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6	Toxicological Reviews of Cyanobacterial
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8	Microcystins LR, RR, YR and LA
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1		LIST OF ACRONYMS
2 3		
4	AFB1	Aflatoxin B1
5	ALP	Alkaline phosphatase
6	ALT	Alanine aminotransferase
7	AST	Aspartate aminotransferase
8	BMD	Benchmark dose
9	BMDL	Benchmark dose, lower confidence limit
10	BMDS	Benchmark dose software
11	BMR	Benchmark response
12	BUN	Blood urea nitrogen
13	CI	Confidence interval
14	DEN	Diethylnitrosamine
15	ELISA	Enzyme-linked immunosorbent assay
16	EPA	Environmental Protection Agency
17	ESR	Electron spin resonance
18	FETAX	Frog Embryo Teratogenicity Assay-Xenopus
19	GD	Gestation day
20	GFR	Glomerular filtration rate
21	GGT	γ-Glutamyltransferase
22	GST-P	Glutathione S-transferase (placental form)
23	HPLC	High pressure liquid chromatography
24	i.p.	Intraperitoneal
25	i.v.	Intravenous
26	LC_{50}	Concentration lethal to 50% of population
27	LD_{50}	Dose lethal to 50% of population
28	LDH	Lactate dehydrogenase
29	LOAEL	Lowest-observed-adverse-effect level
30	MAPK	Mitogen-activated protein kinase
31	MCAR	Microcystin-AR

- 1 MCLA Microcystin-LA
- 2 MCLR Microcystin-LR
- 3 MCRR Microcystin-RR
- 4 MCYM Microcystin-YM
- 5 MCYR Microcystin-YR
- 6 MPT Mitochondrial permeability transition
- 7 MW Molecular weight
- 8 NMR Nuclear magnetic resonance
- 9 NOAEL No-observed-adverse-effect level
- 10 OATP Organic acid transport protein
- 11 PAS Periodic acid-Schiff
- Point of departure
- Protein phosphatase 1
- Protein phosphatase 2A
- 15 RfC Reference concentration
- 16 RfD Reference dose
- 17 ROS Reactive oxygen species
- 18 RVR Renal vascular rate
- 19 SDH Sorbitol dehydrogenase
- 20 SRR Standardized rate ratio
- 21 TDI Tolerable Daily Intake
- 22 TEF Toxicity equivalency factor
- 23 TUNEL Terminal deoxynucleotidyl transferase-mediated dUTP-biotin nick end-labeling
- 24 UF Uncertainty factor
- 25 VAF Virus antibody free
- 26 WHO World Health Organization

PREFACE

 The Safe Drinking Water Act (SDWA), as amended in 1996, requires the Environmental Protection Agency (EPA) to publish a list of contaminants which, at the time of publication, are not subject to any proposed or promulgated national primary drinking water regulations, that are known or anticipated to occur in public water systems, and which may require regulations under SDWA. This list, known as the Contaminant Candidate List (CCL), was first published in 1998 and then again in 2005. The 1998 and 2005 CCLs include "cyanobacteria (blue-green algae), other freshwater algae, and their toxins" as microbial contaminants.

In 2001, a meeting was held among EPA, researchers from the drinking water industry, academia and government agencies with expertise in the area of fresh water algae and their toxins. The goal of this meeting was to convene a panel of scientists to assist in identifying a target list of algal toxins that are likely to pose a health risk in source and finished waters of the drinking water utilities in the U.S. Toxin selection was based on four criteria: health effects, occurrence in the United States, susceptibility to drinking water treatment and toxin stability. Microcystins were identified at this meeting as being toxins of high priority based on those criteria.

The National Center for Environmental Assessment has prepared this Toxicological Review of Cyanobacterial Toxins: Microcystins (LR, RR, YR and LA) as one in a series of doseresponse assessments to support the health assessment of unregulated contaminants on the CCL. The purpose of this document is to compile and evaluate the available data regarding microcystin toxicity to aid the Office of Water in regulatory decision making. It is not intended to be a comprehensive treatise on the chemical or toxicological nature of microcystins.

In Section 6, Major Conclusions in the Characterization of Hazard and Dose Response, EPA has characterized its overall confidence in the quantitative and qualitative aspects of the hazard and dose response by addressing knowledge gaps, uncertainties, quality of data and scientific controversies. The discussion is intended to convey the limitations of the assessment and to aid and guide the Office of Water in the ensuing steps of the human health risk assessment of microcystins.

1	AUTHORS, CONTRIBUTORS AND REVIEWERS
2	
3	
4	AUTHORS
5	
6	Belinda Hawkins, Ph.D. (Chemical Manager)
7	National Center for Environmental Assessment
8	Office of Research and Development
9	U.S. Environmental Protection Agency
10	Cincinnati, OH
11	Hardan Cadan Hamal
12	Heather Carlson-Lynch
13	Syracuse Research Corporation
14 15	Syracuse, NY
16	Marc Odin
17	Syracuse Research Corporation
18	Syracuse, NY
19	Sylucuse, 141
20	Julie Stickney
21	Syracuse Research Corporation
22	Syracuse, NY
23	
24	
25	REVIEWERS
26	
27	INTERNAL EPA REVIEWERS
28	
29	Joyce Donohue, Ph.D.
30	Office of Water
31	Washington, DC
32	
33	Elizabeth Hilborn, D.V.M., M.P.H.
34	National Health and Environmental Effects Research Laboratory
35	Office of Research and Development
36	Research Triangle Park, NC
37	James Cinclair Dh D
38	James Sinclair, Ph.D. Office of Water
39 40	Cincinnati, OH
41	Cilicillian, Off
42	Jeff Swartout
43	National Center for Environmental Assessment
44	Office of Research and Development
45	Cincinnati, OH

1. INTRODUCTION

This toxicological review presents background and justification for hazard and dose-response assessments of microcystins LR, RR, YR and LA. U.S. Environmental Protection Agency (EPA) toxicological reviews may include oral reference dose (RfD) and inhalation reference concentration (RfC) values for chronic and less-than-lifetime exposure durations and a carcinogenicity assessment.

 The RfD and RfC provide quantitative information for use in risk assessments for health effects known or assumed to be produced through a nonlinear (possibly threshold) mode of action. These reference values are defined as an estimate of an exposure, designated by duration and route, to the human population (including susceptible subgroups), that is likely to be without an appreciable risk of adverse effects. Reference values may be derived for acute (≤24 hours), short-term (up to 30 days), subchronic (up to 10% of average lifespan) and chronic (up to lifetime) exposures, all considered to be continuous exposures throughout the duration specified. A reference value is derived from a BMDL (a statistical lower confidence limit on the benchmark dose), a no-observed-adverse effect level (NOAEL), lowest-observed-adverse-effect level (LOAEL) or other suitable point of departure with uncertainty/variability factors applied to reflect limitations of the data used. The RfD is expressed in units of mg/kg-day, and the RfC in units of mg/m³.

The carcinogenicity assessment provides information on the carcinogenic hazard potential of the substance in question and quantitative estimates of risk from oral exposure and inhalation exposure. The information includes a weight-of-evidence judgment of the likelihood that the agent is a human carcinogen and the conditions under which the carcinogenic effects may be expressed. Quantitative risk estimates are presented in three ways. The *slope factor* is the result of application of a low-dose extrapolation procedure and is presented as the risk per mg/kg-day. The *unit risk* is the quantitative estimate in terms of either risk per μ g/L drinking water or risk per μ g/m³ air breathed. Another form in which risk is presented is a drinking water or air concentration providing cancer risks of 1 in 10,000; 1 in 100,000; or 1 in 1,000,000.

Development of these hazard identification and dose-response assessments for microcystins has followed the general guidelines for risk assessment as set forth by the National Research Council (1983). EPA guidelines that were used in the development of this assessment include the following: Guidelines for the Health Risk Assessment of Chemical Mixtures (U.S. EPA, 1986a), Guidelines for Mutagenicity Risk Assessment (U.S. EPA, 1986b), Guidelines for Developmental Toxicity Risk Assessment (U.S. EPA, 1991), Guidelines for Reproductive Toxicity Risk Assessment (U.S. EPA, 1996), Guidelines for Neurotoxicity Risk Assessment (U.S. EPA, 1998a), Guidelines for Carcinogen Risk Assessment (U.S. EPA, 2005a), Supplemental Guidance for Assessing Susceptibility from Early-Life Exposure to Carcinogens (U.S. EPA, 2005b), Recommendations for and Documentation of Biological Values for Use in Risk Assessment (U.S. EPA, 1988), (proposed) Interim Policy for Particle Size and Limit Concentration Issues in Inhalation Toxicity (U.S. EPA, 1994a), Methods for Derivation of Inhalation Reference Concentrations and Application of Inhalation Dosimetry (U.S. EPA, 1994b), Use of the Benchmark Dose Approach in Health Risk Assessment (U.S. EPA, 1995), Science Policy Council

Handbook: Peer Review (U.S. EPA, 1998b, 2000a, 2005c), Science Policy Council Handbook: Risk Characterization (U.S. EPA, 2000b), Benchmark Dose Technical Guidance Document (U.S. EPA, 2000c), Supplemental Guidance for Conducting Health Risk Assessment of Chemical Mixtures (U.S. EPA, 2000d) and A Review of the Reference Dose and Reference Concentration Processes (U.S. EPA, 2002).

Microcystins are a group of at least 80 naturally occurring hepatotoxins produced by freshwater cyanobacteria (blue-green algae) including *Microcystis*, *Anabaena*, *Nodularia*, *Nostoc* and *Oscillatoria* (Duy et al., 2000). Microcystins were first isolated from cyanobacterial extracts in the 1980s (WHO, 1999).

Much of the toxicological research on microcystins has focused on a single congener, microcystin-LR (MCLR). In addition to MCLR, this report focuses on three other major microcystin congeners: microcystin-YR, microcystin-RR and microcystin-LA (abbreviated as MCYR, MCRR and MCLA throughout this document). Literature searches were conducted for studies relevant to the derivation of toxicity and carcinogenicity values for these four microcystin congeners. The following databases were searched: MEDLINE (PubMed), TOXLINE, BIOSIS, CANCERLIT, TSCATS, CCRIS, DART/ETIC, EMIC, GENETOX, HSDB

and RTECS. The relevant literature was reviewed through May 2006.

Microcystins are monocyclic heptapeptide toxins produced by a number of cyanobacterial species, including members of *Microcystis*, *Anabaena*, *Nodularia*, *Nostoc* and *Oscillatoria* (Duy et al., 2000). At least 80 microcystin congeners have been identified. A general structure for microcystins is shown in Figure 2-1.

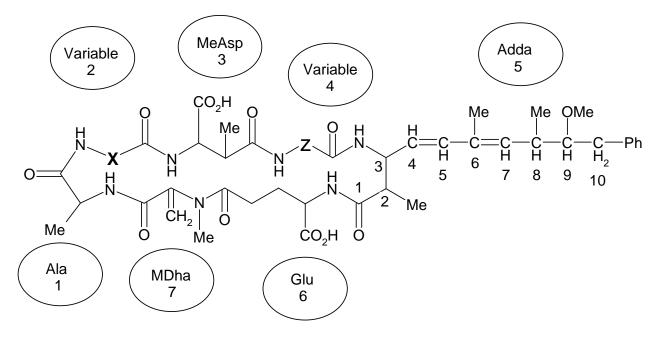


Figure 2-1. General Structure of Microcystins

Microcystins are monocyclic heptapeptides consisting of D-alanine (Ala); two variable amino acids (at positions X and Z in Figure 2-1); D- β -methylaspartic acid (MeAsp); (2S,3S,8S,9S)-3-amino-9-methoxy-2,6,8-trimethyl-10-phenyldeca-4,6-dienoic acid (Adda); isolinked D-glutamic acid (Glu) and *N*-methyl dehydroalanine (MDha). Structural variations occur in all seven of the amino acid peptides, but most commonly in the L-amino acids at positions X and Z in Figure 2-1 (shown as "variable" amino acids in the figure). The most common L-amino acids at position X are leucine (L), arginine (R) and tyrosine (Y), while those at position Z are arginine (R) and alanine (A). The congeners take their names from the L-amino acids in these positions. For example, the microcystin congener with leucine in the X position and arginine in the Z position is microcystin-LR.

 Little information on the chemical and physical properties of microcystins was located. Duy et al. (2000) provided the most thorough (albeit general) review of the properties of microcystins. The microcystins identified to date have molecular weights (MWs) ranging from 900 to 1200. Microcystins are nonvolatile and generally quite hydrophilic, although a few have lipophilic properties. Microcystins are soluble in water, ethanol and methanol, and insoluble in

- acetone, ether, chloroform and benzene. Laboratory studies show microcystins to be stable
- 2 under changes in temperature and pH. Microcystins are stable in sunlight; however, in the
- 3 presence of pigments (type unspecified) and sunlight, microcystins can be decomposed or
- 4 isomerized (Duy et al., 2000). Limited information suggests that microcystins can
- 5 bioaccumulate in aquatic organisms; these toxins have been measured in a number of fish and
- 6 aquatic invertebrates.

3. TOXICOKINETICS

The available information on toxicokinetics of microcystins is primarily focused on exposure via injection routes, either intravenous or intraperitoneal. Few data are available on the oral and inhalation exposure routes. No data on the absorption, metabolism or elimination of microcystins after *in vivo* oral or inhalation exposure in humans or animals were located in the literature. Toxicokinetic data available from parenteral routes of exposure are of uncertain relevance to oral and inhalation exposure routes. The database on microcystins does not contain any toxicokinetic models for microcystins.

A number of studies on the toxicokinetics of microcystins have used ³H-dihydro-MCLR (usually produced by reduction of the MDha moiety in MCLR with tritiated sodium borohydride) as a test material. While there are similarities between the organ distribution, hepatocellular uptake and clinical syndrome after exposure to dihydroMCLR and MCLR (Meriluoto et al., 1990), there are differences in the binding of these compounds to molecular targets. Craig et al. (1996) showed that, while dihydroMCLA was capable of inactivating protein phosphatase 2Ac through a rapid binding mechanism, it did not subsequently form a covalent bond with PP2Ac, while MCLA did. Further information on the structural requirements for microcystin toxicity is provided in Section 4.4.9. Potential differences in the binding of dihydro-microcystin analogs dictates that caution be exercised in generalizing toxicokinetic information derived using these compounds to microcystins as a group. In particular, information on the subcellular localization of dihydromicrocystins may not be applicable to microcystins containing an intact MDha residue.

3.1. ABSORPTION

Pulmonary absorption of MCLR (purified from a bloom sample) was rapid following intratracheal instillation in mice (Ito et al., 2001). Immunostaining of the lung occurred within 5 minutes, followed by a lag period of 60 minutes before staining was observed in the liver. Based on the positive immunostaining of alveoli, it was concluded that absorption occurred at the alveoli. The lethal dose level for intratracheal injection was similar to a lethal dose for intraperitoneal (i.p.) injection (i.e., both about $100~\mu g/kg$). The authors reported that the lungs were not affected by microcystin administration, but it is unclear whether a detailed histopathological evaluation of the lungs was conducted in addition to the immunostaining.

The occurrence of hepatotoxicity and lethality following oral exposure to microcystins (see Section 4.2) is evidence of oral absorption of the toxin. However, quantitative assessments of oral absorption were not located. Ito et al. (1997a) qualitatively studied the oral absorption and distribution of MCLR (purified from a bloom sample) following gavage dosing in mice (500 μ g/kg). Immunostaining techniques indicated that MCLR was absorbed primarily in the small intestine, although some absorption did occur in the stomach. Erosion was observed in the surface epithelial cells and in the lamina propria of the small intestine villi. Erosion may facilitate uptake of the toxin into the bloodstream.

The oral bioavailability of MCLR was indirectly studied in *in situ* experiments using isolated intestinal loops of rats (Dahlem et al., 1989). Rats given an infusion of MCLR (>95% pure by high pressure liquid chromatography (HPLC) with UV detection) into the ileum showed clinical signs (i.e., labored breathing and circulatory shock) and evidence of liver toxicity within 6 hours of a single 5 mg/kg dose. Liver toxicity was assessed as an increase in the liver-to-body weight ratio and the presence of gross and histopathological liver lesions characteristic of microcystin toxicity (i.e., enlarged livers, hepatocyte rounding and disassociation, hemorrhage). Infusion of a similar dose into a jejunal loop produced a lower degree of liver toxicity, as compared to the ileal loop infusion. These results suggest that there could be site-specificity in intestinal absorption of MCLR; however, differences in absorptive surface area were not taken into account in the experiment. *In vitro* experiments reported in this publication indicated that cholestyramine, a bile acid sequestrant, bound MCLR, and the presence of cholestyramine in the ileal loop infusion significantly reduced MCLR liver toxicity (Dahlem et al., 1989).

Oral absorption of ³H-dihydromicrocystin was also demonstrated using ileal loop exposure in swine (Stotts et al., 1997a,b). In the exposed swine, the maximum blood concentration of the toxin occurred 90 minutes after dosing.

3.2. DISTRIBUTION

The distribution of microcystins is limited due to the poor ability of these toxins to cross cell membranes. Microcystins are primarily taken up into the liver by the multispecific active transport system for bile acids. Once inside the cell, these toxins bind covalently to cytosolic proteins, resulting in retention in the liver. The cytosolic proteins bound by microcystins have been identified as the protein phosphatase enzymes (PP1 and PP2A). It should be noted that dihydromicrocystin analogs do not appear to form covalent bonds with PP1 and PP2A, although they are able to rapidly bind and inactivate the enzymes (Craig et al., 1996). Binding to and inhibition of these phosphatase enzymes are directly related to the mechanism of toxicity for microcystins and are further discussed in Section 4.4.7.3. This section will describe the overall organ distribution, cellular uptake, subcellular localization and protein binding of microcystins.

3.2.1. Organ Distribution

The organ distribution of a ¹²⁵I-labelled heptapeptide toxin (MW 1019) isolated from *Microcystis aeruginosa* (while not identified by the study authors as such, probably because the toxin had not yet been named, this is assumed to be a microcystin) was investigated in female rats following intravenous (i.v.) administration (Falconer et al., 1986; Runnegar et al., 1986). The heptapeptide toxin was purified by HPLC prior to reaction with ¹²⁵I in the presence of NaI and lactoperoxidase. Labelling of the toxin was confirmed by HPLC and mouse bioassay. The highest tissue concentrations of microcystins were detected in the liver and kidney. After 30 minutes, 21.7% of the administered dose was present in the liver, 5.6% was present in the kidneys, 7% remained in the gut contents, and 0.9% was cleared in the urine (Falconer et al., 1986). The balance of the administered dose was not reported; however, the authors reported that no significant accumulation was observed in other organs or tissues.

Brooks and Codd (1987) reported extensive liver uptake following i.p. injection of 125 μg/kg of a ¹⁴C-labelled toxin extracted from *M. aeruginosa* strain 7820 (assumed to be a microcystin) in mice. Seventy percent of the radiolabel was found in the liver after 1 minute, increasing to almost 90% after 3 hours. Radiolabel was also found in the lungs, kidneys, heart, large intestine, ileum and spleen.

The distribution of 3 H-MCLR (>95% pure) was evaluated following i.p. injection of a sublethal (45 µg/kg) or lethal (101 µg/kg) dose in mice (Robinson et al., 1989). The tissue distribution of radiolabel was similar after injection of either a lethal or a sublethal dose. Liver accumulation reached a maximal value of 60% by 60 minutes. For the 101 µg/kg dose, the liver, intestine and kidney contained 56, 7 and 0.9% of the radiolabel, respectively. Heart, spleen, lung and skeletal muscle each contained less than 1% of the radiolabel.

Immunostaining methods were used to evaluate the organ distribution following intratracheal instillation of MCLR purified from a bloom sample (Ito et al., 2001). Following instillation of a lethal dose (100 μ g/kg), the lung, liver, small intestine and kidney were positively stained for MCLR. Intense staining was observed in the lung by 5 minutes post-instillation, followed by the kidney (10 minutes), the small intestine (45 minutes) and the liver (60 minutes). After approximately 90 minutes, bleeding began around the hepatic central vein. The authors described the pathological changes in the liver as essentially the same as those seen following oral or i.p. injection exposure routes. Intratracheal instillation of a sublethal dose (50 μ g/kg) resulted in immunostaining of the lung, liver, kidney, cecum and large intestine (Ito et al., 2001). No discernable pathological changes were observed at this dose level. Ito et al. (2002) synthesized glutathione and cysteine conjugates of microsystin-LR and administered them by intratracheal instillation in mice. These conjugates are, according to the authors, known metabolites of microcystins. The metabolites were demonstrated to be less toxic than the parent compound (lethal doses were about 12-fold higher than the MCLR lethal dose) and were distributed primarily to the kidney and intestine, as opposed to the liver.

The distribution of MCLR (purified from a bloom sample) following oral gavage administration to mice ($500 \mu g/kg$) was investigated using immunostaining methods (Ito et al., 1997a). MCLR was detected in large amounts in the villi of the small intestine. Erosion of the villi was observed, which may have enhanced absorption of the toxin into the bloodstream. MCLR was also present in the blood plasma, liver, lungs, kidneys and heart.

The distribution of ³H-dihydroMCLR in mice was shown to differ for the oral and i.p. injection routes of exposure (Nishiwaki et al., 1994). Intraperitoneal injection of ³H-dihydroMCLR resulted in rapid and continuous uptake in the liver, with approximately 72% of the administered dose present in the liver after 1 hour. The ³H-dihydroMCLR was synthesized by reduction of N-methyldehydroalanine from microcystin-LR. Small amounts of radiolabel were found in the small intestine (1.4%), kidney and gallbladder (0.5%), lungs (0.4%) and stomach (0.3%) following i.p. injection. Oral administration of ³H-dihydroMCLR resulted in much lower concentrations in the liver, with less than 1% of the administered dose found in the liver at either 6 hours or 6 days post administration. ³H-DihydroMCLR is rapidly distributed to the liver of swine following i.v. injection or ileal loop infusion (Stotts et al., 1997a,b). Smaller amounts were distributed to the kidneys, lungs, heart, ileum and spleen.

MCLR was not found in the milk of dairy cattle that were exposed to *M. aeruginosa* cells via drinking water (Orr et al., 2001) or ingestion of gelatin capsule containing the cells (Feitz et al., 2002).

3.2.2. Cellular Uptake

The cellular uptake of ³H-dihydroMCLR was evaluated using primary rat hepatocytes in suspension and in isolated perfused rat liver (Eriksson et al., 1990a; Hooser et al., 1991a). The uptake (as measured by scintillation counting of washed cells) of a mixture of unlabelled MCLR and ³H-dihydroMCLR was shown to be specific for freshly isolated rat hepatocytes (Eriksson et al., 1990a). Uptake was negligible in human hepatocarcinoma cells (Hep G2), mouse fibroblast (NIH-3T3) and human neuroblastoma cells (SH-SY5Y). The uptake of ³H-dihydroMCLR was also shown to be inhibited by bile acid transport inhibitors such as antamanide, sulfobromophthalein and rifampicin, and by the bile salts cholate and taurocholate (by competing for the bile acid transporter).

The uptake of ³H-dihydroMCLR was demonstrated to be rapid for the first 5-10 minutes, followed by a plateau, in both rat hepatocyte suspensions and the isolated perfused rat liver (Hooser et al., 1991a). Uptake was measured as radioactivity in fractionated cells versus radioactivity in medium. The uptake of ³H-dihydroMCLR was inhibited by incubation of suspended rat hepatocytes at 0°C, suggesting the involvement of an energy-dependent process (Hooser et al., 1991a). Uptake was also inhibited by preincubation of hepatocytes with rifampicin, presumably via competitive inhibition of the bile acid transporter (Hooser et al., 1991a).

Many studies have demonstrated that inhibition of microcystin uptake at the bile acid transporter reduces or eliminates the liver toxicity observed following *in vitro* or *in vivo* exposures (Runnegar et al., 1981, 1995a; Runnegar and Falconer, 1982; Hermansky et al., 1990a,b; Thompson and Pace, 1992). The human organic acid transport protein (OATP) was shown to mediate the transport of ³H-microcystin (type not specified) in *Xenopus laevis* oocytes, and this uptake was inhibited by sulfobromophthalein and taurocholate. This transport protein is found in the human brain and may be related to the acute neurotoxicity seen in hemodialysis patients exposed to microcystins (see Section 4.4.5.1).

Runnegar et al. (1991) studied the influence of dose level and exposure time on the uptake of ¹²⁵I-microcystin-YM in isolated rat hepatocytes (measured as radioactivity in centrifuged cell pellet). Hepatocyte uptake was initially rapid with a plateau in the uptake rate observed after 10 minutes. The initial uptake rate (in the first minute of exposure) increased with increasing concentration, but cumulative uptake ceased at a dose that resulted in plasma membrane blebbing.

 Microcystin-YM uptake by isolated rat hepatocytes was temperature-dependent and was inhibited 70-80% by the addition of sodium deoxycholate or sulfobromophthalein (Runnegar et al., 1995b). This provides evidence to indicate that microcystin uptake occurs by carrier mediated transport, most likely via the bile acid transporter. Pretreatment of mice with bile acid

transporter inhibitors (cyclosporine A, rifamycin, trypan blue and trypan red) abolished 1 microcystin toxicity, suggesting limited or no uptake of microcystins (Runnegar et al., 1995b). 2 Further, *in vitro* preincubation of hepatocytes with bile acids or bile acid transport inhibitors 3 (taurocholate, trypan blue, cholate, sulfobromophthalein, cyclosporine A, trypan red and 4 rifamycin) each decreased the uptake of microcystin-YM, as measured by assays for protein 5 phosphatase inhibition in cell lysates (Runnegar et al., 1995b). Pretreatment with protein 6 phosphatase inhibitors (i.e., microcystin-YM and calyculin A) also resulted in the inhibition of 7 both microcystin-YM uptake and protein phosphatase inhibition, suggesting that the bile acid 8 transporter is itself regulated by serine/threonine phosphorylation. 9

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Many cell types and established cell lines, including both rodent and some human cells, have been evaluated for potential susceptibility to microcystin uptake and toxicity. Primary isolated hepatocytes have been shown to be the most sensitive to cytotoxicity, due to the presence of the organic ion/bile acid transport system (Eriksson et al., 1990b). In addition, primary cultures of liver cells cease to express these bile acid transport proteins after 2-3 days of being maintained in culture. Therefore, established liver cell lines are generally not useful for evaluating microcystin toxicity (Eriksson and Golman, 1993; Heinze et al., 2001). Chong et al. (2000) evaluated microcystin toxicity in eight permanent cell lines (including rodent, primate and human cell lines), only two of which (human oral epidermoid carcinoma KB cells and rat Reuber H35 hepatoma H-4-II-E cells) showed cytotoxicity following MCLR exposure. The toxic response in these cells was most evident if MCLR was added when the cells were seeded. Established monolayers were more resistant to microcystin toxicity. Mechanistic studies that evaluate organ and cell type specificity for microcystins are further discussed in Section 4.4.7.1.

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3.2.3. Subcellular Localization and Cytosolic Protein Binding

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Tissue distribution was evaluated in mice given i.v. injection of a sublethal dose of ³H-MCLR (Robinson et al., 1991a). The liver contained approximately 67% of the radiolabel by 60 minutes, and the amount of hepatic radioactivity did not change throughout the 6-day study period, despite urinary and fecal elimination of 24% of the administered dose. The subcellular distribution of radioactivity in the liver demonstrated that approximately 70% of the hepatic radiolabel was present in the cytosol. *In vitro* experiments showed that radiolabeled microcystin in the liver was bound to high molecular weight cytosolic proteins (Robinson et al., 1991b). The nature of the binding was demonstrated to be covalent, saturable and specific for a protein with a molecular weight of approximately 40,000. Binding was inhibited by okadaic acid (a potent inhibitor of serine/threonine phosphatases [1 and 2A]), suggesting that the target protein is protein phosphatase 1 or 2A. A discussion of protein phosphatase binding and inhibition by microcystins is provided under mechanistic studies in Section 4.4.7.3, below. Binding proteins for MCLR were found in cytosol derived from several different organs, suggesting that liver specificity is not due to limited distribution of target proteins. Covalent binding to hepatic proteins may be responsible for the long retention of microcystins in the liver. Lin and Chu (1994) evaluated the kinetics of MCLR distribution in serum and liver cytosol derived from mice. Uptake of pure MCLR, as analyzed by direct competitive enzyme-linked immunosorbent assay (ELISA), into the serum was shown to be rapid following an i.p. injection of 35 µg/kg (sublethal dose). The toxin reached a maximum concentration in the serum by 2 hours and in liver cytosol by 12 hours post-injection. MCLR was shown to be bound to liver cytosolic

proteins and the kinetics of binding were correlated with inhibition of protein phosphatase 2A activity. The maximum decrease in enzyme activity was observed 6-12 hours following injection.

Pace et al. (1991) demonstrated significant accumulation of ³H-MCLR in isolated perfused liver despite a low overall extraction ratio (16% in liver, 79% in perfusate). In the liver, radiolabel corresponding to MCLR (15%) and a more polar metabolite (85%) was primarily found in the cytosolic fraction.

The subcellular distribution of ³H-dihydroMCLR was evaluated using primary rat hepatocytes in suspension and the isolated perfused rat liver (Hooser et al., 1991a). ³H-dihydroMCLR was primarily localized in the cytosolic fraction in both the hepatocytes and liver. In the hepatocytes, precipitation with trichloroacetic acid indicated that approximately 50% of the ³H-dihydroMCLR was found as free toxin, while the remaining 50% was bound to cytosolic proteins. Since little of the radiolabel was in the insoluble pellet containing insoluble actin and other elements, the authors suggested that ³H-dihydroMCLR did not bind significantly to actin or other cytoskeletal proteins (Hooser et al., 1991a).

The subcellular protein binding of ³H-dihydroMCLR was evaluated in rat liver homogenates (Toivola et al., 1994). Most of the radiolabeled toxin (80%) was bound to cytosolic proteins. ³H-dihydroMCLR was shown to bind both protein phosphatase 1 (PP1) and protein phosphatase 2A (PP2A); however, PP2A was detected primarily in the cytosol, while PP1 was found in the mitochondrial and post-mitochondrial particulate fraction (membrane proteins). The binding of microcystins to PP1 and PP2A and the inhibition of protein phosphatase activity is further discussed in Section 4.4.7.3 (Mechanistic Studies).

Limited information in humans exposed to microcystins intravenously via dialysate indicates that a large proportion of microcystins in the blood are bound to proteins. Hilborn et al. (2005) compared two techniques for measuring microcystin in the serum of six patients. Use of ELISA, which detects free microcystins, resulted in serum microcystin concentrations ranging from 8 to 51% of the concentrations obtained using gas chromatography/mass spectrometry (GC/MS) detection of 2-methyl-3-methoxy-4-phenylbutyric acid (MMPB, which is derived from both free and protein-bound microcystins by chemical oxidation). These results indicate that microcystins are bound to proteins in human blood, and that analysis for microcystins using ELISA techniques may underestimate total blood concentrations.

3.3. METABOLISM

 Urinary and fecal metabolites of MCLR were analyzed in samples collected 6 and 12 hours following i.v. injection of a sublethal dose of ³H-MCLR in mice (Robinson et al., 1991a). Approximately 60% of the radiolabel in both the urine and the feces was associated with the parent compound. MCLR was metabolized in liver cytosol preparations to a product that binds to a high molecular weight cytosolic protein (Robinson et al., 1991b). The parent compound also binds to this protein, which has been suggested to be the catalytic subunit of protein phosphatase 2A. In isolated perfused rat liver, binding of both the parent toxin (³H-MCLR) and a more polar metabolite to cytosolic proteins was also demonstrated (Pace et al., 1991). Polar

metabolites accounted for 65-85% of the hepatic cytosol radiolabel. Metabolites of MCLR were not further characterized in these studies.

³H-Dihydromicrocystin is not extensively metabolized in swine liver after i.v. injection or ileal loop exposure, and is primarily present in hepatic tissues as the parent compound (Stotts et al., 1997a,b).

Administration of 125 μg/kg of *Microcystis* toxin 7820 to mice resulted in decreased levels of cytochrome b5 and cytochrome P450 (Brooks and Codd, 1987). Pretreatment of mice with microsomal enzyme (mixed function oxidase) inducers (β-naphthoflavone, 3-methylcholanthrene and phenobarbital) was shown to eliminate this effect on hepatic enzymes and to extend survival and reduce liver toxicity (i.e., changes in liver weight). In an *in vitro* study using mouse liver microsomes, cytochrome P450 associated enzyme activity (i.e., metabolism of aminopyrene and p-nitrophenol) was not altered by an unidentified toxin isolated from *M. aeruginosa* (assumed to be a microcystin; Cote et al., 1986).

The hepatic metabolism of MCRR and MCLR (purified from blooms) was studied following i.p. injection in mice and rats (Kondo et al., 1996). Glutathione and cysteine conjugates were identified at 3 and 24 hours in both mouse and rat livers. Structural modification of the 3-amino-9-methoxy-2,6,8,-trimethyl-10-phenyldeca-4,6-dienoic acid (Adda) and methyldehydroalanine (MDha) moieties of the toxins was indicated. Figure 3-1 shows the glutathione and cysteine conjugates of microcystins.

Kondo et al. (1992) demonstrated that glutathione and cysteine conjugates of MCLR and MCYR were less toxic than the parent compounds based on LD₅₀ estimates, but were still significantly toxic (LD₅₀ values ranged from 217 to 630 μg/kg in mice). Glutathione and cysteine conjugates of MCLR were shown to inhibit protein phosphatases 1 and 2A *in vitro* to the same degree as MCLR; however, these metabolites were primarily distributed to the kidney and intestine following intratracheal instillation in mice (Ito et al., 2002). This result suggests that the lower toxicity of glutathione and cysteine conjugates may be related to distribution to excretory organs and elimination of metabolites *in vivo*. Metcalf et al. (2000) also demonstrated that microcystin conjugates with glutathione, cysteine-glycine and cysteine were less toxic in the mouse bioassay; however, these conjugates were also shown to be weaker inhibitors of protein phosphatases 1 and 2A *in vitro*. Takenaka (2001) illustrated that glutathione conjugates of MCLR are formed by glutathione S-transferase enzymes found in both rat liver cytosol and microsomes.

$$\frac{X \quad Y}{\text{Microcystin LR Leu Arg}}$$

$$\frac{X \quad Y}{\text{Microcystin YR Tyr Arg}}$$

Microcystin RR Arg Arg

Cys conjugate
$$Z = H_2N$$
 COOH

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Figure 3-1. Structures of GSH and Cys Conjugates of Microcystins LR, YR and RR (Kondo et al., 1992)

СООН

Several studies have demonstrated an increase in hepatic glutathione levels following exposure to microcystins (Ding et al., 2000a; Bouaïcha and Maatouk, 2004; Gehringer et al., 2004). MCLR was shown to induce the *de novo* synthesis of glutathione in mice exposed to a toxic sublethal dose (75% of the LD₅₀) (Gehringer et al., 2004). Increased transcription of glutathione-S-transferase was also demonstrated in this study.

3.4. ELIMINATION

Limited information on the elimination of microcystins from the human body is available from follow-up of dialysis patients exposed to microcystins intravenously (see Section 4.4.5.1 for further detail). In two separate incidents in Brazil (one in Caruaru, one in Rio de Janeiro), microcystins were detected in patients' serum more than 50 days after documented exposure (Hilborn et al., 2005; Soares et al., 2006).

The excretion of microcystins was investigated in female rats (Falconer et al., 1986). The blood half-life was measured following i.v. administration of a ¹²⁵I-labelled heptapeptide toxin extracted from *M. aeruginosa* (MW 1019, assumed to be a microcystin). A biphasic blood elimination curve was demonstrated, with the first component having a half-life of 2.1 minutes and the second component having a half-life of 42 minutes. After 120 minutes, 9.4% of the administered dose was present in the intestinal contents and 2.9% was present in the urine, suggesting that biliary excretion plays a significant role in elimination of microcystins. Biliary excretion was also demonstrated in isolated perfused rat liver (Pace et al., 1991). In the bile collected 10-20 minutes after toxin exposure, 78% of the radiolabel was associated with the parent toxin, while the remaining radiolabel was associated with more polar metabolites.

 MCLR excretion was also evaluated in mice (Robinson et al., 1991a). A biexponential plasma elimination curve was observed following i.v. injection of a sublethal dose of ³H-MCLR. Plasma half-lives of 0.8 and 6.9 minutes were reported for the first and second phase of elimination, respectively. Approximately 24% of the administered dose was eliminated in the urine (9%) and feces (15%) throughout the 6-day study period. Approximately 60% of the excreted microcystin, measured at 6 and 12 hours following injection, was present as the parent compound.

Ito et al. (1997a) demonstrated that MCLR is secreted in the mucous of goblet cells from both the small and large intestine of mice following administration by oral gavage (500 μ g/kg). MCLR was not detected in urine in this study.

 Stotts et al. (1997a,b) evaluated the toxicokinetics of ³H-dihydroMCLR in swine following i.v. injection and ileal loop exposure. Elimination of ³H-dihydromicrocystin was rapid and followed a biphasic pattern, suggesting that the liver rapidly removes the toxin from the blood. Clearance from the blood is slower at higher dose levels, presumably due to the liver toxicity and circulatory shock observed at high doses. ³H-Dihydromicrocystin was detected in the bile as early as 30 minutes after i.v. injection. Following ileal loop exposure, the concentration of toxin was consistently higher in the portal venous blood as compared to peripheral blood. This suggests that first pass metabolism may play a role in the clearance of dihydroMCLR.

1 2 3.5. PHYSIOLOGICALLY-BASED TOXICOKINETIC MODELS

No physiologically based toxicokinetic models have been developed for microcystins.

4. HAZARD IDENTIFICATION

4.1. STUDIES IN HUMANS – EPIDEMIOLOGY, CASE REPORTS, CLINICAL CONTROLS

A number of case reports and epidemiological investigations have examined the relationship between human exposure to cyanobacteria and various health endpoints. In all of these studies, humans were exposed to blooms of cyanobacteria in environmental settings. As a result, the potential for co-exposure to multiple cyanobacterial toxins and/or other microorganisms or compounds to contribute to observed toxicity cannot be ruled out.

4.1.1. Oral Exposure

4.1.1.1. Short-Term Studies and Case Reports

Dillenberg and Dehnel (1960) reported on a series of animal poisonings and human exposures to cyanobacterial blooms in various lakes of Saskatchewan, Canada during the previous year. Several cases of individual or group human exposures during recreational activities were reported. In general, the symptoms were gastrointestinal in nature, including nausea, stomach pain and diarrhea; headache and muscle weakness were also reported. Stool samples from three of the victims showed evidence of cyanobacteria (*Microcystis* and *Anabaena*). In addition, water samples from the lakes in which the victims had been swimming showed cyanobacteria. At the time of this report, cyanobacterial toxins had not been fully characterized. Thus, no data on the nature or quantity of toxins in the affected waters were provided.

Billings (1981) reported a series of outbreaks of human illness potentially associated with exposure to cyanobacteria in two Pennsylvania lakes. Swimmers in both lakes reported symptoms, including headache, abdominal cramping, nausea, vomiting, diarrhea, hay fever-like symptoms, ear aches, eye irritation, sore throat, sneezing, runny nose and swollen lips within a few hours of swimming in the waters. Investigation by the state Departments of Environmental Resources and Health served to rule out common bacterial, protozoal and viral agents in the outbreaks. In the second lake, a bloom of *Anabaena* was identified. Indirect evidence (rapid onset of symptoms, absence of other potential causative agents and consistency with previous reports of health effects after exposure to cyanobacteria) led the investigators to postulate a role for exposure to *Anabaena* in the health outcomes.

 Turner et al. (1990) reported a similar type of outbreak among army recruits who had consumed reservoir water during canoe exercises. Detailed case reports were presented for two recruits. Both 16-year-old recruits presented with several days' history of malaise, sore throat, blistering around the mouth, dry cough, pleuritic pain and abdominal pain. One also had experienced vomiting and diarrhea. Physical examination revealed fever, left basal pulmonary consolidation (pneumonia) and abdominal tenderness in both patients. Blood tests revealed low platelet counts in both patients. Both were tested for a variety of pathogens, including *Leptospira*, *Legionella*, *Chlamydia*, *Coxiella*, *Mycoplasma* and influenza and adenovirus, all

with negative result. The authors reported similar symptoms (sore throat, headache, abdominal pain, dry cough, diarrhea, vomiting and blistered mouth) in 16 additional soldiers who had taken part in the canoe exercises. The reservoir contained a bloom of cyanobacteria, primarily *M. aeruginosa*. Further, a sample of the bloom taken the day after the patients were admitted into the hospital showed microcystins to be present, including MCLR. High levels of *Escherichia coli* were also found in reservoir water sampled 2 weeks later. The authors suggested that microcystin exposure may have had a role in the pulmonary consolidation and low platelet count of the two patients, citing evidence from studies in mice (the authors cited Falconer et al., 1981 and Slatkin et al., 1983).

Teixeira et al. (1993) characterized an epidemic of gastroenteritis, primarily among children, in the vicinity of the Itaparica Dam in Bahia, Brazil. The authors collected data on the incidence of treatments for diarrhea between February and May of 1988. Timing of 1118 cases of diarrhea in local health units was recorded, showing a spike in the incidence of gastroenteritis coinciding with the flooding of the Itaparica Dam reservoir. Most (about 70%) of the cases involved children under the age of 5 years. Additional data were collected on the age, residence, symptoms, foods consumed, source of drinking water and travel for 76 patients given outpatient treatment for diarrhea. Fecal, blood and urine samples were collected from these patients and analyzed for chemical and biological contaminants (i.e., bacteriologic, virologic, cholinesterase, heavy metals). In addition, water samples were analyzed for chemical and biological contaminants (i.e., organophosphates, carbamates, heavy metals, fecal coliform). Analysis of biological samples showed no contaminants. Untreated water samples showed high levels of fecal coliform, but samples of treated water did not contain significant levels. Untreated water samples also revealed high counts (1104-9755 units per mL) of Anabaena and Microcystis cells, 4-32 times the World Health Organization (WHO) maximum acceptable cell count for untreated water (300 units/mL at the time). No data were provided on levels of cyanobacteria in the treated water. It is not clear from the publication whether affected persons were exposed to treated or untreated water. This study does not provide information on health effects of microcystin exposure, but provides some circumstantial evidence for gastrointestinal effects from exposure to cyanobacteria.

A case control study investigated the incidence of gastrointestinal and dermatological symptoms among persons exposed to Murray River water (Australia) (el Saadi and Cameron, 1993; el Saadi et al., 1995). Physicians in 8 of 11 towns along the Murray River participated in the study, recruiting 102 gastrointestinal and 86 dermatological cases between January and March, 1992. Gastrointestinal cases were patients with abdominal pain, vomiting or diarrhea; dermatological cases had rash, itching or blistering of the mouth. Control patients (132) were selected as the next patient entering the office after each case, when possible. For each study participant, age, sex, primary source of drinking and domestic water (rain/spring, untreated river water or chlorinated river water from a town supply) and recreational water contact (none, river or lake contact, or other, such as pool contact) during the previous week were recorded. River water samples were collected and cyanobacteria identified and quantified. *Anabaena*, *Aphanizomenon* and *Planktothrix* were the most common cyanobacteria identified in the samples; small numbers of *M. aeruginosa* were infrequently identified. Both univariate and multivariate analysis of the data showed the odds of having gastrointestinal symptoms to be raised in persons drinking chlorinated river water or using untreated river water for domestic

purposes. Likewise, both types of statistical analyses showed increased odds of having dermatological symptoms for persons less than 20 years of age and for persons using untreated river water for domestic purposes. The proportion of patients with gastrointestinal symptoms and the proportion with dermatological symptoms both correlated with mean weekly log cyanobacterial cell count in the river, although the correlation was statistically significant only for gastrointestinal symptoms. However, when upper and lower reaches of the river were analyzed separately, nonsignificant correlations were observed.

No data on cyanobacterial toxins in the river water were provided. The symptoms reported in this study cannot be readily attributed to any particular toxin (if they are indeed associated with toxin exposure rather than exposure to the living cyanobacterial cells) due to the absence of toxin analyses, as well as the identification of genera with potential to produce multiple toxins. In addition, the potential for other microbial or chemical contaminants in the untreated river water was not evaluated in this study.

Pilotto et al. (1997) conducted a prospective study of gastrointestinal and dermatological symptoms among people exposed to cyanobacteria at water recreation sites in Australia. Study participants were individuals 6 years of age and older who were present at one of several water bodies that were both used for recreational purposes and expected to have algal blooms during the summer. Interviewers visited these sites on several Sundays and invited all individuals to participate. Participants completed a questionnaire to evaluate health status and the nature and duration of water-contact activities. In addition, subjects were asked whether they had symptoms or recreational water contact in the 5 days prior to study initiation, in order to control for the effects of prior health conditions and prior exposure on study findings. Five hundred and fourteen persons had either pre-existing symptoms or water contact in the days prior to initial interview. Participants were contacted by telephone 2 and 7 days later, at which time the occurrence of diarrhea, flu-like symptoms, skin rashes, mouth ulcers, fevers or eye or ear infections in the intervening time was recorded. Water samples for cyanobacterial cell count and toxin analysis were collected at 10 a.m. and 2 p.m. on the day of initial interview. Toxin presence was assessed by mouse bioassay (i.p. injection of 500 mg freeze-dried cells/kg body weight).

 Of 1029 persons invited to participate, 921 persons participated in the study (Pilotto et al., 1997). Interviewers were able to contact 845 of these persons by telephone 2 days after initial interview, and 852 persons 7 days later. No differences in the reporting of gastrointestinal and dermatological symptoms were found between those who had water contact and those without water contact (on the day of the initial interview) when contacted 2 days later (Pilotto et al., 1997). However, when subjects with water contact or symptoms prior to initial interview were excluded, a significant trend to increasing symptom occurrence with duration of exposure was observed among persons contacted 7 days after initial interview. Cyanobacterial cell count showed some correlation with symptom occurrence, but presence/absence of hepatotoxins did not. The authors postulated that any association between symptoms and exposure resulted from the allergenic nature of the cells rather than exposure to toxins.

Pilotto et al. (1999) evaluated the relationship between cyanobacterial exposure and perinatal outcomes in an ecological study conducted in Australia. Cyanobacterial monitoring

data (cell counts) were collected from raw drinking water supplies in 156 towns. Perinatal outcome data were obtained from several registries (for calendar years 1992-1994) and the following variables assessed: premature birth (<36 weeks), low birth weight (<2.5 kg), very low birth weight (<1.5 kg) and congenital defects (at least one). Maternal residence at birth was used to assess exposure based on cyanobacterial cell counts. Exposure was assessed at various gestational periods, either as the proportion of time with cyanobacterial exposure (proportion of weeks with non-zero levels) or average alert level (alert level 1 = <2000 cells/mL; alert level 2 = 2000-15,000 cells/mL; alert level 3 = <15,000 cells/mL). Data on 32,700 births were collected, although the numbers with exposure data in different gestational periods varied. A significant difference in the incidence of low birth weight and very low birth weight babies was observed between mothers with and without cyanobacterial exposure during the first trimester. Very low birth weight incidences increased with increasing exposure to cyanobacteria (as measured by the proportion of first trimester with non-zero cyanobacterial cell counts). At the highest exposure category (exposure to cyanobacteria during 100% of first trimester), the odds ratio (OR) was 1.42 (95% confidence interval [CI] = 1.00-2.02). When exposure was assessed as the average alert level (cell concentration interval), there was a significant increase in congenital defects at the highest average alert level of 2.5-3.0 (OR = 2.03, 95% CI = 1.37-3.01), but the trend was not significant. This study had a number of limitations, including a lack of individual exposure data and lack of data on cyanobacteria or toxins in the finished water (after various treatment processes). Further, because the measure of exposure was cyanobacteria rather than toxin, it is difficult to interpret this study in the context of microcystin effects.

Falconer et al. (1983) compared the hepatic enzyme levels in patients served by a public water supply contaminated with a bloom of *M. aeruginosa* with levels in patients living in areas served by other water supplies. Enzymes assessed in the study were γ-glutamyltransferase (GGT), aspartate aminotransferase (AST), alanine aminotransferase (ALT) and alkaline phosphatase (ALP). The study population consisted of all patients served by a single hospital laboratory and referred for liver function tests before, during and after a bloom of *M. aeruginosa* in the Malpas Dam reservoir of Australia. Patients were classified either as residents of the city of Armidale, which uses the reservoir for drinking water supply, or residents of neighboring towns with independent water supplies. Liver function test results within each comparison group were further sorted by date into three categories: testing during the 5 weeks before the first signs of the bloom appeared, testing during the 3-week bloom or the 2 weeks following copper sulfate treatment of the bloom (identified as the high-risk time interval due to the cell lysis and subsequent toxin release) or testing during the 5 weeks that followed.

 Results of plasma enzyme analyses were compared before, during and after the bloom among residents of Armidale and surrounding areas (Falconer et al., 1983). Analysis of variance was used to assess differences in enzyme levels between comparison groups and between times within comparison groups. Results of the statistical analysis indicated a significant rise in GGT levels in residents of Armidale during the bloom period. ALT levels in Armidale residents increased during the bloom period, but the change was not statistically significant.

The authors noted substantial variability in enzyme levels, attributing this finding to the imprecise method of selecting study participants (Falconer et al., 1983). It should be noted that several of the enzyme measurements for the referent population were associated with one

individual requiring repeat analysis for chronic kidney disease. Alcoholism, which can increase GGT levels, was reported to occur at about the same proportion (7-10%) in the groups assessed before and during the bloom, although it was substantially lower in the post-bloom group of Armidale residents. The authors concluded that the change in GGT among Armidale residents before and during the bloom period might potentially be associated with exposure to *M. aeruginosa*.

4.1.1.2. Long-Term Studies and Epidemiological Studies

Zhou et al. (2002) conducted a retrospective cohort analysis of colorectal cancer and exposure to microcystins in drinking water in a Chinese province in which an association had been reported previously (Jiao et al., 1985; Chen et al., 1994). Cases of primary colorectal adenocarcinoma between 1977 and 1996 from eight randomly selected towns within Haining City of Zhejiang Province were used as the study population. Cases were identified using the local cancer registry and independently verified by two pathologists. Drinking water source used longest during the lifetime was used as a surrogate measure of exposure to microcystins. Information on drinking water source was obtained by interview of patients or family members of deceased cases. In each of the eight towns, 10 water sources (3 rivers, 3 ponds, 2 wells and 2 taps) were randomly selected and sampled for microcystins twice in each of the months of June through September (total of eight samples from each source). Water samples were analyzed for microcystins by ELISA; the authors did not specify the targeted congeners. The authors do not specify the nature of the "tap" water sources, but the text implies that the tap water derives from one or more treatment plants.

The average incidence rate of colorectal cancer across all of the study areas was 8.37/100,000 per year. The incidence rate was compared among the four different water sources, with well water users serving as the referent population. Compared with the incidence among well water users, the colorectal cancer incidence rates among users of the other water sources were significantly increased. Tap water use was associated with a relative risk of 1.88, while river and pond water use were both associated with a relative risk over 7.0. There was no difference in colorectal cancer incidence between river and pond water users. The authors suggested that exposure to trihalomethane compounds might account for the increase in incidence among tap water users. Table 4-1 shows the incidence rate, relative risk and 95% CIs for these exposure comparisons.

Microcystins were detected at concentrations exceeding 50 pg/mL (considered by the authors to be the limit for positive detection) only in river and pond water, and the average concentrations in these sources were substantially higher (30- to 50-fold) than well or tap water. A similar proportion (about 25%) of the residents in each of the eight towns used river and pond water for drinking water, allowing an analysis comparing the average microcystin concentration in river and pond water in each town with the incidence rate by town. This analysis showed a strong correlation between incidence rate and concentration of microcystin (Spearman correlation

Table 4-1. Relative Risk of Colorectal Cancer and Microcystin Concentration by Drinking Water Source (Zhou et al., 2002)

Water Source	Colorectal Cancer Incidence Rate per 100,000	Relative Risk of Colorectal Cancer	95% CI	Number of Microcystin Samples >50 pg/mL	Mean Microcystin Concentration (pg/mL)	Maximum Microcystin Concentration (pg/mL)
Well water	3.61	-	-	0/ 12	0.73	9.13
Tap water	6.77	1.88	1.39-2.54	0/ 17	4.85	11.34
River water	28.5	7.94	6.11-10.31	25/ 69	141.08	1083.43
Pond water	27.76	7.7	5.75-10.30	6/ 35	106.19	1937.94

coefficient = 0.88, p<0.01). Figure 4-1 shows the relationship between colorectal cancer incidence and average microcystin concentration.

This study provides suggestive evidence for an association between microcystin exposure and colorectal cancer. It is also consistent with earlier reports of an association between drinking river or pond water and incidence of colorectal cancer in the Zhejiang Province of China (Jiao et al., 1985; Chen et al., 1994; studies published in Chinese and not translated for this review). However, because demographic information was not provided in the report, it is not clear whether dietary, genetic and lifestyle factors associated with colorectal cancer have been adequately controlled in the analysis. Further, other potential biological and chemical contaminants in the river and pond water were not considered.

Several epidemiological studies have examined the relationship between drinking water source (well, river, pond or ditch) and liver cancer in Haimen City, Jian-Su Province, China, an area with an elevated hepatocarcinoma incidence (Yu, 1989; Yu et al., 1989). These studies, published in Chinese and not translated for this review, showed an increased risk of primary liver cancer associated with consumption of pond or ditch water (Ueno et al., 1996). According to Health Canada (2002), Yu (1989) showed that consumption of pond or ditch water was associated with an 8-fold increase in liver cancer incidence when compared with well water consumption. Health Canada (2002) reported that a larger study of 65 counties in China, also published in Chinese (Chen et al., 1991), showed the opposite; that consumption of deep well water was associated with an increased risk of liver cancer.

Ueno et al. (1996) conducted a survey of microcystin content in drinking water supplies in Haimen City to test the hypothesis that microcystins in surface drinking water supplies could contribute to the higher incidence of liver cancer. Microcystins were measured by ELISA in shallow and deep wells, as well as in ponds/ditches and river waters. The authors did not indicate which congeners were targeted by the ELISA. Occurrence of microcystins was higher in pond/ditch water (17% reported as positive with concentration >50 pg/mL) and river water (32% positive) samples than in shallow wells (4% positive) or deep wells (no detections >50 pg/mL). Further, microcystin concentrations averaged across the drinking water types were different, averaging 101, 160 and 68 pg/mL in pond/ditch, river and shallow well samples, respectively. These data, while suggestive, do not directly associate exposure to microcystins and liver cancer, since individual exposures were not measured or estimated, and other biological or chemical contaminants in the surface waters have not been ruled out.

In a case-control study of liver cancer in Haimen City, conducted by Yu et al. (2002), a variety of liver cancer risk factors were evaluated, including hepatitis B and C virus infection, aflatoxin B1 or microcystin exposure, smoking, drinking, diet and genetic polymorphisms. From a pool of 248 patients with hepatocellular carcinoma and 248 age-, sex- and residence-matched controls, 134 paired cases and controls assented to blood samples for virus infection and ALDH2 and CYP2E1 gene polymorphism analyses. Data from these analyses were combined with questionnaire information on possible lifestyle and dietary risk factors for liver cancer. Microcystin exposure was assessed categorically based on drinking water supply (tap, deep or

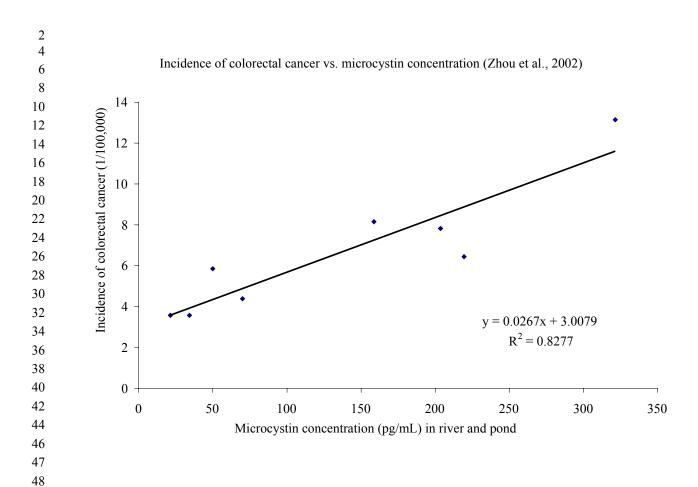


Figure 4-1. Relationship between Colorectal Cancer and Microcystin Concentration in River and Pond Water in Haining City, China (Zhou et al., 2002)

shallow well, river, ditch or pond), as in earlier studies (Yu et al., 1989). Neither univariate nor multivariate analysis of the data indicated an association between consumption of river, pond or ditch water and hepatocellular carcinoma. Hepatitis B virus infection was strongly associated with primary liver cancer, and history of i.v. injection was also identified as a risk factor (Yu et al., 2002).

Fleming et al. (2002) conducted an ecological epidemiological investigation of the relationship between drinking water source and incidence of primary liver cancer in Florida. The study was prompted by data showing cyanobacteria and toxins, especially microcystins, in surface drinking water sources in Florida. The study population consisted of all cases of primary hepatocellular carcinoma reported to the Florida state cancer registry between 1981 and 1988. The study population was divided into comparison groups consisting of those served by surface drinking water supply and those using other sources. Residence at the time of diagnosis was used to place cases into the various comparison groups. Surface water treatment plants and their service areas were geocoded, as were deep groundwater treatment plants. Several comparisons were made. First, incident cases residing in the service area of a surface water treatment plant were compared with those residing in the service area of a deep groundwater treatment plant. Within this comparison, there were several referent groups; one randomly sampled from the available groundwater service areas, one matched on median income and rent, one matched on ethnic makeup and one matched on income, rent and ethnicity. Second, incident cases in the surface water service area were compared with equally-sized buffer areas surrounding the surface water service area, but not served by the treatment plant. Finally, incident cases were compared with the incidence in the general Florida population.

Evaluation of the individual incidence rates in the 18 surface water service areas with the groundwater service areas did not reveal any statistically significant differences among the individual incidence rates. When the service areas were pooled, residence in a surface water service area was associated with a statistically significant reduced risk of hepatocellular carcinoma compared with either groundwater service areas (standardized rate ratios [SRR] ranged from 0.8 to 0.98 for the four groundwater comparison groups) or the general Florida population (SRR of 0.8). It should be noted that the measure of exposure, residence within a surface water service area, was estimated as the average size plus two standard deviations of the service area for this comparison.

When comparisons were made between residence in the actual (i.e., not estimated as above) surface water service areas and residence in the buffer areas surrounding the service areas, a statistically significant increase in the incidence of hepatocellular carcinoma was observed for those residing within the surface water service area (SRR = 1.39, CI = 1.38-1.4). Analyses of 1990 census data suggested that the ethnic and socioeconomic backgrounds of the service areas and buffer areas were similar, although the authors did not report these data. Interestingly, the incidence of hepatocellular carcinoma in the buffer areas was significantly lower than that in the general Florida population (SRR = 0.59).

An ecological study such as this is useful for generating hypotheses, but not for establishing an exposure-response relationship due to the lack of exposure data on individuals. In this case in particular, there is strong potential for misclassification of exposure. Residence in

a surface water service area at the time of diagnosis of hepatocellular carcinoma is a poor measure of potential exposure to cyanobacterial toxins, especially given residential mobility and likely latency time for cancer development. Further, the initial comparisons with groundwater service areas used GIS-generated estimates of surface water service areas rather than actual service areas, leading to greater potential for misclassification.

Fleming et al. (2004) also conducted an ecological study assessing the relationship between incident colorectal cancer and proximity to a surface drinking water treatment plant, with the latter representing a surrogate for exposure to cyanobacteria. Methods for this study were identical to those described above for Fleming et al. (2002) except that colorectal cancer data were abstracted from the Florida Cancer Data System from 1981-1999. As with Fleming et al. (2002), comparisons were made between the colorectal cancer incidence rates in the 18 surface water treatment service areas with several referent groups (a random group of groundwater treatment service areas, a group of groundwater treatment service areas matched on median income and rent, a group of groundwater treatment service areas matched on ethnic makeup, a group of groundwater treatment service areas matched on both median income and ethnicity, groups residing in an equally-sized buffer areas surrounding the surface water service area and, finally, the general Florida population). Mann Whitney rank sum tests of all comparisons did not suggest an association between colorectal cancer and residence at time of diagnosis in a surface water treatment area (details not provided). This ecological study is subject to the same limitations as described above for Fleming et al. (2002).

4.1.2. Inhalation Exposure

No studies of human exposure to microcystins via inhalation were identified in the materials reviewed for this document.

4.2. ACUTE, SHORT-TERM, SUBCHRONIC AND CHRONIC STUDIES AND CANCER BIOASSAYS IN ANIMALS – ORAL AND INHALATION

Early research on cyanobacterial toxins examined the effects of exposure to cyanobacterial organisms rather than on the toxins now known to produce many of the toxicological effects. In the case of microcystins, the isolation and characterization of important toxin(s) did not occur until the 1980s (WHO, 1999). As a result, many studies have used various extracts of cyanobacterial blooms as test substances in toxicological experiments. These studies contribute to the hazard identification for cyanobacterial toxins, but, as discussed below, are not useful for dose-response assessment.

 The quantity of an individual cyanobacterial toxin in different bloom samples and extracts varies widely, being influenced by a number of different factors. Some toxins are produced by more than one genus of cyanobacteria. For example, microcystins have been shown to be produced by *Microcystis*, *Anabaena*, *Planktothrix*, *Nostoc* and others (WHO, 1999). Some species (e.g., *Anabaena*) can produce more than one cyanotoxin (WHO, 1999). Even within a species, different strains produce varying levels of toxin; some produce little or no toxin at all. Growth conditions can also contribute to the level of toxin produced by a given species and strain (WHO, 1999). Finally, any given sample may contain multiple genera, species or strains

of cyanobacteria, as well as other contaminating organisms. Some of these variables will also apply to cyanobacterial cells cultured in a laboratory, although clonal cultures may be characterized as to toxin content. In general, there is no clear means of predicting the toxin content in a given bloom sample or cell extract.

In addition to the variations in toxin production within bloom and/or culture samples, there are variations in toxin concentration depending on the method used for producing a material for toxicological administration. Among the studies reviewed, the administered material included lyophilized bloom samples in solution, cell-free extracts, cell-free lysates, partially purified toxins, purified toxins and others. As endotoxins, microcystins exist primarily within the cyanobacterial cell, and are released when cells are lysed. As a result, studies of extracts obtained by removing intact cells may not contain much, if any microcystin. Purified or partially-purified toxins were used in a number of studies; however, the specific microcystin congener or congeners may not have been identified. As a result, data using bloom samples, cell extracts or partially purified toxins cannot be considered reliable information in relating exposure to a given toxin with toxicological effect.

Giving due consideration to the limitations of algal extract studies, a distinction is made between toxicological data obtained using purified microcystins and toxicological data obtained using a bloom sample or extract. These data are discussed separately in this report, with the latter data being considered supplemental due to the confounding factors outlined above. As a result, studies of cyanobacterial extracts are not reviewed in depth.

4.2.1. Oral Exposure

4.2.1.1.1. Purified Microcystins

4.2.1.1. Acute Studies

Fawell et al. (1999) conducted acute, subchronic and developmental toxicity studies of MCLR in mice and/or rats. In the acute portion of the study, single oral gavage doses of MCLR (purity not specified) in aqueous solution were given to male and female CR1:CD-1(ICR)BR(VAF plus) mice and CR1:CD(SD)BR(VAF plus) rats (five per sex per species). Doses of 500, 1580 and 5000 μ g/kg body weight were administered. Untreated control groups were not included. The animals were observed for up to 14 days prior to sacrifice and necropsy. Microscopic examinations of the lung and liver were conducted. LD₅₀ values were calculated.

Oral LD_{50} values were estimated to be about 5000 µg/kg for mice and over 5000 µg/kg for rats. Animals that died showed clinical signs, including hypoactivity and piloerection; however, clinical signs were absent in survivors. Body weights among surviving animals were not affected during the 14-day follow-up. Necropsy of the animals that died showed darkly discolored and distended livers, as well as pallid kidneys, spleen and adrenals. Livers of all animals that died had moderate or marked centrilobular hemorrhage. The incidence and severity of liver lesions increased in a dose-dependent fashion, as shown in Table 4-2.

Table 4-2. Incidence of Liver Lesions in Mice and Rats Treated with a Single Dose of MCLR							
(Fawell et al., 1999)							

Dose (µg/kg)	Number Animals Treated	Mortality	Diffuse Hemorrhage	Moderate Centrilobular Hemorrhage	Marked Centrilobular Hemorrhage	Centrilobular Necrosis	Cytoplasmic Vacuolation		
Mice									
500	10	0	2	0	0	0	0		
1580	10	1	2	2	1	0	0		
5000	10	5	1	7	0	2	0		
Rats									
500	10	0	8	0	0	0	0		
1580	10	0	7	0	0	0	0		
5000	10	1	8	1	1	1	1		

 Diffuse hemorrhage in the liver was seen in rats and mice of all dose groups, but the incidence was not clearly related to dose. Because an untreated control group was not included in this study, it is not possible to say whether the liver effects were treatment-related; thus, neither a NOAEL nor a LOAEL could be determined. The single mouse death at 1580 μ g/kg indicates that this is a frank effect level (FEL) in this species; the FEL in rats was 5000 μ g/kg with a single rat death at this dose.

 Yoshida et al. (1997) assessed the acute oral toxicity of purified MCLR (>95% pure by HPLC) in female BALB/c mice. Preliminary experiments using doses of 16.8 and 20 mg/kg resulted in death within 160 minutes in two mice; therefore, doses of 8.0, 10.0 and 12.5 were chosen for LD₅₀ determination. MCLR in saline solution was administered via gavage to a total of five 6-week-old mice. Two control mice received saline via gavage. Mortality was observed over a 24-hour period, and dead animals, including those in the preliminary study, were immediately necropsied. One surviving mouse was sacrificed and necropsied 24 hours after treatment; the remainder were sacrificed and necropsied after a week. The liver, kidneys and lung were sectioned and examined by light microscopy. Electron microscopy was used to identify apoptotic cells in the livers of treated mice. The remaining tissues were subjected to histopathological analysis.

Mortality within 24 hours was 0/1 at 8 mg/kg, 0/2 at 10 mg/kg and 2/2 at 12.5 mg/kg. The oral LD₅₀ was calculated to be 10.9 mg/kg. No effects on the stomach, intestine, skin or

organs other than the liver and kidneys were observed. Liver effects were pronounced in animals that died, including centrilobular hemorrhage and hepatocyte degeneration, as well as free hepatocytes in the veins of mice receiving doses in excess of 12.5 mg/kg (in the preliminary experiments). Effects on the kidneys included proteinaceous eosinophilic materials in the Bowman's spaces of mice receiving high doses (>12.5 mg/kg). In a single mouse treated with 10.0 mg/kg and sacrificed after 24 hours, evidence of hepatocellular necrosis was observed in the centrilobular and midzonal regions, and single cell death (possibly apoptotic) was reported in the centrilobular region, as well as surrounding necrotic areas. In the other mouse treated with 10 mg/kg and the two mice treated with 8.0 mg/kg (all sacrificed 1 week after treatment), the livers contained hypertrophic hepatocytes in the centrilobular region and fibrosis in the centrilobular and midzonal regions. A few apoptotic cells were observed in these animals. No kidney effects were reported in animals that survived treatment for at least 24 hours. No control group was included, so neither a NOAEL nor a LOAEL could be determined from this study. The deaths of both mice given 12.5 mg/kg MCLR indicate that this is an FEL.

Ito et al. (1997a) compared the acute effects of MCLR on the livers and gastrointestinal tracts of young and aged mice. Single doses of 500 μg/kg MCLR (purity not specified) dissolved in ethanol and diluted in saline were administered via oral gavage to aged (29 mice aged 32 weeks) and young (12 mice aged 5 weeks) male ICR mice. Three aged and three young untreated mice served as controls. Twenty-two aged mice were sacrificed at 2 hours, five mice at 5 hours, and two mice at 19 hours after treatment; four young mice were sacrificed at each time point. Liver damage and gastrointestinal erosion were evaluated.

The results showed marked differences between young and aged mice in both liver damage and gastrointestinal effects. In young mice, no liver pathology or gastrointestinal changes were reported. In contrast, 18 of 29 aged mice treated with the same dose developed pathological changes of the liver. Among the aged mice, 8 of 29 had liver injury of the highest severity, characterized as bleeding, disappearance of many hepatocytes in the whole liver and friable tissue (severity rating of +4). Five of 29 mice had liver changes characterized by bleeding and disappearance of hepatocytes in centrilobular region (severity rating of +3). Necrosis in the centrilobular region was observed in 4 of 29 mice (severity rating of +2) and eosinophilic changes in the centrilobular region were indicated for one mouse (severity rating of +1).

Gastrointestinal effects observed in the treated aged mice included necrosis to one-third depth of the mucosa and severe duodenal damage (including decreased villi density, separation of epithelial cells from lamina propria and edema of both the submucosa and villi). Details of the incidence of these effects were not reported; however, the authors indicated that the degree of liver injury was related to the severity of gastrointestinal effects. Regeneration of intestinal tissues was evident in some of the mice sacrificed at later time points (5 and 19 hours after treatment). Among untreated aged mice, serum enzyme levels (AST and ALT) were not different, but gastrointestinal condition was somewhat compromised in the aged mice. Aged mice had thinning of surface epithelial cells with consequent exposure of lamina propria and glands in some areas. The authors postulated that the oral uptake of MCLR was dependent on gastrointestinal tract erosion and the loss of permeability in capillaries of the villi. This study

identified a freestanding LOAEL of 500 μ g/kg (in aged mice only) for liver and gastrointestinal effects.

In a poorly described study, Fitzgeorge et al. (1994) administered MCLR via gavage to newly weaned CBA/BALBc mice. The commercially-obtained compound was described only as "suitably purified." The LD₅₀ was estimated to be 3000 μg/kg, and increases in liver (43%) and kidney (5.9%) weights were reported. The authors reported that there was no change in lung or spleen weight; other endpoints were either not examined or not reported.

Rogers et al. (2005) evaluated the potential synergism between MCLR and anatoxin-a administered by gavage to CD-1 mice (sex not specified). A total of 60 fasted mice were given gavage doses of 0, 500 or 1000 μ g/kg MCLR (purity \geq 98%) followed 50 minutes later with gavage doses of 0, 500, 1000 or 2500 μ g/kg anatoxin-a (purity \geq 95%). Controls were given distilled water by gavage. While not specified, group sizes are presumed to have been five animals per treatment. The animals were observed for clinical signs of toxicity, loss of appetite and mortality; body weight was measured before treatment and 3 hours later. The duration of observation was not reported. No deaths, clinical signs of toxicity or differences in body weight were observed. Effect levels cannot be identified from this study due to inadequate reporting and because few toxicological endpoints were evaluated.

4.2.1.1.2. Cyanobacterial Extracts

Rao et al. (2005) compared the acute oral effects of microcystin extract in aged (36 weeks old) and young (6 weeks old) Swiss albino mice. A single LD₅₀ dose of extract was administered to male mice; mortality occurred after 4-5 hours. Both groups of mice had increased relative liver weight and DNA fragmentation compared to control, but there was no difference between the age groups. In contrast, glutathione depletion and lipid peroxidation were significantly greater in the aged mice when compared with young mice. Further, while most serum enzymes were increased over controls in both groups, GGT was increased to a greater extent in aged mice than in young mice.

4.2.1.2. Short-Term Studies

4.2.1.2.1. Purified Microcystins

Heinze (1999) evaluated the effects of MCLR in drinking water on 11-week-old male hybrid rats (F1 generation of female WELS/Fohm x male BDIX). Groups of 10 rats were given doses of 0, 50 or 150 µg/kg body weight for 28 days in drinking water. Water consumption was measured daily and rats were weighed at weekly intervals. Dose estimates provided by the authors were not adjusted to account for incomplete drinking water consumption (3-7% of supplied water was not consumed over the 28-day period). The test material was obtained commercially, but the authors did not report a measure of purity. After 28 days of exposure, rats were sacrificed by exsanguination under ether anesthesia. Organ weights (liver, kidneys, adrenals, thymus and spleen) were recorded and hematology, serum biochemistry and histopathology of liver and kidneys were evaluated.

Hematological evaluation demonstrated an increase in the number of leukocytes in rats in the highest dose group (38% increase). Serum biochemistry showed significantly increased mean levels of ALP and lactate dehydrogenase (LDH) in both treatment groups (84 and 100% increase in LDH, 34 and 33% increase in ALP in low and high doses, respectively) and no changes in mean levels of ALT or AST. A dose-dependent increase in relative liver weights was observed (17 and 26% at the low and high doses, respectively). Table 4-3 shows the mean enzyme levels and relative liver weights.

Table 4-3. Serum Enzyme Levels and Relative Liver Weights (Mean ± Standard Deviation) in Rats Ingesting MCLR in Drinking Water (Heinze, 1999)

Parameter	Control n=10	50 μg/kg n=10	150 μg/kg n=10
Relative liver weight (g/100 g body weight)	2.75 ± 0.29	3.22 <u>+</u> 0.34*	3.47 <u>+</u> 0.49*
Lactate dehydrogenase (microkatals/L)	16.64 <u>+</u> 4.48	30.64 <u>+</u> 5.05*	33.58 <u>+</u> 1.16*
Alkaline phosphatase (microkatals/L)	9.67 <u>+</u> 2.20	13.00 <u>+</u> 3.81*	12.86 ± 1.85*

^{*} p<0.05 when compared with control

The authors also reported a dose-dependent increase in absolute liver weights, although the data were not provided. No statistically significant changes in other organ weights or body weights were observed. In treated animals, histopathological alterations in the liver were classified as toxic hepatosis. The incidence of liver lesions is summarized in Table 4-4. Lesions were spread diffusely throughout the parenchyma and included increased cell volume, increased mitochondria, cell necrosis, activation of Kupffer cells and increased amounts of periodic acid-Schiff (PAS)-positive substances, indicating cell damage. Liver lesions were observed in both treatment groups, but the severity of the damage was increased in the 150 μ g/kg dose group. No effects on the kidneys were observed. A NOAEL could not be determined from this study. The lowest dose tested (50 μ g/kg) represents a LOAEL based on liver lesions, increased relative liver weights and changes in liver enzymes (ALP and LDH).

4.2.1.2.2. Cyanobacterial Extracts

Davidson (1959) treated groups of three mice with drinking water or feed mixed with different extracts (crude, aqueous and filtered) from a *Nostoc rivulare* water bloom for 21 days. The only effects reported were ruffled hair and nervousness in the mice treated with crude extract. Kalbe (1984) observed no change in body weight among juvenile mice and rats given filtrates from two different water blooms of *M. aeruginosa* for 2-8 weeks.

Table 4-4. Incidence of Liver Lesions in Rats Ingesting MCLR in Drinking Water for 28 Days (Heinze, 1999)

Liver Histopathology	Control n=10	50 μg/kg n=10	150 μg/kg n=10						
Degenerative and Necrotic Hepatocytes with Hemorrhage									
Slight	0	4	0						
Moderate	0	6	6						
Intensive damage	0	0	3						
Degenerative and Necrotic Hepatocytes without Hemorrhage									
Slight	0	0	0						
Moderate	0	0	1						
Intensive damage	0	0	0						
PAS-positive Material									
Slight	1	5	0						
Moderate	0	5	8						
Intensive damage	0	0	2						
Activation of Kupffer Cells									
Slight	0	0	0						
Moderate	0	10	10						
Intensive damage	0	0	0						
Lipid Granules and Droplets									
Slight	0	4	0						
Moderate	1	2	1						
Intensive damage	0	0	0						

Orr et al. (2003) exposed yearling beef cattle to live cultures of *M. aeruginosa* in drinking water in an effort to evaluate whether microcystins accumulated in the liver or blood of the animals. Four steers were treated for 28 days, and four untreated steers served as controls. No effects on body weight, weight gain, food or water consumption or plasma enzymes (GGT, glyceraldehydes dehydrogenase, AST or bilirubin) were observed. The authors reported no detectable microcystins (by HPLC and gas chromatography-mass spectrometry) in either plasma samples collected throughout treatment or in samples of the liver collected upon sacrifice at the termination of exposure. Analysis of the liver samples by ELISA showed measurable microcystins; however, the authors indicated that these results likely represented cross-reaction with something besides microcystins, given the failure of the more sensitive HPLC analysis to detect microcystins.

Schaeffer et al. (1999) reported the results of an unpublished 1984 study in which *Aphanizomenon flos-aquae*, a cyanobacterium consumed as a food supplement, was fed to mice in the diet. The authors used recent analysis of the *A. flos-aquae*, which often coexists with *Microcystis* species, to estimate the microcystin content in the material consumed by the mice. Analysis of the *A. flos-aquae* samples used in the feeding study showed an average concentration of 20±5 µg MCLR per gram of *A. flos-aquae*. The authors estimated the daily exposure of MCLR in the exposed mice to range from 43.3 µg/kg body weight per day to 333.3 µg/kg-day. No clinical signs of toxicity were reported, and no effects on mortality, body weight, organ weights or histology were observed in the treated mice. In addition, no effects on reproductive parameters were reported in five treated mice (highest dose group) allowed to breed.

4.2.1.3. Subchronic Studies

4.2.1.3.1. Purified Microcystins

Fawell et al. (1999) conducted acute, subchronic and developmental toxicity studies of MCLR given via gavage to Cr1:CD-1(ICR)BR (VAF plus) mice (age not specified). MCLR was obtained commercially and administered in distilled water. The concentration in the dosing solution was verified by HPLC with UV detection. Daily oral gavage doses of 0, 40, 200 or 1000 μg/kg body weight were given to groups of 15 male and 15 female mice for 13 weeks. Daily clinical observations were made, body weight and food consumption were recorded weekly, and eye examinations were conducted prior to and at the conclusion of treatment. Hematology and serum biochemistry were evaluated for seven mice of each treatment group during the final week of treatment. Upon sacrifice after 13 weeks, gross examination of organs and microscopic evaluation of tissues were performed. All tissues were examined in the control and high dose animals, while only lungs, liver and kidney were examined in the other treated animals.

Mean body weight gain was decreased approximately 15% in all treated male groups. Mean terminal body weights differed from controls by about 7% in these groups. No doserelated trends were evident for body weight gain or body weight in males. The only body weight change observed in females was an increase in body weight gain in the 200 μ g/kg-day group. Hematological evaluation showed slight (10-12%) decreases in mean hemoglobin concentration, red blood cell count and packed cell volume among females receiving 1000 μ g/kg body weight.

ALP, ALT and AST levels were significantly elevated (2- to 6-fold higher) in the high-dose males, and ALP and ALT were likewise elevated (2- and 6-fold higher, respectively) in high dose females. ALT and AST were also elevated (2-fold) in the mid-dose males. GGT was slightly decreased in some treatment groups. Serum albumin and protein were reduced (13%) in males of the mid- and high-dose groups. Table 4-5 shows the blood chemistry results.

Table 4-5. Blood Chemistry Results (Mean <u>+</u> Standard Deviation) for Mice Treated with MCLR for 13 Weeks (Fawell et al., 1999)									
MCLR Dose (μg/kg-day)	Alkaline Phosphatase (ALP) (U/I) Alanine Aminotransferase (ALT) (U/I) Aspartate Aminotransferase (AST) (U/I) Gamma Glutamyl Transaminase (GGT) (U/I)		Total Protein (g%)	Albumin (g%)					
Male									
Control	91 <u>+</u> 22.2	27 <u>+</u> 8.0	68 <u>+</u> 27.7	6 <u>+</u> 1.0	5.5 <u>+</u> 0.32	3.2 ± 0.19			
40	95 <u>+</u> 29.2	37 <u>+</u> 17.2	64 <u>+</u> 12.2	4 <u>+</u> 0.7	5.1 <u>+</u> 0.26	3.0 <u>+</u> 0.13			
200	94 <u>+</u> 32.3	59 ^a + 28.0	121 ^b ± 43.7	3° ± 0.4	4.8 ^b ± 0.29	$2.8^{\circ} \pm 0.13$			
1000	232 ^b ± 103.2	159° ± 75	121 ^b ± 26.3	4 <u>+</u> 0.4	4.8° ± 0.21	$2.8^{\circ} \pm 0.11$			
Female									
Control	167 <u>+</u> 24.6	32 <u>+</u> 11.3	101 <u>+</u> 38.3	4 <u>+</u> 1.0	5.1 <u>+</u> 0.30	3.1 <u>+</u> 0.14			
40	187 <u>+</u> 76.2	25 <u>+</u> 7.8	74 <u>+</u> 13.2	3 <u>+</u> 0.5	5.2 <u>+</u> 0.28	3.2 <u>+</u> 0.16			
200	156 <u>+</u> 33.4	27 <u>+</u> 9.4	74 <u>+</u> 22.1	3 <u>+</u> 0.0	5.3 <u>+</u> 0.31	$3.4^{a} \pm 0.14$			
1000	339 ^b <u>+</u> 123.7	220 ^b <u>+</u> 149.1	144 <u>+</u> 71.7	3 <u>+</u> 0.4	5.1 <u>+</u> 0.22	3.1 <u>+</u> 0.18			

^a Significantly different from controls at p<0.05

Histopathological changes in the liver were reported in the males and females of the midand high-dose groups, with a dose-related increase in incidence and intensity. The liver lesions were multifocal and observed throughout the liver lobule. Table 4-6 summarizes the incidence of liver histopathological changes. Sex-related differences in liver pathology were not apparent. No lesions were found in other tissues.

The authors characterized the 40 $\mu g/kg$ body weight dose as a clear NOAEL and indicated that histopathological changes observed in the 200 $\mu g/kg$ dose group were not severe. The mid dose (200 $\mu g/kg$ -day) represents a LOAEL based on the liver histopathological changes and statistically significant blood chemistry changes. The WHO (1999) used the NOAEL value of 40 $\mu g/kg$ -day from this study as the basis for its provisional Tolerable Daily Intake for MCLR.

^b Significantly different from controls at p<0.01

^c Significantly different from controls at p<0.001

Table 4-6. Incidence of Liver Histopathology in Mice Treated with MCLR for 13 Weeks (Fawell et al., 1999)

Liver Histopathology	Control	40 μg/kg-day	200 μg/kg-day	1000 μg/kg-day
Male	n=15	n=15	n=15	n=15
Acute inflammation	0	1	0	0
Chronic inflammation	1	2	4	15
Congestion	3	0	0	1
Hepatocyte vacuolation	5	5	6	3
Hemosiderin deposits	0	0	0	15
Hepatocyte degeneration	0	0	1	14
Female	n=15	n=15	n=15	n=15
Autolysis	0	0	0	1
Chronic inflammation	5	8	8	14
Congestion	0	0	0	1
Hepatocyte vacuolation	5	5	11	8
Hemosiderin deposits	0	0	1	14
Hepatocyte degeneration	0	0	1	9

4.2.1.3.2. Cyanobacterial Extracts

Falconer et al. (1994) administered dried bloom materials in the drinking water of pigs for 44 days. Plasma samples collected over 56 days showed dose- and time-dependent increases in GGT, ALP and total bilirubin, as well as a decrease in plasma albumin. Dose-related changes in the incidence and severity of histopathological changes of the liver were also observed, including cytoplasmic degeneration, hepatic cord disruption, single cell necrosis, periacinar degeneration, congestion and Kupffer cell proliferation.

4.2.1.4. Chronic Studies

4.2.1.4.1. Purified Microcystins

Ueno et al. (1999) evaluated the toxicity of MCLR in mice chronically exposed via drinking water. Two hundred 6-week-old female BALB/c mice were randomly assigned to receive either no treatment or drinking water (*ad libitum*) containing 20 μ g/L MCLR for 7 days/week. The MCLR had been isolated from lyophilized algal bloom materials from Lake Suwa in Nagano, Japan and had been characterized as 95% pure by HPLC. Twenty animals from each group were sacrificed at 3, 6 and 12 months, while the remaining 40 animals were retained for chronic toxicity evaluation and sacrificed at 18 months.

Weekly estimates of food and water consumption and daily observations for clinical signs of toxicity, morbidity and mortality were recorded. Body weights were recorded weekly for the first 2 months, biweekly up until the first year and monthly until sacrifice. At 3, 6, 12 and 18 months, blood was obtained from 20 animals from each group. Samples from 10 animals per group were used for hematological evaluation, and samples from 10 additional animals were used for serum biochemistry evaluation. At each scheduled sacrifice time, complete necropsy of 10 animals per group was conducted. Animals from the chronic toxicity group were necropsied when moribund or dead (prior to scheduled sacrifice) or upon sacrifice at 18 months. Relative and absolute organ weights (liver, kidneys, spleen, thymus, adrenal, ovaries, brain, heart and uterus) were recorded for 9-10 animals per group at each scheduled sacrifice, and histopathological evaluation of these and numerous other organs was conducted. Finally, three to five animals per group were subjected to immunohistochemistry of the liver upon sacrifice to determine the distribution of MCLR in the liver.

Based on weekly estimates of water consumption, the authors calculated the average total intake of MCLR over 18 months to be 35.5 μ g/mouse. No clinical signs of toxicity were observed in either of the groups, and survival in the control and chronic treatment groups was similar. No statistically significant differences in body weight, food consumption, water consumption or hematology were observed; however, hematology data from the 3-month sacrifice were lost due to sampling errors. Treated mice were reported to have a statistically significant decrease in ALP at month 12 (13%) and a significant increase in cholesterol at month 18 (22%). Neither effect was considered by the authors to be toxicologically significant in the absence of other treatment-related effects; however, the increase in cholesterol could be related to the interaction of MCLR with bile acid transport in the liver.

A decrease in heart weight among treated mice sacrificed at month 12 was not considered treatment-related in the absence of histopathological changes. Sporadic changes in absolute and relative thymus weight in treated mice were observed, but histological and morphometric evaluation of the thymus revealed no abnormalities attributable to exposure. In contrast to other studies, the authors observed no difference in the incidence of liver histopathology between treated and control mice. Immunohistochemistry of the liver revealed no accumulation of MCLR. This study identifies a free-standing NOAEL of 2.7 µg/kg-day in female mice (calculated assuming a 24.5 g body weight and an exposure duration of 548 days).

Ito et al. (1997b) evaluated the carcinogenicity and liver toxicity of chronic gavage doses of MCLR. A water bloom from Lake Suwa, Japan served as the source of the MCLR, which was isolated and dissolved in ethanol and saline for dosing. The purity of the isolated MCLR was not specified. Twenty-two ICR mice (13 weeks old) were given either 80 or 100 gavage doses of 80 μg/kg MCLR over the course of 28 weeks. Ten mice were sacrificed after 80 treatments, five were sacrificed after 100 treatments and seven were withdrawn from treatment after 100 doses and sacrificed 2 months later. Three mice served as untreated control. Although the authors did not specify the nature of the postmortem examinations, it appears that the liver was the only organ examined. No change in mean liver weight was observed in the MCLR-treated animals compared with controls. The authors reported "light" injuries to hepatocytes in the vicinity of the central vein in 8 of 15 mice sacrificed immediately after treatment, and in 5 of 7 mice that were withdrawn from treatment for 2 months after exposure. No fibrous changes or neoplastic nodules were observed. Analysis for MCLR and its metabolites by immunohistochemistry failed to detect either the parent compound or any metabolites in the livers of mice sacrificed immediately after treatment.

Thiel (1994) briefly reported the results of a chronic toxicity study of MCLA in vervet monkeys. The report is a brief summary published in the proceedings of an international workshop; a published version of this study was not located. According to the summary, three monkeys were given increasing intragastric doses of MCLA for 47 weeks, while three other monkeys served as controls. Doses increased from 20 µg/kg at the commencement of the study to 80 µg/kg at study termination. The rate of dosage increase was not reported. Monthly measures of body weight and clinical signs (respiration, pulse, temperature) showed no effect of treatment. Blood was withdrawn monthly; hematological parameters examined were hematocrit, bilirubin, hemoglobin, erythrocyte and leukocyte count and platelet count. No statistically significant changes in hematological parameters were observed. No changes were observed in serum biochemistry analyses (albumin, globulins and electrolytes, as well as AST, LDH, ALP, ALT and GGT). Histopathological examination of the liver and other organs (not specified) did not show any differences in treated monkeys when compared with controls.

4.2.1.4.2. Cyanobacterial Extracts

Falconer et al. (1988) conducted a chronic exposure experiment using an extract of a *M. aeruginosa* water bloom in Swiss Albino mice. A concentration-dependent increase in mortality, reduced body weight and a concentration-dependent increase in ALT levels were observed among groups of mice receiving serial dilutions of the extract as their drinking water for a year. There was some evidence that bronchopneumonia incidence was related to concentration of

extract. No significant differences in liver histopathology were observed, although the observed liver changes were slightly more prevalent in treated animals. The data showed some indication of sex differences in susceptibility; male mice showed effects (including mortality and enzyme level increases) at lower concentrations than females.

4.2.1.5. Initiation/Promotion Studies – Cyanobacterial Extracts

Falconer (1991) and Falconer and Buckley (1989) reported evidence of skin tumor promotion by extracts of *Microcystis*. *Microcystis* extract was administered via drinking water to mice pretreated topically with an initiating dose of dimethylbenzanthracene (DMBA). The total skin tumor weight in mice drinking *Microcystis* extract was significantly higher than that of mice receiving only water after initiation. The number of tumors per mouse was only slightly increased in mice receiving extract; the weight difference was largely due to the weight of individual tumors (Falconer and Buckley, 1989). The total weight of tumors in this group also exceeded that of mice pretreated with DMBA and subsequently treated with topical croton oil, with or without concurrent consumption of *Microcystis* extract. Details of the tumor incidence in the mice were not provided by the authors. When *Microcystis* extract was provided in the drinking water of mice pretreated with two oral doses of N-methyl-N-nitroso-urea, no evidence of promotion of lymphoid or duodenal adenomas and adenocarcinomas was observed. No primary liver tumors were observed (Falconer and Humpage, 1996).

Humpage et al. (2000) administered *M. aeruginosa* extract in drinking water to mice pretreated with azoxymethane. Mice were sacrificed at intervals up to 31 weeks after commencement of extract exposure. Enzyme analysis showed a concentration-dependent increase in ALP and decrease in albumin in mice treated with extract. The authors observed a concentration-dependent increase in the mean area of aberrant crypt foci of the colon, although the number of foci per colon and the number of crypts per focus were not different among the groups. The authors proposed that increased cell proliferation caused the increase in size of foci. Histological examination of the livers of mice treated with extract showed more leukocyte infiltration in animals treated with the highest concentration of extract compared to those receiving a low concentration.

4.2.2. Inhalation Exposure

All available studies of inhalation exposure used MCLR, and there were no inhalation studies using cyanobacterial extracts.

4.2.2.1. Acute Studies

Fitzgeorge et al. (1994) conducted experiments in CBA/BALBc mice with MCLR administered via intranasal instillation and inhalation. This study is poorly described, giving few details of study design and findings. A single experiment with mice (number unspecified) inhaling a fine aerosol (particle size 3-5 μ m) with 50 μ g/L MCLR for an unspecified duration of time did not result in any deaths, clinical signs of toxicity or histopathological changes. The nature of the examinations was not reported. The authors estimated the delivered dose of MCLR to be very small (about 0.0005 μ g/kg).

4.2.2.2. Short-Term Studies

those in mice dosed intravenously with MCLR.

Benson et al. (2005) exposed groups of six male BALB/c mice to monodisperse submicron aerosols of MCLR via nose-only inhalation for 30, 60 or 120 minutes each day for 7 consecutive days. The concentration of MCLR was 260-265 µg/m³ and doses deposited in the respiratory tract were estimated to be 3, 6 and 12.5 µg/kg body weight. Control mice were exposed to the aerosolized vehicle (20% ethanol in water). Clinical signs were recorded daily. Sacrifice by injection of Euthasol occurred the day after the last exposure. Blood was collected by cardiac puncture and serum was subjected to clinical chemistry analysis (blood urea nitrogen [BUN], creatinine, total bilirubin, ALP, AST, ALT, total protein, albumin and globulin). Organ weights (adrenals, lung, liver, kidney, spleen and thymus) were recorded and histopathological examinations of the liver, respiratory tract tissues, adrenals, kidney, spleen, thymus, gastrointestinal tract and testes were conducted.

A brief abstract describes a study of acute MCLR exposure via inhalation (Creasia,

1990). Details of study design and results were not reported. The LC₅₀ for mice exposed to a

MCLR aerosol (nose only) for 10 minutes was reported to be 18 µg/L (mg/m³) air with a 95%

exposure of mice to the LC₅₀ concentration, an LD₅₀ of 43 µg/kg body weight was estimated.

The authors reported that histological lesions in mice killed by aerosol exposure were similar to

confidence interval of 15.0-22.0 µg/L (mg/m³). Based on studies of lung deposition after

No clinical signs or effects on body weight or organ weights were observed following exposure to MCLR aerosol. Histopathological examination revealed treatment-related lesions in the nasal cavity only. Lesions were not observed in the liver, other organs or in other parts of the respiratory tract. As indicated in Table 4-7, the incidence and severity of nasal lesions increased with length of the daily exposure period. The lesions consisted primarily of necrosis or inflammation of respiratory epithelial cells and degeneration, necrosis and atrophy of olfactory epithelial cells. Necrotic lesions of olfactory epithelial cells were generally larger patches, while few cells were involved in respiratory epithelial cell necrosis.

4.2.2.3. Subchronic and Chronic Studies

No subchronic or chronic animal studies evaluating the inhalation route of exposure were identified in the materials reviewed for this document.

Table 4-7. Incidence and Severity of Nasal Cavity Lesions in Mice Inhaling Microcystin Aerosol for 7 Days (Benson et al., 2005)

Lesion	Severity	Control	Daily Exp	Daily Exposure Period (minutes)					
	J		30	60	120				
Respiratory Epithelial Necrosis									
Turbinate 1	Minimal	0/6	1/6	0/6	0/6				
	Mild	0/6	0/6	6/6	0/6				
	Moderate	0/6	0/6	0/6	2/6				
Turbinate 2	Mild	0/6	0/6	6/6	3/6				
	Moderate	0/6	0/6	0/6	3/6				
Respiratory Epithelial Inflammation									
Turbinate 1	Mild	0/6	1/6	0/6	1/6				
Turbinate 2	Mild	0/6	1/6	0/6	0/6				
Olfactory Epithelial Degeneration, N	Vecrosis and A	trophy							
Turbinate 1	Mild	0/6	0/6	0/6	4/6				
	Moderate	0/6	0/6	0/6	1/6				
Turbinate 2	Mild	0/6	0/6	6/6	0/6				
	Moderate	0/6	0/6	0/6	6/6				
Turbinate 3	Mild	0/6	0/6	6/6	0/6				
	Moderate	0/6	0/6	0/6	4/6				
	Marked	0/6	0/6	0/6	2/6				

4.3. REPRODUCTIVE/DEVELOPMENTAL STUDIES - ORAL AND INHALATION

4.3.1. Oral Exposure

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4.3.1.1. Purified Microcystins

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Fawell et al. (1999) conducted acute, subchronic and developmental toxicity studies of MCLR given via gavage to Cr1:CD-1(ICR)BR (VAF plus) mice (age not specified). MCLR (0, 200, 600 or 2000 µg/kg) was administered to groups of 26 mice on days 6-15 of pregnancy. The mice were sacrificed on day 18 and necropsied. Weight and sex of the fetuses were recorded, and external, visceral and skeletal examinations performed. Seven of 26 dams receiving 2000 ug/kg died and 2 others were sacrificed prematurely due to morbidity. Altered liver appearance was noted during gross examination of these animals. Surviving dams in this group did not display any clinical signs of toxicity or differences in body weight or food consumption. The authors reported that fetal body weight was significantly lower than controls and there was delayed skeletal ossification at the highest dose; however, the data were not presented in the publication. These effects may be associated with maternal toxicity. According to the authors, no effects on resorption or litter size were observed, nor were there increases in external, visceral or skeletal abnormalities in fetuses of any treatment group. Data on reproductive and developmental parameters were not provided. Based on the authors' description of the findings, it seems evident that the 600 µg/kg-day dose represents a NOAEL for both maternal and developmental toxicity in mice. Further, the deaths of seven dams receiving 2000 µg/kg-day clearly identify this dose as a maternal FEL. However, in the absence of the data showing incidences of reproductive and developmental parameters, it is difficult to determine conclusively whether the high dose also represents a LOAEL for developmental toxicity.

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4.3.1.2. Cyanobacterial Extracts

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Falconer et al. (1988) conducted a limited study of reproductive effects using an extract from an *M. aeruginosa* bloom sample. Eight female mice that had been given 1/4th dilution of the extract as drinking water (estimated to contain 14 µg/mL of unspecified microcystin toxin) since weaning were mated with similarly treated males. No difference in number of litters, pups per litter, sex ratio or litter weight were observed. Reduced brain size was reported to occur in 7 of 73 pups from treated parents and in none of 67 pups from controls. The litter distribution of the affected pups was not reported by the authors. One of the small brains was examined histologically, revealing extensive damage to the hippocampus.

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4.3.2. Inhalation Exposure

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No reports of developmental or reproductive toxicity by the inhalation route of exposure were identified in the materials reviewed for this document.

4.4. OTHER STUDIES

4.4.1. Neurological Effects

 Neurological effects have been reported to occur with acute lethal doses of MCLR. Clinical signs such as hypoactivity and piloerection have been observed in mice and rats exposed to lethal doses of MCLR by oral or i.p. administration (Hooser et al., 1989a; Fawell et al., 1999). These signs were observed in animals that subsequently succumbed to the lethal exposures. No reports of neurotoxicity by the inhalation route of exposure were identified in the materials reviewed for this document.

 Maidana et al. (2006) reported that long-term memory retrieval (as assessed by step-down inhibitory avoidance task) was impaired in rats receiving intrahippocampal injection of 0.01 or 20 μ g/L of a microcystin extract from *Microcystis* strain RST 9501. Exposure at 0.01 μ g/L also impaired spatial learning (as assessed by performance on the radial arm maze), but exposure at the higher concentration did not. The authors indicated that the primary microcystin produced by this strain is [D-Leu¹] MCLR, a variant of MCLR. Oxidative damage, as measured by lipid peroxides and DNA damage, was increased in tissue homogenates of the hippocampus from treated animals.

Foxall and Sasner (1981) conducted limited *in vitro* studies on the neurological effects of a crude extract from a bloom of *M. aeruginosa*. Little detail on experiment design was reported; frog and mouse heart, frog sartorius muscle, frog sciatic nerve and mouse ileum were used in the experiments. The authors reported that the extract had no effect on electrical or mechanical events.

4.4.2. Immunological Effects

Evaluation of the immunotoxicity of microcystins *in vivo* was reported in only two studies, both of which used a cyanobacterial cell extract rather than purified microcystins. Shirai et al. (1986) reported that mice, immunized i.p. with either live or sonicated cells from a *Microcystis* water bloom, developed delayed-type hypersensitivity when challenged 2 weeks later with a subcutaneous injection of sonicated cells. Delayed hypersensitivity was assessed by footpad swelling, which was increased approximately 2-fold over controls at the highest doses of cells. It is not clear whether an endotoxin in the bloom sample was responsible for the development of hypersensitivity, or whether the antigenic epitope existed on other components of the sample.

Shen et al. (2003) assessed the effect of cyanobacterial cell extract on immune function. Mice received 14 daily i.p. injections containing a cell-free extract from a water bloom dominated by *M. aeruginosa*. Doses were reported as 16, 32 and 64 mg lyophilized cells/kg body weight or as 4.97, 9.94 and 19.88 µg/kg microcystin equivalents. HPLC analysis indicated that the microcystin content of the extract was 79.53%, although specific congeners in the extract were not reported. The following immunotoxicity endpoints were examined: phagocytosis, lymphocyte proliferation and antibody production in response to sheep red blood cells. Phagocytic capacity was reduced at the two highest doses, but percentage phagocytosis was not

affected. B-lymphocyte proliferation was significantly reduced (33% compared to controls at 32 mg/kg), while changes in T-lymphocyte proliferation were mild, and deemed biologically insignificant. Finally, humoral immune response, as measured by antibody-forming plaques, was reduced in a dose-dependent manner in treated mice.

Chen et al. (2004, 2005) evaluated the role of nitric oxide generation and macrophage related cytokines on the reduced phagocytic capacity induced by pure MCLR. A dose-dependent inhibition of nitric oxide production was observed in activated macrophages, and a repressive effect was seen in cytokine formation at the mRNA level (e.g., IL-1β, TNF-α, GM-CSF, IFN-γ) after either 24 hour (Chen et al., 2004) or 6 hour treatment (Chen et al., 2005). Hernandez et al. (2000) indicated that MCLR enhanced the early spontaneous polymorphonuclear leukocyte (PMN) adherence (not late or PMN stimulated early or late) at low concentrations, suggesting that microcystins may affect the immune system.

4.4.3. Hematological Effects

Several studies have noted thrombocytopenia in laboratory animals treated with microcystins or bloom extracts purportedly containing microcystins (Slatkin et al., 1983; Adams et al., 1985, 1988; Takahashi et al., 1995). Early investigations explored whether microcystins had a direct effect on platelets, and whether platelets might be responsible for pulmonary thrombi (Slatkin et al., 1983; Jones, 1984). However, *in vitro* studies have shown that MCLR neither induces nor impedes the aggregation of platelets (Adams et al., 1985). Pulmonary thrombi apparently consist of necrotic hepatocytes circulating in the blood (see Section 4.4.5.2.1). More recent information supports the hypothesis that hematological effects observed in animals acutely exposed to microcystins are secondary effects of liver hemorrhage (Takahashi et al., 1995).

Takahashi et al. (1995) reported dose-dependent reductions in erythrocyte count, leukocyte count, hemoglobin concentration, hematocrit and coagulation parameters 1 hour after rats were exposed to MCLR (100 and 200 μ g/kg i.p). None of these parameters changed until after massive liver hemorrhage commenced. Further, hematological changes such as increased prothrombin time and fibrin deposition in the renal glomeruli were not observed. The authors concluded that the depletion of blood components occurred as a result of liver hemorrhage.

Interestingly, mild thrombocytopenia was reported in 1-week-old mice treated with a large i.p. dose of MCLR, even though none of these mice died (Adams et al., 1985).

4.4.4. Effects by Dermal Exposure

No animal studies evaluating the effects in animals of dermal exposure to purified microcystins were identified in the materials reviewed for this document; only one study using a cyanobacterial extract was located. Davidson (1959) applied a crude extract from a bloom of *N. rivulare* to the shaved backs of three mice every 2 hours for a total of 12 hours. Heavy scales were observed on the treated areas; the scales were gone within 4 days and hair regrowth occurred in the following weeks. Dermal application of an aqueous extract or aqueous filtrate did not result in any effects.

4.4.5. Effects by Parenteral Exposure

4.4.5.1. Effects in Humans after Parenteral Exposure

In February of 1996, unfinished water from a reservoir with a cyanobacterial bloom was used at a hemodialysis center in Caruaru Brazil, leading to numerous deaths among the patients treated with the water (Jochimsen et al., 1998; Pouria et al., 1998; Carmichael et al., 2001; Azevedo et al., 2002). The first report of this incident was published by Jochimsen et al. (1998), and Pouria et al. (1998) published follow-up information on the status of the patients. Azevedo et al. (2002) provided the most up-to-date information on patient status after the incident. Using water samples from the reservoir, serum and tissue samples from patients, and a variety of methods, each publication identified microcystins as the primary causative factor in the deaths and disease among patients. Carmichael et al. (2001) used analytical results from tissue samples with dialysate volumes to estimate the concentration of microcystins in the water to which patients were exposed, as a direct measure of exposure concentration was not available.

Of the 131 patients undergoing dialysis during the 4 days when unfinished water was supplied to the center, 116 experienced symptoms, including visual disturbances, nausea, vomiting and muscle weakness; 100 developed acute liver failure; and 52 had died as of December, 1996 (Azevedo et al., 2002). The acute presentation of the disease included malaise, weakness, dizziness, vertigo, visual disturbances and blindness, nausea, vomiting and abdominal pain. Clinical signs included hepatomegaly and jaundice. Biochemistry showed high concentrations of bilirubin and ALP, moderate increases in AST and ALT, hypoglycemia, hypoalbuminemia and severe hypertrigliceridemia. Major hematology findings were slightly low platelet count (within normal range) and reduced platelet aggregation, as well as red blood cell abnormalities (anisocytosis, acanthocytosis and schistocytosis) in some patients. Liver biopsy and autopsy samples showed severe, diffuse individual hepatocyte necrosis throughout the liver lobule, with cell-plate disruption and apoptosis; however, no intrahepatic hemorrhage was observed. Leukocyte infiltration and canalicular cholestasis were also observed.

 Microcystins were implicated as the major contributing factor to patient death and morbidity (Jochimsen et al., 1998; Pouria et al., 1998; Carmichael et al., 2001; Azevedo et al., 2002). Evidence for the role of microcystins was derived from a variety of sources discussed by Azevedo et al. (2002) and Carmichael et al. (2001). Quantitative analysis of the phytoplankton in the reservoir from which the water was supplied to the dialysis center showed that cyanobacteria represented about 99% of the phytoplankton in the reservoir, although the species present during the outbreak were not identified. Analyses of filter systems in the dialysis center showed microcystins; the carbon filter also had cylindrospermopsin. Analyses of patient sera and liver samples provided additional evidence. Patient sera were analyzed for other potential toxins (chlorines, chloramines, trace elements, heavy metals, agricultural compounds and pesticides), but none were found (Pouria et al., 1998). Both serum and liver analyses for microcystins revealed MCYR, MCLR and MCAR. None of the biological samples contained cylindrospermopsin. Finally, physiological effects observed in the patients closely mirrored effects observed in laboratory animals exposed to microcystins. Specifically, the liver damage observed in patients was similar to that observed in mice (Jochimsen et al., 1998). Using

microcystin concentrations measured in patients' livers and typical dialysis volume, Carmichael et al. (2001) estimated the average concentration of microcystins in the dialysate to be 19.5 μ g microcystins/L.

Soares et al. (2006) reported another incident involving human exposure to microcystins via dialysate. In November, 2001, 44 patients of a renal dialysis center in Rio de Janeiro, Brazil, were assumed to be exposed to microcystins after a bloom of *Microcystis and Anabaena* occurred in the reservoir supplying water to the center. The concentration of microcystins in the drinking water was 0.4 µg/L by ELISA. In the water used at the center (after treatment by activated carbon column), the concentration was 0.32 µg/L. Of the 44 patients exposed, 90% had serum microcystin concentrations above the limit of detection (0.16 ng/mL). Twelve of the patients were selected for 2-month follow-up monitoring of serum levels. Over the follow-up period, serum concentrations ranged from <0.16 to 0.96 ng/mL. The highest serum concentrations occurred 1 month after initial exposure. The authors did not provide any information on health effects from the exposure.

Pilotto et al. (2004) reported that about 20% of 114 volunteers subjected to skin-patch testing using cell suspensions and extracts of various cyanobacterial cultures (including two strains of *M. aeruginosa*) experienced a mild dermal skin reaction (erythema). The reaction did not vary with cyanobacterial species.

4.4.5.2. Effects in Animals after Parenteral Exposure

4.4.5.2.1. Acute Studies with Parenteral Exposure

The acute toxicity of microcystins administered i.p. has been extensively studied. The primary target organ for acute microcystin toxicity is the liver; effects have also been observed in the kidney, lungs and gastrointestinal tract. Some of the effects observed in organs other than the liver are believed to be secondary to liver effects.

A number of references report LD_{50} estimates for injected MCLR (Slatkin et al., 1983; LeClaire et al., 1988; Lovell et al., 1989a; Hermansky et al., 1990c; Miura et al., 1991; Stotts et al., 1993; Gupta et al., 2003); two report LD_{50} values for MCRR and MCYR (Stotts et al., 1993; Gupta et al., 2003). Table 4-8 summarizes the available estimates of microcystin LD_{50} values after injection exposure to purified microcystins.

 As the table shows, the LD_{50} for MCLR in mice ranges between 30 and 60 µg/kg. The acute lethality of MCYR is slightly lower than MCLR; LD_{50} estimates for MCYR were 111 and 171 µg/kg (Gupta et al., 2003 and Stotts et al., 1993, respectively). The LD_{50} for MCRR is higher still, with LD_{50} values estimated as 235 µg/kg (Gupta et al., 2003) and 650 µg/kg (Stotts et al., 1993). In rats, the LD_{50} for MCLR was similar to that in mice. There is some evidence that

	Table 4-8. LD ₅₀ Values of Purified Microcystin Congeners by Intraperitoneal Administration									
Sex/Strain	Purity	Purity Vehicle LD ₅₀ (95% CI) Duration Comments								
MCLR-Mice	MCLR-Mice									
Male/Balb/C	75%	NS*	32.6 (±1.2) μg/kg	24 hours	Impurities tested for toxicity at high doses with negative results	Lovell et al., 1989a				
Female/Swiss albino Hale- Stoner	NS	NS	60 μg/kg	NS		Slatkin et al., 1983				
Female/NIH non-Swiss	>95%	Distilled water	61 μg/kg	NS		Hermansky et al., 1990c				
Male/Swiss Webster	NA	0.09% saline	<100 μg/kg	NS	LD ₅₀ by up and down method	Stotts et al., 1993				
Not specified/ CBA/Balbc	NS	NS	250 μg/kg	NS		Fitzgeorge et al., 1994				
Female/Swiss albino	NS	NS	43.0 μg/kg (37.5-49.4)	24 hours	LD ₅₀ by up and down method	Gupta et al., 2003				
Male/Swiss albino	NS	Methanol and PBS	43 μg/kg (37.5-49.4)	24 hours	LD ₅₀ by up and down method	Rao et al., 2005				

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Table 4-8 cont.								
Sex/Strain Purity Vehicle LD ₅₀ (95% CI) Duration Comments Refe								
MCLR-Rats						•		
Male/Fischer	NS	NS	50 μg/kg (36-68)	72 hours	Intraarterial injection. Abstract only	LeClaire et al., 1988		
Male/Fischer 344	>95%	saline	Fed rats: 122 μg/kg (106-141) Fasted rats: 72 μg/kg (60-83)	25 hours		Miura et al., 1991		
MCRR-Mice	•							
Male/Swiss Webster	NS	0.09% saline	~650 μg/kg	NS	LD ₅₀ by up and down method	Stotts et al., 1993		
Female/Swiss albino	NS	NS	235.40 µg/kg (202.3-272.8)	24 hours	LD ₅₀ by up and down method	Gupta et al., 2003		
MCYR-Mice								
Male/Swiss Webster	NS	0.09% saline	~171 µg/kg	NS	LD ₅₀ by up and down method	Stotts et al., 1993		
Female/Swiss albino	NS	NS	110.6 μg/kg (81.7-149.6)	24 hours	LD ₅₀ by up and down method	Gupta et al., 2003		

^{*} Not specified.

the LD_{50} for MCLR was higher in fed rats (122 μ g/kg) than in fasted rats (72 μ g/kg) (Miura et al., 1991).

In general, death occurs quickly in animals receiving a lethal injected dose of MCLR. Mice typically die within 1-2 hours of a lethal i.p. dose of MCLR (Adams et al., 1988; Gupta et al., 2003). Mean time to death for mice exposed to a lethal dose of MCRR or MCYR is also within 2 hours (Gupta et al., 2003). In mice, an i.p. dose of 100 μg/kg MCLR typically results in 100% mortality (Adams et al., 1988; Hooser et al., 1989a; Hermansky et al., 1990c). Hooser et al. (1989a) compared the effects of MCLR in mice and rats and observed significant differences in survival time; both male and female rats given less than 240 μg/kg survived between 20 and 32 hours; rats receiving higher doses died within 8 hours. In contrast, female mice receiving 100 μg/kg died within 1.5 hours (Hooser et al., 1989a). Miura et al. (1991) administered MCLR i.p. to fed and fasted rats and reported a protective effect of feeding. Median time to death for rats given 100 μg/kg MCLR was 32 hours in fed rats and less than 2 hours in fasted rats.

The sequence of events leading to death in laboratory rodents has been extensively studied (Slatkin et al., 1983; Adams et al., 1988; Hooser et al., 1989a,b, 1990; Takahashi et al., 1995). In general, similar effects have been reported in both rats and mice, but effects occur later in rats than in mice (Hooser et al., 1989a). Gross and microscopic changes in swine treated intravenously are similar to those observed in rodents (Lovell et al., 1989b).

Ten minutes after mice received a lethal i.p. dose, clinical signs, enzyme changes and liver weight changes were generally absent (Slatkin et al., 1983; Adams et al., 1988). Beginning approximately 20 minutes after dosing in mice, liver weights increased as the livers became suffused with blood (Slatkin et al., 1983; Adams et al., 1988; Hermansky et al., 1990c). Adams et al. (1988) estimated that as much as 44% of the total blood volume was located in the liver 30 minutes after a lethal dose of MCLR in mice. Similarly, in swine treated intravenously, blood volume lost to the liver was estimated to be about 40% (Beasley et al., 2000). At 30 minutes, there were isolated areas of hepatic necrosis, and at 45 minutes, there was marked liver congestion and widespread hepatic necrosis (Hermansky et al., 1990c). Pulmonary thrombi observed at the time of death were generally believed to result from necrotic hepatocytes (Adams et al., 1988). Other pulmonary effects observed at 30-60 minutes after exposure included congestion, bronchial epithelial hyperplasia and necrosis, edema and hemorrhage (Gupta et al., 2003). In general, hepatic enzyme levels show little or no change until 20-30 minutes after dosing, when hepatic hemorrhage is beginning (Adams et al., 1988; Hooser et al., 1989a; Hermansky et al., 1990c; Takahashi et al., 1995).

At 60 minutes, there was severe disassociation of hepatocytes, hepatocyte loss and hemorrhage, with disintegration of liver architecture (Hooser et al., 1991b; Guzman et al., 2003). Thrombocytopenia coincided with hepatic hemorrhage as blood accumulated in the liver (Slatkin et al., 1983; Adams et al., 1988; Takahashi et al., 1995).

The available studies demonstrate a very steep dose-response curve for MCLR acute toxicity. In female NIH non-Swiss outbred mice, the only change observed after i.p. administration of 50 µg MCLR/kg was Kupffer-cell hyperplasia, while all mice receiving 100

μg/kg died (Hermansky et al., 1990c). Hooser et al. (1989a) reported that male rats dosed i.p. with 20, 40 or 80 μg/kg and females dosed with 40 μg/kg MCLR showed no clinical signs of toxicity nor gross or microscopic lesions in the liver or other organs examined, while 120 μg/kg was lethal in some rats and 160 μg/kg was consistently lethal. Lovell et al. (1989a) administered a sublethal i.p. dose (about 25 μg/kg) of MCLR to male mice and reported a significant increase in liver weight (8.7%), but no clinical signs or hepatic lesions.

Induction of apoptosis is believed to be one mechanism for liver damage in acutely exposed animals (Hooser, 2000). Yoshida et al. (1998) treated mice with single sublethal doses of MCLR (20, 30 and 45 μ g/kg i.p.) and observed them for 7 days. Multiple apoptotic cells were noted in the centrilobular regions of the livers of these mice. Hooser (2000) reported widespread apoptosis in most hepatocytes after rats were treated with a single i.p. dose of 500 μ g/kg MCLR. Yoshida et al. (1998) reported the occurrence of two types of apoptotic hepatocytes in mice given sublethal doses of MCLR; one that revealed MCLR by immunohistochemistry and one that did not. The authors suggested that the latter type may contain MCLR that had lost the antigenic epitope, or may have become apoptotic via other means, including ischemia or hypoxia. Apoptosis induction is not restricted to the liver; Botha et al. (2004) reported significantly increased apoptotic indices in the gastrointestinal tract of BALB/c mice as early as 8 hours after a single 75% LD₅₀ dose (specific dose not reported) of MCLR i.p. The authors observed immunohistochemical evidence of MCLR in the lamina propria and postulated that MCLR was involved in the induction of apoptosis. The role of apoptosis in microcystin-induced liver toxicity is further discussed in Section 4.4.7.5 (Mechanistic Studies).

Effects outside the liver have been reported after acute injection exposure to MCLR. It remains unclear whether such effects may be indirectly related to hepatotoxicity. Some studies have shown increases in kidney weight (Hooser et al., 1989a; Lovell et al., 1989a) or other signs of kidney damage (LeClaire et al., 1988; Zhang et al., 2002) in rats and mice after injection of sublethal doses of MCLR. Lovell et al. (1989a) observed dilation of cortical tubules and eosinophilic granular or fibrillar material in the cortical tubules after MCLR administration. In addition to reports of kidney effects, there are scattered reports of cardiac effects, including degeneration and necrosis of myocardial cells after i.p. or i.v. exposure to MCLR (LeClaire et al., 1988; Zhang et al., 2002). LeClaire et al. (1995) reported that mechanisms such as reflex tachycardia and increased cardiac output, which typically would allow the heart to compensate for the acute hypotension caused by blood pooling in the liver, were impaired in rats given a lethal dose of MCLR. The authors suggested a cardiogenic component to the toxicity of MCLR. Oishi and Watanabe (1986) observed tachycardia in mice 20 minutes after i.p. injection of lyophilized cells from *M. aeruginosa*. Finally, a few studies suggest that acute i.p. exposure to MCLR can affect lipid peroxidation levels in both the intestinal mucosa (Moreno et al., 2003) and liver (Towner et al., 2002) of rats.

Two injection studies support the finding by Ito et al. (1997a) by oral exposure that young animals are not as susceptible to the acute toxic effects of microcystins as older animals. Adams et al. (1985) administered lethal doses of MCLR via i.p. injection to 1-, 2- and 3-week-old mice. None of the 1- or 2-week-old mice died, whereas 23 of 31 mice aged 3 weeks died within 2 hours. The 3-week-old mice that survived were rechallenged with MCLR a week later,

and all died. Rao et al. (2005) reported that time to death decreased with age in mice treated i.p. with MCLR.

Guzman and Solter (2002) evaluated the acute effects of repeated injection of MCLR. Male BALB/c mice were injected with 45 μ g/kg MCLR daily for 2, 4 or 7 days. Livers of mice receiving four or seven daily doses were pale and moderately enlarged, with an accentuated reticular pattern; absolute and relative liver weights were statistically increased over controls. Histopathology revealed apoptotic hepatocytes in the centrilobular region of mice receiving two doses, and marked hepatocytomegaly, disorganized hepatocytes, multinucleated hepatocytes and cytoplasmic vacuolation in mice receiving four or seven doses. Guzman et al. (2003) reported immunostaining of some centrilobular hepatocytes following two doses of 45 μ g/kg; however, protein phosphatase activity was not affected.

Acute toxicity of bloom extracts is highly variable, likely reflecting the variable toxin content of algal blooms. LD₅₀ values estimated for various bloom extracts range from 14 mg dry weight cells/kg body weight to 1924 mg dry weight cells/kg (see Table 4-9). These studies used a variety of test materials, including lyophilized cyanobacterial cells, cell-free lysates, etc.

Early studies of purified but unidentified toxins from *M. aeruginosa* show essentially the same pattern of acute hepatotoxicity and time to death after i.p. injection of lethal doses in rats and mice that is seen with purified MCLR (Elleman et al., 1978; Foxall and Sasner, 1981; Falconer et al., 1981; Theiss et al., 1984, 1985, 1988; Jones and Carmichael, 1984; Siegelman et al., 1984; Dabholkar and Carmichael, 1987). In addition, many studies of cyanobacterial bloom extracts (primarily *M. aeruginosa*) administered via i.p. injection to laboratory rodents show similar effects (Ashworth and Mason, 1945; Ohtake et al., 1989; Rao et al., 1994; Porfino et al., 1999; Sabour et al., 2002).

Jackson et al. (1984) also observed liver pathology in sheep exposed intraruminally to a bloom sample identified as *M. aeruginosa*. Time to death ranged from 18 to 48 hours post exposure. In animals that succumbed, the livers were hemorrhagic and necrotic.

Effects on organs other than the liver have been reported in some studies of bloom extracts. Bhattacharya et al. (1997) observed changes suggesting distal tubular dysfunction (proteinurea, as well as decreases in kidney LDH and AST levels) in rats injected with LD₅₀ doses of *M. aeruginosa* extract, but no histopathological changes. Dose-dependent increases in urea and creatinine and decreases in total protein and albumin were also observed. Picanco et al. (2004) reported that i.p. injection of an extract from a culture of *M. aeruginosa* (strain NPJB-1) into either young or mature mice resulted in increased alveolar collapse and increased number of polymorphonuclear and mononuclear cell infiltrations when compared with saline-treated controls. There was a very low concentration of contaminating bacteria in the culture (i.e., the culture was not axenic) used in this study, and the authors acknowledged that materials in the extract other than microcystins may have contributed to the pulmonary effects.

Table 4-9. Intraperitoneal LD ₅₀ Values for Bloom Extracts								
Microcystin Source	Species	Sex/Strain	Vehicle	LD ₅₀ (95% CI)	Duration	Comments	Reference	
Solution of lyophilized cells of <i>M. aeruginosa</i>	Rats	Male/Jcl: Wistar	Saline	67.4 mg dry weight cells/kg, 2 hours	2 hours		Oishi and Watanabe, 1986	
Solution of lyophilized cells of <i>M. aeruginosa</i>	Mice	Male/Jcl:ICR	Saline	14.4 mg dry weight cells/kg, 1 hour	1 hour	None of mice surviving past 1 hour died within 1 week	Oishi and Watanabe, 1986	
Solution of lyophilized cells of <i>M. ichthyoblabe</i>	Mice	Male/Swiss	0.9% saline	502-1924 mg cell dry weight/ kg body weight	Not specified	Microcystin content ranged from 0.73-0.78 µg/g	Sabour et al., 2002	
Cell-free lysate of <i>M</i> . <i>aeruginosa</i>	Mice	Male/Swiss	0.9% saline	431 mg/kg dry weight, 24 hours	24 hours	LD ₅₀ by up and down method	Rao et al., 1994	
Solution of lyophilized bloom sample	Mice	Not specified/ Balb/C	Saline	25-250 mg dry weight phytoplankton sample/kg	Not specified	Bloom dominated by <i>M. aeruginosa</i> . MCLR content of samples ranged from 53-952 µg/g DW biomass	Tarczynska et al., 2000	
Solution of lyophilized bloom sample	Mice	Male/Swiss albino	0.9% saline	154.28 mg algae/kg, 48 hours	48 hours	M. aeruginosa was 95% of bloom biomass	Porfino et al., 1999	
Cell-free lysate of <i>M</i> . aeruginosa	Mice	Male/Swiss albino	Not specified	3.5 g extract/kg, 24 hours	24 hours	MCLR dose approximately 9.625 mg/kg	Rao et al., 2005	
Purified toxin from <i>M</i> . aeruginosa	Mice	Not specified/ white, strain unspecified	Ethanol and water	466 <u>+</u> 13 μg/kg	Not specified	Congener not identified/specified	Bishop et al., 1959	
purified toxin of <i>M</i> . aeruginosa	Mice	Male/white, strain unspecified	Not specified	56 μg/kg (43-60)	Not specified	Congener not identified/specified	Elleman et al., 1978	

In further support for the findings of age-dependent liver effects from purified microcystins, Foxall and Sasner (1981) reported that neonatal and juvenile (age not specified) mice had no clinical symptoms or liver pathology after i.p. injections of a crude lysate from a bloom of M. *aeruginosa*. Details of the experiment and results were not provided. The authors reported that the mice did not die until they were 20 days of age, but it is not clear from the publication whether the mice treated as neonates died later or only mice that were at least 20 days old when treated died. Mature mice were also treated for comparison, but the results were not reported. The authors concluded that "young animals were not sensitive to the toxin but developed sensitivity as they matured".

4.4.5.2.2. Short-Term Studies with Parenteral Exposure

Guzman and Solter (1999) and Solter et al. (1998, 2000) evaluated the effects on rats of short-term administration of MCLR via continuous i.p. infusion. After 28 days of exposure at 16, 32 or 48 µg/kg-day, there were dose-dependent increases in serum levels of sorbitol dehydrogenase (SDH), AST, GGT, ALP and bile acids, while a dose-dependent decrease in serum albumin and a decrease in ALT synthesis were also observed (Solter et al., 1998, 2000).

Immunohistochemistry on the liver showed evidence of bioaccumulation of MCLR in liver cytosol, with measured liver concentrations increasing at a greater rate than the administered dose (Solter et al., 1998). Apoptotic cells and cytoplasmic vacuolation were observed in the livers of rats receiving 32 and 48 μ g/kg-day. A later study exposing rats at the same doses showed evidence for oxidative damage in the liver, as measured by dose-dependent increases in malondialdehyde, a lipid peroxidation byproduct (Guzman and Solter, 1999). This observation is consistent with evidence for oxidative stress after short-term exposure to MCLR.

4.4.5.2.3. Subchronic Studies with Parenteral Exposure

Shi et al. (2002; Chinese publication, only abstract reviewed) reported oxidative stress in rats injected with i.p. doses of 4, 8 or 12 μ g/kg-day MCLR for 35 days. Serum GGT and whole blood glutathione were decreased, while LDH and AST increased after exposure, with no change in ALT levels. Hepatocyte proliferation and apoptosis were also observed. Oxidative stress and apoptosis are discussed further in Section 4.4.7 (Mechanistic Studies).

Elleman et al. (1978) administered daily i.p. injections of a purified toxin from a bloom of M. aeruginosa to white male rats (strain unspecified) for 6 weeks. Doses were reported as fractions of the LD_{100} (0.75, 0.5, 0.25 and 0). These doses correspond to 52.5, 35 and 17.5 μ g/kg based on the reported LD_{100} (70 μ g/kg). Two mice from each group were sacrificed weekly for necropsy and histopathological examination of the liver, kidney, heart, lung, spleen and brain. Eleven of 16 high-dose and 5 of 14 mid-dose mice died prior to scheduled sacrifice; none of the mice receiving the low dose died early. Of the 11 high dose animals that died prematurely, nine died during the first week with symptoms of acute toxicity and liver hemorrhage. In the remaining groups, progressive liver changes were seen with each week, and dose-dependent pathology was observed. The authors noted numerous mitotic figures in hepatocytes of the low-

dose mice early on. Other histopathological findings in the liver were hepatocyte degeneration,

scattered necrosis, fibrosis and mononuclear cellular infiltration; details of the severity of these findings in each group were not reported.

4.4.5.2.4. Chronic Studies with Parenteral Exposure

Milutinovic et al. (2002, 2003) evaluated the kidney effects of chronic i.p. administration of MCLR and MCYR in rats. Doses of 10 μg/kg were administered to groups of five male Wistar rats every other day for 8 months. After sacrifice under CO₂ anesthesia, the kidneys were removed, fixed, sectioned and stained. During exposure, the treated rats exhibited clinical signs of toxicity and reduced body weight. Microscopic examination of the kidneys of treated animals showed collapsed glomeruli and dilated tubules with eosinophilic casts and some cytoplasmic vacuolation. The interstitial space was infiltrated with lymphocytes. More renal corpuscles were significantly damaged in the MCLR-treated group than the MCYR-treated group. Cytoskeletal abnormalities and DNA damage typical of apoptosis or necrosis were also observed in tubular epithelial cells (Milutinovic et al., 2002, 2003). Although they did not report details of the liver pathology in this study, the authors noted that the kidneys were more damaged than the livers, suggesting that adaptation to exposure may have occurred in the livers.

4.4.5.2.5. Initiation/Promotion Studies with Parenteral Exposure

Nishiwaki-Matsushima et al. (1992) demonstrated that i.p administration of MCLR could enhance the number and area of glutathione S-transferase (placental form; GST-P) positive foci in a medium-term rat liver bioassay. In male F344 rats pretreated with 200 mg/kg diethylnitrosamine (DEN) and partially hepatectomized, the number of GST-P positive foci was significantly increased when the rats were subsequently treated with 10 μ g/kg MCLR i.p. twice a week. In a follow-up experiment, rats were pretreated with DEN and then given twice weekly i.p. doses of 10, 25 or 50 μ g/kg MCLR. A dose-dependent increase in the number and area of GST-P positive foci was observed in the animals treated with MCLR (Nishiwaki-Matsushima et al., 1992).

Ohta et al. (1994) also used the two-stage rat liver bioassay model to evaluate the promotion capability of MCLR in rats pretreated with DEN, but not subjected to partial hepatectomy. After treatment with 200 mg/kg DEN, rats given twice weekly i.p. injections of 25 μ g/kg MCLR had significantly more GST-P positive foci and significant increases in the area of such foci when compared with DEN pretreatment alone. MCLR alone had negligible initiating capability.

 Hu et al. (2002) reported significant enhancement of gamma-GT foci in a two-stage medium-term rat bioassay. Microcystin treatment (congener not specified) in DEN-pretreated rats resulted in 100% incidence of gamma-GT foci, while DEN treatment alone resulted in foci in only 22% of rats. Immunohistochemistry showed that microcystin exposure reduced expression of the bax gene and increased expression of the bcl-2 gene. The Bax protein induces apoptosis, while the Bcl-2 protein inhibits apoptosis (Klassen, 2001). This finding suggests that apoptosis is inhibited by low doses of microcystin; in contrast, higher doses appear to induce apoptosis (see Section 4.4.7.5 below).

Sekijima et al. (1999) conducted a similar experiment using either DEN or aflatoxin B1 (AFB1) as an initiator (with partial hepatectomy) and MCLR or a combination of AFB1 and MCLR for promotion. In rats pretreated with 200 mg/kg DEN and subsequently given 10 µg/kg MCLR i.p., there were increases in both the number and area of GST-P positive foci, but this difference did not reach statistical significance. In rats pretreated with DEN and subsequently given either AFB1 alone or a combination of MCLR and AFB1, a statistically significant increase in number and area of foci was observed. The effect on GST-P positive foci of combined treatment with MCLR and AFB1 was not synergistic, however; the number and area of foci in animals treated with both was not larger than the sum of the foci induced by each compound individually. When altered hepatic foci of all types (including basophilic and eosinophilic/clear, rather than only GST-P positive foci) were analyzed, the number of foci was significantly greater in rats treated with both MCLR and AFB1 than in those treated with either toxin alone. To assess the effect of MCLR on initiation by AFB1, Sekijima et al. (1999) pretreated rats with AFB1 followed by twice weekly i.p. injections of 1 or 10 µg/kg MCLR for 6 weeks. The number and area of GST-P positive foci were significantly increased in animals given MCLR compared with controls. There was no difference in number or area of foci between the two doses of MCLR. The authors suggested that the higher dose may have had cytotoxic effects on hepatocytes.

Ito et al. (1997b) treated 13 ICR mice with 100 i.p. injections (5 times/week) of 20 μg/kg MCLR over 28 weeks. Five mice were sacrificed immediately after the last injection, while eight mice were withdrawn from treatment and sacrificed 2 months later. Three untreated mice served as controls. Relative liver weights were 4.7% of body weight in the control mice, 9% in mice sacrificed immediately after the last treatment, and 6.8% in those withdrawn from treatment and sacrificed later. Statistical comparisons among the relative liver weights were not provided, nor were the data with which to perform these comparisons. Neoplastic nodules were observed in the livers of all mice of both treatment groups. The nodules ranged in size up to 5 mm in diameter. The mean numbers of nodules in the treated animals (7.7 and 9.9 nodules per cm² area in the groups sacrificed immediately and 2 months later, respectively) were not significantly different from the controls (control data not provided). The incidence of nodule development in the few control mice was not reported. The types of nodules were characterized as A) weakly staining with hematoxylin and eosin and with small nuclei; B) intensely staining with eosin but not with PAS or C) mainly occupied with fat droplets. The small number of animals in the treatment and control groups limits the usefulness of these data.

4.4.5.2.6. Developmental/Reproductive Studies with Parenteral Exposure

Chernoff et al. (2002) investigated the developmental toxicity of MCLR in CD-1 mice. Pregnant mice were treated with i.p. or subcutaneous doses of MCLR (95% pure) on gestation days (GDs) 7-8, 9-10 or 11-12. Doses of 0, 32, 64 and 128 µg/kg were administered i.p., while only the 128 µg/kg dose level was administered subcutaneously. Mice were sacrificed by CO₂ inhalation on GD 17. After litter and gravid uterus weights were recorded, fetuses were examined for gross malformations and preserved for skeletal examination. Livers of the dams were examined grossly and subjected to histopathology. No effects on maternal weight gain, litter size, average fetal weight or incidence of gross or skeletal abnormalities were observed. Histological examination of the maternal livers showed no effects of treatment.

In a separate experiment by the same researchers (Chernoff et al., 2002), pregnant mice treated with 32-128 $\mu g/kg$ MCLR (via i.p. injection on GD 7-8, 9-10 or 11-12) were allowed to give birth, and the growth and viability of the offspring were followed for 5 days. A different, apparently more potent, lot of MCLR was used in this experiment. Mortality exceeded 50% (19 of 35 dams) in the 64 $\mu g/kg$ dose group and only 1 of 34 animals in each of the 96 and 128 $\mu g/kg$ dose groups survived treatment. Among the surviving animals, there were no effects on viability, birth weight or growth of litters during the brief follow-up period.

Experiments with rabbit whole embryo cultures *in vitro* suggest that low concentrations of MCLR (10-20 μ M) can alter the organization of actin filaments and microtubules, although cell morphology is not significantly affected (Frangez et al., 2003; Zuzek et al., 2003). At high concentrations (100 μ M), MCLR causes cell rounding and loss of adhesion properties, with consequent cell detachment and dispersion. Frangez et al. (2003) showed that the zona pellucida (a glycoprotein envelope surrounding the ovum) forms an effective barrier against the effects of MCLR, as rabbit whole embryos embedded in zona pellucida were not affected even at high concentrations of MCLR.

Development of mouse embryos in culture was inhibited by a purified toxin from a bloom dominated by *Microcystis* (Sepulveda et al., 1992). At a concentration of 120 μ g/mL of toxin in the culture medium, development of two-cell embryos was halted and cytolysis occurred in some embryos. Disruption of the actin cortex was also observed in these embryos. At 60 μ g/mL, two-thirds of the embryos divided once more, and the remainder did not develop further. In eight-cell embryos treated with 120 μ g/mL toxin, compaction was prevented or reversed. Embryos at this stage treated with 240 μ g/mL did not develop further; cells were rounded and lysed. The specific toxin used in this experiment was not identified.

Conflicting results have been observed in teratogenicity testing of purified microcystins in the Frog Embryo Teratogenicity Assay-Xenopus (FETAX) assay with *X. laevis* embryos. In one study, MCLR at concentrations of 25-250 µg/L was shown to induce both skeletal and soft tissue malformations in *X. laevis* embryos (Dvoráková et al., 2002). In contrast, Fischer and Dietrich (2000) reported no effects on mortality, malformation or growth in these embryos after exposure to either MCLR or MCRR at concentrations up to 2000 µg/L. Dvoráková et al. (2002) attributed the differing results to interlaboratory variability or variability in sensitivity of the embryos. O'Brien et al. (2003) reported no effects of MCYR on *X. laevis* embryo mortality, malformation or growth.

Interestingly, Dvoráková et al. (2002) demonstrated that biomass from a *Microcystis* species (*wesenbergii*) that does not produce microcystins could induce malformations in *Xenopus* embryos. O'Brien et al. (2003) also tested two extracts each from *Plantothrix rubescens* and *M. aeruginosa*, reporting that all four extracts resulted in facial narrowing and growth retardation in 96-hour *Xenopus* embryos, while purified MCYR had no effects.

4.4.6. Effects by Intratracheal or Intranasal Instillation

Ito et al. (2001) evaluated the distribution of MCLR after intratracheal instillation of lethal doses in male ICR mice and included a limited description of toxic effects. MCLR in saline solution was instilled at various doses (50, 75, 100, 150 and 200 μ g/kg) into 34 mice; 3 mice were sham-exposed as controls. Mortality was 100% in 12 mice receiving doses of 100 μ g/kg and greater. At 75 μ g/kg, two of four mice died, while no deaths occurred in 18 mice given 50 μ g/kg intratracheally. The time course of hepatotoxicity was further evaluated in eight mice given intratracheal doses of 100 μ g/kg. One mouse was sacrificed at each of 5, 10, 20, 30, 45, 60, 90 and 120 minutes. Immunostaining for MCLR showed the toxin in the lungs within 5 minutes and in the liver after 60 minutes. Hemorrhage in the liver was observed after 90 minutes and became severe by 120 minutes.

Fitzgeorge et al. (1994) conducted experiments in CBA/BALBc mice with MCLR administered via intranasal instillation and inhalation. This study is poorly described, giving few details of study design and findings. The LD₅₀ for intranasal instillation of MCLR was equal to the i.p. LD₅₀ (250 µg/kg). Liver and kidney weights were increased in the animals receiving MCLR intranasally (41.6 and 7.5% respectively). The authors further evaluated the relationship between dose and liver weight increase after intranasal instillation of MCLR. At single intranasal doses of 31.3, 62.5, 125, 250 and 500 µg/kg, liver weight increased proportionally (0, 1.5, 24.4, 37.4 and 87%). Seven daily intranasal doses of 31.3 µg/kg, a dose that had produced no liver weight change after a single dose, resulted in a liver weight increase of 75%. Fitzgeorge et al. (1994) reported histopathological findings, but failed to specify which findings resulted from single doses and which resulted from the multiple-dose experiment reported in the same publication. Findings included necrosis of respiratory and olfactory epithelium in the nasal mucosa and centrilobular necrosis with hemorrhage in the liver. Early changes in the liver included vacuolar degeneration and necrosis of hepatocytes near the central vein. The adrenal glands showed effects as well, with vacuolation and necrosis of the inner cortex, as well as congestion of medullary blood vessels. No histopathological changes were observed in the trachea, lungs, esophagus, pancreas, spleen, lymph nodes, kidneys or brain.

4.4.7. Mechanistic Studies

Many mechanistic studies have been conducted to characterize the toxicology of microcystins. These studies include *in vivo* investigations in laboratory animals, *in situ* studies in isolated perfused organ systems and *in vitro* assays in isolated cell preparations. Mechanistic studies have evaluated many aspects of microcystin toxicity, including: 1) the reason for target organ and cell type specificity of microcystins, 2) description of the subcellular effects that occur in susceptible cells, 3) interaction with serine and threonine protein phosphatases (i.e., PP1 and PP2A) as the molecular target for microcystins, 4) the role of cytoskeletal effects, 5) the importance of oxidative stress and apoptosis as a mode of toxic action and 6) the use of chemoprotectants to reduce toxicity. Each of these topics is discussed in further detail below. Mechanistic data related to the genotoxicity of microcystins is presented below in Section 4.4.8 (Genotoxicity and Cell Proliferation).

4.4.7.1. Target Organ/Cell Type Specificity

Oral and injection studies in laboratory animals have demonstrated that the liver is the primary target organ for microcystin toxicity (see Section 4.2). Mechanistic studies suggest that the target organ specificity is directly related to the limited ability of microcystins to cross cell membranes in the absence of an active transport system, such as the bile acid transporter in hepatocytes. Evidence of the importance of the bile acid transporter to liver toxicity is provided by studies that used bile acids and bile acid transport inhibitors (Runnegar et al., 1981, 1993, 1995b; Runnegar and Falconer, 1982; Eriksson et al., 1990a; Hermansky et al., 1990a,b; Hermansky et al., 1991). These studies demonstrated that the liver toxicity produced by *in vitro* or *in vivo* exposures to microcystins was reduced or eliminated by inhibition of hepatocellular uptake using bile acid transport inhibitors (e.g., antamanide, sulfobromophthalein and rifampicin) and bile salts (i.e., cholate and taurocholate). Additional discussion of the cellular uptake of microcystins is provided in Section 3.2 (Distribution).

Runnegar et al. (1993) demonstrated that i.p. injection of mice with MCYM or MCLR caused inhibition of liver protein phosphatase activity followed by evidence of liver toxicity (i.e., increased liver weight). Kidney protein phosphatase activity was unchanged following the *in vivo* exposure even at lethal doses. *In vitro* exposure of kidney extracts to microcystins did result in a decrease in kidney phosphatase activity, and no difference in sensitivity was observed between liver and kidney phosphatase inhibition. This result suggests that target organ specificity is most likely due to slower intercellular uptake of microcystins in the kidney.

The cell type specificity of microcystins was investigated using isolated rat hepatocytes, rat renal epithelial cells (ATCC 1571) and rat skin fibroblasts (ATCC 1213) (Khan et al., 1995; Wickstrom et al., 1995). The time course of light microscopic and ultrastructural effects was examined following in vitro exposure to MCLR (Khan et al., 1995). Effects were noted after 4 minutes in hepatocytes, 1 hour in renal cells and 8 hours in fibroblasts. Similar lesions observed in all cell types included blebbing, loss of cell-cell contact, clumping and rounding, cytoplasmic vacuolization and redistribution of cellular organelles. Effects that were seen only in hepatocytes include loss of microvilli, whirling of rough ER, dense staining and dilated cristae of mitochondria and pinching off of membrane blebs. The nuclear changes typical of apoptosis were seen in renal cells and fibroblasts. Cell type differences may be related to the specific proteins that were overphosphorylated within each cell type. The authors postulated that the lack of apoptotic changes in hepatocytes in this study might be related to the short exposure duration or the failure of their transmission electron microscopic method to examine severely damaged cells that had detached from the coverslips. Wickstrom et al. (1995) evaluated the changes in cytoskeletal morphology after MCLR exposure in these cell types. High concentrations and long incubation times were required for cytoskeletal changes in kidney and skin cells; however, the nature of the changes was similar in all cell types (e.g., actin aggregation). Wickstrom et al. (1995) suggested that microcystins may enter kidney cells and fibroblasts via pinocytosis.

McDermott et al. (1998) treated several cell types with MCLR (primary rat hepatocytes, human fibroblasts, human endothelial cells, human epithelial cells and rat promyelocytes). Hepatocytes underwent membrane blebbing, cell shrinkage, organelle redistribution, chromatin

condensation and, in some cells, DNA fragmentation. Similar changes were observed in the other cell types, but a longer duration of exposure was required.

Matsushima et al. (1990) demonstrated that injection of MCYR into mouse skin epithelial cells and human fibroblasts resulted in morphological changes in cell shape (i.e., spindle shape to round forms). These cells were thought to be resistant to microcystins; however, microcystins YR, LR and RR produced a dose-dependent inhibition of protein phosphatase activity using the partially purified enzyme derived from mouse skin cell cytosol. The authors suggest that absence of a direct effect in these cells is due to lack of penetration through the cell membrane.

Many cell types and established cell lines have been evaluated for potential susceptibility to microcystin uptake and toxicity. Primary isolated hepatocytes have been shown to be the most sensitive to cytoxicity, due to the presence of the organic ion/bile acid transport system (Eriksson et al., 1987, 1990b). Uptake was negligible in human hepatocarcinoma cells (Hep G2), mouse fibroblasts (NIH-3T3), erythrocytes and human neuroblastoma cells (SH-SY5Y). Hepatic endothelial cells have also been shown to be resistant to microcystin toxicity (Solow et al., 1989; Runnegar et al., 1994). Primary cultures of liver cells cease to express these bile acid transport proteins after 2-3 days of being maintained in culture. Therefore, established liver cell lines are not generally useful for evaluating microcystin toxicity (Eriksson et al., 1994; Battle et al., 1997; Heinze et al., 2001).

4.4.7.2. Characterization of Subcellular Effects in the Liver

The liver effects that occur following *in vivo* exposures to microcystins are generally discussed in Section 4.2 (Animal Studies). Many additional mechanistic studies describe liver histopathology, ultrastructural changes and biomarkers of cytotoxicity in either isolated perfused rat liver (Berg et al., 1988; Pace et al., 1991; Runnegar et al., 1995b) or primary isolated hepatocytes (Runnegar et al., 1981; Runnegar and Falconer, 1982; Aune and Berg, 1986; Runnegar and Falconer, 1986; Berg and Aune, 1987; Runnegar et al., 1987; Falconer and Runnegar, 1987a,b; Thompson et al., 1988; Solow et al., 1989; Mereish et al., 1989; Mereish and Solow, 1990; Eriksson et al., 1990a; Boe et al., 1991; Khan et al., 1995; Runnegar et al., 1995b; Yea et al., 2001; Batista et al., 2003).

Toxicological effects of microcystins in the isolated perfused rat liver were similar to those demonstrated following *in vivo* exposure (Pace et al., 1991). During a 60-minute exposure, MCLR caused liver engorgement and cessation of bile flow. Electron microscopy revealed loss of sinusoidal architecture, dilation of bile canaliculi and the space of Disse and decreased intracellular contact. Mitochondrial swelling, disruption of endoplasmic reticulum and formation of whorls and loss of desmosomal intermediate filaments were also observed. Mitochondrial function was impaired, with inhibition of state 3 respiration and a decrease in the respiratory control index.

Runnegar et al. (1995b) demonstrated a decrease in protein phosphatase activity in perfused rat liver exposed to MCYM. Cessation of bile flow, increased perfusion pressure, decreased protein secretion and decreased glucose secretion were also observed. Histological

changes included hepatocytes swelling, loss of sinusoidal architecture, pyknotic nuclei and extensive necrosis. Exposure to high concentrations of toxin extracts in the isolated perfused liver produced loss of cord architecture due to hepatocyte disassociation, membrane damage and cytolysis and nuclear effects (pyknosis, karyokinesis, karyolysis) (Berg et al., 1988). Ultrastructural effects included swollen mitochondria, vacuoles, necrosis, abnormal nuclei, bile canaliculi lacking microvilli and whorls of rough endoplasmic reticulum.

Studies in primary isolated hepatocytes have described the morphological and histopathological changes induced by microcystins that relate to loss of sinusoidal architecture and cytotoxicity (Runnegar et al., 1981; Runnegar and Falconer, 1982; Aune and Berg, 1986; Runnegar and Falconer, 1986; Berg and Aune, 1987; Runnegar et al., 1987; Falconer and Runnegar, 1987a,b; Thompson et al., 1988; Solow et al., 1989; Mereish et al., 1989; Mereish and Solow, 1990; Eriksson et al., 1990a; Boe et al., 1991; Runnegar et al., 1995b; Khan et al., 1995; Yea et al., 2001; Batista et al., 2003). Microcystin exposure to hepatocytes in suspension or cultured in a monolayer results in membrane blebbing that becomes more pronounced and localized in one region of the cell surface. Cells are observed to be rounded in appearance and become dissociated from one another. Microfilaments are reorganized as a compact spherical body in the vicinity of the blebbing, while the rest of cell is depleted of filamentous actin. MCLR disrupts hepatocellular morphology within minutes, leading to loss of sinusoidal architecture and hemorrhage. Morphological changes in hepatocytes (i.e., blebbing, rounding) have been shown to occur prior to any effect on cell membrane integrity (measured as LDH leakage or release of radiolabeled adenine nucleotides) or cell viability (generally measured as decreased trypan blue exclusion) (Runnegar et al., 1981; Runnegar and Falconer, 1982; Aune and Berg, 1986; Ding et al., 2000a).

Thompson et al. (1988) described the time course of cellular effects of microcystins (type not specified) on cultured rat hepatocytes. Disintegration of attachment matrix occurred by 15 minutes, followed by cells clustered in groups with no extracellular material at 1 hour, and release of cells from plates between 2 and 4 hours. LDH release did not occur until after these visual effects, but was dose-related.

Similar toxicological effects were observed in isolated human hepatocytes (Yea et al., 2001; Batista et al., 2003). MCLR produced blebbing, fragmentation and hepatocyte disassociation. Cytotoxicity, as measured by LDH leakage, occurred after morphological changes were evident. Yea et al. (2001) indicated that cytotoxicity in human hepatocytes was observed at a concentration (1 μ M) that did not affect rat hepatocytes. Batista et al. (2003) also reported a slightly higher susceptibility to microcystin-induced morphological change in human hepatocytes as compared to rat hepatocytes.

 The ultrastructural effects observed following microcystin exposure in isolated rat hepatocytes (i.e., condensation of chromatin, segregation of organelles, separated by apoptotic microbodies, decrease in cell volume and increase in cell density) suggest that hepatocyte cell death is apoptosis and not necrosis (Boe et al., 1991). Microcystin exposure in hepatocytes causes cell shrinkage, externalization of membrane phosphatidylserine, DNA fragmentation and chromatin condensation, indicating a rapid apoptosis (Ding et al., 2000b). Apoptosis is discussed in further detail below as a possible mode of action for microcystin liver toxicity.

Several studies have suggested that microcystins may increase the release and decrease the reincorporation of arachadonic acid into cellular membranes (Adams et al., 1985; Naseem et al., 1990, 1991; Nobre et al., 2001).

4.4.7.3. Molecular Target: Inhibition of Type 1 and 2A Protein Phosphatases

The primary molecular target of microcystins has been identified as serine and threonine protein phosphatases PP1/PP2A. Protein phosphatases dephosphorylate proteins while protein kinases phosphorylate them. Together, protein kinases and phosphatases maintain the balance of phosphorylation and dephosphorylation of key proteins involved in cell cycle regulation. Because more than 97% of protein phosphates occur at serine and threonine residues (Gehringer, 2004), the PP1 and PP2A protein phosphatases are particularly important. Inhibition of these enzymes results in the increased phosphorylation of a number of regulatory proteins. Importantly, PP1 is believed to be the major phosphorylase *a* phosphatase in the liver (Runnegar et al., 1993). PP2A, the major soluble serine/threonine phosphatase, regulates several mitogenactivated protein kinases (Gehringer, 2004).

Microcystins bind to these enzymes under both *in vivo* and *in vitro* study conditions, resulting in an inhibition of enzyme activity leading to an increase in protein phosphorylation. Microcystins have been shown to directly inhibit the activity of PP1 and PP2A derived from several different species (i.e., fish, mammals, plants) and cell types (Honkanen et al., 1990; MacKintosh et al., 1990; Matshushima et al., 1990; Yoshizawa et al., 1990; Sim and Mudge, 1993; Xu et al., 2000; Leiers et al., 2000; Becchetti et al., 2002). Microcystins do not alter protein kinase activity, suggesting the balance of phosphorylation and dephosphorylation is related to protein phosphatase inhibition alone. Microcystins have been used as a tool to investigate the importance of serine and threonine phosphorylation to specific cellular functions. The regulatory effects of phosphorylation on sodium channel opening in renal cells (Becchetti et al., 2002), smooth and skeletal muscle contraction (Hayakawa and Kohama, 1995; Knapp et al., 2002) and insulin secretion (Leiers et al., 2000) have been studied.

Runnegar et al. (1993) demonstrated the inhibition of PP1 and PP2A activity in the liver following i.p. injection of MCYM and MCLR. Increased protein phosphorylation preceded the observed increase in liver weight and was correlated with hepatotoxicity. Decreased phosphatase activity was also demonstrated in the isolated perfused liver (Runnegar et al., 1995b).

The relationship between phosphatase inhibition by microcystins and changes in cytoskeletal structure and cell morphology has been reviewed (Eriksson and Golman, 1993). Inhibition of protein phosphatase activity by MCLH and 7-dmMCRR was associated with an increased phosphorylation of cytoskeletal and cytosolic proteins (Eriksson et al., 1990b). Concentrations that produce a marked increase in protein phosphorylation were accompanied by a complete reorganization of microfilament network. The cytoskeletal effects of microcystins are discussed in further detail below. Microcystin LH and 7-dmMCRR were equipotent inhibitors of purified PP1 and PP2A; however, higher concentrations of 7-dmMCRR were required to increase protein phosphorylation. Table 4-10 shows studies with comparative data on inhibition of protein phosphatases (IC₅₀s) by MCLR, MCYR, MCRR and MCLA.

Table 4-10. Studies Comparing Protein Phosphatase Inhibition Activity of Microcystin Congeners

Reference	IC ₅₀ (nM)							
1101010100	MCLR	MCLA	MCYR	MCRR				
PP2A Inhibition								
Craig et al., 1996	0.15	0.16						
Nishiwaki-Matsushima et al., 1991	0.28			0.78				
Matsushima et al., 1990	7.6		4.5	5.8				
PP1 Inhibition								
MacKintosh et al., 1995	0.2		0.2					
Mixture of PPs								
Yoshizawa et al., 1990	1.6		1.4	3.4				

The molecular interaction between microcystins and protein phosphatases has been evaluated using immunoprecipitation, autoradiography, reverse phase liquid chromatography, X-ray crystallography, nuclear magnetic resonance (NMR) solution structures, and molecular dynamics simulation (Runnegar et al., 1995; MacKintosh et al., 1995; Goldberg et al., 1995; Craig et al., 1996; Bagu et al., 1997; Mattila et al., 2000; Mikhailov et al., 2003; Maynes et al., 2004, 2006). Molecular modeling and molecular dynamics simulations have indicated that microcystins bind in a Y-shaped groove containing the catalytic site on the surface of PP1 (Mattila et al., 2000). Studies with PP1 suggest that the C-terminal β12-β13 loop of PP1 (containing residues 268-281) is important for microcystin-protein phosphatase interactions as well as for substrate recognition (Maynes et al., 2004, 2006). Information available to date indicates that the binding process primarily involves the amino acids Glu, Adda, Leu and MDha of microcystins. Figure 4-2 shows a schematic representation of the interactions between microcystin-LR and protein phosphatase 1; these interactions are discussed further below.

Microcystins LR, LA and LL interact with the catalytic subunits of PP1 and PP2A in two phases. The first phase occurs within minutes and consists of rapid inactivation of the phosphatase. The second, slower phase of interaction represents a covalent interaction that takes place within several hours (Craig et al., 1996). The initial binding and inactivation of protein phosphatases appears to result from several non-covalent interactions that are still being elucidated. Mattila et al. (2000) demonstrated an interaction of the Glu amino acid (reported as

IGlu in the publication) carboxyl group of MCLR with a metal ion (Fe, Mn) in the PP1 catalytic 1 site. Glu appears to be an important component because esterification eliminates toxicity 2 (Namikoshi et al., 1993; Rinehart et al., 1994). Herfindal and Selheim (2006), in a review of the 3

mechanisms of microcystin toxicity, indicated that the Adda side chain is involved in a 4

hydrophobic interaction between the Trp 206 and Ile130 residues in the hydrophobic groove of 5

PP1. Mattila et al. (2000) suggested that the long side chain of the Adda residue may contribute 6 7

to orienting the toxin into the catalytic site. The Adda amino acid residue of microcystins plays

an important role in the inhibition of protein phosphatase activity (Nishiwaki-Matsushima et al., 8

1991; Gulledge et al., 2002, 2003a,b). Isomerization of the diene from 4E,6E to 4E,6Z on the 9 Adda chain (see Figure 2-1) eliminates the toxic activity of microcystins (Harada et al., 1990; 10

Nishiwaki-Matsushima et al., 1991; Stotts et al., 1993). Microcystin analogues containing only 11 12

Adda and one additional amino acid are capable of substantial inhibition of PP1 and PP2A, while modifications to the Adda structure abolished the inhibition (Gulledge et al., 2003b). Finally,

Herfindal and Selheim (2006) indicated that the L-Leu of MCLR participates in a hydrophobic

interaction with Tyr 272 of PP1 (on the β12-β13 loop).

The second phase of interaction between microcystins and protein phosphatase consists of covalent bonding (Craig et al., 1996). Immunoprecipitation and autoradiography methods indicate that covalent bonds result from the interaction between the methylene of the MDha residue of microcystins and the thiol of Cys273 located at the C-terminal of PP1. NMR solution structures and X-ray crystallography data on the MCLR/PP1 complex illustrate the covalent linkage at Cys-273 (Goldberg et al., 1995; Bagu et al., 1997). Site-directed mutagenesis replacing Cys273 in PP1 results in a loss of microcystin binding (MacKintosh et al., 1995; Maynes et al., 2004). Based on sequence similarity between PP1 and PP2A, it has been suggested that Cys-266 is the site of equivalent covalent linkage between PP2A and microcystins (Craig et al., 1996).

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Microcystin analogues containing a reduced MDha residue are not capable of covalent binding to protein phosphatases. MacKintosh et al. (1995) reported that a reduction of the MDha residue of MCYR by ethanethiol abolished covalent binding to PP1. Likewise, Craig et al. (1996) showed that reduction of the MDha residue of MCLA abolished the covalent binding phase with PP2A. Maynes et al. (2006) confirmed the lack of covalent interaction by determining the crystal structure of dihydroMCLA bound to PP1. Their work showed that the β12-β13 loop of PP1 takes on a different conformation when the covalent bond is absent, and that other interactions (including hydrogen bonding) are responsible for the bond between dihydroMCLA and PP1.

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The importance of covalent bonding between microcystins and protein phosphatases to toxicity resulting from the enzyme inhibition is uncertain, as other interactions are apparently responsible for the rapid inactivation of the enzymes (Herfindal and Selheim, 2006). Modifications to either molecule (microcystin or protein phosphatase) to prevent covalent bonding generally decrease, but do not eliminate, the toxic action (Meriluoto et al., 1990;

MacKintosh et al., 1995; Hastie et al., 2005).

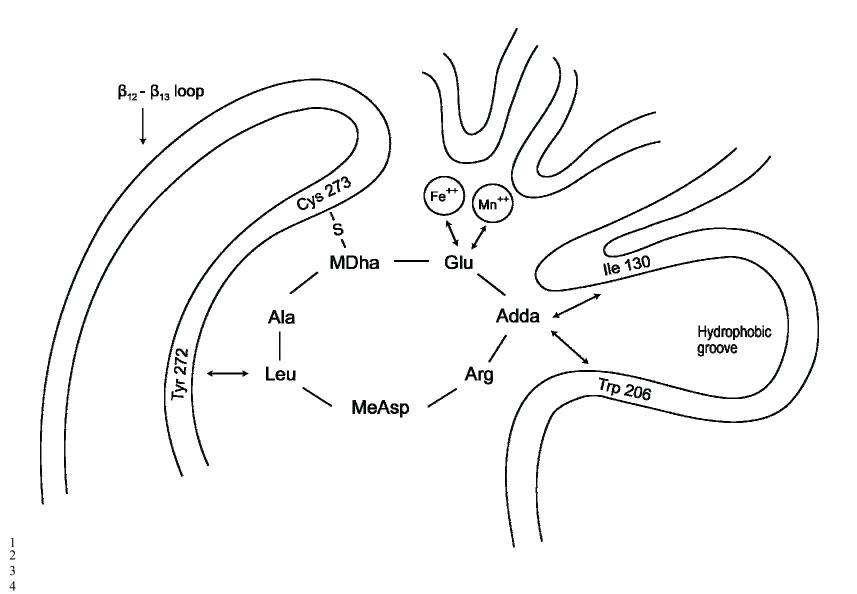


Figure 4-2. Schematic Representation of Interactions between Microcystin-LR and the Catalytic Site of Protein Phosphatase 1

Microcystins may also bind to other molecular targets in addition to protein phosphatases. Chen et al. (2006) used bioinformatic approaches to identify human liver aldehyde dehydrogenase 2 (ALDH2) as a potential molecular target of MCLR. After screening a phage display library to identify potential ligands specific for MCLR, Chen et al. (2006) used molecular docking studies to show that MCLR could bind to ALDH2. The authors postulated that this interaction could lead to aldehyde-induced reactive oxygen species (ROS) and apoptosis.

4.4.7.4. Cytoskeletal Effects

The cytoskeletal effects of microcystins in the liver have been visually demonstrated in several studies using light, electron and fluorescent microscopy (Runnegar and Falconer, 1986; Eriksson et al., 1989; Hooser et al., 1989a,b, 1991b; Falconer and Yeung, 1992). Ultrastructural changes in rats given a lethal dose of microcystin A¹ include the following: a widening of intracellular spaces; progressive disassociation followed by rounding, blebbing and invagination of hepatocytes; loss of microvilli in the space of Disse; breakdown of the endothelium; hemorrhage; and loss of lobular architecture (Hooser et al., 1989b). No effects were noted in endothelial cells or Kupffer cells. In isolated hepatocytes, actin aggregates were seen at the base of the membrane blebs. As membrane blebs grew larger and were drawn toward one pole of the cell, the microfilaments were organized toward the same pole, resulting in rosette formation with a condensed band of microfilaments at the center (Runnegar and Falconer, 1986; Eriksson et al., 1989; Hooser et al., 1991b; Falconer and Yeung, 1992; Wickstrom et al., 1995; Ding et al., 2000a). Frangez et al. (2003) also demonstrated cytoskeletal changes in rabbit primary whole embryo cultured cells. Actin and microtubule disorganization was demonstrated to lead to detachment and cellular toxicity.

The observed reorganization of microfilaments that leads to alteration of hepatocyte morphology was not shown to be due to effects on actin polymerization (Runnegar and Falconer, 1986; Eriksson et al., 1989; Falconer and Yeung, 1992). Instead, microcystins cause an increase in the phosphorylation of cytokeratin intermediate filament proteins (Falconer and Yeung, 1992): Ohta et al., 1992; Wickstrom et al., 1995; Blankson et al., 2000). Toivola et al. (1997) evaluated the effects of MCLR on hepatic keratin intermediate filaments in primary hepatocytes cultures. A disruption of the desmoplakin² organization at the cell surface (disorganization of desmosomes) is followed by a dramatic reorganization of the intermediate filament and microfilament networks, resulting in intermediate filaments being organized around a condensed actin core. The major target proteins for microcystin-induced hyperphosphorylation include keratins 8 and 18 and desmoplakin (DP) I/II. Keratins 8 and 18 are the major proteins of intermediate filaments in hepatocytes; DP I and II attach keratin filaments in epithelial cells to desmosomes. Hyperphosphorylation of DPI/II leads to loosening of cell junction and loss of interactions with cytoplasmic intermediate filaments. The hyperphosphorylation of keratin proteins leads to increased solubility (caused by disassembly or prevention of subunit polymerization), leading to the observed morphological effects. Phosphopeptide mapping shows

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¹ The authors refer to the test compound as microcystin-A, which may reflect an old nomenclature no longer in use. Available information is insufficient to identify the congener with current nomenclature.

² Desmoplakin is the principal plaque protein in a desmosome, which is a localized thickening of membrane that serves as an adhesion junction connecting contiguous cells.

four specific tryptic peptides in soluble keratin 18 that are highly phosphorylated; however, no specific phosphorylation sites have been identified for keratin 8. A Ca²⁺/calmodulin-dependent kinase may be involved in regulating the serine-specific phosphorylation of keratin proteins 8 and 18.

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Some investigators have suggested that generation of reactive oxygen species may play a role in the cytoskeletal changes induced by microcystins. Ding et al. (2001) illustrated generation of superoxide and hydrogen peroxide radicals preceding microfilament disorganization and cytotoxicity. Hepatocellular glutathione levels were affected by microcystins, and administration of N-acetylcysteine was shown to protect against cytoskeletal alterations (Ding et al., 2000a).

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4.4.7.5. Apoptosis

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The ultrastructural changes observed in hepatocytes after microcystin exposure suggest that cell death is related to apoptosis and not necrosis. These changes include cell shrinkage (decreased volume and increased density), condensation of chromatin and segregation of organelles separated by apoptotic microbodies (Boe et al., 1991; Fladmark et al., 1998; McDermott et al., 1998; Ding et al., 2000b; Mankiewicz et al., 2001). The effects of microcystins on the signaling pathways involved in rapid apoptosis have been investigated in several studies (Ding et al., 1998a,b, 2000b, 2001, 2002; Ding and Ong, 2003). Mitochondrial permeability transition (MPT) is considered to be a critical rate-limiting event in apoptosis. Oxidative stress may play a role in the induction of MPT and the onset of apoptosis. In cultured hepatocytes exposed to microcystins, an increase in the generation of ROS preceded the onset of MPT, mitochondrial depolarization and apoptosis. A dose- and time-dependent increase in ROS and lipid peroxidation, measured as malondialdehyde formation, was shown to precede morphological changes in hepatocytes and release of LDH. The addition of deferoxamine or cyclosporine A inhibited the formation of ROS and delayed the onset of MPT and cell death. The addition of superoxide dismutase prevented collapse of cytoskeleton and release of LDH from isolated hepatocytes. An early surge of mitochondrial Ca²⁺ was shown to occur prior to MPT and cell death. Prevention of this Ca²⁺ surge by one of several methods (i.e., chelation of intracellular Ca²⁺, blockage of the mitochondrial Ca²⁺ uniporter or use of mitochondrial uncoupler) prevented MPT and cell death. Electron transport chain inhibitors (e.g., rotenone, actinomycin A, oligomycin or carbonyl cyanide m-chlorophenylhydrazone) also inhibited the onset of MPT. MCLR caused the release of cytochrome c through MPT, which is considered universal in mitochondrial apoptosis; however, caspases -9 and -3 were not activated. The increase in intracellular Ca²⁺ may instead facilitate the activation of calpain, which occurred following exposure to microcystins (Ding and Ong, 2003). Botha et al. (2004) demonstrated that apoptosis and oxidative stress can be induced in nonhepatic cells by microcystins. LDH leakage and increased apoptotic indices were observed in the human colon carcinoma cell line (CaCo2) and MCF-7 cells (deficient in pro-caspase-3). These changes were accompanied by increased H₂O₂ formation and increased calpain activity.

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45 46 Western blot analysis has been used to show that MCLR increases the expression of p53 and the pro-apoptotic Bax protein in both cultured rat hepatocytes treated with MCLR *in vitro* and rat liver after *in vivo* exposure (Fu et al., 2005). Expression of the anti-apoptotic protein

Bcl-2 was decreased in vitro, but in vivo MCLR treatment did not lead to a difference in the expression of this protein. This finding suggests that MCLR may induce apoptosis through other mechanisms in addition to the inhibition of protein phosphatases.

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4.4.7.6. Lipid Peroxidation

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Several studies have investigated the role of glutathione homeostasis and lipid peroxidation in microcystin-induced liver toxicity (Runnegar et al., 1987; Eriksson et al., 1989; Bhattacharya et al., 1996; Ding et al., 2000a; Towner et al., 2002; Gehringer et al., 2003a,b, 2004; Bouaïcha and Maatouk, 2004). Ding et al. (2000a) indicated that microcystin exposure in isolated hepatocytes resulted in an initial increase in glutathione synthesis followed by a later depletion of glutathione. MCLR was shown to induce the *de novo* synthesis of glutathione in mice exposed to a toxic sublethal dose (75% of the LD_{50}) (Gehringer et al., 2004). Increased transcription of glutathione-S-transferase was also demonstrated in his study. Gehringer et al. (2004) suggest that increased lipid peroxidation induced by microcystins is accompanied by an increase in glutathione peroxidase, transcriptional regulation of glutathione-S-transferase and glutathione peroxidase and de novo synthesis of glutathione. Bouaïcha and Maatouk (2004) also reported that a low noncytotoxic concentration (2 ng/mL) of MCLR in primary rat hepatocytes caused an initial increase in ROS formation and an increase in glutathione; however, a decrease in lipid peroxidation was observed in this study. Electron spin resonance (ESR) spin trapping techniques have demonstrated the formation of two possible lipid-derived free radical metabolites in rat liver following in vivo exposure to MCLR (Towner et al., 2002). Vitamin E and selenium supplementation in mice provided some protection against liver toxicity and lethality by MCLR (Gehringer et al., 2003a,b). Measures of liver toxicity included serum enzyme determination, lipid peroxidation, glutathione levels and histopathology. Hermansky et al. (1991) reported that membrane active antioxidants (i.e., vitamin E, silymarin and glutathione) provided some protection from microcystin toxicity (i.e., LDH leakage) and lethality; however, free radical scavengers and water soluble antioxidants were ineffective (see below).

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Several studies have reported MCLR-induced increases in lipid peroxidation as well as decreases in antioxidant enzymes (Moreno et al., 2005; Jayaraj et al., 2006). Jayaraj et al. (2006) measured oxidative stress in mice treated i.p. with an LD₅₀ dose of MCLR. Significant increases in heat shock protein-70 and hepatic lipid peroxidation were observed. Further, GSH was depleted, and there were decreases in the activity of glutathione peroxidase, superoxide dismutase, catalase, glutathione reductase and glutathione-S-transferase in the animals treated at the LD₅₀. Similarly, Moreno et al. (2005) reported significant reductions in glutathione peroxidase, glutathione reductase, superoxide dismutase and catalase, along with increases in lipid peroxidation, in both the liver and kidney of rats treated intraperitoneally with single doses of MCLR.

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Some studies report the absence of lipid peroxidation during microcystin-induced hepatotoxicity. A time-dependent leakage of LDH, ALT and AST was observed in liver slices with no change observed for glutathione content or lipid peroxidation (Bhattacharya et al., 1996). In addition, Runnegar et al. (1987) indicated that glutathione depletion did not occur until after morphological changes (i.e., blebbing) were observed. Eriksson et al. (1989) indicated that rapid

deformation of isolated rat hepatocytes by MCLR was not associated with alterations in glutathione homeostasis.

4.4.7.7. Prevention of Liver Toxicity and Lethality

Several types of agents have been evaluated as potential chemoprotectants against microcystin toxicity, including inhibitors of bile acid transport, microsomal enzyme inducers, calcium channel blockers, free radical scavengers, water-soluble antioxidants and membrane active antioxidants. It was initially reported that preincubation of hepatocytes with chemicals that interfere with uptake of bile acids (sulfobromophthalein, rifampicin, sodium cholate and sodium deoxycholate) also prevents hepatocyte deformation (Runnegar et al., 1981; Runnegar and Falconer, 1982).

Hermansky et al. (1991) evaluated several possible chemoprotectants by measuring LDH leakage and lethality following i.p. injection of MCLR. No protective effect was observed using calcium channel blockers, free-radical scavengers or water-soluble antioxidants administered prior to MCLR. Membrane active antioxidants, such as vitamin E, silymarin and glutathione, provided some protection from microcystin toxicity and lethality. Phenobarbital provided partial protection; however, tetrachlorodibenzo-*p*-dioxin did not afford protection, suggesting that the protective effect may not be related to microsomal enzyme induction. Rifampicin and cyclosporine A provided complete protection by blocking uptake of microcystins at the bile acid transporter. Hermansky et al. (1990a,b) reported that rifampicin can be given 15 minutes after MCLR injection and still prevent lethality, while cyclosporine A prevents lethality only if given 0.5-3 hours before MCLR injection.

Silymarin and dithioerythritol, both antioxidants, were shown to reduce MCLR toxicity, as measured by LDH and adenine nucleotide release and light microscopy in primary cultures of adult rat hepatocytes (Mereish and Solow, 1990). Dithioerythritol and silymarin have both been shown to increase the content of reduced thiols (i.e., glutathione). Silymarin has additionally been shown to stabilize membranes, inhibit lipoxygenase, reduce leukotrienes, scavenge free radicals and increase protein synthesis.

Mereish et al. (1991) indicated that silymarin pretreatment inhibited microcystin liver toxicity in mice (evidenced by histopathology and serum enzyme levels) following i.p. injection, but not oral administration. As discussed above, vitamin E and selenium supplementation in mice provided some protection against liver toxicity and lethality by MCLR (Gehringer et al., 2003a,b).

Thompson and Pace (1992) evaluated several types of agents for their ability to protect against MCLR toxicity in cultured hepatocytes. Toxicity was measured as morphology under light microscopy, LDH release and protein synthesis inhibition. The uptake of MCLR into hepatocytes was also measured. Cytochalasins D&E, fungal metabolites that interfere with actin polymerization into microfilaments, were shown to protect against LDH release and provided moderate protection from rounding and clustering of cells; however, these compounds produced cytotoxicity themselves at concentrations that were required for protection against microcystin toxicity. Cholate and deoxycholate are competitive inhibitors for the bile acid transporter.

These compounds provided some protection against LDH leakage, but were also cytotoxic. Trypan blue and trypan red also provided some protection related to blocking microcystin uptake into hepatocytes. The antibiotic rifampicin was shown to prevent microcystin uptake and toxicity at low non-cytotoxic concentrations, suggesting a possible therapeutic use in microcystin poisoning.

Rao et al. (2004) confirmed that pretreatment with cyclosporine-A, rifampin and silymarin each provided 100% protection against a lethal dose of MCLR (although the route was not specified, it is assumed to be i.p., based on the lethal dose). Protected animals had significantly reduced glutathione and increased hepatic lipid peroxidation up to 7 days after treatment, but levels were returned to normal by 14 days.

Adams et al. (1985) demonstrated that MCLR lethality following i.p. injection in mice was reduced by pretreatment with a single subcutaneous injection of carbon tetrachloride. Lethality was also reduced in young mice in this study (no deaths at 1 and 2 weeks of age, 23/31 deaths at 3 weeks of age), suggesting that normal hepatic function is necessary for the uptake and hepatotoxicity of MCLR. The microsomal enzyme inhibitors SKF525A and cobalt chloride produced no effect on MCLR lethality, indicating that microsomal metabolism is not critical for MCLR toxicity. The administration of hydrocortisone was also shown to protect against MCLR lethality in mice, possibly due to a decrease in the release of arachadonic acid from membrane phospholipids. Naseem et al. (1990) demonstrated that pretreatment of cultured rat hepatocytes with glucocorticoids (flucinolone, dexamethasone and hydrocortisone) reduced the release of arachadonic acid and metabolites caused by MCLR.

β-Carotene and lutein inhibited the effect of MCLR on hepatocyte morphology in mouse primary hepatocyte cultures (Matsushima-Nishiwaki et al., 1995). β-Carotene protected the cytokeratin network from disassembly and suppressed the hyperphosphorylation of cytokeratins 8 and 18. Several carotenoid analogs were evaluated, and the protective effect appeared to be related to the number of *trans* configured double-bonds in the carotenoid.

4.4.7.8. Extra-Hepatic Effects of Microcystins

 An isolated perfused kidney model was used to evaluate the kidney toxicity of MCLR (Nobre et al., 1999, 2001). MCLR produced vascular, glomerular and tubular effects in the exposed kidney. An increase in perfusion pressure was followed by an increase in the glomerular filtration rate (GFR), increased urinary flow rate and a reduction in tubular transport at the proximal tubules. Histopathological changes included protein in the urinary spaces, but were not further described. Dexamethazone and indomethacin were shown to antagonize the effects of MCLR on perfusion pressure, renal vascular resistance (RVR), GFR and urinary flow. These results suggest a role for phospholipase A2 and cyclooxygenase in the kidney toxicity of microcystins. Nobre et al. (2003) utilized rat peritoneal macrophages exposed to MCLR to further investigate the role of inflammatory mediators in the isolated perfused kidney model. Macrophage supernatants from exposed rats caused an increase in RVR, GFR and urinary flow and reduced Na⁺ transport. These effects were reduced by cyclohexamide, dexamethasone and quinacrine, further suggesting the involvement of PLA₂ and other inflammatory mediators in microcystin-induced kidney toxicity.

 Moreno et al. (2003) investigated the effects of MCLR on intestinal physiology following an i.p. injection of 100 µg/kg in rats. Lipid peroxidation was increased in both the serum and the intestinal mucosa of treated rats. With the exception of sucrase, intestinal brush border enzymes were unaffected by MCLR exposure. An increase in the specific activity of acid phosphatase and succinate dehydrogenase in intestinal homogenates suggests an effect of MCLR on lysosomal and mitochondrial membranes, respectively. Nobre et al. (2004) used perfused rat ileal segments and ligated intestinal loops to evaluate the effect of MCLR on electrolyte and water secretion. MCLR caused significant secretion of water, sodium, potassium and chloride. Aziz (1974) observed that a dialyzable component of whole cell lysate from *M. aeruginosa* caused fluid accumulation in the ligated small intestine of guinea pigs.

Sicińska et al. (2006) evaluated the effects of MCLR on human erythrocytes *in vitro*. MCLR exposure resulted in the formation of echinocytes, hemolysis, conversion of oxyhemoglobin to methemoglobin, and a decrease in membrane fluidity. In addition, measures of oxidative stress were affected in treated erythrocytes; glutathione reductase and superoxide dismutase activity were decreased, while ROS and lipid peroxidation were increased.

Several studies have evaluated the effects of MCLR on immune system components *in vitro* (Lankoff et al., 2004; Teneva et al., 2005; Chen et al., 2005; Kujbida et al., 2006). Lankoff et al. (2004) reported that MCLR inhibited B-cell proliferation in human and chicken peripheral blood lymphocytes at all concentrations tested, and decreased T-cell proliferation only at the highest concentration. Apoptosis was enhanced in both human and chicken lymphocytes (Lankoff et al., 2004). Similarly, MCLR was cytotoxic to mouse splenocytes, and caused apoptosis in B-cells but not in T-cells (Teneva et al., 2005).

Kujbida et al. (2006) assessed the effects of MCLR and [Asp3]-MCLR on human polymorphonuclear lymphocytes (PMNs) *in vitro*. Both compounds caused migration of neutrophils in a chemotaxis chamber, suggesting that PMNs may migrate from the blood stream to organs that concentrate microcystins, such as the liver. In addition, both caused a dose-related increase in ROS production as measured by chemiluminescence of PMN degranulation products that accompany ROS production. The phagocytosis of *Candida albicans* by PMNs was increased after exposure to either compound, but only MCLR increased the intracellular killing of *C. albicans*. These findings suggest the possibility that PMNs may mediate some of the toxic effects of microcystins.

4.4.8. Genotoxicity and Cell Proliferation

Available data give conflicting results when purified MCLR has been tested for mutagenicity. Pure MCLR did not induce mutations in the Ames assay either with or without metabolic activation, although microcystin-containing extracts did induce mutations (Ding et al., 1999). A crude toxin extracted from *M. aeruginosa* did not induce mutations in the Ames assay (Grabow et al., 1982). In contrast, Suzuki et al. (1998) observed increased ouabain resistance mutation frequency in human embryo fibroblast cells treated with MCLR (purity not specified). Similarly, Zhan et al. (2004) reported a 5-fold increase over control in the frequency of thymidine kinase mutations when human lymphoblastoid TK6 cells were treated with

commercially-obtained MCLR. More slow-growing mutants were observed than fast-growing mutants, suggesting that the mutation damage was larger than the TK locus, and that MCLR induced large deletions, recombinations or rearrangements. Repavich et al. (1990) reported that Ames assays (using strains TA98, TA100 and TA102) of a purified hepatotoxin (supplied by Wright State University and presumed to be microcystin) were negative, as were *Bacillus subtilis* multigene sporulation assays.

The conflicting information on mutagenicity may be related to differences in the cell uptake of MCLR. For example, the failure of MCLR to induce mutations in bacterial cells may be related to poor uptake. Zhan et al. (2004) reported that MCLR is not taken up by many cell types, including bacteria; however, the authors did not provide references to support this assertion. While hepatocytes take up MCLR at a significant rate, other cell types show limited or no uptake unless measures are taken to enhance the penetration of the cells by MCLR. The cellular uptake of microcystins is discussed in detail in Sections 3.2 (Distribution) and 4.4.7 (Mechanistic Studies).

A number of studies have reported DNA damage after MCLR treatment *in vivo* (Rao and Bhattacharya, 1996), and in primary rat hepatocytes (Ding et al., 1999) and human hepatoma cells (Zegura et al., 2003, 2004). Recent studies suggest that apoptosis may be intimately linked to observations of DNA damage in cells treated with MCLR. Lankoff et al. (2004) showed a strong correlation between DNA damage, as measured by the comet assay, and the induction of apoptosis, as measured by the terminal deoxynucleotidyl transferase-mediated dUTP-biotin nick end-labeling (TUNEL) assay, in human lymphocytes. Other evidence has suggested that the comet assay can give a false positive measure of DNA damage when apoptosis is induced, as DNA fragmentation is one consequence of apoptosis induction (Lankoff et al., 2004). The authors postulated that earlier reports of DNA damage measured by the comet assay may have been related to early stages of apoptosis due to cytotoxicity rather than a direct effect on DNA. The induction of apoptosis appears to be dose-related. Humpage and Falconer (1999) showed that low (picomolar) concentrations of commercially-obtained MCLR induced cytokinesis and inhibited apoptosis in primary mouse hepatocytes, while higher (nanomolar) concentrations resulted in the inverse effects.

Bouaïcha et al. (2005) reported that noncytotoxic concentrations of MCLR did not cause the formation of hydrophobic DNA adducts in primary cultured rat hepatocytes treated *in vitro*, but did decrease the amount of endogenous hydrophobic adducts. MCLR was also shown to cause a dose- and time-dependent increase in the formation of 8-oxo-7,8-dihydro-2'-deoxyguanosine (a measure of oxidative DNA damage) both in cultured hepatocytes and in rat liver cells after *in vivo* treatment via i.p. injection (Maatouk et al., 2004; Bouaïcha et al., 2005).

Conflicting results have been reported in studies of MCLR-induced clastogenicity. MCLR (commercially-obtained) has induced micronuclei in human lymphoblastoid cells and mouse bone marrow erythrocytes (Ding et al., 1999; Zhan et al., 2004). Lankoff et al. (2004) observed no effect of MCLR on the incidence of chromosomal aberrations in human peripheral blood lymphocytes. Observations of polyploidy in MCLR-treated cells (Humpage and Falconer, 1999; Lankoff et al., 2003) may be related to its effects on cytokinesis. Lankoff et al. (2003) showed that MCLR, through its effect on microtubules, damages the mitotic spindle, leading to

the formation of polyploid cells. Repavich et al. (1990) reported a dose-related increase in chromosome breakage in human lymphocytes exposed to a purified hepatotoxin (presumed to be a microcystin).

Mechanistic evidence provides support for the hypothesis that MCLR can act as a promoter at low doses. Zhu et al. (2005) reported that MCLR can transform immortalized colorectal crypt cells, resulting in anchorage-independent growth and enhanced proliferation. MCLR has been shown to increase the expression of the bcl-2 protein (that inhibits apoptosis) and decrease the expression of the bax protein (that induces apoptosis) (Hu et al., 2002). Further, MCLR upregulates the transcription factors c-fos and c-jun, leading to abnormal proliferation (Zhao and Zhu, 2003). Gehringer (2004) reviewed the molecular mechanisms leading to promotion by MCLR and the related tumor promoter, okadaic acid. Gehringer (2004) reported that MCLR inhibits protein phosphatase PP2A, which regulates several mitogenactivated protein kinases (MAPK). The MAPK cascade regulates transcription of genes required for cell proliferation, including c-jun and c-fos. In addition, activation of the MAPK cascade has been postulated to inhibit apoptosis and thus increase cell proliferation. Finally, Gehringer (2004) noted that MCLR has been reported to increase phosphorylation of p53, which is involved in the regulation of the cell cycle and apoptosis as well as the carcinogenic process.

4.4.9. Structure-Activity Relationships

With a few exceptions, microcystin congeners exhibit i.p. LD_{50} values between 50 and 300 µg/kg in mice (Rinehart et al., 1994; WHO, 1999). MCLR is one of the most potent congeners (i.p. LD_{50} approximately 50 µg/kg). Limited comparative testing of *in vitro* protein phosphatase inhibition (IC₅₀) of MCLR, -RR and -YR resulted in IC₅₀ values of 1.6, 3.4 and 1.4 nM, respectively (Yoshizawa et al., 1990), indicating that microcystin congeners may be relatively similar in protein inhibition potency. Pharmacokinetic differences among the various microcystin congeners may be at least partially responsible for observed variations in lethal potency (Ito et al., 2002). Microcystin congeners of varying hydrophobicity were shown to interact differently with lipid monolayers (Vesterkvist and Meriluoto, 2003). Effects on membrane fluidity could alter the cellular uptake of these toxins.

Wolf and Frank (2002) proposed toxicity equivalency factors (TEFs) for the four major microcystin congeners based on LD_{50} values obtained after i.p. administration. The proposed TEFs, using MCLR as the index compound (TEF=1.0) were 1.0 for MCLA and MCYR and 0.1 for MCRR. The application of TEFs based on i.p. LD_{50} values to assessment of risk from oral or dermal exposure is questionable given that differences in liphophilicity and polarity of the congeners may lead to variable absorption by non-injection routes of exposure.

The molecular interaction between microcystins and the catalytic subunits of protein phosphatases has been extensively studied (see Section 4.4.7.3 for more detail). The interaction was shown to occur in two phases. The first phase occurs within minutes and results in rapid inactivation of the phosphatase (Craig et al., 1996). The amino acids Glu and Adda appear to be important for the rapid inactivation of the protein phosphatases and for subsequent toxicity (Harada et al., 1990; Nishiwaki-Matsushima et al., 1991; Namikoshi et al., 1993; Rinehart et al., 1994; Gulledge et al., 2002, 2003a,b). The carboxyl group of the Glu residue in MCLR

apparently interacts with a metal (Fe, Mn) ion in the PP1 catalytic site (Mattila et al., 2000). The Adda side chain is involved in a hydrophobic interaction between the Trp 206 and Ile130 residues in the hydrophobic groove of PP1 (Herfindal and Selheim, 2006). The few apparently non-toxic (i.p. LD_{50} values >1000 μ g/kg) microcystin congeners exhibit structural alterations in the Adda or Glu regions (Harada et al., 1990; Stotts et al., 1993; Rinehart et al., 1994).

The second, slower phase of interaction represents a covalent interaction between the methylene of the MDha residue of microcystins and the thiol of Cys273 located at the C-terminal of PP1 that takes place over several hours (Craig et al., 1996). Microcystin analogues containing a reduced MDha residue are not capable of covalent binding to protein phosphatases (MacKintosh et al., 1995; Craig et al., 1996; Maynes et al., 2006). The importance of covalent bonding to the toxic effect of microcystins is uncertain, as other interactions are apparently responsible for the rapid inactivation of the enzymes (Herfindal and Selheim, 2006). Modifications to either molecule (microcystin or protein phosphatase) to prevent covalent bonding generally decrease, but do not eliminate the toxic action (Meriluoto et al., 1990; MacKintosh et al., 1995; Hastie et al., 2005; Herfindal and Selheim, 2006).

4.5. MODE OF ACTION – NONCANCER AND CANCER

Microcystins appear to result in different cellular effects depending on dose. In a review of the mechanistic data on microcystins, Gehringer (2004) postulated a dualistic response for microcystins and okadaic acid (another potent inhibitor of PP1 and PP2A). Gehringer (2004) outlined evidence suggesting that at high doses, microcystins cause alterations in cellular structure and function that may lead to cell death via apoptosis or necrosis, while at low doses, microcystins inhibit apoptosis and cause cell proliferation. The high-dose effects are likely to be responsible for the acute toxicity and lethality of microcystins and are discussed below.

4.5.1. Target Organ Specificity

The liver is the primary site of toxicological action for microcystins after oral, i.v., i.p. and intranasal instillation exposure. Acute and short-term exposures to microcystins have resulted in intrahepatic hemorrhage in both rats and mice (Ito et al., 1997a; Fawell et al., 1999; Heinze, 1999). One subchronic study in mice also showed liver effects (Fawell et al., 1999), including hepatocyte degeneration, chronic inflammation and hemosiderin deposits, but no hemorrhage. Liver hemorrhage, resulting from apoptosis and necrosis of hepatocytes leading to disintegration of hepatic architecture, appears to be the most prominent effect observed in available toxicological studies. The main reason for this target organ specificity is the greater cellular uptake of microcystins by hepatocytes compared with other cells. Microcystins are actively transported into hepatocytes by the bile acid transporter system, while uptake by other cell types is limited by the lack of an active transport system. *In vitro* studies demonstrate that preincubation of hepatocytes with compounds that block the uptake of bile acids prevent damage to the hepatocyte from microcystin exposure (Runnegar et al., 1981, 1993, 1995a; Runnegar and Falconer, 1982; Eriksson et al., 1990a; Hermansky et al., 1990a,b, 1991). Treatment of various cell types with microcystins results in rapid damage to hepatocytes, while changes in other cell types occur after much longer exposure (Khan et al., 1995; Wickstrom et al., 1995; McDermott et al., 1998).

Available mechanistic data indicate that the hepatotoxic effects of microcystins begin with one or two molecular effects: inhibition of protein phosphatases and induction of oxidative stress. At present, it is not clear whether the two effects are linked, whether they lead independently to similar cellular effects (e.g., cytoskeletal damage and apoptosis), or whether one effect is the predominant cause of hepatocellular damage. In both cases, the initial cellular effects are associated with cytoskeletal changes and the induction of apoptosis. These alterations in the structural integrity and function of hepatocytes lead to profound hepatotoxicity and hemorrhage. In summary, the key events in the hepatotoxicity of microcystins appear to be:

- 1. Molecular events (inhibition of protein phosphatase and/or induction of oxidative stress);
- 2. Cellular effects (cytoskeletal damage and/or apoptosis); and
- 3. Tissue damage (altered liver structure and function, and intrahepatic hemorrhage).

The molecular events and cellular effects leading to tissue damage are discussed further below.

4.5.2.1. Molecular Events

cytokeratin intermediate filament proteins.

Protein Phosphatase Inhibition. As discussed above in Section 4.4.7.3, microcystins are potent inhibitors of serine and threonine PP1 and PP2A. Inhibition of these protein phosphatases results in aberrant phosphorylation of a number of cellular proteins, with the potential for multiple effects on the cell. Current data suggest that the inhibition of PP1/PP2A by microcystins can trigger cytoskeletal damage and apoptosis.

One outcome of microcystin-induced inhibition of PP1 and PP2A is the hyperphosphorylation of cytokeratin intermediate filament proteins (Falconer and Yeung, 1992; Ohta et al., 1992; Wickstrom et al., 1995; Blankson et al., 2000). Specifically, microcystin exposure results in hyperphosphorylation of keratins 8 and 18 and desmoplakin I/II (Toivola et al., 1997). It has been suggested that protein kinases PKC, PKA or the calcium/calmodulin-dependent kinase may play a role in the hyperphosphorylation of these proteins (Gehringer, 2004). The hyperphosphorylation of desmoplakin I/II results in the loosening of cell junctions and loss of interactions with cytoplasmic intermediate filaments, while the hyperphosphorylation of keratins 8 and 18 leads to increased solubility. Some of the morphological changes observed in hepatocytes (e.g., blebbing, rounding) may result from the hyperphosphorylation of

Guzman et al. (2003) reported an increase in the phosphorylation of p53 in rat livers after i.p. exposure to MCLR. PP1 and PP2A help to regulate the activity of p53 through dephosphorylation. Thus, inhibition of these enzymes can result in hyperphosphorylation of p53. Increases in the phosphorylation of p53 can cause an increase in the transcription of p21^{WAF1}, which in turn inhibits cyclin D-, E- and A-dependent kinases. The result of the latter inhibition is to stall the cell cycle in G1 phase (Gehringer, 2004). This disruption of cell cycling can allow for repair of DNA damage or for apoptosis to occur. For MCLR, there is evidence of

hyperphosphorylation of p53, but other steps in this cascade of events have not yet been investigated.

Oxidative Stress. A number of studies indicate that oxidative stress may play a role in microcystin-induced hepatotoxicity. As noted above in Section 4.4.7.5, studies of chemoprotectants show that several antioxidants can provide protection against the toxicity of microcystins. Among the antioxidants shown to protect against the effects of microcystin, either *in vitro* or *in vivo*, are vitamin E, silymarin, dithioerythritol, desferoxamine, N-acetylcysteine, superoxide dismutase and glutathione. Further, dose and time-dependent increases in reactive oxygen species have been shown to precede morphological changes in hepatocytes, and the addition of superoxide dismutase prevents the cytoskeletal collapse caused by microcystins. Ding and Ong (2003) have proposed two primary pathways by which microcystins increase oxidative stress leading to cell death. First, microcystins may deplete glutathione, leading to oxidative damage and cell death. Second, microcystins may increase the production of ROS by disrupting the mitochondrial electron transport chain, leading to mitochondrial permeability transition and apoptosis.

Microcystins may enhance oxidative stress by altering glutathione homeostasis; however, the importance of glutathione homeostasis in MCLR-induced hepatotoxicity is not clear. Glutathione serves as an intracellular antioxidant, by scavenging free radicals, by serving as a substrate for the reduction of hydrogen peroxide by glutathione peroxidase and by detoxifying xenobiotics. In addition, depletion of glutathione can disrupt microfilament structures in some cell types (Ding and Ong, 2003). MCLR lethality has been prevented in mice by pretreatment with glutathione (Hermansky et al., 1991), and hepatocytes pretreated with a glutathione precursor were likewise protected from MCLR toxicity. Some studies have shown glutathione depletion after microcystin exposure (Runnegar et al., 1987); however, depletion did not occur until after membrane blebbing had been observed. Other studies have reported an increase in glutathione after MCLR exposure (Ding et al., 2000a; Bouïacha and Maatouk, 2004). Finally, Eriksson et al. (1989) indicated that the rapid deformation of rat hepatocytes after MCLR exposure was not associated with changes in glutathione levels. Thus, the role of glutathione homeostasis in MCLR-induced hepatotoxicity has not yet been determined.

A variety of studies have demonstrated the importance of mitochondrial permeability transition in the apoptotic cascade induced by MCLR (see Ding and Ong, 2003; Gehringer, 2004). In particular, studies have shown that pretreating hepatocytes with cyclosporin A, a specific inhibitor of MPT, prevented cell death from microcystin exposure (Ding and Ong, 2003; Gehringer, 2004). Ding and Ong (2003) outlined the following pathways for MCLR-induced apoptosis via MPT. First, microcystin disrupts the mitochondrial electron transport chain, leading to the release of reactive oxygen species from mitochondria and mitochondrial permeability transition. MPT triggers a release of cytochrome c and mitochondrial calcium. Cytochrome c may activate one or more caspases that trigger apoptosis, although neither caspase-9 nor caspase-3 appear to be involved based on current information (Ding and Ong, 2003; Gehringer, 2004). The release of mitochondrial calcium activates calpain and calcium/calmodulin-dependent protein kinase II, both of which also lead to apoptosis.

4.5.2.2. Cellular Effects

Cytoskeletal Changes. Morphological changes observed in hepatocytes treated *in vitro* with MCLR include membrane blebbing, cell rounding and dissociation. Membrane blebs become localized in one region of the cell, and microfilaments are reorganized as a compact spherical body near the blebbing (Runnegar and Falconer, 1986; Eriksson et al., 1989; Hooser et al., 1991b; Falconer and Yeung, 1992; Wickstrom et al., 1995; Ding et al., 2000a). These morphological changes occur before cell viability or cell membrane integrity is affected. Electron microscopy of isolated perfused rat liver showed that these cellular effects led to loss of sinusoidal architecture, dilation of bile canaliculi and the space of Disse and decreased intercellular contact (Pace et al., 1991). Intrahepatic hemorrhage results from the breakdown of liver structure, and the liver is rapidly engorged with blood.

Apoptosis. Membrane blebbing is also a characteristic of the apoptotic process (Gehringer, 2004). A growing body of evidence indicates that microcystin exposure can trigger apoptosis. Hooser (2000) used several visualization (light and electron microscopy) and analytical techniques (TUNEL and electrophoresis to evaluate DNA laddering) to demonstrate widespread apoptosis in the livers of rats 3 hours after an i.p. dose of MCLR. Characteristic apoptotic changes including cell rounding, shrinkage, disassociation, loss of microvilli and chromatin margination and condensation were observed in a majority of hepatocytes. The author postulated that the rapidity with which the apoptotic process occurred overwhelmed the phagocytic capacity of the liver, such that apoptotic hepatocytes depleted their energy stores and later underwent necrosis. In mice, apoptotic hepatocytes have also been observed, but not to the degree reported in rats. Hooser (2000) postulated that intrahepatic hemorrhage and death occurred so quickly in mice that cellular changes characteristic of apoptosis did not have time to develop.

4.5.3. Conclusion

The mechanisms by which microcystins induce hepatic damage have not been fully elucidated. Available evidence suggests roles for both protein phosphatase inhibition and oxidative stress as important molecular events, since chemoprotectant studies show that pretreatment with compounds that inhibit either of these effects can protect against hepatotoxicity in MCLR-treated animals. It is possible, even likely, that microcystin exposure triggers a series of independent or linked events that cause cytoskeletal damage and/or apoptosis, given the numerous cellular functions controlled by PP1 and PP2A, as well as the number of effects triggered by oxidative stress. These cytoskeletal and apoptotic changes apparently lead to the altered hepatic structure/function and intrahepatic hemorrhage observed in animal studies.

4.6. SYNTHESIS AND EVALUATION OF MAJOR NONCANCER EFFECTS

The preponderance of toxicological data on the effects of microcystins is restricted to the MCLR congener. A single, poorly-described study, reported only in a secondary source is available for the LA congener. Data on the YR and RR congeners are limited to i.p. LD_{50} values and measures of relative inhibition of protein phosphatases. As a result, this section largely describes the available information on the toxic effects of MCLR, with limited reference to other congeners.

Anecdotal reports indicate that, in humans, exposure to cyanobacterial blooms (including microcystin-producing genera) can result in neurological, gastrointestinal and dermatological symptoms, such as headache; muscle weakness; eye, ear and throat irritation; nausea; stomach pain; diarrhea; blistering around the mouth; and hay-fever like symptoms (Dillenberg and Dehnel, 1960; Billings, 1981; Turner et al., 1990; Teixeira et al., 1993; el Saadi and Cameron, 1993). Effects were reported in persons exposed via recreational contact (swimming, boating) and drinking water. Turner et al. (1990) also reported pneumonia in army recruits exposed to a cyanobacterial bloom. Symptoms occurring after exposure to cyanobacteria cannot be directly attributed to microcystin toxins (or other endotoxins); some effects may result from exposure to the cyanobacterial cells themselves, or from exposure to multiple toxins in the bloom.

The primary noncancer health effect of exposure to MCLR is liver damage. The liver is targeted largely because hepatocytes are among only a few cell types that actively take up microcystins, which do not readily cross the cell membrane. Severe liver damage (diffuse individual hepatocyte necrosis, cell-plate disruption and apoptosis) occurred in dialysis patients exposed to microcystins³ in dialysate (Jochimsen et al., 1998; Pouria et al., 1998; Carmichael et al., 2001; Azevedo et al., 2002). At high acute doses in laboratory animals, MCLR caused potentially fatal hemorrhaging within the liver. While the liver is the usual target of microcystin toxicity, there have been some reports of effects in other systems, including hematological, kidney, cardiac, neurological and gastrointestinal effects. It has been suggested that some effects in other organs observed after high doses of MCLR may result from ischemia or hypoxia caused by hepatic hemorrhage. However, some effects outside the liver have been observed in the absence of hemorrhage.

Much of the toxicological data on microcystins are limited to reports of liver effects after single lethal or sublethal doses administered via i.p. injection. These studies indicate that injected doses of 50-200 µg/kg MCLR or MCYR are usually lethal in mice and rats within a few hours (Adams et al., 1988; Hooser et al., 1989a; Hermansky et al., 1990c; Stotts et al., 1993; Gupta et al., 2003; Rao et al., 2005). Miura et al. (1991) showed that the median time to death is greatly increased in fed rats (32 hours) when compared with fasted rats (less than 2 hours). The authors suggest that fasting may increase the sensitivity of animals to the mitochondrial toxicity of microcystins, although this could not be conclusively demonstrated. In the liver, MCLR destroys the cytoskeleton of hepatocytes, leading to hepatocyte disassociation, degeneration,

³ Exposure was to untreated water containing cyanobacteria. The presence of microcystins was confirmed in patient biopsy samples; however, it is possible that the patients may also have been exposed to other microbial or chemical contaminants.

apoptosis and necrosis (Hermansky et al., 1990c; Hooser et al., 1991b). Hepatic hemorrhage and disintegration of the liver architecture follow quickly (Hooser et al., 1991b). Effects reported to occur outside the liver include pulmonary thrombi derived from necrotic hepatocytes, kidney effects such as dilation of cortical tubules and eosinophilic material in the cortical tubules, and degeneration and necrosis of myocardial cells (Adams et al., 1988; LeClaire et al., 1988; Zhang et al., 2002). As previously stated, some of these effects may occur secondary to hepatic hemorrhage.

Injection studies suggest a very steep dose-response curve for acute liver effects from microcystin exposure. In several studies, mice and rats receiving single i.p. doses of 20-40 μ g/kg MCLR showed no clinical toxicity and few or no gross or microscopic effects in the liver or other organs (Hooser et al., 1989a; Lovell et al., 1989a; Hermansky et al., 1990c), while i.p. doses of 50-200 μ g/kg are usually lethal within a few hours (Hooser et al., 1989a; Hermansky et al., 1990c; Stotts et al., 1993; Gupta et al., 2003).

4.6.1. Oral

Table 4-11 provides a summary of the noncancer effects from repeated-dose oral studies of MCLR toxicity in laboratory animals. The table includes all of the studies that used purified microcystins as the test substance. As the table indicates, the toxicological database for effects of microcystins after oral exposure is limited.

 Liver Effects. One study of human exposure to drinking water before, during and after a bloom of *M. aeruginosa* reported a significant increase in GGT levels during the bloom compared with levels before the bloom (Falconer et al., 1983). The study population consisted of all persons subjected to liver function tests in the area served by the affected drinking water supply; as such, it is not representative of the general population. The liver is the primary target organ when laboratory animals are exposed to high doses of MCLR. Oral exposure to single 500 μg/kg doses of MCLR caused diffuse hemorrhage in the liver of mice and rats; more pronounced liver damage occurred at higher doses (Ito et al., 1997a; Fawell et al., 1999). Young mice (5 weeks old) did not develop signs of hepatotoxicity at 500 μg/kg MCLR, while aged mice (32 weeks old) developed clear signs (Ito et al., 1997a). This difference may result in part from differences in gastrointestinal absorption of microcystins, but cannot be entirely explained by absorption differences, since similar age-dependent effects were reported after i.p. exposure (Adams et al., 1985; Rao et al., 2005).

A single 28-day study of oral exposure to 50 or 150 μ g/kg MCLR in drinking water showed increased liver weight, slight to moderate liver lesions with hemorrhages and increased ALP and LDH in rats exposed at 50 μ g/kg-day (Heinze, 1999). A subchronic study in mice using a similar dose range identified a LOAEL of 200 μ g/kg (Fawell et al., 1999). At this dose, mild liver lesions including chronic inflammation, hemosiderin deposits and single hepatocyte degeneration were observed, as well as increased ALT and AST in male animals. The 40 μ g/kg dose was identified as a NOAEL. Mild hepatocyte injury was reported in mice given 80 or 100 gavage doses of 80 μ g/kg each over 28 weeks, corresponding to time-weighted average doses of 33-41 μ g/kg-day (Ito et al., 1997b). Based on the report, it appears that a limited postmortem

	Table 4-11. Summary Noncancer Results in All Animal Studies of Oral Exposure to Purified Microcystin-LR								
Species	Sex	Average Daily Dose (µg/kg-day)	Exposure	NOAEL (µg/kg- day)	LOAEL (µg/kg- day)	Responses	Comments	Reference	
Acute Exposure									
Rat	M/F	500, 1580, 5000	Single gavage	ND	5000*	Mortality; diffuse hepatic hemorrhage at lower doses	No untreated controls. Dose- dependent increase in hepatotoxicity	Fawell et al., 1999	
Mouse	M/F	500, 1580, 5000	Single gavage	ND	1580*	Mortality; diffuse hepatic hemorrhage at lower dose	No untreated controls. Dose- dependent increase in hepatotoxicity	Fawell et al., 1999	
Mouse	F	8000, 10000, 12500	Single gavage	ND	12500*	Mortality (2/2); hypertrophic hepatocytes, fibrosis in centrilobular and midzonal regions at lower doses	No untreated controls. 1-2 animals/dose group.	Yoshida et al., 1997	
Mouse	М	0, 500	Single gavage	ND	500 (aged mice only)	Centrilobular hepatic hemorrhage and necrosis; necrosis of intestinal mucosa and duodenal damage	Effects observed in aged (32 week-old) mice; no effects on liver or gastrointestinal tract in young (5 week-old) mice	Ito et al., 1997a	
Short-Tei	Short-Term Exposure								
Rat	М	0, 50, 150	Drinking water, 28 day	ND	50	Slight to moderate degenerative and necrotic hepatocytes with hemorrhages; increased serum enzymes (ALP and LDH)		Heinze, 1999	

	Table 4-11. cont.								
Species	Sex	Average Daily Dose (µg/kg-day)	Exposure	NOAEL (μg/kg- day)	LOAEL (µg/kg- day)	Responses	Comments	Reference	
Subchron	Subchronic Exposure								
Mouse	M/F	0, 40, 200, 1000	Daily gavage, 13 weeks	40	200	Minimal/slight chronic inflammation with hemosiderin deposits and single hepatocyte degeneration; increased serum enzymes (ALT and AST)		Fawell et al., 1999	
Chronic l	Exposure								
Mouse	F	0, 3	Drinking water, 18 months	3	ND	No effects on survival, body weight, hematology, serum biochemistry, organs or histopathology	Minor changes in ALP and cholesterol not considered toxicologically significant by researchers	Ueno et al., 1999	
Mouse	Not given	Not available	Gavage, 80 μg/kg, 80-100 times over 28 weeks	ND	ND	Light injuries to hepatocytes in the vicinity of the central vein	Only liver examined; only three control animals; dosing frequency unclear	Ito et al., 1997b	
Developr	nental To	xicity							
Mouse	F	0, 200, 600, 2000	Gavage, GD 6-15	600	2000*	Maternal mortality (7/26) and morbidity (2/26 humanely sacrificed); reduced fetal body weight, delayed skeletal ossification	Authors defined 600 µg/kg-day as NOAEL but did not present data on reproductive or developmental parameters to support identification of LOAEL	Fawell et al., 1999	

^{*} Frank Effect Level (FEL)

examination was conducted in this study, which was primarily aimed at evaluating carcinogenicity. No liver or other toxicity was reported in female mice given approximately 3 µg/kg-day MCLR in drinking water for 18 months (Ueno et al., 1999).

Neurological Effects. The database contains scattered reports of neurological symptoms after exposure to high doses of MCLR. Dialysis patients exposed to microcystins in dialysate reported symptoms such as visual disturbance, blindness, vertigo, headache and muscle weakness (Jochimsen et al., 1998). Clinical signs in mice and rats orally exposed to lethal doses (about 5000 μ g/kg) include hypoactivity and piloerection (Fawell et al., 1999).

Other Organs. Gastrointestinal effects (necrosis, duodenal damage) were observed in aged mice exposed orally to single 500 μ g/kg doses of MCLR (Ito et al., 1997a). Kidney effects including eosinophilic materials in the Bowman's spaces were observed in two mice exposed to a lethal dose of 12.5 mg/kg (Yoshida et al., 1997). Female mice exposed subchronically to 1000 μ g/kg had slight increases in hemoglobin concentration, erythrocyte count and packed cell volume (Fawell et al., 1999). Milutinovic et al. (2002, 2003) briefly reported that kidney effects are more pronounced than liver effects in rats chronically exposed to i.p. doses of MCLR and MCYR (time weighted average dose, 5 μ g/kg for 8 months). Details of the liver examinations were not reported in this study, limiting the usefulness of these data.

Developmental Effects. A single oral study of developmental toxicity in mice reported maternal toxicity, liver effects and deaths in some dams treated at the highest dose of MCLR (2000 µg/kg during GD 6-15), along with reduced fetal body weight and delayed skeletal ossification. No effects on reproductive or developmental parameters were observed in other treatment groups, and 600 µg/kg was identified as a NOAEL for developmental toxicity (Fawell et al., 1999). One study of developmental toxicity after i.p. injection of 32-238 µg/kg MCLR in mice confirmed the lack of developmental or reproductive effects in the absence of maternal toxicity (Chernoff et al., 2002). A study in which an extract of *M. aeruginosa* (estimated to contain about 14 µg/L unspecified toxin) was administered in the drinking water to mice before and during pregnancy revealed small brains in 7 of 73 pups from treated parents and none in untreated controls (Falconer et al., 1988). The litter distribution of the affected pups was not reported by the authors. It is not possible to attribute this effect to microcystin exposure, as the extract may have contained other compounds.

In vitro studies suggest that MCLR can disrupt the cytoskeleton of embryonic cells, causing cell detachment, retarding division or causing cytolysis (Sepulveda et al., 1992; Frangez et al., 2003; Zuzek et al., 2003). MCLR effects on these and other cell types may be limited by the degree of uptake. Frangez et al. (2003) showed that an intact zona pellucida prevented effects in rabbit whole embryo cultures.

4.6.2. Inhalation

Very limited information is available on the toxicity of MCLR via inhalation exposure. The available data indicate that short-term inhalation of a low concentration of MCLR can cause local damage to the epithelial cells of the nasal cavity. A single study of inhalation exposure in

mice revealed dose-dependent damage to the respiratory and olfactory epithelial cells of the nasal cavity (Benson et al., 2005). Exposure occurred over 7 days at 260 $\mu g/m^3$ for 30, 60 and 120 minutes per day (authors estimated the deposited doses as 3, 6 and 12 $\mu g/kg$). No effects on the liver or other organs were observed.

Several limited lines of evidence suggest that high doses of MCLR via respiratory exposure routes can lead to systemic uptake with subsequent liver effects. Systemic uptake of MCLR by respiratory routes of exposure has been demonstrated in studies of acute, high-dose exposure (Creasia, 1990; Fitzgeorge et al., 1994; Ito et al., 2001). Importantly, the LD50 for MCLR given via either intranasal or intratracheal instillation is similar to that of MCLR given via i.p. injection (Fitzgeorge et al., 1994; Ito et al., 2001). As with i.p. and oral exposure, liver hemorrhage is the proximate cause of death in animals lethally dosed via intranasal or intratracheal instillation (Fitzgeorge et al., 1994; Ito et al., 2001). Further evidence of systemic effects comes from a brief abstract describing lethality in mice exposed via inhalation (nose only) to MCLR aerosols. Creasia (1990) reported an LC50 for MCLR of 18 mg/m³ air for 10 minutes (authors estimated the deposited dose as 45 μ g/kg), and indicated that histopathological findings in deceased mice were similar to those reported after i.v. dosing. Ito et al. (2001) suggested that MCLR could enter the bloodstream either via local damage to the nasal mucosa leading to exposure of the nasal blood vessels, or through transport to the lung and absorption into alveolar capillaries.

4.7. WEIGHT-OF-EVIDENCE EVALUATION AND CANCER CHARACTERIZATION

4.7.1. Summary of Overall Weight of Evidence

Applying the Guidelines for Carcinogen Risk Assessment (U.S. EPA, 2005a), there is inadequate information to assess carcinogenic potential of microcystins by the oral, dermal or inhalation routes of exposure. One poorly-described long-term carcinogenicity bioassay found no increase in neoplastic liver nodules after gavage administration in mice (Ito et al., 1997b). The few available epidemiological studies that suggest a positive association between liver or colorectal cancers and microcystins are limited by ecological study design, poor measures of exposure, potential coexposure to other microbial or chemical contaminants and, in some cases, failure to control for known liver and colorectal risk factors. MCLR has been shown to have a promotional effect in two-stage rat liver bioassays using i.p. administration; however, the relevance of this effect to environmental exposures is uncertain. Mechanistic information provides some support for a possible promotional effect of MCLR.

4.7.2. Synthesis of Human, Animal and Other Supporting Evidence

Several human epidemiological studies have reported an association between consumption of drinking water containing cyanobacteria and microcystins and liver or colon cancer in certain areas of China (Yu et al., 1989 and Yu, 1989 as cited in Ueno et al., 1996; Zhou et al., 2002). In all of these studies, the use of a surface drinking water supply was used as a surrogate for exposure to microcystins. Individual exposure to microcystins was not estimated.

Further, it is not clear whether these studies adequately controlled for confounding factors, such as hepatitis infection or aflatoxin exposure.

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Ito et al. (1997b) conducted the only study of oral carcinogenicity of a purified microcystin. In this study, chronic gavage doses of MCLR over 28 weeks failed to induce neoplastic nodules of the liver in mice. Limited information from two-stage, medium-term rat liver bioassays where MCLR was administered i.p. suggest that MCLR can act as a promoter, increasing the number and/or size of GST (placental form) positive foci in livers of rats pretreated with an initiating agent (Nishiwaki-Matsushima et al., 1992; Ohta et al., 1994; Sekijima et al., 1999; Hu et al., 2002). In one such study, MCLR alone showed no initiating activity (Ohta et al., 1994). Ito et al. (1997b) observed an increase in the size of neoplastic liver nodules in mice given 100 i.p. injections of MCLR without an initiating agent; however, the numbers of treated and control animals were small.

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Studies of cyanobacterial extract also suggest a possible promotional effect. In mice given an extract of *M. aeruginosa* in drinking water, the mean area of aberrant crypt foci of the colon was significantly increased, although the number of foci was not affected (Humpage et al., 2000). Similarly, the total weight of skin tumors was increased in mice given an extract of Microcystis in drinking water after topical DMBA pretreatment (Falconer and Buckley, 1989; Falconer, 1991). It is not possible to determine whether the observed effects resulted from exposure to microcystins or to other contaminants in the extracts.

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Mechanistic data indicate that at low doses, MCLR may increase cell proliferation. MCLR has been shown to increase the expression of the bcl-2 protein (that inhibits apoptosis) and decrease the expression of the bax protein (that induces apoptosis) (Hu et al., 2002). Further, MCLR upregulates the transcription factors c-fos and c-jun, leading to abnormal proliferation (Zhao and Zhu, 2003). Gehringer (2004) reviewed the molecular mechanisms leading to promotion by MCLR and the related tumor promoter, okadaic acid. Gehringer (2004) reported that MCLR inhibits protein phosphatase PP2A, which regulates several MAPKs. The MAPK cascade regulates transcription of genes required for cell proliferation, including c-iun and c-fos. In addition, activation of the MAPK cascade has been postulated to inhibit apoptosis and thus increase cell proliferation. Finally, Gehringer (2004) noted that MCLR has been reported to increase phosphorylation of p53, which is involved in the regulation of the cell cycle and apoptosis.

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Genotoxicity studies of MCLR have given conflicting results, with negative findings in Ames assays (Grabow et al., 1982; Ding et al., 1999) while positive results were observed with human cell lines (Suzuki et al., 1998; Zhan et al., 2004). Evidence for MCLR-induced DNA damage as measured by the comet assay has been called into question by the finding that apoptosis can lead to false positive findings in this assay (Lankoff et al., 2004). There is some evidence for a clastogenic effect of MCLR (Ding et al., 1999; Zhan et al., 2004).

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4.8. SUSCEPTIBLE POPULATIONS AND LIFE STAGES

Little information is available on potentially susceptible populations. Studies in laboratory rodents suggest that the acute effects of MCLR may be more pronounced in adult or aged animals than in juvenile animals (Adams et al., 1985; Ito et al., 1997a; Rao et al., 2005). In these studies, young animals showed little or no effect at MCLR doses lethal to adult animals. Age-dependent differences in toxicity were observed after both oral and i.p. exposure, suggesting that differences in gastrointestinal uptake were not entirely responsible for the effect of age. The relevance of these age-related differences to acute toxicity in humans is unclear.

Available information does not suggest any pronounced gender differences in response to microcystins. Studies with algal extracts suggest the possibility that male mice may be more sensitive than female mice to oral exposure to algal extracts (Falconer et al., 1988). However, the relevance of this finding to human microcystin exposure is uncertain given the potential for coexposure to other contaminants in algal extracts.

Because microcystins inhibit the action of protein phosphatases (PP1 and PP2A), coexposure to other compounds that inhibit these enzymes (for example, okadaic acid) may enhance the toxicological effects of microcystins.

5. DOSE RESPONSE ASSESSMENTS

5.1. NARRATIVE DESCRIPTION OF THE EXTENT OF THE DATABASE

The available information on the toxicokinetic behavior of microcystins in humans or animals after oral or inhalation exposure is limited to a single study of the organ distribution of dihydro-MCLR. No other data are available on the absorption, distribution, metabolism or elimination of microcystins via environmentally relevant exposure routes. Acute lethality data show a significant difference in lethal doses via injected and oral routes of exposure, suggesting that the toxicokinetic behavior of microcystins is an important determinant of health effects after oral exposure; thus, the deficiencies in this category of data are significant.

The mode by which microcystins affects its primary target organ, the liver, is remarkably well-studied. There are abundant mechanistic data ranging from target organ specificity down to molecular targets. *In vitro* studies using human hepatocytes show effects similar to those in animal hepatocytes, indicating that the mode of toxicological action is similar. Human hepatocytes appear to be more susceptible to the action of MCLR than rat hepatocytes (Yea et al., 2001; Batista et al., 2003). Further evidence for the relevance of this mode of toxicological action to humans comes from reports of human exposure. Liver histopathology on humans exposed to MCLR via dialysate showed effects similar to those seen in animals, although intrahepatic hemorrhage was not observed (Azevedo et al., 2002).

The toxicological database for microcystins is almost exclusively limited to data on a single congener, MCLR. Data on the other congeners is restricted to *in vitro* studies of protein phosphatase inhibition, i.p. LD₅₀ measures and a single, poorly-described toxicological evaluation of MCLA in primates. The database on the oral toxicity of MCLR is adequate to support the derivation of RfD values. Human data on the oral toxicity of MCLR are limited by potential co-exposure to other cyanobacterial toxins and microorganisms. There are three studies of acute oral exposure to MCLR in two laboratory animal species; however, none of these identified a NOAEL, and the minimum dose tested was the same for all three. Further, neither of the two experiments published in Fawell et al. (1999) nor the experiment by Yoshida et al. (1997) used an untreated control group. One animal study evaluated the oral toxicity of MCLR after short-term (<30 days) exposure and and one after subchronic (30-90 days) exposure. The short-term study (Heinze, 1999) used a small number of animals (10/dose) and did not identify a NOAEL, but was otherwise of good study quality. The subchronic study (Fawell et al., 1999) used an adequate number of animals (30/dose) and identified both a NOAEL and LOAEL. Two chronic exposure studies are available, but one of these (Ito et al., 1997b) apparently conducted only a limited examination of the liver for toxicity, and the other (Ueno et al., 1999) used a single dose and did not identify a LOAEL. A single, well-conducted developmental toxicity study in the mouse is available; however, the results are presented only briefly and without any supporting data in the publication (Fawell et al., 1999). The noncancer database is missing a chronic toxicity study in a second species, as well as a multigeneration reproductive toxicity study and neurotoxicity study.

The database on the inhalation toxicity of MCLR is inadequate for the derivation of any RfC. There are no human data on the inhalation of MCLR. There is a single well-reported animal study addressing inhalation exposure to MCLR for 7 days (Benson et al., 2005). This study used only one exposure concentration with daily exposure for 30, 60 or 120 minutes and, as such, is not adequate for short-term RfC derivation.

The available data on carcinogenicity are inadequate for carcinogenicity assessment. There is no well-conducted long-term carcinogenicity bioassay for microcystin. Several studies using an initiation-promotion protocol are available, as are limited mechanistic data suggesting a potential promoting effect of microcystins.

5.2. ORAL REFERENCE DOSE

Data considered in deriving oral reference dose for each exposure duration are summarized in the following exposure-response array (Figure 5-1) as well as in Table 5-1 below. Due to the limited toxicological database for microcystins, both the table and the figure include all studies in the published literature that examined the oral toxicity of purified MCLR in laboratory animals, with one exception. The publication by Ito et al. (1997b) did not provide the dosing frequency or information to estimate an average daily dose; thus, this study is not included in the table or figure.

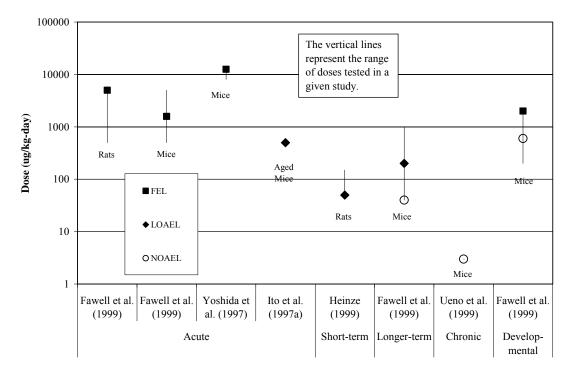


Figure 5-1. Exposure Response Array for Oral Exposure: All Studies of Purified Microcystin-LR

Table 5-1. Available Dose-Response Information for Oral Exposure to Purified MCLR							
Species	Sex	Average Daily Dose (μg/kg-day)	Exposure	NOAEL (μg/kg- day)	LOAEL or FEL (µg/kg-day)	Reference	
Acute Ex	posure						
Rat	M/F	500, 1580, 5000	Single gavage	ND	5000*	Fawell et al., 1999	
Mouse	M/F	500, 1580, 5000	Single gavage	ND	1580*	Fawell et al., 1999	
Mouse	F	8000, 10000, 12500	Single gavage	ND	12500*	Yoshida et al., 1997	
Mouse	M	0, 500	Single gavage	ND	500 (aged mice only)	Ito et al., 1997a	
Short-Tei	rm Expo	sure					
Rat	M	0, 50, 150	Drinking water, 28 d	ND	50	Heinze, 1999	
Subchron	ic Expo	sure					
Mouse	M/F	0, 40, 200, 1000	Daily gavage, 13 weeks	40	200	Fawell et al., 1999	
Chronic I	Exposure	e					
Mouse	F	0, 3	Drinking water, 18 months	3	ND	Ueno et al., 1999	
Developr	nental T	oxicity					
Mouse	F	0, 200, 600, 2000	Gavage, GD 6-15	600	2000*	Fawell et al., 1999	

^{*} Frank Effect Level (FEL)

5.2.1. Acute Oral RfD

The acute oral data for MCLR are inadequate for the derivation of an acute RfD. There are four studies of acute exposure to MCLR (Yoshida et al., 1997; Ito et al., 1997a; Fawell et al., 1999); however, none of the available studies identified a NOAEL. Yoshida et al. (1997) conducted a LD₅₀ determination using a small number of mice (5) treated with a single gavage dose, and no untreated controls. Histopathology of surviving mice (one at 8 mg/kg and two at 10 mg/kg) showed liver lesions (hypertrophic hepatocytes with centrilobular and midzonal fibrosis). The high dose in this study was an FEL based on deaths of the 2 treated animals. Fawell et al.

(1999) administered single gavage doses of 500, 1580 or 5000 μg/kg MCLR to groups of rats and mice (10/dose/species). There were no untreated control groups in this study. In both species, diffuse hepatic hemorrhage was observed at the low dose (500 μg/kg); however, in the absence of untreated controls for comparison, this dose cannot be identified as a LOAEL. Ito et al. (1997a) administered single gavage doses of 500 μg/kg to young (5 weeks old) and aged (32 weeks old) mice. Centrilobular hemorrhage and hepatocyte necrosis, as well as gastrointestinal lesions, were observed in the aged mice, while no effect was observed in the young mice. This study identifies a freestanding LOAEL. The absence of an acute study of adequate quality precludes derivation of an acute oral RfD.

5.2.2. Short-Term Oral RfD

5.2.2.1. Choice of Principal Study and Critical Effect

A single short-term study and a single developmental toxicity study of orally-administered MCLR are available. Heinze (1999) evaluated the effects of MCLR in drinking water in 11-week-old male hybrid rats. Groups of 10 rats (5 of each sex) were given approximate doses of 0, 50 or 150 μ g/kg body weight for 28 days. Serum biochemistry showed significantly increased mean levels of ALP and LDH in both treatment groups (84 and 100% increase in LDH, 34 and 33% increase in ALP in low and high doses, respectively). A dose-dependent increase in relative liver weights was observed at both dose levels (17 and 26% at the low and high doses, respectively). Liver lesions were observed in both treatment groups, but the severity of the damage was increased in the 150 μ g/kg dose group. Moderate to severe degenerative and necrotic hepatocytes with hemorrhage was observed in 0 of 10 controls, 6 of 10 low-dose and 9 of 10 high-dose rats.

Fawell et al. (1999) evaluated the developmental toxicity of MCLR administered via gavage to mice at doses of 0, 200, 600 and 2000 μ g/kg on GDs 6-15. Seven of 26 dams receiving 2000 μ g/kg died and two others were sacrificed prematurely due to morbidity. At this dose, fetal body weight was significantly lower than controls and there was delayed skeletal ossification; these effects may have been associated with maternal toxicity. Data on reproductive and developmental parameters were not provided in the reference; thus, a LOAEL for developmental toxicity could not be determined. This study identified a NOAEL of 600 μ g/kg-day for both developmental and maternal effects. The high dose of 2000 μ g/kg was an FEL based on maternal mortality. The study by Heinze (1999) identified a lower LOAEL and more sensitive effect (hepatotoxicity) than the developmental toxicity study did; thus, this study was chosen as the basis for the short-term RfD.

5.2.2.2. Methods of Analysis

Liver toxicity observed by Heinze (1999) included liver lesions, serum enzyme changes, and changes in relative liver weight. All three of these endpoints were considered for determining the point of departure for RfD derivation.

All quantal models in U.S. EPA's Benchmark Dose (BMD) software (version 1.3.2) were fit to the incidence data for liver lesions in rats (males and females combined) exposed to MCLR in the drinking water for 28 days (Heinze, 1999). The incidence data for liver lesions are reported in Table 4-4 (Section 4.2.1.2.1). Of the liver lesions observed, the category of degenerative and necrotic hepatocytes with hemorrhage showed a strong dose-related trend with greater incidence and severity with higher dose, and no control animals were affected. As the table shows, at 50 μ g/kg, 4/10 rats had slight lesions and 6/10 had lesions of moderate severity. At 150 μ g/kg, 6/10 rats had moderate lesions and 3/10 had intensive damage. For BMD modeling, the moderate and severe lesion categories were collapsed into one. The data modeled are shown in Table 5-2.

Table 5-2. Incidence of Liver Lesions Used for BMD Modeling (Heinze, 1999)						
	150 μg/kg-day					
Lesion incidence	0/10	6/10	9/10			

In accordance with the U.S. EPA (2000c) BMD methodology, the default benchmark response (BMR) of 10% increase in extra risk was used. The high response rate (60%) at the lowest dose with a positive response precludes the use of a lower BMR for this analysis. Models were run using the default restrictions on parameters built into the BMDS. The modeling results are shown in Table 5-3. Adequate fits were achieved with all models, except the quantal quadratic. While the gamma, multistage, quantal linear and Weibull all converged on the same model, the log probit model gave the best fit, as assessed by AIC. Figure 5-2 shows the fit of the log probit model to the data. Appendix A contains the full model outputs. The BMD and BMDL estimated by the log probit model for the liver lesion data are 11.0 and 6.4 μ g/kg-day, respectively.

The linear model for continuous data was fit to the increased LDH and ALP levels reported in Heinze (1999). These data are shown in Table 4-3 (Section 4.2.1.2.1). With only three observations in each of these datasets, there were not enough data points to use the remaining models (polynomial, power, or Hill), which each have more than three parameters.⁴

The linear model did not provide adequate fit to either dataset as measured by goodness-of-fit criteria (see Appendix A for model outputs). The linear model was also fit to the relative liver weight changes (also three observations) reported in Heinze (1999). These data are shown in Table 4-3 (Section 4.2.1.2.1). In accordance with the U.S. EPA methodology, the default BMR of one standard deviation change from the control mean was used, and the polynomial coefficients were restricted to be positive. The linear model provided an adequate fit to the data (see Appendix A for model output). The BMD and BMDL estimated by the linear model for the relative liver weight data are 85 and 58 μ g/kg-day, respectively.

⁴ The number of parameters describing the shape of the dose-response curve cannot exceed the number of dose groups (U.S. EPA, 2000c).

Table 5-3. BMD Modeling Results for Heinze (1999) Liver Lesion Data							
Model	Degrees of Freedom	χ^2	χ^2 Goodness of Fit p -Value	AIC	BMD (µg/kg- day)	BMDL (μg/kg-day)	
Log-probit (slope ≥1)	2	0.01	0.99	21.97	11.04	6.38	
Gamma (power ≥1)	2	0.09	0.96	22.05	6.31	3.92	
Multistage (degree=1)*	2	0.09	0.96	22.05	6.31	3.92	
Quantal Linear	2	0.09	0.96	22.05	6.31	3.92	
Weibull (power ≥1)	2	0.09	0.96	22.05	6.31	3.92	
Log-logistic (slope ≥1)	1	0.00	1	23.96	10.14	1.24	
Quantal Quadratic	2	5.77	0.06	26.17	24.81	19.01	
Logistic	1	3.43	0.06	28.31	19.43	11.40	
Probit	1	3.50	0.06	28.43	19.69	12.31	

 ^{*} Degree of polynomial initially set to (n-1) where n= number of dose groups including control;
 model selected is lowest degree model providing adequate fit. Betas restricted to ≥0.

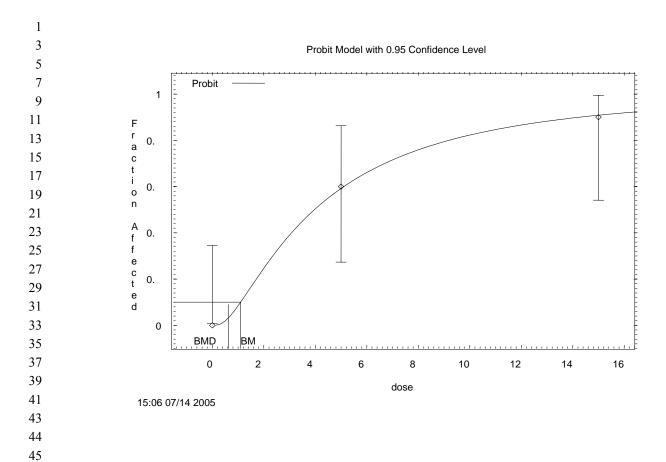


Figure 5-2. Probit Model Fit to Liver Lesion Incidence Data from Heinze (1999)

5.2.2.3. RfD Derivation

The BMDL of 6 μ g/kg-day from the Heinze (1999) data is the lowest BMDL among the modeled datasets and was used as the point of departure (POD) for the short-term RfD. Dividing the BMDL of 6 μ g/kg-day by a composite uncertainty factor (UF) of 1000 results in a short-term RfD for MCLR of 6×10^{-6} mg/kg-day.

Short-term RfD = BMDL ÷ UF = $6 \mu g/kg$ -day ÷ 1000 = **0.000006 mg/kg-day or 6x10**-6 mg/kg-day

The composite UF of 1000 includes a factor of 10 for interspecies extrapolation, a factor of 10 to account for interindividual variability in the human population and a factor of 10 for database limitations, as follows.

- A default 10-fold UF for intraspecies differences was used to account for potentially susceptible individuals in the human population. There is insufficient information on the toxicity of microcystins in exposed humans. Cases of human poisoning have been attributed to ingestion of water containing microcystin-producing cyanobacteria, but no dose-response information is available. There is no information on the degree to which humans of varying gender, age, health status or genetic makeup might vary in the disposition of, or response to, ingested microcystins. There are some data to suggest that adult or aged rodents may be more susceptible than young rodents to the acute toxicity of MCLR (Ito et al., 1997a). Further, studies with algal extracts suggest the possibility that male mice may be more sensitive than female mice to oral exposure to algal extracts (Falconer et al., 1988); however, the relevance of this finding to human microcystin exposure is unclear.
- An interspecies UF of 10 was used to account for differences in response between laboratory rodents and humans. No information is available on the toxicity of purified microcystins in humans, and data on toxicokinetic differences between animals and humans in the disposition of ingested microcystins are not available. Limited data from *in vitro* studies suggests that human hepatocytes may be more susceptible to the effects of MCLR than rat hepatocytes (Yea et al., 2001; Batista et al., 2003), supporting the use of a full 10-fold UF.
 - A 10-fold UF is used to account for deficiencies in the database. Database deficiencies include the lack of a detailed developmental toxicity study, a neurotoxicity study, a multi-generation reproductive toxicity study and supporting information on systemic toxicity in a second species.

5.2.3. Subchronic Oral RfD

5.2.3.1. Choice of Principal Study and Critical Effect

A single subchronic oral toxicity study is available for MCLR. Fawell et al. (1999) identified both a NOAEL (40 μ g/kg-day) and a LOAEL (200 μ g/kg-day for slight liver injury) after subchronic exposure of male and female mice. Fawell et al. (1999) administered daily oral gavage doses of 0, 40, 200 or 1000 μ g MCLR per kg body weight to groups of 15 male and 15 female mice for 13 weeks. Histopathological changes in the liver and serum enzyme changes were reported in the mid- and high-dose groups. Both the histopathology and the serum enzymes showed dose-dependent changes. The authors considered the liver changes in the 200 μ g/kg-day dose group to represent a minimal effect.

The NOAEL from this 90-day study (40 µg/kg-day) is only slightly lower than the LOAEL identified in the 28-day study above (50 µg/kg-day). Further, Heinze (1999) observed more severe effects in rats exposed to 50 µg/kg-day for 28 days in drinking water than Fawell et al. (1999) observed in mice exposed via gavage doses of 200 µg/kg-day for 90 days. The reason for this difference in response is not clear. Both studies used a commercially-produced test material from the same manufacturer. In the drinking water study, MCLR was dissolved in ethanol and diluted to a stock solution that was subsequently used to prepare drinking water. It is possible that the small intake of ethanol may have potentiated the hepatic effects of MCLR; however, there are no data to determine whether this is likely or not. In the gavage study, test solutions were prepared with distilled water and the concentration was confirmed by HPLC with UV detection (Fawell et al., 1999). The accuracy of dosing in the gavage study was likely to be greater than in the drinking water study. The authors of the drinking water study indicated that the MCLR solution was prepared daily, and water consumption was measured daily. Between 3 and 7% of the water solution administered over the 28 days was not consumed, and the dose estimates were not corrected for this loss (Heinze, 1999). This loss of administered dose would lead to a small overestimate of the LOAEL in the drinking water study, leading to a further discrepancy in the results of the two studies.

The drinking water study used smaller group sizes (10 males/dose) than the gavage study (15/sex/dose or 30/dose). However, the incidence of liver lesions (with necrosis and hemorrhage) in the drinking water study increased from 0% to 100% (including slight, moderate, and intensive lesions) between the control and low-dose group, and there was a dose-related change in the severity of the lesions, leaving little question that the effect was treatment-related.

These studies appear to contradict evidence from acute parenteral studies indicating that mice are more sensitive to the acute effects of MCLR. Typically, mice die within a few hours of a lethal injected dose, while rats may survive 24-48 hours. Species-specific differences in oral absorption of MCLR do not appear to account for the discrepancy between these studies; in an acute study of orally-administered MCLR using both mice and rats, mortality occurred at a lower dose in mice (1580 $\mu g/kg$) than in rats (5000 $\mu g/kg$; Fawell et al., 1999). It is possible that the more mild effects in the mice in the subchronic study resulted from an adaptive response to MCLR exposure. The longer exposure duration may have allowed for liver regeneration and

repair that was not possible in the shorter-term study; however, there is no information to support this hypothesis. In fact, Ito et al. (1997) reported that light injuries to hepatocytes were still evident in five of seven mice 2 months after treatment with MCLR had ceased.

5.2.3.2 Methods of Analysis

The data from both Heinze (1999) and Fawell et al. (1999) were considered for identifying the point of departure for the subchronic RfD derivation. Results of the modeling for Heinze (1999) are reported in Section 5.2.2.2. Among the liver lesions reported by Fawell et al. (1999), only chronic inflammation showed evidence of a dose-response relationship. Consequently, all quantal models in U.S. EPA's Benchmark Dose Software (BMDS) were fit to the incidence data for chronic liver inflammation in male and female mice reported by Fawell et al. (1999). The incidence data are reported in Table 4-6. In accordance with the U.S. EPA methodology, the default BMR of 10% increase in extra risk was used. Models were run using the default restrictions on parameters built into the BMDS. Adequate fits were achieved with all models. For both male and female mice, the probit model provided the best fit, as assessed by AIC. Table 5-4 gives the results for the best fit models; Appendix A contains the full outputs for all models.

Table 5-4. BMD Modeling Results for Fawell et al. (1999) Chronic Liver Inflammation Data

	Degrees of Freedom	X^2	χ^2 Goodne ss of Fit <i>p</i> -Value	AIC	BMD (µg/kg- day)	BMDL (µg/kg- day)
Male: Probit Model	2	0.21	0.90	40.75	107.59	66.45
Female: Probit Model	2	0.94	0.63	72.84	86.34	56.92

Fawell et al. (1999) also reported significant increases in ALP, AST, and ALT in high-dose animals. These data are shown in Table 4-5 (Section 4.2.1.3.1). Of these, only the increase in ALT in male mice showed a dose-response trend amenable to modeling. All continuous models but the Hill model⁵ were fit to the ALT data for male mice reported in Fawell et al. (1999). In accordance with the U.S. EPA methodology, the default BMR of one standard deviation change from the control mean was used, and the polynomial coefficients were restricted to be positive. Only the linear model with a nonhomogenous variance provided an adequate fit to the data. Table 5-5 gives the results from the linear model (see Appendix A for model output). The BMD and BMDL estimated by the linear model (nonhomogenous variance) for the ALT increases in male mice are 82 and 58 μg/kg-day, respectively.

⁵ There were too few dose groups to apply the Hill model.

Table 5	Table 5-5. BMD Modeling Results for Fawell et al. (1999) ALT Data in Male Mice								
	Degrees of Freedom	Goodness of Fit p-Value	AIC	BMD (µg/kg- day)	BMDL (μg/kg-day)				
Linear model, non- homogenous variance	2	0.10	447.68	81.84	58.37				

The BMDL from the 28-day drinking water study (6 μ g/kg-day) is approximately an order of magnitude lower than any of the BMDL values from the 90-day gavage study (57-66 μ g/kg-day). Details of the BMD modeling and analysis of the data from Heinze (1999) are provided above in Section 5.2.2.2, Method of Analysis, under Short-Term Oral RfD.

5.2.3.3. RfD Derivation

The BMDL of 6 μ g/kg-day from the Heinze (1999) data is used as the POD for the subchronic RfD. A composite UF of 1000 is used to derive the subchronic RfD, including a factor of 10 for interspecies extrapolation, a factor of 10 for interindividual variability, and a factor of 10 for database limitations (see Section 5.2.2.3 above for details). Although the BMDL comes from a 28-day study, a UF for exposure duration is not proposed, based on the lower toxicity observed in the 90-day gavage study conducted by Fawell et al. (1999). The subchronic RfD is, therefore, set equal to the short-term RfD of 0.006 μ g/kg-day or $6x10^{-6}$ mg/kg-day.

Subchronic RfD = BMDL \div UF = $6 \mu g/kg-day \div 1000$ = $0.006 \mu g/kg-day$ or $6x10^{-6} mg/kg-day$

5.2.4. Chronic Oral RfD

5.2.4.1. Choice of Principal Study and Critical Effect

 Two chronic studies of the oral toxicity of MCLR were identified. Ito et al. (1997b) conducted a chronic gavage study in mice with 80 to 200 doses (80 μ g/kg-day) given over 28 weeks; however, the control group was very small (3 animals) and the postmortem examination was apparently limited to the liver. This study was not given further consideration for RfD development given these study quality concerns.

Ueno et al. (1999) evaluated the toxicity of MCLR in female mice chronically exposed via drinking water 7 days/week. The authors conducted a comprehensive postmortem examination. No treatment-related effects were identified, and the authors observed no

difference in the incidence of liver histopathology between treated and control mice. It is important to note that immunohistochemistry of the liver revealed no accumulation of MCLR. This study identified a free-standing NOAEL of approximately 3 µg/kg-day in female mice.

Although Ueno et al. (1999) used only a single dose level and identified a freestanding NOAEL, it was chosen for RfD derivation because it was a well-conducted chronic study using a relevant exposure route (drinking water). The BMDL of 6 μ g/kg-day from modeling (see Section 5.2.2.2) of Heinze (1999) compares favorably with the free-standing NOAEL of 3 μ g/kg-day reported by Ueno et al. (1996), providing support for the use of the NOAEL from Ueno et al. (1999).

5.2.4.2. RfD Derivation

As noted above, the NOAEL of 3 μ g/kg-day from the study by Ueno et al. (1999) is used as the POD for the chronic RfD. Because this study used a single dose, it was not possible to use BMD modeling to identify the POD. A composite UF of 1000 is used to derive the chronic RfD, including a factor of 10 for interspecies extrapolation, a factor of 10 for interindividual variability and a factor of 10 for database limitations (see Section 5.2.2.3 above for details). Dividing the NOAEL of 3 μ g/kg-day by a composite UF of 1000 results in a chronic RfD for MCLR of 3×10^{-6} mg/kg-day.

```
Chronic RfD = NOAEL \div UF
= 3 \mug/kg-day \div 1000
= 0.003 \mug/kg-day or 3x10<sup>-6</sup> mg/kg-day
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In 1999, the WHO published a provisional Tolerable Daily Intake (TDI) for MCLR based on the subchronic gavage study later published by Fawell et al. (1999). The WHO used the NOAEL of 40 μg/kg-day with a composite UF of 1000 to derive a TDI of 0.04 μg/kg-day or 4×10^{-5} mg/kg-day. The composite uncertainty factor included UFs of 10-fold each for interindividual variability, interspecies extrapolation, and database deficiencies (WHO specifically cited the lack of chronic toxicity and carcinogenicity studies). WHO (1999) did not evaluate either Ueno et al. (1999) or Heinze (1999), which may not have been published at the time.

Table 5-6 provides a summary of the RfD values derived for MCLR in this report.

5.3. INHALATION REFERENCE CONCENTRATION

The available data do not provide adequate information for the derivation of inhalation RfCs for MCLR. Two acute inhalation studies were identified in the literature. In a poorly described study, Fitzgeorge et al. (1994) conducted a single experiment with mice (number unspecified) inhaling a fine aerosol (particle size 3-5 μ m) with 50 μ g/L MCLR for an unspecified duration of time. There were apparently no deaths, clinical signs of toxicity or histopathological changes; however, the authors gave few details of study design and findings. A brief abstract describes a study of acute microcystin exposure via inhalation (Creasia, 1990). The LC₅₀ for

Table 5-6. Summary of Reference Dose Values							
	RfD (mg/kg-day) Critical Effect Principal Study						
Acute	NA						
Short-term	6 x 10 ⁻⁶	Hepatotoxicity	Heinze, 1999				
Subchronic 6 x 10 ⁻⁶ Hepatotoxicity Heinze, 1999							
Chronic	3 x 10 ⁻⁶	No effects observed	Ueno et al., 1999				

mice exposed to a MCLR aerosol (nose only) for 10 minutes was reported to be 18 μ g/L (mg/m³) air with a 95% confidence interval of 15.0-22.0 μ g/L (mg/m³). The authors reported that histological lesions in mice killed by aerosol exposure were similar to those in mice dosed intravenously with MCLR. Neither of these studies provides adequate basis for an acute RfC.

Only one well-conducted study of inhalation exposure to MCLR was identified. Benson et al. (2005) exposed groups of six male BALB/c mice to monodisperse submicron aerosols of MCLR via nose-only inhalation for 30, 60 or 120 minutes each day for 7 consecutive days. The concentration of MCLR was 260-265 μ g/m³. Histopathological examination revealed treatment-related lesions in the nasal cavity only. The incidence and severity of nasal lesions increased with daily exposure duration. This study used only one exposure concentration, and as such, the data are of limited utility for RfC derivation. Further, extrapolation of the effects from this study for the purpose of deriving a short-term RfC would be associated with substantial uncertainty given the brief exposure time (30-120 minutes/day) and duration (7 days). There are no subchronic or chronic animal studies evaluating the inhalation route of exposure.

Route-to-route extrapolation is not considered appropriate for microcystins based on current data. Limited available information indicates that inhalation exposure to microcystins may cause point-of-entry effects (Benson et al., 2005), while oral exposure leads to hepatotoxicity. Data from intratracheal and intranasal instillation studies show hepatic effects after exposure via these routes; however, the relevance of this information to inhalation exposures is uncertain.

5.4. CANCER ASSESSMENT

No dose-response or other information is available regarding the carcinogenicity of pure microcystins.

6. MAJOR CONCLUSIONS IN THE CHARACTERIZATIONS OF HAZARD AND DOSE-RESPONSE

6.1. HUMAN HAZARD POTENTIAL

Microcystins are a group of naturally occurring hepatotoxins produced by freshwater cyanobacteria. No studies of the absorption, distribution, metabolism or elimination of microcystins LR, YR, RR or LA *in vivo* have been conducted. Acute lethality data suggest a significant difference in the lethal dose after oral exposure to MCLR when compared with injection routes of exposure, suggesting low absorption of orally-administered MCLR. The cellular uptake and distribution of MCLR has been extensively studied, showing preferential uptake of MCLR by hepatocytes due to the presence of a bile acid transporter.

Data on human exposures to microcystin-producing cyanobacteria have shown gastrointestinal and dermal effects; however, it is not clear whether these are effects of microcystins, other endotoxins, or the microorganisms themselves. The preponderance of the available toxicological studies in animals employed the MCLR congener. Both oral and parenteral exposure studies in laboratory animals point to the liver as the primary target organ of MCLR, and parenteral exposure studies suggest a steep dose-response curve for the hepatotoxic effects. The toxicological database is limited to only a few studies using oral exposure to purified microcystins; all of these used the MCLR congener, and most were conducted in mice. The database includes four studies of acute exposure, one 28-day and one 90-day study, and two chronic studies. In all of the studies where a toxicological effect was observed, the primary target organ was the liver (one study also identified gastrointestinal lesions). The liver effects included hepatocyte degeneration and necrosis, inflammation, fibrosis, hypertrophy and hemorrhage. Humans exposed to microcystins via dialysate suffered acute liver failure and, in many cases, death. Liver biopsies conducted on the decedents showed hepatocyte necrosis and apoptosis, but no intrahepatic hemorrhage.

A single well-described study of short-term (7 days) inhalation exposure identified the upper respiratory tract as a target organ. Damage to the respiratory and olfactory epithelial cells of the nasal cavity was observed in mice in this study. Studies using intratracheal or intranasal instillation have shown systemic effects, including liver toxicity, after these exposures.

A single oral developmental toxicity study with inadequate data reporting indicated that developmental effects in the absence of maternal toxicity are not likely, and that developmental effects, if any, would occur at much higher doses than liver effects.

Mechanistic studies indicate that the inhibition of protein phosphatases 1 and 2A (a well-established molecular effect of microcystin exposure) and/or oxidative stress play a role in the hepatotoxicity of MCLR. Cellular effects in hepatocytes include membrane blebbing, cell rounding and dissociation, and apoptosis. These cellular effects lead to alterations in the liver structure and function as well as intrahepatic hemorrhage.

6.2. DOSE RESPONSE

The available oral data were sufficient for derivation of short-term, subchronic and chronic oral RfDs for MCLR. Based on a BMDL of 6 μ g/kg-day for hepatotoxicity in rats exposed to MCLR in drinking water for 28 days (Heinze, 1999), an RfD of 0.006 μ g/kg-day (6x10⁻⁶ mg/kg-day) was derived for short-term and subchronic exposure durations. A UF of 1000 was used to derive the RfD. The UF comprises component factors of 10 for interspecies extrapolation, 10 for interindividual variability and 10 for database deficiencies. The subchronic RfD did not include an additional UF for extrapolating from a 28-day study because a subchronic (90-day) gavage study identified a higher NOAEL. A chronic RfD of 0.003 μ g/kg-day (3x10⁻⁶ mg/kg-day) was derived from a free-standing NOAEL of 3 μ g/kg-day in female mice chronically exposed via drinking water. A composite UF of 1000 was used, with factors of 10 each for interindividual variability, interspecies extrapolation and database deficiencies. Inhalation RfCs were not derived since there were no studies of adequate quality for this purpose. There is inadequate evidence to evaluate the carcinogenicity of microcystins LR, RR, YR and LA.

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1	APPENDIX A
2	
3	
4	Contents: Benchmark Dose Modeling Output Files for:
5	
6	1) Heinze, 1999 Liver Lesions
7	2) Fawell et al., 1999 Male Chronic Liver Inflammation
8	3) Fawell et al., 1999 Female Chronic Liver Inflammation
9	4) Fawell et al., 1999 Male and Female (combined) Chronic Liver Inflammation
10	5) Heinze, 1999 Relative Liver Weight Changes
11	6) Heinze, 1999 Lactate Dehydrogenase Changes
12	7) Heinze, 1999 Alkaline Phosphatase Changes
13	8) Fawell et al. 1999 Male Alanine Aminotransferase Changes

```
1
    Heinze,1999. Liver Lesions
2
     ______
             $Revision: 2.2 $ $Date: 2001/03/14 01:17:00 $
Input Data File: C:\BMDS\HEINZE_MOD_AND_INT_WITH_HEMORRHAGE.(d)
3
4
5
6
7
             Gnuplot Plotting File: C:\BMDS\HEINZE MOD AND INT WITH HEMORRHAGE.plt
                                                 Thu \overline{J}ul \overline{14} 1\overline{5}:03:08 2\overline{0}05
      ______
8
9
      BMDS MODEL RUN
10
11
12
        The form of the probability function is:
13
14
        P[response] = background+(1-background)*CumGamma[slope*dose,power],
15
        where CumGamma(.) is the cummulative Gamma distribution function
16
17
18
        Dependent variable = COLUMN2
19
        Independent variable = COLUMN1
20
        Power parameter is restricted as power >=1
21
22
23
        Total number of observations = 3
        Total number of records with missing values = 0
24
25
26
27
28
29
        Maximum number of iterations = 250
        Relative Function Convergence has been set to: 1e-008
        Parameter Convergence has been set to: 1e-008
30
                       Default Initial (and Specified) Parameter Values
31
                           Background =
                                           0.0454545
32
                                Slope =
                                           0.0153804
33
                                Power =
                                             1.02976
34
35
36
37
                Asymptotic Correlation Matrix of Parameter Estimates
38
                ( *** The model parameter(s) -Background
39
                      have been estimated at a boundary point, or have been
40
    specified by the user,
41
                      and do not appear in the correlation matrix )
42
43
                       Slope
44
45
          Slope
                            1
46
47
48
49
                                Parameter Estimates
50
                                                      Std. Err.
51
            Variable
                                Estimate
52
          Background
                                        0
                                                        NA
53
               Slope
                                0.0166997
                                                    0.00500499
54
               Power
55
56
    NA - Indicates that this parameter has hit a bound
57
          implied by some inequality constraint and thus
58
          has no standard error.
59
60
61
62
                              Analysis of Deviance Table
63
64
            Model
                       Log(likelihood) Deviance Test DF
                                                                P-value
```

Full model -9.98095
Fitted model -10.0255 0.089063 2 0.9564
Reduced model -20.7944 21.6269 2 <.0001

AIC: 22.051

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000	0.0000	0.000	0	10	0
50.0000	0.5661	5.661	6	10	0.2162
150.0000	0.9183	9.183	9	10	-0.2115
Chi-square =	0.09	DF = 2	P-value	= 0.9553	

Benchmark Dose Computation

Specified effect = 0.1

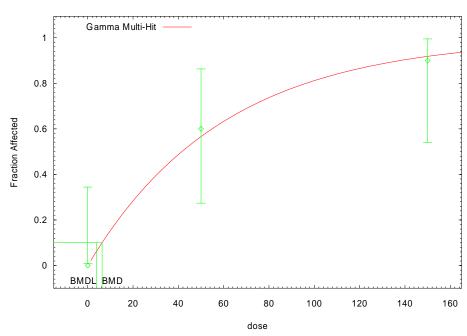
Risk Type = Extra risk

Confidence level = 0.95

BMD = 6.30914

BMDL = 3.92229

Gamma Multi-Hit Model with 0.95 Confidence Level



15:03 07/14 2005

```
1
2
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4
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6
7
    ______
            Logistic Model $Revision: 2.1 $ $Date: 2000/02/26 03:38:20 $
             Input Data File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY DOCUMENTS\ CYANO
    TOX REV\HEINZE LIVER LESIONS.(d)
    Gnuplot Plotting File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY DOCUMENTS\_CYANO TOX REV\HEINZE_LIVER_LESIONS.plt
 8
                                               Thu Mar 09 11:41:20 2006
9
      ______
10
11
     BMDS MODEL RUN
12
13
14
       The form of the probability function is:
15
16
       P[response] = 1/[1+EXP(-intercept-slope*dose)]
17
18
19
       Dependent variable = COLUMN2
20
       Independent variable = COLUMN1
21
22
23
24
25
26
       Slope parameter is not restricted
       Total number of observations = 3
       Total number of records with missing values = 0
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
27
28
29
30
       Parameter Convergence has been set to: 1e-008
31
                       Default Initial Parameter Values
32
                          background =
                                                 Ω
                                                      Specified
33
                           intercept =
                                          -2.28075
34
                                         0.0300564
                               slope =
35
36
37
               Asymptotic Correlation Matrix of Parameter Estimates
38
39
                ( *** The model parameter(s) -background
40
                     have been estimated at a boundary point, or have been
41
     specified by the user,
42
                      and do not appear in the correlation matrix )
43
44
                   intercept
                                    slope
45
46
     intercept
                          1
                                   -0.75
47
48
         slope
                      -0.75
49
50
51
52
53
                               Parameter Estimates
54
           Variable
                              Estimate
                                                    Std. Err.
55
                                                   0.772069
                               -2.02314
          intercept
56
57
58
                                                   0.0123743
              slope
                               0.0344016
59
60
                            Analysis of Deviance Table
61
62
           Model
                      Log(likelihood) Deviance Test DF
                                                              P-value
63
         Full model
                           -9.98095
64
       Fitted model
                            -12.1529
                                          4.34392 1
                                                                  0.03714
```

Reduced model -20.7944 21.6269 2 <.0001

AIC: 28.3058

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000	0.1168	1.168	0	10	-1.15
50.0000	0.4248	4.248	6	10	1.121
150.0000	0.9584	9.584	9	10	-0.9248

Chi-square = 3.43 DF = 1 P-value = 0.0639

Benchmark Dose Computation

Specified effect = 0.1

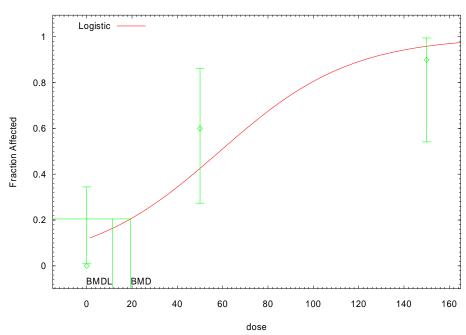
Risk Type = Extra risk

Confidence level = 0.95

BMD = 19.4327

BMDL = 11.4032

Logistic Model with 0.95 Confidence Level



11:41 03/09 2006

```
______
           Logistic Model $Revision: 2.1 $ $Date: 2000/02/26 03:38:20 $
2
3
4
5
           Input Data File: C:\BMDS\HEINZE MOD AND INT WITH HEMORRHAGE.(d)
           Thu Jul 14 15:04:46 2005
6
7
     ______
8
     BMDS MODEL RUN
9
    10
11
       The form of the probability function is:
12
13
       P[response] = background+(1-background)/[1+EXP(-intercept-slope*Log(dose))]
14
15
16
       Dependent variable = COLUMN2
17
       Independent variable = COLUMN1
18
       Slope parameter is restricted as slope >= 1
19
20
       Total number of observations = 3
21
22
23
24
25
26
27
28
29
30
       Total number of records with missing values = 0
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
       Parameter Convergence has been set to: 1e-008
      User has chosen the log transformed model
31
                    Default Initial Parameter Values
32
                       background =
                                             0
33
34
                        intercept =
                                      -5.97477
                            slope =
                                       1.63093
35
36
37
              Asymptotic Correlation Matrix of Parameter Estimates
38
39
              ( *** The model parameter(s) -background
40
                   have been estimated at a boundary point, or have been
41
    specified by the user,
42
                   and do not appear in the correlation matrix )
43
44
                 intercept
                                slope
45
46
     intercept
                        1
                                -0.99
47
48
                    -0.99
        slope
49
50
51
52
53
                           Parameter Estimates
54
          Variable
                                               Std. Err.
                           Estimate
55
        background
                                   0
                                                 NA
56
         intercept
                            -5.97477
                                                4.77026
57
58
             slope
                             1.63093
59
    NA - Indicates that this parameter has hit a bound
60
         implied by some inequality constraint and thus
61
        has no standard error.
62
63
     Warning: Likelihood for the fitted model larger than the Likelihood for the
64
    full model.
```

1
2
4 5
6
8
10
12
13 14
15 16
17 18
19
21
23
25
26 27
28 29
30 31
32 33
34 35
1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37
51

Analysis of Deviance Table

Model Log(likelihood) Deviance Test DF P-value

Full model

-9.98095 -9.98095 -3.55271e-015 1 -1 -20.7944 21.6269 2 <.0001 Fitted model Reduced model

AIC: 23.9619

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000	0.0000	0.000	0	10	0
50.0000	0.6000	6.000	6	10	-3.44e-015
150.0000	0.9000	9.000	9	10	-1.123e-014
Chi-square =	0.00	DF = 1	P-value	= 1.0000	

Benchmark Dose Computation

Specified effect = 0.1

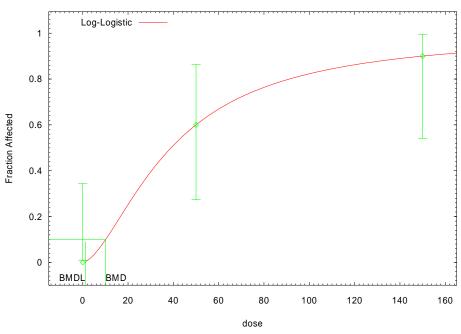
Risk Type Extra risk

Confidence level = 0.95

> BMD = 10.137

BMDL = 1.23783

Log-Logistic Model with 0.95 Confidence Level



15:04 07/14 2005

```
______
2
3
4
5
            Multistage Model. $Revision: 2.1 $ $Date: 2000/08/21 03:38:21 $
             Input Data File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY DOCUMENTS\ CYANO
    TOX REV\HEINZE LIVER LESIONS.(d)
            Gnuplot Plotting File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY
6
7
    DOCUMENTS\ CYANO TOX REV\HEINZE LIVER LESIONS.plt
                                               Thu Mar 09 11:37:17 2006
8
     ______
9
10
     BMDS MODEL RUN
11
12
13
       The form of the probability function is:
14
15
        P[response] = background + (1-background) * [1-EXP(
    -beta1*dose^1-beta2*dose^2)]
16
17
18
        The parameter betas are restricted to be positive
19
20
21
       Dependent variable = COLUMN2
\overline{22}
        Independent variable = COLUMN1
23
24
25
26
     Total number of observations = 3
     Total number of records with missing values = 0
     Total number of parameters in model = 3
Total number of specified parameters = 0
27
28
     Degree of polynomial = 2
29
30
31
     Maximum number of iterations = 250
32
     Relative Function Convergence has been set to: 1e-008
33
     Parameter Convergence has been set to: 1e-008
34
35
36
37
                       Default Initial Parameter Values
38
                          Background = 0.0617654
39
                             Beta(1) =
                                           0.015138
40
                             Beta(2) =
                                                  0
41
42
43
               Asymptotic Correlation Matrix of Parameter Estimates
44
45
      ( *** The model parameter(s) -Background
                                                   -Beta(2)
46
     have been estimated at a boundary point, or have been specified by the
47
    user, and do not appear in the correlation matrix )
48
49
                    Beta(1)
50
51
       Beta(1)
                           1
52
53
54
55
                               Parameter Estimates
56
57
           Variable
                                                    Std. Err.
                               Estimate
58
         Background
                                                       NA
59
            Beta(1)
                               0.0166997
                                                  0.00582148
60
            Beta(2)
61
62
    NA - Indicates that this parameter has hit a bound
63
          implied by some inequality constraint and thus
64
         has no standard error.
```

Analysis of Deviance Table

Model	Log(likelihood)	Deviance	Test DF	P-value
Full model	-9.98095			
Fitted model	-10.0255	0.089063	3 2	0.9564
Reduced model	-20.7944	21.6269	9 2	<.0001

AIC: 22.051

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Chi^2 Res.
i: 1 0.0000 i: 2	0.0000	0.000	0	10	-1.000
50.0000 i: 3	0.5661	5.661	6	10	0.138
150.0000	0.9183	9.183	9	10	-0.244
Chi-square =	0.09	DF = 1	P-value	= 0.7623	

Benchmark Dose Computation

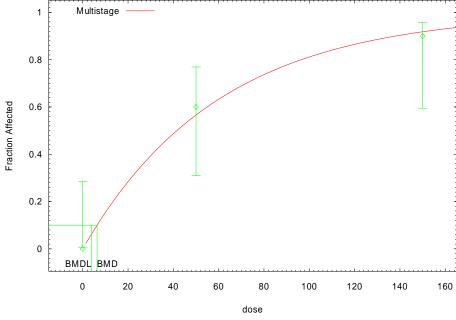
Specified effect = 0.1

Risk Type = Extra risk

Confidence level = 0.95

BMD = 6.30914

BMDL = 3.92229



11:37 03/09 2006

```
______
2
           Multistage Model. $Revision: 2.1 $ $Date: 2000/08/21 03:38:21 $
           Input Data File: C:\BMDS\HEINZE MOD AND INT WITH HEMORRHAGE.(d)
           4
5
                                           Thu Jul 14 15:05:31 2005
6
7
     ______
8
     BMDS MODEL RUN
9
    10
11
       The form of the probability function is:
12
13
       P[response] = background + (1-background) * [1-EXP(
14
    -beta1*dose^1)]
15
16
       The parameter betas are restricted to be positive
17
18
19
       Dependent variable = COLUMN2
20
       Independent variable = COLUMN1
21
\overline{22}
     Total number of observations = 3
23
     Total number of records with missing values = 0
24
25
26
     Total number of parameters in model = 2
Total number of specified parameters = 0
     Degree of polynomial = 1
27
28
29
     Maximum number of iterations = 250
30
     Relative Function Convergence has been set to: 1e-008
31
     Parameter Convergence has been set to: 1e-008
32
33
34
35
                     Default Initial Parameter Values
36
                       Background =
                                     0.0617654
37
                          Beta(1) =
                                       0.015138
38
39
40
              Asymptotic Correlation Matrix of Parameter Estimates
41
42
              ( *** The model parameter(s) -Background
43
                    have been estimated at a boundary point, or have been
    specified by the user,
44
45
                    and do not appear in the correlation matrix )
46
47
                   Beta(1)
48
49
       Beta(1)
50
51
52
53
                            Parameter Estimates
54
55
          Variable
                                               Std. Err.
                            Estimate
56
        Background
                                                  NΑ
57
58
           Beta(1)
                            0.0166997
                                             0.00582148
59
    NA - Indicates that this parameter has hit a bound
60
         implied by some inequality constraint and thus
61
        has no standard error.
62
```

Analysis of Deviance Table

Model	Log(likelihood)	Deviance	Test DF	P-value
Full model	-9.98095			
Fitted model	-10.0255	0.089063	3 2	0.9564
Reduced model	-20.7944	21.6269	2	<.0001

AIC: 22.051

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Chi^2 Res.
i: 1 0.0000 i: 2	0.0000	0.000	0	10	0.000
50.0000 i: 3	0.5661	5.661	6	10	0.138
150.0000	0.9183	9.183	9	10	-0.244
Chi-square =	0.09	DF = 2	P-value	= 0.9553	

Benchmark Dose Computation

Specified effect = 0.1

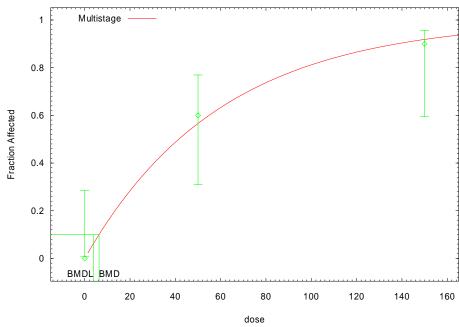
Risk Type = Extra risk

Confidence level = 0.95

BMD = 6.30914

BMDL = 3.92229

Multistage Model with 0.95 Confidence Level



15:05 07/14 2005

```
1
2
3
4
5
6
7
    ______
             Probit Model $Revision: 2.1 $ $Date: 2000/02/26 03:38:53 $
             Input Data File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY DOCUMENTS\ CYANO
    TOX REV\HEINZE LIVER LESIONS.(d)
    Gnuplot Plotting File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY DOCUMENTS\_CYANO TOX REV\HEINZE_LIVER_LESIONS.plt
8
                                               Thu Mar 09 11:49:47 2006
9
     ______
10
11
     BMDS MODEL RUN
12
13
14
       The form of the probability function is:
15
16
       P[response] = CumNorm(Intercept+Slope*Dose),
17
18
       where CumNorm(.) is the cumulative normal distribution function
19
20
21
22
23
24
25
26
       Dependent variable = COLUMN2
       Independent variable = COLUMN1
       Slope parameter is not restricted
       Total number of observations = 3
       Total number of records with missing values = 0
27
28
29
30
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
       Parameter Convergence has been set to: 1e-008
31
32
33
34
                       Default Initial (and Specified) Parameter Values
                          background =
                                                      Specified
                                                0
35
                           intercept =
                                          -1.57069
36
                               slope =
                                          0.0201403
37
38
39
               Asymptotic Correlation Matrix of Parameter Estimates
40
41
                ( *** The model parameter(s) -background
42
                     have been estimated at a boundary point, or have been
43
    specified by the user,
44
                      and do not appear in the correlation matrix )
45
46
                  intercept
                                    slope
47
48
     intercept
                                   -0.75
                          1
49
50
         slope
                      -0.75
                                        1
51
52
53
54
55
                               Parameter Estimates
56
           Variable
                               Estimate
                                                    Std. Err.
57
58
                                                    0.433188
           intercept
                               -1.209
              slope
                               0.0190194
                                                  0.00553748
59
60
61
62
                            Analysis of Deviance Table
63
64
                       Log(likelihood) Deviance Test DF
           Model
                                                              P-value
```

Full model -9.98095
Fitted model -12.2154 4.46887 1 0.03452
Reduced model -20.7944 21.6269 2 <.0001

AIC: 28.4308

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000	0.1133	1.133	0	10	-1.131
50.0000	0.3982	3.982	6	10	1.304
150.0000	0.9499	9.499	9	10	-0.7234
Chi-square =	3.50	DF = 1	P-value	= 0.0613	

Benchmark Dose Computation

Specified effect = 0.1

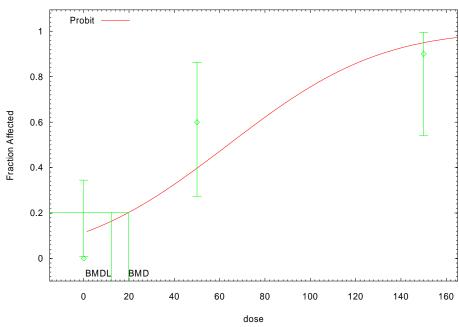
Risk Type = Extra risk

Confidence level = 0.95

BMD = 19.6901

BMDL = 12.3138

Probit Model with 0.95 Confidence Level



11:49 03/09 2006

```
1
    ______
           Probit Model $Revision: 2.1 $ $Date: 2000/02/26 03:38:53 $
2
3
4
5
           Input Data File: C:\BMDS\HEINZE MOD AND INT WITH HEMORRHAGE.(d)
           Thu Jul 14 15:06:15 2005
6
7
     ______
8
     BMDS MODEL RUN
9
    10
11
       The form of the probability function is:
12
13
       P[response] = Background
14
                 + (1-Background) * CumNorm(Intercept+Slope*Log(Dose)),
15
16
       where CumNorm(.) is the cumulative normal distribution function
17
18
19
       Dependent variable = COLUMN2
20
       Independent variable = COLUMN1
21
22
23
24
25
26
       Slope parameter is restricted as slope >= 1
       Total number of observations = 3
       Total number of records with missing values = 0
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
27
28
29
30
       Parameter Convergence has been set to: 1e-008
31
      User has chosen the log transformed model
32
33
34
                    Default Initial (and Specified) Parameter Values
35
                       background =
36
                        intercept =
                                      -3.68466
37
                           slope =
                                            1
38
39
40
              Asymptotic Correlation Matrix of Parameter Estimates
41
42
              ( *** The model parameter(s) -background
                                                      -slope
43
                   have been estimated at a boundary point, or have been
    specified by the user,
44
45
                   and do not appear in the correlation matrix )
46
47
                intercept
48
49
     intercept
50
51
52
53
                           Parameter Estimates
54
55
          Variable
                                              Std. Err.
                           Estimate
56
        background
                                  Ω
                                                 NA
57
58
         intercept
                            -3.68338
                                              0.323658
             slope
59
60
    NA - Indicates that this parameter has hit a bound
61
        implied by some inequality constraint and thus has no standard error.
62
```

Analysis of Deviance Table

Model	Log(likelihood)	Deviance	Test DF	P-value
Full model	-9.98095			
Fitted model	-9.98638	0.0108688	3 2	0.9946
Reduced model	-20.7944	21.6269	2	<.0001

AIC: 21.9728

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000	0.0000	0.000	0	10	0
50.0000	0.5904	5.904	6	10	0.06155
150.0000	0.9078	9.078	9	10	-0.08513
Chi-square =	0.01	DF = 2	P-value	= 0.9945	

Benchmark Dose Computation

Specified effect = 0.1

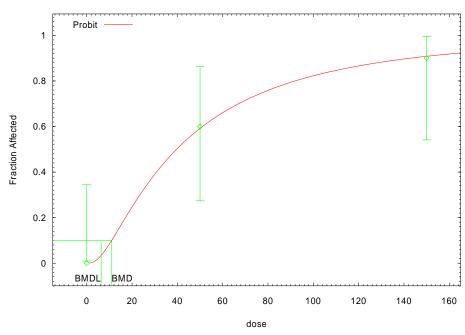
Risk Type = Extra risk

Confidence level = 0.95

BMD = 11.0433

BMDL = 6.37572

Probit Model with 0.95 Confidence Level



15:06 07/14 2005

```
______
           Quantal Linear Model $Revision: 2.2 $ $Date: 2000/03/17 22:27:16 $
2
3
4
5
           Input Data File: C:\BMDS\HEINZE MOD AND INT WITH HEMORRHAGE.(d)
           Thu Jul 14 15:07:01 2005
6
7
     ______
8
     BMDS MODEL RUN
9
    10
11
       The form of the probability function is:
12
13
       P[response] = background + (1-background)*[1-EXP(-slope*dose)]
14
15
16
       Dependent variable = COLUMN2
17
       Independent variable = COLUMN1
18
19
       Total number of observations = 3
20
       Total number of records with missing values = 0
21
22
23
24
25
26
27
28
29
30
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
       Parameter Convergence has been set to: 1e-008
                    Default Initial (and Specified) Parameter Values
                                     0.0454545
                       Background =
                           Slope =
                                     0.0129727
                            Power =
                                                Specified
31
32
33
              Asymptotic Correlation Matrix of Parameter Estimates
34
35
              ( *** The model parameter(s) -Background
36
                   have been estimated at a boundary point, or have been
37
    specified by the user,
38
                   and do not appear in the correlation matrix )
39
40
                    Slope
41
42
        Slope
                        1
43
44
45
46
                           Parameter Estimates
47
48
          Variable
                                              Std. Err.
                           Estimate
49
        Background
                                                 NA
50
                           0.0166997
                                             0.00500498
             Slope
51
52
    NA - Indicates that this parameter has hit a bound
53
        implied by some inequality constraint and thus
54
        has no standard error.
55
56
57
58
                          Analysis of Deviance Table
59
          Model
                    Log(likelihood) Deviance Test DF
                                                      P-value
60
        Full model
                         -9.98095
61
      Fitted model
                         -10.0255
                                     0.089063
                                                            0.9564
                                                  2
                         -20.7944
                                      21.6269
62
      Reduced model
                                                  2
                                                           < .0001
63
64
              AIC:
                          22.051
```

0.09 DF = 2

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000	0.0000	0.000	0	10	0
50.0000	0.5661	5.661	6	10	0.2162
150.0000	0.9183	9.183	9	10	-0.2115

P-value = 0.9553

Benchmark Dose Computation

Chi-square =

Specified effect = 0.1

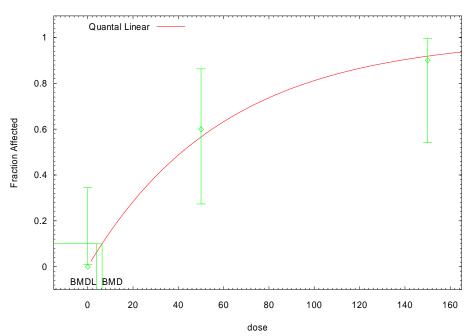
Risk Type = Extra risk

Confidence level = 0.95

BMD = 6.30914

BMDL = 3.92229

Quantal Linear Model with 0.95 Confidence Level



15:07 07/14 2005

```
______
           Quantal Quadratic Model $Revision: 2.2 $ $Date: 2000/03/17 22:27:16 $
2
3
4
5
           Input Data File: C:\BMDS\HEINZE MOD AND INT WITH HEMORRHAGE.(d)
           Thu Jul 14 15:07:40 2005
6
7
     ______
8
     BMDS MODEL RUN
9
    10
11
       The form of the probability function is:
12
13
       P[response] = background + (1-background)*[1-EXP(-slope*dose^2)]
14
15
16
       Dependent variable = COLUMN2
17
       Independent variable = COLUMN1
18
19
       Total number of observations = 3
20
       Total number of records with missing values = 0
21
22
23
24
25
26
27
28
29
30
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
       Parameter Convergence has been set to: 1e-008
                    Default Initial (and Specified) Parameter Values
                       Background =
                                   0.0454545
                           Slope = 8.64849e-005
                           Power =
                                                Specified
31
32
33
              Asymptotic Correlation Matrix of Parameter Estimates
34
35
              ( *** The model parameter(s) -Background
36
                   have been estimated at a boundary point, or have been
37
    specified by the user,
38
                   and do not appear in the correlation matrix )
39
40
                    Slope
41
42
        Slope
                        1
43
44
45
                           Parameter Estimates
46
47
48
          Variable
                                              Std. Err.
                           Estimate
49
        Background
                                                NA
50
                          0.000171114
                                           5.74454e-005
             Slope
51
52
    NA - Indicates that this parameter has hit a bound
53
        implied by some inequality constraint and thus
54
        has no standard error.
55
56
57
58
                         Analysis of Deviance Table
59
          Model
                    Log(likelihood) Deviance Test DF
                                                      P-value
60
        Full model
                         -9.98095
61
      Fitted model
                         -12.0873
                                                            0.1217
                                      4.21262
                                                  2
                         -20.7944
                                      21.6269
62
      Reduced model
                                                  2
                                                           < .0001
63
64
                         26.1745
              AIC:
```

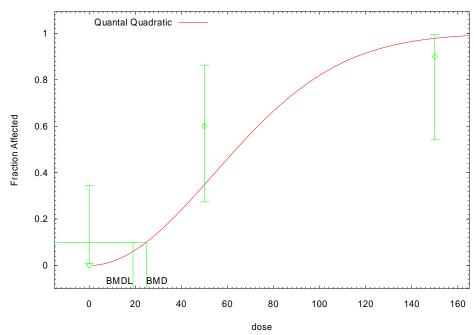
Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000	0.0000	0.000	0	10	0
50.0000	0.3480	3.480	6	10	1.673
150.0000	0.9787	9.787	9	10	-1.725

Chi-square = 5.77 DF = 2 P-value = 0.0558

Benchmark Dose Computation

Specified effect	=	0.1
Risk Type	=	Extra risk
Confidence level	=	0.95
BMD	=	24.814
BMDL	=	19.01

Quantal Quadratic Model with 0.95 Confidence Level



15:07 07/14 2005

```
______
2
           Weibull Model $Revision: 2.2 $ $Date: 2000/03/17 22:27:16 $
            Input Data File: C:\BMDS\HEINZE MOD AND INT WITH HEMORRHAGE.(d)
4
           5
                                           Thu Jul 14 15:08:20 2005
6
7
8
     BMDS MODEL RUN
9
    10
11
       The form of the probability function is:
12
13
       P[response] = background + (1-background)*[1-EXP(-slope*dose^power)]
14
15
16
       Dependent variable = COLUMN2
17
       Independent variable = COLUMN1
18
       Power parameter is restricted as power >=1
19
20
       Total number of observations = 3
21
22
       Total number of records with missing values = 0
       Maximum number of iterations = 250
23
24
25
26
27
28
29
30
       Relative Function Convergence has been set to: 1e-008
       Parameter Convergence has been set to: 1e-008
                     Default Initial (and Specified) Parameter Values
                        Background =
                                       0.0454545
                            Slope =
                                       0.0129727
31
                            Power =
32
33
34
              Asymptotic Correlation Matrix of Parameter Estimates
35
36
              ( *** The model parameter(s) -Background
37
                    have been estimated at a boundary point, or have been
38
    specified by the user,
39
                    and do not appear in the correlation matrix )
40
41
                     Slope
42
43
         Slope
                        1
44
45
46
47
                            Parameter Estimates
48
49
          Variable
                            Estimate
                                                Std. Err.
50
         Background
                                    Ω
                                                  NA
51
             Slope
                            0.0166997
                                              0.00500498
52
             Power
53
54
    NA - Indicates that this parameter has hit a bound
55
         implied by some inequality constraint and thus
56
         has no standard error.
57
58
59
60
                          Analysis of Deviance Table
61
62
          Model
                     Log(likelihood) Deviance Test DF
                                                         P-value
63
         Full model
                         -9.98095
64
                                       0.089063
                                                    2
                                                              0.9564
       Fitted model
                          -10.0255
```

Reduced model -20.7944 21.6269 2 <.0001

AIC: 22.051

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000 50.0000	0.0000 0.5661	0.000 5.661	0	10 10	0 0.2162
150.0000	0.9183	9.183	9	10	-0.2115

Chi-square = 0.09 DF = 2 P-value = 0.9553

Benchmark Dose Computation

Specified effect = 0.1

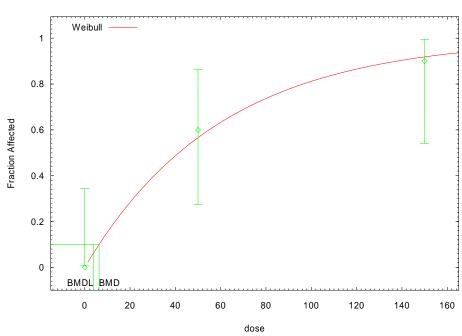
Risk Type = Extra risk

Confidence level = 0.95

BMD = 6.30914

BMDL = 3.92229

Weibull Model with 0.95 Confidence Level



15:08 07/14 2005

```
1
2 3 4 5 6 7 8 9
    Fawell et al., 1999 Male Chronic Inflammation
     ______
              $Revision: 2.2 $ $Date: 2001/03/14 01:17:00 $
            Input Data File: C:\BMDS\FAWELL MALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_MALE_CHRONIC_INFLAMMATION.plt
                                               Wed Dec 28 15:45:24 2005
     ______
10
11
     BMDS MODEL RUN
12
13
14
        The form of the probability function is:
15
16
        P[response] = background+(1-background)*CumGamma[slope*dose,power],
17
       where CumGamma(.) is the cummulative Gamma distribution function
18
19
20
       Dependent variable = COLUMN2
21
22
23
        Independent variable = COLUMN1
        Power parameter is restricted as power >=1
24
25
26
27
28
29
30
        Total number of observations = 4
        Total number of records with missing values = 0
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
        Parameter Convergence has been set to: 1e-008
31
32
                       Default Initial (and Specified) Parameter Values
33
                          Background =
                                            0.09375
34
                               Slope =
                                         0.00193589
35
                               Power =
                                                1.3
36
37
38
                Asymptotic Correlation Matrix of Parameter Estimates
39
40
                  Background
                                    Slope
                                                 Power
41
42
    Background
                                    0.037
                                                 0.041
43
44
          Slope
                       0.037
                                        1
                                                     1
45
46
         Power
                       0.041
                                        1
                                                     1
47
48
49
50
                               Parameter Estimates
51
52
           Variable
                               Estimate
                                                    Std. Err.
53
         Background
                               0.0999826
                                                    0.054818
54
               Slope
                               0.0276638
                                                    0.488219
55
               Power
                                 8.09035
                                                     116.456
56
57
58
                             Analysis of Deviance Table
59
60
                       Log(likelihood) Deviance Test DF
           Model
                                                             P-value
61
         Full model
                           -18.2628
62
        Fitted model
                                         0.376765
                                                                   0.5393
                            -18.4512
                                                      1
63
      Reduced model
                            -39.4295
                                           42.3333
                                                        3
                                                                  < .0001
```

AIC: 42.9024

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000	0.1000	1.500	1	15	-0.4301
40.0000	0.1000	1.500	2	15	0.4304
200.0000	0.2667	4.001	4	15	-0.0002949
1000.0000	1.0000	15.000	15	15	0.00692

Chi-square = 0.37 DF = 1 P-value = 0.5429

Benchmark Dose Computation

Specified effect = 0.1

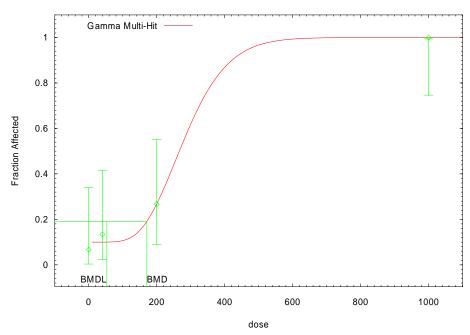
Risk Type = Extra risk

Confidence level = 0.95

BMD = 170.825

BMDL = 53.1183

Gamma Multi-Hit Model with 0.95 Confidence Level



15:45 12/28 2005

```
1
    ______
            Logistic Model $Revision: 2.1 $ $Date: 2000/02/26 03:38:20 $
2
3
4
5
            Input Data File: C:\BMDS\FAWELL MALE CHRONIC INFLAMMATION.(d)
            Gnuplot Plotting File: C:\BMDS\FAWELL MALE CHRONIC INFLAMMATION.plt
                                             Thu \overline{M}ar 0\overline{9} 11:56:\overline{5}8 2006
6
7
     ______
8
     BMDS MODEL RUN
9
    10
11
       The form of the probability function is:
12
13
       P[response] = 1/[1+EXP(-intercept-slope*dose)]
14
15
16
       Dependent variable = COLUMN2
17
       Independent variable = COLUMN1
18
       Slope parameter is not restricted
19
20
       Total number of observations = 4
21
22
       Total number of records with missing values = 0
       Maximum number of iterations = 250
23
24
25
26
27
28
29
       Relative Function Convergence has been set to: 1e-008
       Parameter Convergence has been set to: 1e-008
                     Default Initial Parameter Values
                        background =
                                                   Specified
                                               Ω
30
                         intercept =
                                         -2.07771
31
                                       0.00552538
                             slope =
32
33
34
               Asymptotic Correlation Matrix of Parameter Estimates
35
36
               ( *** The model parameter(s) -background
37
                    have been estimated at a boundary point, or have been
38
    specified by the user,
39
                    and do not appear in the correlation matrix )
40
41
                  intercept
                                  slope
42
43
     intercept
                         1
                                  -0.72
44
45
         slope
                     -0.72
46
47
48
49
                             Parameter Estimates
50
51
           Variable
                             Estimate
                                                 Std. Err.
52
53
          intercept
                              -2.49527
                                                 0.612166
                            0.00805129
                                               0.00337645
              slope
54
55
56
57
58
                           Analysis of Deviance Table
59
           Model
                     Log(likelihood) Deviance Test DF
                                                          P-value
60
         Full model
                          -18.2628
                          -18.4278
                                        0.329996
61
       Fitted model
                                                     2
                                                                0.8479
      Reduced model
                          -39.4295
                                        42.3333
                                                               <.0001
62
                                                     3
63
64
              AIC:
                           40.8556
```

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000	0.0762	1.143	1	15	-0.139
40.0000	0.1022	1.533	2	15	0.3983
200.0000	0.2921	4.382	4	15	-0.217
1000.0000	0.9962	14.942	15	15	0.2408
Chi-square :	= 0.28	DF = 2	P-value	= 0.8680	

Benchmark Dose Computation

Specified effect = 0.1

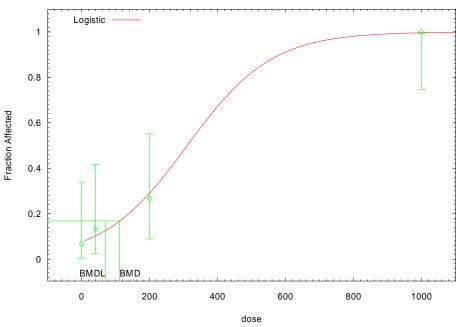
Risk Type = Extra risk

Confidence level = 0.95

BMD = 111.719

BMDL = 70.686

Logistic Model with 0.95 Confidence Level



11:56 03/09 2006

```
1
2
3
4
5
    ______
            Logistic Model $Revision: 2.1 $ $Date: 2000/02/26 03:38:20 $
            Input Data File: C:\BMDS\FAWELL MALE CHRONIC INFLAMMATION.(d)
            Gnuplot Plotting File: C:\BMDS\FAWELL MALE CHRONIC INFLAMMATION.plt
                                             Wed \overline{D}ec 2\overline{8} 15:47:\overline{1}1 2005
6
7
     ______
8
     BMDS MODEL RUN
9
    10
11
       The form of the probability function is:
12
13
       P[response] = background+(1-background)/[1+EXP(-intercept-slope*Log(dose))]
14
15
16
       Dependent variable = COLUMN2
17
       Independent variable = COLUMN1
18
       Slope parameter is restricted as slope >= 1
19
20
       Total number of observations = 4
21
22
23
24
25
26
27
28
29
30
       Total number of records with missing values = 0
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
       Parameter Convergence has been set to: 1e-008
       User has chosen the log transformed model
31
                      Default Initial Parameter Values
32
                         background =
                                       0.0666667
33
34
                                         -10.0178
                          intercept =
                              slope =
                                          1.86367
35
36
37
               Asymptotic Correlation Matrix of Parameter Estimates
38
39
                 background
                              intercept
                                               slope
40
41
    background
                          1
                                 0.0022
                                             -0.0023
42
43
                     0.0022
     intercept
                                      1
                                                  -1
44
45
         slope
                    -0.0023
                                     -1
                                                   1
46
47
48
49
                             Parameter Estimates
50
51
           Variable
                             Estimate
                                                  Std. Err.
52
53
         background
                             0.0999997
                                                 0.0547725
                              -65.9875
                                                   5466.64
          intercept
54
55
              slope
                               12.1748
                                                   1031.77
56
57
58
                           Analysis of Deviance Table
59
                      Log(likelihood) Deviance Test DF
           Model
                                                          P-value
60
         Full model
                           -18.2628
61
       Fitted model
                           -18.4512
                                        0.376844
                                                                0.5393
                                                      1
      Reduced model
                                         42.3333
                                                      3
62
                           -39.4295
                                                                < .0001
63
64
                           42.9024
               AIC:
```

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000	0.1000	1.500	1	15	-0.4303
40.0000	0.1000	1.500	2	15	0.4303
200.0000	0.2667	4.000	4	15	-4.206e-005
1000.0000	1.0000	15.000	15	15	0.0004285

Chi-square = 0.37 DF = 1 P-value = 0.5428

Benchmark Dose Computation

Specified effect = 0.1

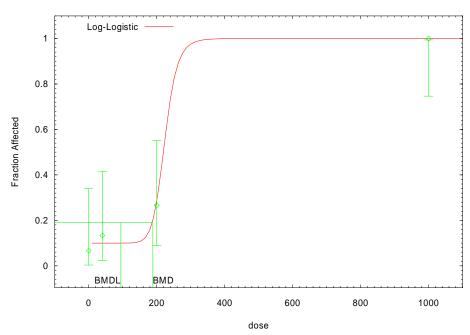
Risk Type = Extra risk

Confidence level = 0.95

BMD = 188.582

BMDL = 94.487

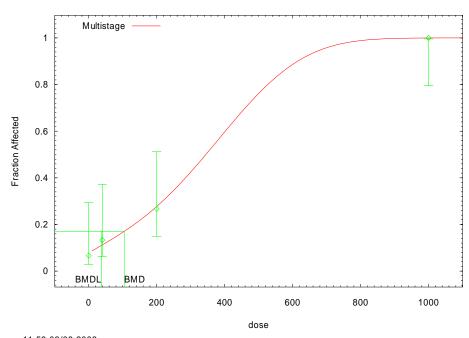
Log-Logistic Model with 0.95 Confidence Level



15:47 12/28 2005

```
______
             Multistage Model. $Revision: 2.1 $ $Date: 2000/08/21 03:38:21 $
2
3
4
5
             Input Data File: C:\BMDS\FAWELL MALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_MALE_CHRONIC_INFLAMMATION.plt
                                                Thu \overline{M}ar 0\overline{9} 11:59:\overline{4}4 2006
6
7
     ______
8
     BMDS MODEL RUN
9
     10
11
        The form of the probability function is:
12
13
        P[response] = background + (1-background) * [1-EXP(
     -beta1*dose^1-beta2*dose^2-beta3*dose^3)]
14
15
16
        The parameter betas are restricted to be positive
17
18
19
        Dependent variable = COLUMN2
20
        Independent variable = COLUMN1
21
\overline{22}
     Total number of observations = 4
23
     Total number of records with missing values = 0
24
25
26
     Total number of parameters in model = 4
Total number of specified parameters = 0
     Degree of polynomial = 3
27
28
29
     Maximum number of iterations = 250
30
     Relative Function Convergence has been set to: 1e-008
31
     Parameter Convergence has been set to: 1e-008
32
33
34
                       Default Initial Parameter Values
35
                          Background =
36
                             Beta(1) =
                                                   0
37
                             Beta(2) =
                                                   0
38
                             Beta(3) = 1.00264e+011
39
40
41
                Asymptotic Correlation Matrix of Parameter Estimates
42
43
                ( *** The model parameter(s) -Beta(2)
44
                      have been estimated at a boundary point, or have been
45
    specified by the user,
46
                      and do not appear in the correlation matrix )
47
48
                  Background
                                  Beta(1)
                                                Beta(3)
49
50
    Background
                                    -0.65
                                                   0.14
51
52
53
       Beta(1)
                       -0.65
                                        1
                                                  -0.42
54
        Beta(3)
                       0.14
                                     -0.42
55
56
57
58
                               Parameter Estimates
59
           Variable
                               Estimate
                                                     Std. Err.
60
         Background
                               0.0782274
                                                     0.200883
                                                  0.00219636
61
            Beta(1)
                             0.000933123
            Beta(2)
62
                                                        NΑ
63
             Beta(3)
                            7.10432e-009
                                                 1.57018e-008
64
```

1 2 3 4 5 6 7	NA - Indicates implied by has no sta					
5		A	nalysis of	Deviance Tab	le	
7				eviance Tes	t DF P-	value
8 9 10	Full model Fitted model Reduced model	-1: -1: -3:	8.2628 8.3175 9.4295	0.10941 42.3333	1 3	0.7408 <.0001
11 12 13	AIC:		42.635			
14 15 16		Good	ness of F	it		
17 18	Dose E	stProb.	Expected	Observed	Size	Chi^2 Res.
19 20	i: 1 0.0000	0.0782	1.173	1	15	-0.160
21 22	i: 2 40.0000	0.1124	1.686	2	15	0.210
23 24	i: 3 200.0000	0.2774	4.161	4	15	-0.054
25 26	i: 4 1000.0000	0.9997	14.996	15	15	1.000
27 28	Chi-square =	0.11	DF = 1	P-val	ue = 0.7438	
29 30		<u>.</u>				
31 32	Benchmark Do	se Computa	tion			
33 34	Specified effec	t =	0.1			
35 36	Risk Type	= E:	xtra risk			
37 38	Confidence leve	1 =	0.95			
39	ВМ	D =	104.279			
40 41	BMD	L =	37.4815			



11:59 03/09 2006

```
______
             Multistage Model. $Revision: 2.1 $ $Date: 2000/08/21 03:38:21 $
2
3
4
5
             Input Data File: C:\BMDS\FAWELL MALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_MALE_CHRONIC_INFLAMMATION.plt
                                                Thu \overline{M}ar 0\overline{9} 12:00:\overline{4}8 2006
6
7
     ______
8
     BMDS MODEL RUN
9
     10
11
        The form of the probability function is:
12
13
        P[response] = background + (1-background) * [1-EXP(
     -beta1*dose^1-beta2*dose^2)]
14
15
16
        The parameter betas are restricted to be positive
17
18
19
        Dependent variable = COLUMN2
20
        Independent variable = COLUMN1
21
\overline{22}
     Total number of observations = 4
23
     Total number of records with missing values = 0
24
25
26
     Total number of parameters in model = 3
Total number of specified parameters = 0
     Degree of polynomial = 2
27
28
29
     Maximum number of iterations = 250
30
     Relative Function Convergence has been set to: 1e-008
31
     Parameter Convergence has been set to: 1e-008
32
33
34
35
                       Default Initial Parameter Values
36
                          Background =
                                                   0
37
                             Beta(1) =
                                                   0
38
                             Beta(2) = 1.01264e+014
39
40
41
                Asymptotic Correlation Matrix of Parameter Estimates
42
43
                  Background
                                   Beta(1)
                                                Beta(2)
44
45
    Background
                                     -0.65
                                                   0.33
46
47
       Beta(1)
                       -0.65
                                        1
                                                  -0.75
48
49
       Beta(2)
                       0.33
                                     -0.75
                                                      1
50
51
52
53
                               Parameter Estimates
54
55
            Variable
                               Estimate
                                                     Std. Err.
56
57
58
          Background
                               0.0889744
                                                     0.205309
                                                   0.00292558
             Beta(1)
                            8.80277e-005
             Beta(2)
                            5.76916e-006
                                                 6.80122e-006
59
60
61
62
                             Analysis of Deviance Table
63
64
                       Log(likelihood) Deviance Test DF
           Model
                                                                P-value
```

Full model -18.2628
Fitted model -18.4529 0.380206 1 0.5375
Reduced model -39.4295 42.3333 3 <.0001

AIC: 42.9058

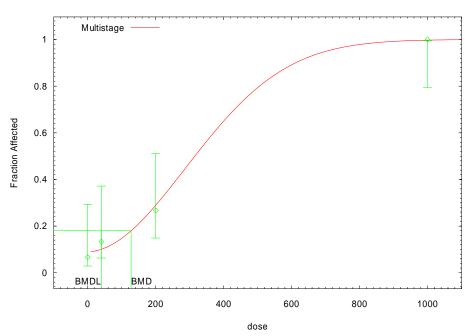
Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Chi^2 Res.
i: 1 0.0000	0.0890	1.335	1	15	-0.275
i: 2 40.0000 i: 3	0.1005	1.508	2	15	0.363
200.0000 i: 4	0.2893	4.340	4	15	-0.110
1000.0000	0.9974	14.961	15	15	1.003
Chi-square =	0.35	DF = 1	P-value	= 0.5556	

Benchmark Dose Computation

Specified effect = 0.1
Risk Type = Extra risk
Confidence level = 0.95
BMD = 127.726
BMDL = 39.8667

Multistage Model with 0.95 Confidence Level



12:00 03/09 2006

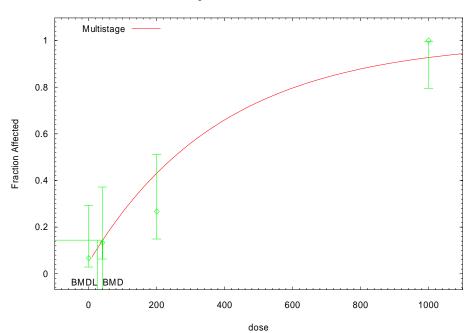
```
______
             Multistage Model. $Revision: 2.1 $ $Date: 2000/08/21 03:38:21 $
2
3
4
5
             Input Data File: C:\BMDS\FAWELL MALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_MALE_CHRONIC_INFLAMMATION.plt
                                               Wed \overline{D}ec 2\overline{8} 15:47:\overline{4}3 2005
6
7
     ______
8
     BMDS MODEL RUN
9
     10
11
       The form of the probability function is:
12
13
        P[response] = background + (1-background) * [1-EXP(
14
     -beta1*dose^1)]
15
16
        The parameter betas are restricted to be positive
17
18
19
       Dependent variable = COLUMN2
20
        Independent variable = COLUMN1
21
\overline{22}
     Total number of observations = 4
23
     Total number of records with missing values = 0
24
25
26
     Total number of parameters in model = 2
Total number of specified parameters = 0
     Degree of polynomial = 1
27
28
29
     Maximum number of iterations = 250
30
     Relative Function Convergence has been set to: 1e-008
31
     Parameter Convergence has been set to: 1e-008
32
33
34
35
                       Default Initial Parameter Values
36
                          Background =
37
                             Beta(1) = 1.04991e+017
38
39
40
                Asymptotic Correlation Matrix of Parameter Estimates
41
42
                  Background
                                  Beta(1)
43
44
    Background
                                    -0.37
45
46
       Beta(1)
                       -0.37
                                        1
47
48
49
50
                               Parameter Estimates
51
52
           Variable
                               Estimate
                                                     Std. Err.
53
         Background
                                                     0.16314
                                0.048937
54
             Beta(1)
                               0.0025684
                                                  0.000904753
55
56
57
58
                             Analysis of Deviance Table
59
                       Log(likelihood) Deviance Test DF
           Model
                                                             P-value
60
         Full model
                            -18.2628
                            -20.3189
61
       Fitted model
                                                                     0.128
                                           4.11219
                                                         2
      Reduced model
                                           42.3333
62
                            -39.4295
                                                         3
                                                                   < .0001
63
64
                             44.6378
               AIC:
```

Dose	EstProb.	Expected	Observed	Size	Chi^2 Res.
i: 1 0.0000 i: 2	0.0489	0.734	1	15	0.381
40.0000 i: 3	0.1418	2.127	2	15	-0.070
200.0000 i: 4	0.4310	6.465	4	15	-0.670
1000.0000	0.9271	13.906	15	15	1.079
Chi-square =	2.94	DF = 2	P-value	= 0.2298	

Benchmark Dose Computation

Specified effect = 0.1 Risk Type Extra risk Confidence level = 0.95 41.0219 BMD = BMDL = 25.943

Multistage Model with 0.95 Confidence Level



15:47 12/28 2005

```
1
    ______
2
            Probit Model $Revision: 2.1 $ $Date: 2000/02/26 03:38:53 $
            Input Data File: C:\BMDS\FAWELL MALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_MALE_CHRONIC_INFLAMMATION.plt
4
5
                                             Thu Mar 09 12:02:26 2006
6
7
     ______
8
     BMDS MODEL RUN
9
     10
11
       The form of the probability function is:
12
13
       P[response] = CumNorm(Intercept+Slope*Dose),
14
15
       where CumNorm(.) is the cumulative normal distribution function
16
17
18
       Dependent variable = COLUMN2
19
       Independent variable = COLUMN1
20
       Slope parameter is not restricted
21
22
       Total number of observations = 4
23
24
25
26
       Total number of records with missing values = 0
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
       Parameter Convergence has been set to: 1e-008
27
28
29
30
                      Default Initial (and Specified) Parameter Values
31
                         background =
                                                0
                                                    Specified
32
                                         -1.32259
                          intercept =
33
                              slope =
                                        0.00345221
34
35
36
               Asymptotic Correlation Matrix of Parameter Estimates
37
38
               ( *** The model parameter(s) -background
39
                     have been estimated at a boundary point, or have been
40
    specified by the user,
41
                     and do not appear in the correlation matrix )
42
43
                  intercept
                                   slope
44
45
     intercept
                                   -0.65
46
47
                      -0.65
         slope
48
49
50
51
                              Parameter Estimates
52
53
           Variable
                                                  Std. Err.
                             Estimate
54
          intercept
                              -1.42508
                                                  0.306506
55
                             0.00435347
                                                0.00171942
              slope
56
57
58
59
                            Analysis of Deviance Table
60
61
           Model
                      Log(likelihood) Deviance Test DF
                                                           P-value
         Full model
                           -18.2628
62
63
       Fitted model
                           -18.3773
                                        0.229031
                                                      2
                                                                 0.8918
      Reduced model
                           -39.4295
                                         42.3333
64
                                                      3
                                                                < .0001
```

AIC: 40.7546

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000 40.0000 200.0000 1000.0000	0.0771 0.1055 0.2897 0.9983	1.156 1.582 4.345 14.974	1 2 4 15	15 15 15 15	-0.151 0.3512 -0.1963 0.16
Chi-square =	- 0.21	DF = 2	P-value	= 0.9002	

Benchmark Dose Computation

Specified effect = 0.1

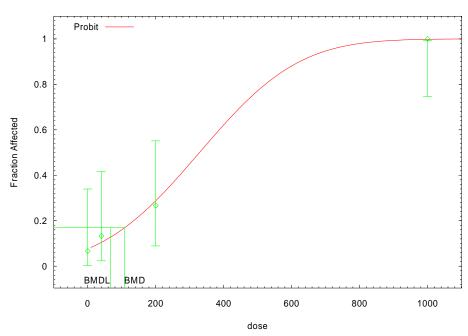
Risk Type = Extra risk

Confidence level = 0.95

BMD = 107.589

BMDL = 66.4468

Probit Model with 0.95 Confidence Level



12:02 03/09 2006

```
1
    ______
            Probit Model $Revision: 2.1 $ $Date: 2000/02/26 03:38:53 $
2
3
4
5
            Input Data File: C:\BMDS\FAWELL MALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_MALE_CHRONIC_INFLAMMATION.plt
                                              Wed Dec 28 15:48:13 2005
6
7
     ______
8
     BMDS MODEL RUN
9
     10
11
       The form of the probability function is:
12
13
       P[response] = Background
14
                   + (1-Background) * CumNorm(Intercept+Slope*Log(Dose)),
15
16
       where CumNorm(.) is the cumulative normal distribution function
17
18
19
       Dependent variable = COLUMN2
20
       Independent variable = COLUMN1
21
22
23
24
25
26
27
28
29
30
       Slope parameter is restricted as slope >= 1
       Total number of observations = 4
       Total number of records with missing values = 0
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
       Parameter Convergence has been set to: 1e-008
31
       User has chosen the log transformed model
32
33
34
                      Default Initial (and Specified) Parameter Values
35
                         background =
                                        0.0666667
36
                          intercept =
                                          -5.60926
37
                                           1.03389
                              slope =
38
39
40
               Asymptotic Correlation Matrix of Parameter Estimates
41
42
                 background
                               intercept
                                                slope
43
44
    background
                                  0.0018
                                               -0.002
45
46
     intercept
                     0.0018
                                       1
                                                   -1
47
48
         slope
                     -0.002
                                      -1
                                                    1
49
50
51
52
53
                              Parameter Estimates
54
           Variable
                              Estimate
                                                  Std. Err.
55
                                                  0.0547723
         background
                                    0.1
56
57
58
                                                    1087.37
          intercept
                               -22.1534
              slope
                                4.01215
                                                    205.229
59
60
61
                            Analysis of Deviance Table
62
63
           Model
                      Log(likelihood) Deviance Test DF
                                                           P-value
64
                           -18.2628
         Full model
```

Fitted model -18.4512 0.376844 1 0.5393 Reduced model -39.4295 42.3333 3 <.0001

AIC: 42.9024

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000	0.1000	1.500	1	15	-0.4303
40.0000	0.1000	1.500	2	15	0.4303
200.0000	0.2667	4.000	4	15	-1.174e-005
1000.0000	1.0000	15.000	15	15	0.0004249
Chi-square =	0.37	DF = 1	P-value	= 0 5428	

Benchmark Dose Computation

Specified effect = 0.1

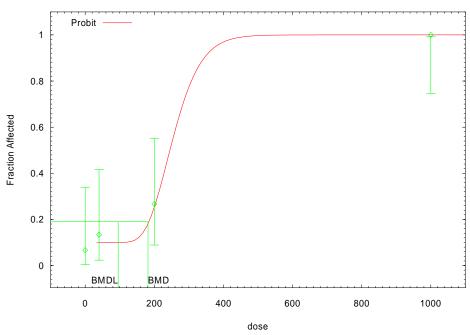
Risk Type = Extra risk

Confidence level = 0.95

BMD = 181.665

BMDL = 95.2612

Probit Model with 0.95 Confidence Level



15:48 12/28 2005

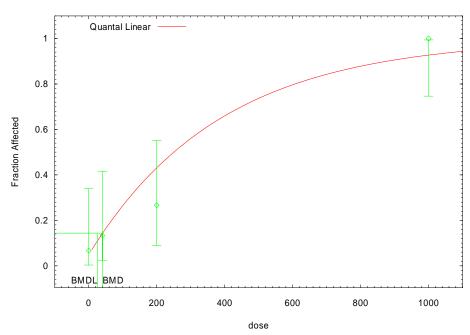
```
______
            Quantal Linear Model $Revision: 2.2 $ $Date: 2000/03/17 22:27:16 $
2
3
4
5
            Input Data File: C:\BMDS\FAWELL MALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_MALE_CHRONIC_INFLAMMATION.plt
                                             Wed Dec 28 15:48:41 2005
6
7
     ______
8
     BMDS MODEL RUN
9
    10
11
       The form of the probability function is:
12
13
       P[response] = background + (1-background)*[1-EXP(-slope*dose)]
14
15
16
       Dependent variable = COLUMN2
17
       Independent variable = COLUMN1
18
19
       Total number of observations = 4
20
       Total number of records with missing values = 0
21
22
23
24
25
26
27
28
29
30
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
       Parameter Convergence has been set to: 1e-008
                      Default Initial (and Specified) Parameter Values
                                      0.09375
                         Background =
                             Slope =
                                        0.0033673
                             Power =
                                                    Specified
31
32
33
               Asymptotic Correlation Matrix of Parameter Estimates
34
35
               ( *** The model parameter(s) -Power
36
                     have been estimated at a boundary point, or have been
37
    specified by the user,
38
                     and do not appear in the correlation matrix )
39
40
                 Background
                                  Slope
41
42
    Background
                         1
                                  -0.23
43
44
         Slope
                      -0.23
45
46
47
48
                             Parameter Estimates
49
50
           Variable
                             Estimate
                                                  Std. Err.
51
         Background
                              0.048937
                                                 0.0439067
52
53
54
55
              Slope
                             0.0025684
                                               0.000746108
56
                           Analysis of Deviance Table
57
58
           Model
                     Log(likelihood) Deviance Test DF
59
         Full model
                          -18.2628
60
       Fitted model
                                         4.11219
                                                      2
                                                                 0.128
                          -20.3189
                                                     3
61
      Reduced model
                                         42.3333
                                                               < .0001
                          -39.4295
62
63
               AIC:
                           44.6378
64
```

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000 40.0000 200.0000 1000.0000	0.0489 0.1418 0.4310 0.9271	0.734 2.127 6.465 13.906	1 2 4 15	15 15 15 15	0.3183 -0.09393 -1.285 1.086
Chi-square =	= 2.94	DF = 2	P-value	= 0.2298	

Benchmark Dose Computation

Specified effect	=	0.1
Risk Type	=	Extra risk
Confidence level	=	0.95
BMD	=	41.0219
BMDL	=	25.943

Quantal Linear Model with 0.95 Confidence Level



15:48 12/28 2005

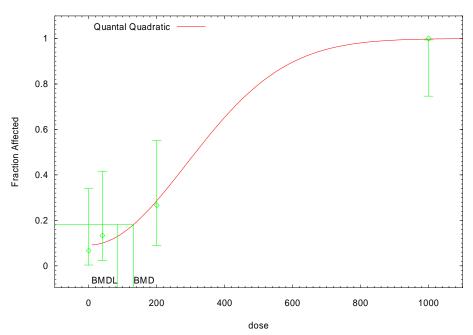
```
______
            Quantal Quadratic Model $Revision: 2.2 $ $Date: 2000/03/17 22:27:16 $
2
3
4
5
            Input Data File: C:\BMDS\FAWELL MALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_MALE_CHRONIC_INFLAMMATION.plt
                                            Wed Dec 28 15:49:14 2005
6
7
     ______
8
     BMDS MODEL RUN
9
    10
11
       The form of the probability function is:
12
13
       P[response] = background + (1-background)*[1-EXP(-slope*dose^2)]
14
15
16
       Dependent variable = COLUMN2
17
       Independent variable = COLUMN1
18
19
       Total number of observations = 4
20
       Total number of records with missing values = 0
21
22
23
24
25
26
27
28
29
30
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
       Parameter Convergence has been set to: 1e-008
                      Default Initial (and Specified) Parameter Values
                        Background = 0.09375
                             Slope = 3.3673e-006
                             Power =
                                                    Specified
31
32
33
               Asymptotic Correlation Matrix of Parameter Estimates
34
35
               ( *** The model parameter(s) -Power
36
                     have been estimated at a boundary point, or have been
37
    specified by the user,
38
                     and do not appear in the correlation matrix )
39
40
                 Background
                                  Slope
41
42
    Background
                         1
                                  -0.32
43
44
         Slope
                     -0.32
45
46
47
48
                             Parameter Estimates
49
50
           Variable
                             Estimate
                                                 Std. Err.
51
         Background
                              0.091177
                                                0.0532315
52
53
54
55
              Slope
                          6.03482e-006
                                              3.53595e-006
56
                           Analysis of Deviance Table
57
58
           Model
                     Log(likelihood) Deviance Test DF
59
         Full model
                          -18.2628
                                       0.382862
60
       Fitted model
                                                     2
                                                               0.8258
                          -18.4542
                                                     3
61
      Reduced model
                                        42.3333
                                                               < .0001
                          -39.4295
62
63
              AIC:
                           40.9085
64
```

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000 40.0000 200.0000 1000.0000	0.0912 0.0999 0.2861 0.9978	1.368 1.499 4.291 14.967	1 2 4 15	15 15 15 15	-0.3298 0.4317 -0.1665 0.1808
Chi-square	= 0.36	DF = 2	P-value	= 0.8372	

Benchmark Dose Computation

Specified effect = 0.1 Risk Type Extra risk Confidence level = 0.95 BMD = 132.132 BMDL = 85.4708

Quantal Quadratic Model with 0.95 Confidence Level



15:49 12/28 2005

```
______
2
3
4
5
            Weibull Model $Revision: 2.2 $ $Date: 2000/03/17 22:27:16 $
            Input Data File: C:\BMDS\FAWELL MALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_MALE_CHRONIC_INFLAMMATION.plt
                                             Wed Dec 28 15:49:46 2005
6
7
     ______
8
     BMDS MODEL RUN
9
     10
11
       The form of the probability function is:
12
13
       P[response] = background + (1-background)*[1-EXP(-slope*dose^power)]
14
15
16
       Dependent variable = COLUMN2
17
       Independent variable = COLUMN1
18
       Power parameter is restricted as power >=1
19
20
       Total number of observations = 4
21
22
23
24
25
26
27
28
29
30
       Total number of records with missing values = 0
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
       Parameter Convergence has been set to: 1e-008
                      Default Initial (and Specified) Parameter Values
                                         0.09375
                         Background =
                              Slope = 4.78181e-007
31
                              Power =
                                          2.28256
32
33
34
               Asymptotic Correlation Matrix of Parameter Estimates
35
36
                 Background
                                  Slope
                                               Power
37
38
    Background
                                  -0.33
                                                0.31
39
40
         Slope
                     -0.33
                                      1
                                                  -1
41
42
         Power
                      0.31
                                     - 1
                                                   1
43
44
45
46
                             Parameter Estimates
47
48
           Variable
                                                 Std. Err.
                             Estimate
49
         Background
                             0.0959027
                                                0.0577365
50
              Slope
                           1.27341e-006
                                              1.17833e-005
51
              Power
                               2.27222
                                                   1.67024
52
53
54
55
                           Analysis of Deviance Table
56
57
           Model
                      Log(likelihood) Deviance Test DF
                                                          P-value
58
         Full model
                           -18.2628
59
       Fitted model
                                                                0.5649
                           -18.4284
                                        0.331283
                                                     1
60
      Reduced model
                                         42.3333
                                                      3
                                                                < .0001
                          -39.4295
61
62
                           42.8569
               AIC:
63
```

1
2
- 3
3 4
5
5
5 6
7
8
9 10
10
11
12
12
13
14
15
16
13 14 15 16 17 18 19 20
18
10
19
20
21
22
44
23 24
24
27
25

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000	0.0959	1.439	1	15	-0.3845
40.0000	0.1009	1.514	2	15	0.4168
200.0000	0.2712	4.067	4	15	-0.03917
1000.0000	0.9998	14.997	15	15	0.05665

Chi-square = 0.33 DF = 1P-value = 0.5678

Benchmark Dose Computation

Specified effect = 0.1

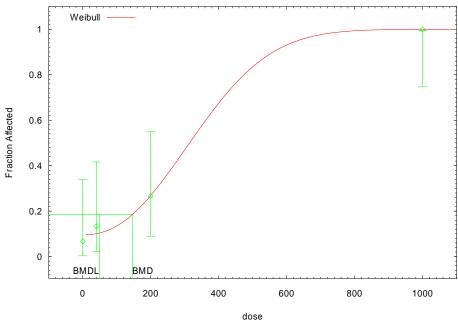
Risk Type Extra risk

Confidence level = 0.95

> BMD = 145.973

BMDL = 48.796

Weibull Model with 0.95 Confidence Level



15:49 12/28 2005

```
Fawell et al., 1999 Female Chronic Inflammation
1
2
3
4
5
    ______
             $Revision: 2.2 $ $Date: 2001/03/14 01:17:00 $
            Input Data File: C:\BMDS\FAWELL_FEMALE_CHRONIC_INFLAMMATION.(d)
            Gnuplot Plotting File: C:\BMDS\FAWELL_FEMALE_CHRONIC_INFLAMMATION.plt
6
7
                                              Wed Dec 28 15:50:58 2005
8
     ______
9
10
     BMDS MODEL RUN
11
12
13
       The form of the probability function is:
14
15
       P[response] = background+(1-background)*CumGamma[slope*dose,power],
16
       where CumGamma(.) is the cummulative Gamma distribution function
17
18
19
       Dependent variable = COLUMN2
20
       Independent variable = COLUMN1
21
22
       Power parameter is restricted as power >=1
23
24
25
26
       Total number of observations = 4
       Total number of records with missing values = 0
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
27
28
29
       Parameter Convergence has been set to: 1e-008
30
31
                      Default Initial (and Specified) Parameter Values
32
                         Background =
                                           0.34375
33
                              Slope =
                                        0.00408479
34
                              Power =
                                           2.14573
35
36
37
               Asymptotic Correlation Matrix of Parameter Estimates
38
39
               ( *** The model parameter(s)
                                            -Power
40
                     have been estimated at a boundary point, or have been
41
    specified by the user,
42
                     and do not appear in the correlation matrix )
43
44
                 Background
                                   Slope
45
46
    Background
                          1
                                   -0.42
47
48
         Slope
                      -0.42
                                       1
49
50
51
52
53
                              Parameter Estimates
54
           Variable
                              Estimate
                                                  Std. Err.
55
                                                 0.0913888
         Background
                               0.385764
56
                                                0.000852726
              Slope
                              0.0020737
57
              Power
58
59
    NA - Indicates that this parameter has hit a bound
60
         implied by some inequality constraint and thus
61
         has no standard error.
62
```

Analysis of Deviance Table

Model	Log(likelihood)	Deviance	Test DF	P-value
Full model	-33.9494			
Fitted model	-34.4584	1.01803	1 2	0.6011
Reduced model	-40.7516	13.6045	5 3	0.003496

AIC: 72.9167

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000	0.3858 0.4347	5.786 6.520	5 8	15 15	-0.4172
200.0000	0.5943	8.914	8	15	-0.4808
1000.0000 Chi-square =	0.9228	13.842	14	15 = 0.5996	0.1531
cmi-square =	1.02	DF = 2	P-value	= 0.5996	

Benchmark Dose Computation

Specified effect = 0.1

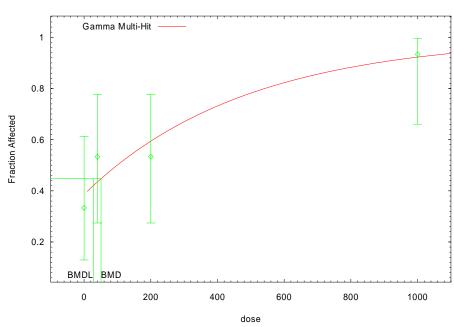
Risk Type = Extra risk

Confidence level = 0.95

BMD = 50.8081

BMDL = 27.7368

Gamma Multi-Hit Model with 0.95 Confidence Level



A-49

15:50 12/28 2005

```
1
    ______
            Logistic Model $Revision: 2.1 $ $Date: 2000/02/26 03:38:20 $
2
3
4
5
            Input Data File: C:\BMDS\FAWELL FEMALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_FEMALE_CHRONIC_INFLAMMATION.plt
                                             Thu Mar 09 12:16:37 2006
6
7
     ______
8
     BMDS MODEL RUN
9
     10
11
       The form of the probability function is:
12
13
       P[response] = 1/[1+EXP(-intercept-slope*dose)]
14
15
16
       Dependent variable = COLUMN2
17
       Independent variable = COLUMN1
18
       Slope parameter is not restricted
19
20
       Total number of observations = 4
21
22
       Total number of records with missing values = 0
       Maximum number of iterations = 250
23
24
25
26
27
28
29
       Relative Function Convergence has been set to: 1e-008
       Parameter Convergence has been set to: 1e-008
                      Default Initial Parameter Values
                         background =
                                                    Specified
                                                Ω
30
                          intercept =
                                         -0.342416
31
                              slope =
                                        0.00261455
32
33
34
               Asymptotic Correlation Matrix of Parameter Estimates
35
36
               ( *** The model parameter(s) -background
37
                     have been estimated at a boundary point, or have been
38
    specified by the user,
39
                     and do not appear in the correlation matrix )
40
41
                  intercept
                                   slope
42
43
     intercept
                          1
                                   -0.51
44
45
         slope
                      -0.51
46
47
48
49
                              Parameter Estimates
50
51
           Variable
                             Estimate
                                                  Std. Err.
52
          intercept
                             -0.374017
                                                  0.335179
53
                             0.00300128
                                                0.00110665
              slope
54
55
56
57
58
                            Analysis of Deviance Table
59
           Model
                      Log(likelihood) Deviance Test DF
                                                           P-value
60
         Full model
                           -33.9494
61
       Fitted model
                                        0.944503
                                                                 0.6236
                           -34.4216
                                                      2
      Reduced model
                           -40.7516
                                         13.6045
62
                                                               0.003496
63
64
                           72.8432
               AIC:
```

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000	0.4076	6.114	5	15	-0.5851
40.0000	0.4368	6.553	8	15	0.7534
200.0000	0.5563	8.345	8	15	-0.1792
1000.0000	0.9326	13.989	14	15	0.0114

Chi-square = 0.94 DF = 2 P-value = 0.6243

Benchmark Dose Computation

Specified effect = 0.1

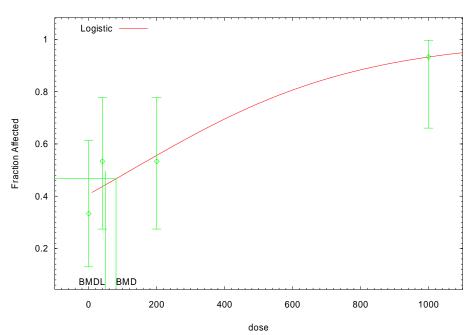
Risk Type = Extra risk

Confidence level = 0.95

BMD = 80.3245

BMDL = 48.7137

Logistic Model with 0.95 Confidence Level



12:16 03/09 2006

```
1
2
3
4
5
     ______
             Logistic Model $Revision: 2.1 $ $Date: 2000/02/26 03:38:20 $
             Input Data File: C:\BMDS\FAWELL_FEMALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_FEMALE_CHRONIC_INFLAMMATION.plt
6
7
                                                Wed Dec 28 15:51:45 2005
     ______
8
9
     BMDS MODEL RUN
10
11
12
        The form of the probability function is:
13
14
        P[response] = background+(1-background)/[1+EXP(-intercept-slope*Log(dose))]
15
16
17
        Dependent variable = COLUMN2
18
        Independent variable = COLUMN1
19
        Slope parameter is restricted as slope >= 1
20
21
22
23
24
25
26
27
28
29
30
        Total number of observations = 4
        Total number of records with missing values = 0
       Maximum number of iterations = 250
        Relative Function Convergence has been set to: 1e-008
        Parameter Convergence has been set to: 1e-008
       User has chosen the log transformed model
31
32
                       Default Initial Parameter Values
33
34
                          background = 0.333333
                           intercept =
                                            -5.18041
35
                               slope =
                                                   1
36
37
38
                Asymptotic Correlation Matrix of Parameter Estimates
39
40
                  background
                                intercept
                                                  slope
41
42
    background
                                    -0.51
                                                  0.47
                           1
43
44
     intercept
                       -0.51
                                                  -0.99
45
46
          slope
                        0.47
                                    -0.99
                                                      1
47
48
49
50
                               Parameter Estimates
51
52
           Variable
                               Estimate
                                                    Std. Err.
53
         background
                                                    0.0985863
                                0.426119
54
           intercept
                                -12.3243
                                                      8.90473
55
                                 2.07233
                                                      1.36766
               slope
56
57
58
                             Analysis of Deviance Table
59
60
                       Log(likelihood) Deviance Test DF
           Model
                                                               P-value
61
          Full model
                            -33.9494
       Fitted model
62
                            -34.5387
                                           1.17872
                                                         1
                                                                    0.2776
63
      Reduced model
                            -40.7516
                                           13.6045
                                                         3
                                                                  0.003496
64
```

1000.0000

AIC: 75.0774

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Scaled Residual	
0.0000	0.4261	6.392	5	15 15	-0.7267 0.7971	
40.0000 200.0000	0.4314 0.5448	6.471 8.172	8 8	15 15	-0.08904	

Chi-square = 1.17 DF = 1P-value = 0.2789

13.966

Benchmark Dose Computation

Specified effect = 0.1

0.9310

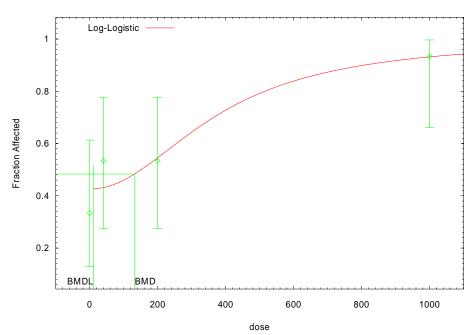
Risk Type Extra risk

Confidence level = 0.95

> BMD = 132.532

BMDL = 11.3311

Log-Logistic Model with 0.95 Confidence Level



15:51 12/28 2005

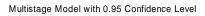
15

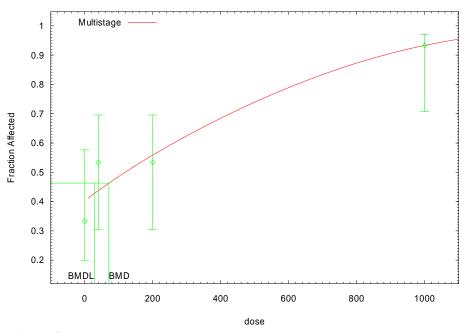
0.03513

```
______
            Multistage Model. $Revision: 2.1 $ $Date: 2000/08/21 03:38:21 $
2
3
4
5
            Input Data File: C:\BMDS\FAWELL FEMALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_FEMALE_CHRONIC_INFLAMMATION.plt
                                               Thu Mar 09 12:17:59 2006
6
7
     ______
8
     BMDS MODEL RUN
9
     10
11
       The form of the probability function is:
12
13
       P[response] = background + (1-background) * [1-EXP(
    -beta1*dose^1-beta2*dose^2-beta3*dose^3)]
14
15
16
       The parameter betas are restricted to be positive
17
18
19
       Dependent variable = COLUMN2
20
       Independent variable = COLUMN1
21
\overline{22}
     Total number of observations = 4
23
     Total number of records with missing values = 0
24
25
26
     Total number of parameters in model = 4
Total number of specified parameters = 0
     Degree of polynomial = 3
27
28
29
     Maximum number of iterations = 250
30
     Relative Function Convergence has been set to: 1e-008
31
     Parameter Convergence has been set to: 1e-008
32
33
34
                      Default Initial Parameter Values
35
                          Background =
                                           0.417955
36
                             Beta(1) =
                                         0.00124572
37
                             Beta(2) =
38
                             Beta(3) = 9.2085e-010
39
40
41
               Asymptotic Correlation Matrix of Parameter Estimates
42
43
                ( *** The model parameter(s) -Beta(2)
44
                     have been estimated at a boundary point, or have been
45
    specified by the user,
46
                      and do not appear in the correlation matrix )
47
48
                 Background
                                  Beta(1)
                                               Beta(3)
49
50
    Background
                                    -0.66
                                                  0.55
51
52
53
       Beta(1)
                      -0.66
                                       1
                                                 -0.93
54
       Beta(3)
                       0.55
                                    -0.93
55
56
57
58
                               Parameter Estimates
59
           Variable
                               Estimate
                                                    Std. Err.
60
         Background
                                0.403311
                                                    0.162784
61
            Beta(1)
                              0.00148059
                                                 0.00267062
            Beta(2)
62
                                                       NΑ
63
            Beta(3)
                            7.06214e-010
                                                2.70652e-009
64
```

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1 2 3 4	implied b	NA - Indicates that this parameter has hit a bound implied by some inequality constraint and thus has no standard error.					
5 6		Į	Analysis of De	viance Table			
7 8	Model	Log(lik	celihood) Dev	riance Test 1	DF P-v	alue	
9 10	Full mode Fitted mode	1 -3 1 -3	33.9494 34.4013 0 40.7516	.903921	1	0.3417	
11 12	Reduced mode	1 -4	10.7516	13.6045	3	0.003496	
13 14	AIC	: 7	74.8026				
15 16		Cood	lnoss of Fit				
17	Goodness of Fit Dose EstProb. Expected Observed Size Chi^2 Res.						
18 19	Dose	EstProb. 	Expected	Observed	Size 	Chi 2 Res.	
20 21		0.4033	6.050	5	15	-0.291	
22 23	i: 2 40.0000	0.4376	6.565	8	15	0.389	
24 25		0.5587	8.381	8	15	-0.103	
26 27	i: 4 1000.0000	0.9330	13.995	14	15	0.005	
28 29 30	Chi-square =	0.90	DF = 1	P-value	= 0.3421		
31	D 1 D						
32 33	Benchmark D	_					
34 35	Specified effe	ct =	0.1				
36 37	Risk Type	= E	Extra risk				
38 39	Confidence lev	el =	0.95				
40 41	В	MD =	70.9904				
41	BM	DL =	28.0638				





2 12:17 03/09 2006

```
______
            Multistage Model. $Revision: 2.1 $ $Date: 2000/08/21 03:38:21 $
2
3
4
5
            Input Data File: C:\BMDS\FAWELL FEMALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_FEMALE_CHRONIC_INFLAMMATION.plt
                                              Wed Dec 28 15:52:16 2005
6
7
     ______
8
     BMDS MODEL RUN
9
     10
11
       The form of the probability function is:
12
13
       P[response] = background + (1-background)*[1-EXP(
    -beta1*dose^1-beta2*dose^2)]
14
15
16
       The parameter betas are restricted to be positive
17
18
19
       Dependent variable = COLUMN2
20
       Independent variable = COLUMN1
21
22
     Total number of observations = 4
23
     Total number of records with missing values = 0
24
25
26
     Total number of parameters in model = 3
Total number of specified parameters = 0
     Degree of polynomial = 2
27
28
29
     Maximum number of iterations = 250
30
     Relative Function Convergence has been set to: 1e-008
31
     Parameter Convergence has been set to: 1e-008
32
33
34
35
                      Default Initial Parameter Values
36
                         Background =
                                          0.417495
37
                            Beta(1) =
                                         0.00110779
38
                            Beta(2) = 1.05844e-006
39
40
41
               Asymptotic Correlation Matrix of Parameter Estimates
42
43
                 Background
                                 Beta(1)
                                              Beta(2)
44
45
    Background
                                   -0.67
                                                 0.58
46
47
       Beta(1)
                      -0.67
                                       1
                                                -0.95
48
49
       Beta(2)
                       0.58
                                   -0.95
                                                     1
50
51
52
53
54
                              Parameter Estimates
55
                                                    Std. Err.
           Variable
                              Estimate
56
57
58
                                                    0.165802
         Background
                               0.402286
            Beta(1)
                             0.00142259
                                                 0.00323206
            Beta(2)
                           7.46169e-007
                                               3.28803e-006
59
60
61
62
                            Analysis of Deviance Table
63
64
                      Log(likelihood) Deviance Test DF
           Model
                                                             P-value
```

```
1 Full model -33.9494
2 Fitted model -34.4149 0.93098 1
3 Reduced model -40.7516 13.6045 3
4
5 AIC: 74.8297
6
7
8 Goodness of Fit
9
10 Dose Est. Prob. Expected Observed S:
```

Dose	EstProb.	Expected	Observed	Size	Chi^2 Res.
i: 1 0.0000 i: 2	0.4023	6.034	5	15	-0.287
40.0000 i: 3	0.4360	6.540	8	15	0.396
200.0000 i: 4	0.5635	8.453	8	15	-0.123
1000.0000	0.9317	13.975	14	15	0.026
Chi-square =	0.93	DF = 1	P-value	= 0.3347	

Benchmark Dose Computation

Specified effect = 0.1

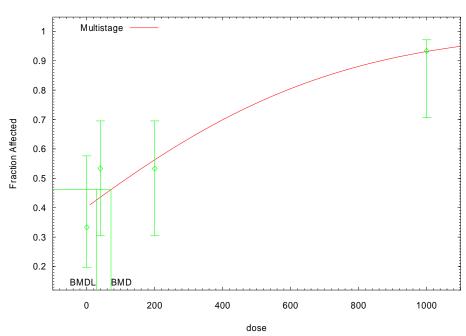
Risk Type = Extra risk

Confidence level = 0.95

BMD = 71.3892

BMDL = 27.9852

Multistage Model with 0.95 Confidence Level



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0.3346

0.003496

```
______
            Multistage Model. $Revision: 2.1 $ $Date: 2000/08/21 03:38:21 $
2
3
4
5
            Input Data File: C:\BMDS\FAWELL FEMALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_FEMALE_CHRONIC_INFLAMMATION.plt
                                              Thu Mar 09 12:20:10 2006
6
7
     ______
8
     BMDS MODEL RUN
9
     10
11
       The form of the probability function is:
12
13
       P[response] = background + (1-background) * [1-EXP(
14
    -beta1*dose^1)]
15
16
       The parameter betas are restricted to be positive
17
18
19
       Dependent variable = COLUMN2
20
       Independent variable = COLUMN1
21
\overline{22}
     Total number of observations = 4
23
     Total number of records with missing values = 0
24
25
26
     Total number of parameters in model = 2
Total number of specified parameters = 0
     Degree of polynomial = 1
27
28
29
     Maximum number of iterations = 250
30
     Relative Function Convergence has been set to: 1e-008
31
     Parameter Convergence has been set to: 1e-008
32
33
34
35
                      Default Initial Parameter Values
36
                         Background = 0.377465
37
                            Beta(1) =
                                        0.00221127
38
39
40
               Asymptotic Correlation Matrix of Parameter Estimates
41
42
                 Background
                                 Beta(1)
43
44
    Background
                                   -0.45
45
46
       Beta(1)
                      -0.45
                                       1
47
48
49
50
                              Parameter Estimates
51
52
           Variable
                              Estimate
                                                   Std. Err.
53
         Background
                                                     0.1375
                               0.385756
54
            Beta(1)
                             0.00207378
                                                0.000944904
55
56
57
58
                            Analysis of Deviance Table
59
60
                      Log(likelihood) Deviance Test DF
           Model
                                                            P-value
61
         Full model
                           -33.9494
       Fitted model
                                          1.01801
                                                       2
62
                           -34.4584
                                                                  0.6011
63
      Reduced model
                           -40.7516
                                          13.6045
                                                       3
                                                                0.003496
64
```

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AIC: 72.9167

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Chi^2 Res.
i: 1 0.0000	0.3858	5.786	5	15	-0.221
i: 2 40.0000 i: 3	0.4347	6.520	8	15	0.402
200.0000 i: 4	0.5943	8.914	8	15	-0.253
1000.0000	0.9228	13.842	14	15	0.148
Chi-square =	1.02	DF = 2	P-value	= 0.5996	

Benchmark Dose Computation

Specified effect = 0.1

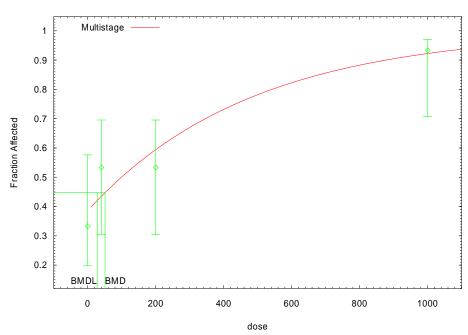
Risk Type = Extra risk

Confidence level = 0.95

BMD = 50.8061

BMDL = 27.7368

Multistage Model with 0.95 Confidence Level

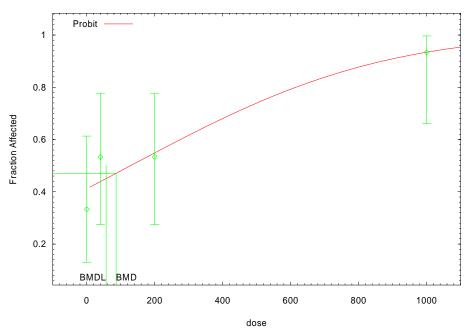


12:20 03/09 2006

```
1
    ______
            Probit Model $Revision: 2.1 $ $Date: 2000/02/26 03:38:53 $
2
3
4
            Input Data File: C:\BMDS\FAWELL FEMALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_FEMALE_CHRONIC_INFLAMMATION.plt
5
                                             Thu Mar 09 12:21:25 2006
6
7
     ______
8
     BMDS MODEL RUN
9
     10
11
       The form of the probability function is:
12
13
       P[response] = CumNorm(Intercept+Slope*Dose),
14
15
       where CumNorm(.) is the cumulative normal distribution function
16
17
18
       Dependent variable = COLUMN2
19
       Independent variable = COLUMN1
20
       Slope parameter is not restricted
21
22
       Total number of observations = 4
23
24
25
26
       Total number of records with missing values = 0
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
       Parameter Convergence has been set to: 1e-008
27
28
29
30
                      Default Initial (and Specified) Parameter Values
31
                         background =
                                                0
                                                    Specified
32
                                        -0.216871
                          intercept =
33
                              slope =
                                        0.00162326
34
35
36
               Asymptotic Correlation Matrix of Parameter Estimates
37
38
               ( *** The model parameter(s) -background
39
                     have been estimated at a boundary point, or have been
40
    specified by the user,
41
                     and do not appear in the correlation matrix )
42
43
                  intercept
                                  slope
44
45
     intercept
                                   -0.53
46
47
                      -0.53
         slope
48
49
50
51
                              Parameter Estimates
52
53
           Variable
                                                  Std. Err.
                             Estimate
54
          intercept
                             -0.223386
                                                   0.20693
55
                             0.00172875
                                              0.000560405
              slope
56
57
58
59
                           Analysis of Deviance Table
60
61
           Model
                      Log(likelihood) Deviance Test DF
                                                           P-value
         Full model
                          -33.9494
62
63
       Fitted model
                           -34.421
                                        0.943277
                                                      2
                                                                  0.624
      Reduced model
                           -40.7516
                                         13.6045
64
                                                      3
                                                               0.003496
```

1 2 3 4 5 6 7 8 9	<i>2</i> 4	AIC:	72.842			
5		G	oodness of I	Fit		
6 7						Scaled
8	Dose	EstPro	b. Expected	d Observed	Size	Residual
	0.0000	0.4116	6.1		15	-0.6161
11	40.0000	0.4387			15	
12	200.0000					-0.1196
13	1000.0000	0.9339	14.00	08 14	15	-0.00859
14 15	Chi-square	= 0.	94 DF = 2	P-valu	e = 0.6252	
16						
17 18	Renchmark	Dose Comp	utation			
19	Bellelimari	r bobe comp	46461011			
20	Specified ef	fect =	0.1			
21						
22 23	Risk Type	=	Extra risk			
23 24	Confidence l	evel =	0.95			
25	confidence i		0.33			
26		BMD =	86.341			
27 28		BMDL =	56.9234			

Probit Model with 0.95 Confidence Level



1 12:21 03/09 2006

```
1
    ______
2
3
4
5
            Probit Model $Revision: 2.1 $ $Date: 2000/02/26 03:38:53 $
            Input Data File: C:\BMDS\FAWELL FEMALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_FEMALE_CHRONIC_INFLAMMATION.plt
                                              Wed Dec 28 15:52:51 2005
6
7
     ______
8
     BMDS MODEL RUN
9
     10
11
       The form of the probability function is:
12
13
       P[response] = Background
14
                   + (1-Background) * CumNorm(Intercept+Slope*Log(Dose)),
15
16
       where CumNorm(.) is the cumulative normal distribution function
17
18
19
       Dependent variable = COLUMN2
20
       Independent variable = COLUMN1
21
22
23
24
25
26
       Slope parameter is restricted as slope >= 1
       Total number of observations = 4
       Total number of records with missing values = 0
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
27
28
29
30
       Parameter Convergence has been set to: 1e-008
31
       User has chosen the log transformed model
32
33
34
                      Default Initial (and Specified) Parameter Values
35
                         background =
                                          0.333333
36
                          intercept =
                                          -5.62294
37
                              slope =
38
39
40
               Asymptotic Correlation Matrix of Parameter Estimates
41
42
                 background
                               intercept
                                                slope
43
44
    background
                                   -0.46
                                                0.43
45
46
     intercept
                      -0.46
                                                -0.99
                                       1
47
48
                       0.43
                                   -0.99
         slope
                                                    1
49
50
51
52
53
                              Parameter Estimates
54
           Variable
                              Estimate
                                                  Std. Err.
55
                                                  0.0937984
         background
                               0.431473
56
57
58
                                 -7.732
          intercept
                                                    4.97551
              slope
                                 1.2906
                                                    0.75516
59
60
61
                            Analysis of Deviance Table
62
63
           Model
                      Log(likelihood) Deviance Test DF
                                                            P-value
64
                           -33.9494
         Full model
```

Fitted model -34.5602 1.22166 1 0.269 Reduced model -40.7516 13.6045 3 0.003496

AIC: 75.1204

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000	0.4315 0.4323	6.472 6.485	5	15 15	-0.7674 0.7897
200.0000	0.5370 0.9327	8.055 13.991	8 14	15 15	-0.02871 0.009746
Chi-square =	= 1.21	DF = 1	P-value	= 0.2706	

Benchmark Dose Computation

Specified effect = 0.1

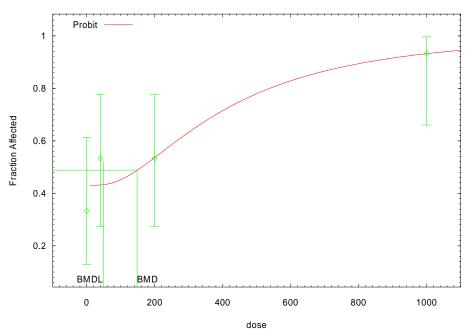
Risk Type = Extra risk

Confidence level = 0.95

BMD = 148.118

BMDL = 48.5214

Probit Model with 0.95 Confidence Level



15:52 12/28 2005

```
______
            Quantal Linear Model $Revision: 2.2 $ $Date: 2000/03/17 22:27:16 $
2
3
4
5
            Input Data File: C:\BMDS\FAWELL FEMALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_FEMALE_CHRONIC_INFLAMMATION.plt
                                             Wed Dec 28 15:54:53 2005
6
7
     ______
8
     BMDS MODEL RUN
9
     10
11
       The form of the probability function is:
12
13
       P[response] = background + (1-background)*[1-EXP(-slope*dose)]
14
15
16
       Dependent variable = COLUMN2
17
       Independent variable = COLUMN1
18
19
       Total number of observations = 4
20
       Total number of records with missing values = 0
21
22
23
24
25
26
27
28
29
30
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
       Parameter Convergence has been set to: 1e-008
                      Default Initial (and Specified) Parameter Values
                         Background = 0.34375
                              Slope =
                                       0.00194591
                              Power =
                                                    Specified
31
32
33
               Asymptotic Correlation Matrix of Parameter Estimates
34
35
               ( *** The model parameter(s) -Power
36
                     have been estimated at a boundary point, or have been
37
    specified by the user,
38
                     and do not appear in the correlation matrix )
39
40
                 Background
                                  Slope
41
42
    Background
                         1
                                  -0.42
43
44
         Slope
                      -0.42
45
46
47
48
                             Parameter Estimates
49
50
           Variable
                             Estimate
                                                  Std. Err.
51
         Background
                              0.385756
                                                 0.0913888
52
53
54
55
              Slope
                            0.00207382
                                               0.000852767
56
                           Analysis of Deviance Table
57
58
           Model
                      Log(likelihood) Deviance Test DF
                                                          P-value
59
         Full model
                          -33.9494
60
       Fitted model
                                         1.01801
                                                      2
                                                                 0.6011
                          -34.4584
61
      Reduced model
                                         13.6045
                                                     3
                                                              0.003496
                          -40.7516
62
63
               AIC:
                           72.9167
64
```

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000	0.3858	5.786	5	15	-0.4171
40.0000	0.4347	6.520	8	15	0.771
200.0000	0.5943	8.914	8	15	-0.4808
1000.0000	0.9228	13.842	14	15	0.153

Chi-square = 1.02 DF = 2 P-value = 0.5996

Benchmark Dose Computation

Specified effect = 0.1

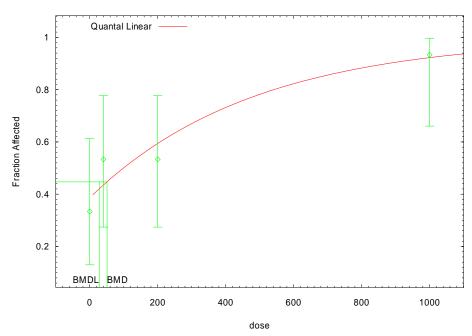
Risk Type = Extra risk

Confidence level = 0.95

BMD = 50.8051

BMDL = 27.7368

Quantal Linear Model with 0.95 Confidence Level



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15:54 12/28 2005

```
1
    ______
            Quantal Quadratic Model $Revision: 2.2 $ $Date: 2000/03/17 22:27:16 $
2
3
4
5
            Input Data File: C:\BMDS\FAWELL FEMALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_FEMALE_CHRONIC_INFLAMMATION.plt
                                             Wed Dec 28 15:55:22 2005
6
7
     ______
8
     BMDS MODEL RUN
9
     10
11
       The form of the probability function is:
12
13
       P[response] = background + (1-background)*[1-EXP(-slope*dose^2)]
14
15
16
       Dependent variable = COLUMN2
17
       Independent variable = COLUMN1
18
19
       Total number of observations = 4
20
       Total number of records with missing values = 0
21
22
23
24
25
26
27
28
29
30
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
       Parameter Convergence has been set to: 1e-008
                      Default Initial (and Specified) Parameter Values
                         Background = 0.34375
                              Slope = 1.94591e-006
                              Power =
                                                    Specified
31
32
33
               Asymptotic Correlation Matrix of Parameter Estimates
34
35
               ( *** The model parameter(s) -Power
36
                     have been estimated at a boundary point, or have been
37
    specified by the user,
38
                     and do not appear in the correlation matrix )
39
40
                 Background
                                  Slope
41
42
    Background
                         1
                                  -0.23
43
44
         Slope
                      -0.23
45
46
47
48
                             Parameter Estimates
49
50
           Variable
                             Estimate
                                                 Std. Err.
51
         Background
                              0.449003
                                                 0.0779228
52
53
54
55
              Slope
                           2.16129e-006
                                              1.00801e-006
56
                           Analysis of Deviance Table
57
58
           Model
                      Log(likelihood) Deviance Test DF
                                                          P-value
59
         Full model
                          -33.9494
60
       Fitted model
                                         1.33385
                                                      2
                                                                 0.5133
                          -34.6163
61
      Reduced model
                                         13.6045
                                                     3
                                                              0.003496
                          -40.7516
62
63
               AIC:
                           73.2326
64
```

1
2
3
4

Goodness of Fit

Dose	EstProb.	Expected	Observed	Size	Scaled Residual
0.0000	0.4490	6.735	5	15	-0.9007
	0.4509	6.764	8	15	0.6416
200.0000	0.4946	7.420	8	15	0.2998
	0.9365	14.048	14	15	-0.05091
Chi-square =	1.32	DF = 2	P-value	= 0.5181	

Benchmark Dose Computation

Specified effect = 0.1

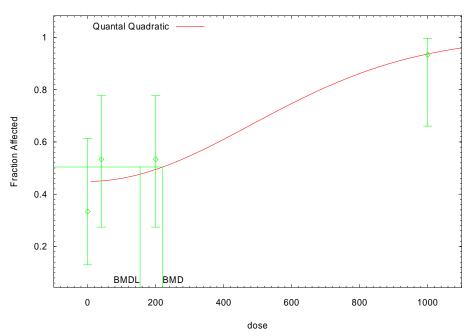
Risk Type = Extra risk

Confidence level = 0.95

BMD = 220.792

BMDL = 153.94

Quantal Quadratic Model with 0.95 Confidence Level



26 15:55 12/28 2005

```
1
2
3
4
5
    ______
            Weibull Model $Revision: 2.2 $ $Date: 2000/03/17 22:27:16 $
            Input Data File: C:\BMDS\FAWELL_FEMALE_CHRONIC_INFLAMMATION.(d)
Gnuplot Plotting File: C:\BMDS\FAWELL_FEMALE_CHRONIC_INFLAMMATION.plt
6
7
                                               Wed Dec 28 15:55:56 2005
     ______
8
9
     BMDS MODEL RUN
10
11
12
       The form of the probability function is:
13
14
       P[response] = background + (1-background)*[1-EXP(-slope*dose^power)]
15
16
17
       Dependent variable = COLUMN2
18
       Independent variable = COLUMN1
19
        Power parameter is restricted as power >=1
20
21
22
23
24
25
26
27
28
29
30
        Total number of observations = 4
       Total number of records with missing values = 0
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
       Parameter Convergence has been set to: 1e-008
                       Default Initial (and Specified) Parameter Values
                          Background =
                                            0.34375
31
                               Slope = 0.000217656
32
                                            1.31712
                               Power =
33
34
35
               Asymptotic Correlation Matrix of Parameter Estimates
36
37
                  Background
                                    Slope
                                                 Power
38
39
    Background
                           1
                                    -0.77
                                                  0.76
40
41
         Slope
                       -0.77
                                       1
                                                    -1
42
43
         Power
                       0.76
                                       -1
                                                     1
44
45
46
47
                               Parameter Estimates
48
49
           Variable
                              Estimate
                                                    Std. Err.
50
         Background
                               0.396812
                                                    0.136488
51
               Slope
                              0.00106435
                                                  0.00719029
52
53
54
              Power
                                 1.09925
                                                    0.999439
55
56
                             Analysis of Deviance Table
57
58
           Model
                       Log(likelihood) Deviance Test DF
                                                             P-value
59
         Full model
                            -33.9494
60
       Fitted model
                                          1.00699
                                                     1
                                                                    0.3156
                            -34.4529
                                                        3
61
      Reduced model
                                           13.6045
                                                                 0.003496
                            -40.7516
62
63
               AIC:
                            74.9057
64
```

Goodness of Fit

	Dose	EstProb.	Expected	Observed	Size	Scaled Residual
(0.000	0.3968	5.952	5	15	-0.5025
4	0.000	0.4327	6.491	8	15	0.7864
20	0.000	0.5792	8.689	8	15	-0.3601
100	0.000	0.9271	13.906	14	15	0.09327

Chi-square = 1.01 DF = 1 P-value = 0.3151

Benchmark Dose Computation

Specified effect = 0.1

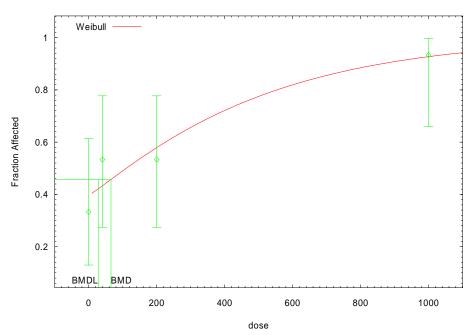
Risk Type = Extra risk

Confidence level = 0.95

BMD = 65.3741

BMDL = 27.7679

Weibull Model with 0.95 Confidence Level



15:55 12/28 2005

```
Heinze, 1999 Relative Liver Weight Changes
1
2
3
4
5
    ______
            Polynomial Model. Revision: 2.2 Date: 9/12/2002
            Input Data File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY DOCUMENTS\ CYANO
6
7
    TOX REV\MODELING\HEINZE ENZYMES AND LIVER WT. (d)
    Gnuplot Plotting File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY DOCUMENTS\_CYANO TOX REV\MODELING\HEINZE_ENZYMES_AND_LIVER_WT.plt
8
9
                                             Thu May 18 09:37:38 2006
10
     11
12
     BMDS MODEL RUN
13
    14
15
       The form of the response function is:
16
17
       Y[dose] = beta 0 + beta 1*dose + beta 2*dose^2 + ...
18
19
20
       Dependent variable = MEAN
21
22
       Independent variable = COLUMN1
       rho is set to 0
23
24
25
26
       The polynomial coefficients are restricted to be positive
       A constant variance model is fit
       Total number of dose groups = 3 Total number of records with missing values = 0
27
28
29
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
30
       Parameter Convergence has been set to: 1e-008
31
32
33
34
                      Default Initial Parameter Values
35
                              alpha =
                                      0.1466
36
                                rho =
                                                0
                                                    Specified
37
                             beta 0 =
                                          2.84857
38
                             beta 1 =
                                       0.00447143
39
40
41
42
                                     Parameter Estimates
43
44
                                                            95.0% Wald Confidence
45
    Interval
46
                                           Std. Err.
                           Estimate
                                                        Lower Conf. Limit
           Variable
47
    Upper Conf. Limit
48
                           0.143276
                                          0.0369936
                                                              0.0707695
              alpha
49
    0.215782
50
             beta 0
                           2.84857
                                           0.101163
                                                                  2.6503
51
    3.04685
52
             beta_1
                         0.00447143
                                         0.00110819
                                                              0.00229942
53
    0.00664343
54
55
56
               Asymptotic Correlation Matrix of Parameter Estimates
57
58
                      alpha
                                 beta 0
                                              beta 1
59
         alpha
                        1
                                5.6e-009
                                            1.9e-009
60
                   5.6e-009
                                               -0.73
        beta 0
                                   1
61
        beta 1
                   1.9e-009
                                  -0.73
                                                   1
62
63
```

Table of Data and Estimated Values of Interest

```
2
     Dose
               N Obs Mean Obs Std Dev Est Mean Est Std Dev Chi^2
    Res.
4
                      _____
                                   _____
                                                 _____
                                                             _____
                                                                            _____
5
                     2.75
                                                             0.379
                                                2.85
3.07
3.52
6
7
        0
              10
                                    0.29
                                                                             -0.824
       50
              10
                       3.22
                                     0.34
                                                  3.07
                                                              0.379
                                                                               1.24
8
                                                                             -0.412
       150
              10
                       3.47
                                     0.49
                                                              0.379
9
10
11
12
      Model Descriptions for likelihoods calculated
13
14
                Yij = Mu(i) + e(ij)

Var\{e(ij)\} = Sigma^2
15
     Model A1:
16
17
                Yij = Mu(i) + e(ij)
Var\{e(ij)\} = Sigma(i)^2
18
19
20
21
     Model R:
                        Yi = Mu + e(i)
22
                 Var\{e(i)\} = Sigma^2
23
24
25
26
                            Likelihoods of Interest
27
28
                 Model
                            Log(likelihood)
                                               \mathsf{DF}
                                                         AIC
                               15.381120
                                                      -22.762240
                 A1
                                               4
29
                  A2
                               16.880747
                                                      -21.761494
                                               6
                               14.144766
30
                fitted
                                               2
                                                      -24.289533
31
                                 7.133405
                                                      -10.266809
32
33
     Test 1: Does response and/or variances differ among dose
34
    levels
35
               (A2 vs. R)
36
      Test 2: Are Variances Homogeneous (A1 vs A2)
      Test 3: Does the Model for the Mean Fit (A1 vs. fitted)
37
38
39
                          Tests of Interest
40
41
               -2*log(Likelihood Ratio) Test df
       Test
                                                      p-value
42
43
                                                         <.0001
                             19.4947
                                              4
       Test 1
       Test 2
44
                             2.99925
                                              2
                                                          0.2232
45
       Test 3
                             2.47271
                                                          0.1158
46
47
    The p-value for Test 1 is less than .05. There appears
48
    to be a
49
    difference between response and/or variances among the
50
    dose levels.
51
    It seems appropriate to model the data
52
53
    The p-value for Test 2 is greater than .05. A
54
    homogeneous variance
55
    model appears to be appropriate here
56
57
58
    The p-value for Test 3 is greater than .05. The model
59
    chosen appears
60
    to adequately describe the data
61
62
63
```

Benchmark Dose Computation

Specified effect =

Estimated standard deviations from the control mean Risk Type

Confidence level = 0.95

> BMD = 84.6525

BMDL = 57.9321

Linear Model with 0.95 Confidence Level

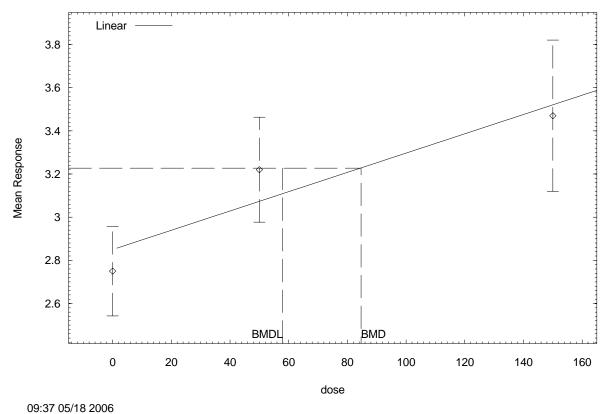


Table of Data and Estimated Values of Interest

```
1
 2
     Dose
                N Obs Mean Obs Std Dev Est Mean Est Std Dev Chi^2
     Res.
 4
                       _____
                                    _____
                                                   _____
                                                               _____
                                                                              -----
 5
                                                               5.39
                                                  20.2
                      16.6
 6
7
        0
              10
                                     4.48
                                                                                  -2.1
        50
              10
                        30.6
                                      5.05
                                                    25.3
                                                                 5.39
                                                                                 3.15
8
                                                                 5.39
                                                                                -1.05
                                                    35.4
       150
              10
                        33.6
                                      1.16
9
10
11
12
       Model Descriptions for likelihoods calculated
13
14
                Yij = Mu(i) + e(ij)

Var\{e(ij)\} = Sigma^2
15
      Model A1:
16
17
                Yij = Mu(i) + e(ij)
Var\{e(ij)\} = Sigma(i)^2
18
19
20
21
      Model R:
                         Yi = Mu + e(i)
22
                 Var\{e(i)\} = Sigma^2
23
24
25
26
                             Likelihoods of Interest
27
28
                 Model
                             Log(likelihood)
                                                \mathsf{DF}
                                                          AIC
                               -54.666589
                                                       117.333177
                  A1
                                                4
29
                  A2
                               -46.093905
                                                       104.187810
                                                 6
30
                 fitted
                               -65.523922
                                                2
                                                       135.047844
31
                               -78.954450
                                                       161.908899
32
33
      Test 1: Does response and/or variances differ among dose
34
     levels
35
               (A2 vs. R)
36
      Test 2: Are Variances Homogeneous (A1 vs A2)
Test 3: Does the Model for the Mean Fit (A1 vs. fitted)
37
38
39
                           Tests of Interest
40
41
        Test -2*log(Likelihood Ratio) Test df
                                                        p-value
42
43
                                                           <.0001
        Test 1
                             65.7211
                                               4
        Test 2
44
                             17.1454
                                               2
                                                        0.0001892
45
        Test 3
                             21.7147
                                               1
                                                           < .0001
46
47
     The p-value for Test 1 is less than .05. There appears
48
     to be a
49
     difference between response and/or variances among the
50
     dose levels.
51
     It seems appropriate to model the data
52
53
     The p-value for Test 2 is less than .05. Consider
54
     running a
55
     non-homogeneous variance model
56
57
     The p-value for Test 3 is less than .05. You may want
58
     to try a
59
     different model
60
61
62
     Benchmark Dose Computation
63
     Specified effect =
64
```

1 2 3	Risk Type		=	Estimated	standard	deviations	from	the	control	mean
4 5	Confidence	level	=	0.95						
6 7		BMD	=	53.3442						
8 9		BMDL	=	39.9199						

```
1
2
3
4
5
    ______
            Polynomial Model. Revision: 2.2 Date: 9/12/2002
            Input Data File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY DOCUMENTS\ CYANO
    TOX REV\MODELING\HEINZE ENZYMES AND LIVER WT.(d)
            Gnuplot Plotting File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY
6
7
    DOCUMENTS\_CYANO TOX REV\MODELING\HEINZE_ENZYMES_AND_LIVER_WT.plt
                                              Thu May 18 09:42:43 2006
8
     ______
9
10
     BMDS MODEL RUN
11
12
13
       The form of the response function is:
14
15
       Y[dose] = beta 0 + beta 1*dose + beta 2*dose^2 + ...
16
17
18
       Dependent variable = MEAN
19
       Independent variable = COLUMN1
20
       The polynomial coefficients are restricted to be positive
21
22
       The variance is to be modeled as Var(i) = alpha*mean(i)^rho
23
24
25
26
       Total number of dose groups = 3
       Total number of records with missing values = 0
       Maximum number of iterations = 250
       Relative Function Convergence has been set to: 1e-008
27
28
29
30
       Parameter Convergence has been set to: 1e-008
31
                      Default Initial Parameter Values
32
                              alpha =
                                          15.6395
33
                                rho =
34
                                             20.22
                             beta_0 =
35
                             beta 1 =
                                             0.101
36
37
38
39
                                     Parameter Estimates
40
41
                                                             95.0% Wald Confidence
42
    Interval
43
                                            Std. Err.
           Variable
                           Estimate
                                                         Lower Conf. Limit
44
    Upper Conf. Limit
45
              alpha
                        2.44639e+006
                                         4.30425e+006
                                                            -5.98979e+006
46
    1.08826e+007
47
                rho
                            -3.55294
                                            0.556161
                                                                   -4.643
48
    -2.46289
49
             beta 0
                            21.3722
                                              1.64609
                                                                   18.146
50
    24.5985
51
                           0.0863674
                                            0.0134905
                                                               0.0599266
             beta 1
52
53
    0.112808
54
55
               Asymptotic Correlation Matrix of Parameter Estimates
56
57
58
                                     rho
                                               beta 0
                      alpha
                                                            beta 1
                                                             -0.\overline{2}3
         alpha
                                                0.\overline{1}9
                         1
                                   -0.99
59
           rho
                      -0.99
                                      1
                                                -0.21
                                                             0.25
60
                      0.19
        beta 0
                                   -0.21
                                                 1
                                                             -0.89
61
        beta 1
                      -0.23
                                    0.25
                                                -0.89
                                                                 1
62
63
```

Table of Data and Estimated Values of Interest

```
2
       Dose
                    N Obs Mean Obs Std Dev Est Mean Est Std Dev Chi^2
      Res.
 4
                             _____
                                             _____
                                                                _____
                                                                               _____
 5
                                                              21.4
25 7
                           16.6
                                                                                6.79
                                             4.48
 6
7
           0
                  10
                                                                                                    -2.2
3.2
          50
                  10
                              30.6
                                                5.05
                                                                                   4.9
 8
                                                                 34.3
                                                                                 2.93
                                                                                                   -0.808
         150
                  10
                              33.6
                                                1.16
 9
10
11
12
       Model Descriptions for likelihoods calculated
13
14
                     \label{eq:condition} \begin{array}{rcl} \text{Yij} &=& \text{Mu(i)} + \text{e(ij)} \\ \text{Var}\{\text{e(ij)}\} &=& \text{Sigma^2} \end{array}
15
       Model A1:
16
17
18
                     \label{eq:continuous} \begin{array}{rcl} \mbox{Yij} &=& \mbox{Mu(i)} + \mbox{e(ij)} \\ \mbox{Var} \left\{ \mbox{e(ij)} \right\} &=& \mbox{Sigma(i)} \mbox{^2} \end{array}
19
20
                     Yij = Mu(i) + e(ij)

Var\{e(ij)\} = alpha*(Mu(i))^rho
21
       Model A3:
22
23
24
25
26
                      Yi = Mu + e(i)

Var\{e(i)\} = Sigma^2
       Model R:
27
28
29
                                     Likelihoods of Interest
30
                      Model
                                     Log(likelihood) DF
                                                                         AIC
                                       -54.666589 4 117.333177
-46.093905 6 104.187810
-54.022155 5 118.044309
-61.092641 4 130.185281
-78.954450 2 161.908899
31
                       A1
32
                       A2
33
                       A3
34
                     fitted
35
                       R
36
37
38
                                Explanation of Tests
39
40
       Test 1: Does response and/or variances differ among Dose
41
      levels?
42
                   (A2 vs. R)
43
       Test 2: Are Variances Homogeneous? (A1 vs A2)
       Test 3: Are variances adequately modeled? (A2 vs. A3)
Test 4: Does the Model for the Mean Fit? (A3 vs. fitted)
44
45
46
47
                                  Tests of Interest
48
49
                    -2*log(Likelihood Ratio) Test df
          Test
                                                                           p-value
50
51
          Test 1
                                     65.7211
                                                          4
                                                                           < .0001
52
                                                         2
          Test 2
                                     17.1454
                                                                       0.0001892
53
          Test 3
                                     15.8565
                                                            1
                                                                          < .0001
54
          Test 4
                                      14.141
                                                            1
                                                                       0.0001696
55
56
      The p-value for Test 1 is less than .05. There appears
57
58
      difference between response and/or variances among the
59
      dose levels
60
      It seems appropriate to model the data
61
62
      The p-value for Test 2 is less than .05. A
63
      non-homogeneous variance
64
      model appears to be appropriate
```

```
1
2
3
4
5
6
7
8
9
10
     The p-value for Test 3 is less than .05. You may want
     to consider a
     different variance model
     The p-value for Test 4 is less than .05. You may want to try a different \,
     model
11
      Benchmark Dose Computation
12
     Specified effect =
13
14
                                 Estimated standard deviations from the control mean
     Risk Type
15
16
17
     Confidence level =
                                       0.95
18
19
20
21
22
23
24
25
26
27
                     BMD =
                                   78.6091
                    BMDL =
                                 63.5158
```

```
Heinze, 1999 Alkaline Phosphatase Changes
1
2
3
4
5
    ______
           Polynomial Model. Revision: 2.2 Date: 9/12/2002
            Input Data File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY DOCUMENTS\ CYANO
6
7
    TOX REV\MODELING\HEINZE_ENZYMES_AND_LIVER_WT.(d)
8
            Gnuplot Plotting File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY
9
    DOCUMENTS\ CYANO TOX REV\MODELING\HEINZE ENZYMES AND LIVER WT.plt
10
                                            Thu May 18 09:46:03 2006
11
     ______
12
13
     BMDS MODEL RUN
14
    15
16
       The form of the response function is:
17
18
       Y[dose] = beta 0 + beta 1*dose + beta 2*dose^2 + ...
19
20
21
       Dependent variable = MEAN
\overline{22}
       Independent variable = COLUMN1
23
       rho is set to 0
24
25
26
       The polynomial coefficients are restricted to be positive
       A constant variance model is fit
27
28
       Total number of dose groups = 3
       Total number of records with missing values = 0
29
       Maximum number of iterations = 250
30
       Relative Function Convergence has been set to: 1e-008
31
       Parameter Convergence has been set to: 1e-008
32
33
34
35
                     Default Initial Parameter Values
36
                             alpha =
                                         7.59287
37
                              rho =
                                              0
                                                  Specified
38
                            beta 0 =
                                         10.6414
39
                            beta 1 =
40
41
42
43
                                   Parameter Estimates
44
45
                                                          95.0% Wald Confidence
46
    Interval
47
                                         Std. Err.
                                                      Lower Conf. Limit
          Variable
                          Estimate
48
    Upper Conf. Limit
49
             alpha
                            7.93453
                                            2.04869
                                                              3.91918
50
    11.9499
51
            beta 0
                           10.6414
                                            0.75283
                                                               9.16591
52
    12.117
53
            beta_1
                                         0.00824683
                                                            0.00186506
                          0.0180286
54
    0.0341921
55
56
57
58
              Asymptotic Correlation Matrix of Parameter Estimates
59
                     alpha
                                beta 0
                                            beta 1
60
                               1.1e-006
                                          -4.7e-007
        alpha
61
        beta 0
                  1.1e-006
                                 1
                                             -0.73
62
        beta 1
                 -4.7e-007
                                 -0.73
63
```

```
Table of Data and Estimated Values of Interest
2
3
4
     Dose
               N
                      Obs Mean
                                  Obs Std Dev Est Mean
                                                            Est Std Dev
                                                                           Chi^2
    Res.
5
6
7
        0
              10
                      9.67
                                     2.2
                                                  10.6
                                                               2.82
                                                                              -1.09
8
       50
              10
                        13
                                     3.81
                                                  11.5
                                                               2.82
                                                                              1.64
9
                                    1.85
                                                 13.3
                                                               2.82
       150
              10
                       12.9
                                                                             -0.545
10
11
12
13
      Model Descriptions for likelihoods calculated
14
15
                16
     Model A1:
17
18
19
                       Yij = Mu(i) + e(ij)
     Model A2:
20
                Var\{e(ij)\} = Sigma(i)^2
21
\overline{22}
                        Yi = Mu + e(i)
     Model R:
23
24
25
26
                 Var\{e(i)\} = Sigma^2
                            Likelihoods of Interest
27
28
29
                 Model
                            Log(likelihood)
                                               DF
                                                         AIC
                 A1
                              -43.827730
                                               4
                                                       95.655461
30
                  A2
                              -40.832314
                                                6
                                                       93.664628
31
                fitted
                              -46.068366
                                               2
                                                       96.136733
32
                              -48.794188
                                                2
                  R
                                                      101.588375
33
34
     Test 1: Does response and/or variances differ among dose
35
     levels
36
               (A2 vs. R)
37
     Test 2: Are Variances Homogeneous (A1 vs A2)
38
     Test 3: Does the Model for the Mean Fit (A1 vs. fitted)
39
40
                          Tests of Interest
41
42
               -2*log(Likelihood Ratio) Test df
       Test
                                                      p-value
43
44
                            15.9237
                                              4
                                                      0.0003485
       Test 1
45
        Test 2
                            5.99083
                                              2
                                                        0.05002
46
       Test 3
                            4.48127
                                                        0.03427
                                              1
47
48
    The p-value for Test 1 is less than .05. There appears
49
50
    difference between response and/or variances among the
51
    dose levels.
52
    It seems appropriate to model the data
53
54
    The p-value for Test 2 is greater than .05. A
55
    homogeneous variance
56
    model appears to be appropriate here
57
58
59
    The p-value for Test 3 is less than .05. You may want
60
    to try a
61
    different model
62
63
64
     Benchmark Dose Computation
```

1 2 3 4 5 6 7	Specified effect	=	1						
	Risk Type	=	Estimated	standard	deviations	from	the	control	mean
	Confidence level	=	0.95						
8 9	BMD	=	156.243						
10 11 12 13 14 15	BMDL	=	87.6609						

```
Fawell et al., 1999 Male Alanine Aminotransferase Changes
1
2
3
    ______
4
           Polynomial Model. Revision: 2.2 Date: 9/12/2002
5
           Input Data File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY
6
    DOCUMENTS\_CYANO TOX REV\MODELING\FAWELL_MALE_ALT.(d)
7
           Gnuplot Plotting File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY
8
    DOCUMENTS\_CYANO TOX REV\MODELING\FAWELL_MALE_ALT.plt
9
                                          Tue May 09 12:59:55 2006
10
     ______
11
12
    BMDS MODEL RUN
13
14
15
      The form of the response function is:
16
17
      Y[dose] = beta_0 + beta_1*dose + beta_2*dose^2 + ...
18
19
20
      Dependent variable = MEAN
21
      Independent variable = COLUMN1
22
      rho is set to 0
23
      The polynomial coefficients are restricted to be positive
24
      A constant variance model is fit
25
26
      Total number of dose groups = 4
27
      Total number of records with missing values = 0
28
      Maximum number of iterations = 250
29
      Relative Function Convergence has been set to: 1e-008
30
      Parameter Convergence has been set to: 1e-008
31
32
33
34
                    Default Initial Parameter Values
                                      1
35
                           alpha =
36
                             rho =
                                           0
                                                Specified
                          beta_0 = 30.4717
37
38
                          beta 1 =
                                    0.129124
39
40
41
42
                                  Parameter Estimates
43
44
                                                       95.0% Wald
45
    Confidence Interval
46
          Variable
                      Estimate Std. Err. Lower Conf. Limit
47
    Upper Conf. Limit
48
             alpha
                       1584.79
                                           289.34
                                                            1017.69
49
    2151.88
50
                      30.4717
                                        6.47011
            beta_0
                                                           17.7905
51
    43.1529
52
                        0.129124
                                      0.0126792
                                                           0.104273
            beta_1
53
    0.153974
54
55
56
              Asymptotic Correlation Matrix of Parameter Estimates
57
```

```
beta_0 beta_1 -6.1e-008 9.2e-007
1
                    alpha
                  1
2
        alpha
3
                             1
       beta_0
                 -6.1e-008
                                            -0.61
4
                 9.2e-007
                                -0.61
       beta_1
                                              1
5
6
7
        Table of Data and Estimated Values of Interest
8
9
              N
                 Obs Mean Obs Std Dev Est Mean Est Std Dev Chi^2
    Dose
10
    Res.
11
12
13
                               8
17.2
14
      0
          15
                     27
                                           30.5
                                                       39.8
                                                                    -0.338
15
      40
           15
                     37
                                           35.6
                                                       39.8
                                                                     0.133
                                28
75
                     59
                                           56.3
                                                        39.8
                                                                     0.263
16
      200
            15
                                            160
                                                     39.8
17
     1000
          15
                   159
                                                                  -0.0579
18
19
20
21
     Model Descriptions for likelihoods calculated
22
23
24
     Model A1:
                   Yij = Mu(i) + e(ij)
25
             Var\{e(ij)\} = Sigma^2
26
27
     Model A2:
                   Yij = Mu(i) + e(ij)
28
              Var\{e(ij)\} = Sigma(i)^2
29
30
               Yi = Mu + e(i)
31
               Var\{e(i)\} = Sigma^2
32
33
34
                         Likelihoods of Interest
35
36
                        Log(likelihood) DF
               Model
                                                  AIC
                                               511.887866
37
                         -250.943933 5
               A1
                                       8
38
                A2
                          -216.540867
                                                449.081734
                         -251.046212
-281.663312
                                         2
39
              fitted
                                               506.092423
40
                                         2
                                                567.326624
41
42
    Test 1: Does response and/or variances differ among dose
43
    levels
44
             (A2 vs. R)
45
     Test 2: Are Variances Homogeneous (A1 vs A2)
46
    Test 3: Does the Model for the Mean Fit (Al vs. fitted)
47
48
                       Tests of Interest
49
50
      Test -2*log(Likelihood Ratio) Test df
                                               p-value
51
                                       6
52
                         130.245
      Test 1
                                                  <.0001
53
                         68.8061
                                       3
      Test 2
                                                  <.0001
                                     2
54
      Test 3
                        0.204557
                                                   0.9028
55
56
    The p-value for Test 1 is less than .05. There appears
```

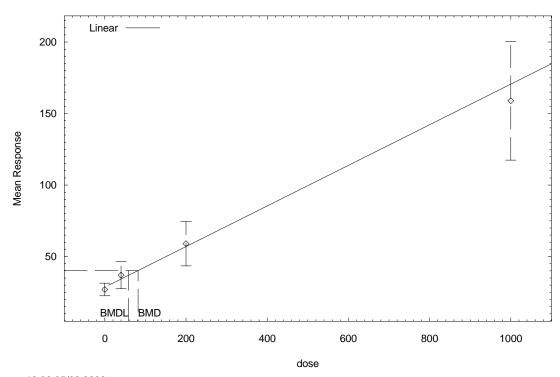
to be a

```
1
    difference between response and/or variances among the
2
    dose levels.
3
    It seems appropriate to model the data
4
5
    The p-value for Test 2 is less than .05. Consider
6
    running a
7
    non-homogeneous variance model
8
9
    The p-value for Test 3 is greater than .05. The model
10
    chosen appears
11
    to adequately describe the data
12
13
14
15
     Benchmark Dose Computation
16
    Specified effect =
                                     1
17
18
    Risk Type
                            Estimated standard deviations from the control mean
19
20
21
    Confidence level =
                                 0.95
22
23
                  BMD =
                              308.304
24
25
                 BMDL =
                              252.245
26
27
```

```
1
2
    ______
3
            Polynomial Model. Revision: 2.2 Date: 9/12/2002
4
            Input Data File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY
5
    DOCUMENTS\ CYANO TOX REV\MODELING\FAWELL MALE ALT.(d)
6
            Gnuplot Plotting File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY
7
    DOCUMENTS\ CYANO TOX REV\MODELING\FAWELL MALE ALT.plt
8
                                           Tue May 09 13:00:39 2006
9
     ______
10
11
     BMDS MODEL RUN
12
13
14
       The form of the response function is:
15
16
       Y[dose] = beta_0 + beta_1*dose + beta_2*dose^2 + ...
17
18
19
       Dependent variable = MEAN
20
       Independent variable = COLUMN1
21
       The polynomial coefficients are restricted to be positive
22
       The variance is to be modeled as Var(i) = alpha*mean(i)^rho
23
24
       Total number of dose groups = 4
25
       Total number of records with missing values = 0
26
       Maximum number of iterations = 250
27
       Relative Function Convergence has been set to: 1e-008
28
       Parameter Convergence has been set to: 1e-008
29
30
31
32
                     Default Initial Parameter Values
33
                            alpha =
                                              1
34
                                               0
                              rho =
35
                           beta 0 =
                                        30.4717
36
                           beta 1 =
                                       0.129124
37
38
39
40
                                   Parameter Estimates
41
42
                                                          95.0% Wald
43
    Confidence Interval
44
                                         Std. Err.
                                                      Lower Conf. Limit
          Variable
                          Estimate
    Upper Conf. Limit
45
                                                            -0.129982
46
             alpha
                          0.098324
                                         0.116485
47
    0.32663
48
                          2.15231
                                          0.291636
                                                              1.58072
               rho
49
    2.72391
50
                          28.6439
            beta_0
                                           2.33865
                                                              24.0602
51
    33.2276
52
                          0.141692
                                          0.0184799
                                                              0.105472
            beta_1
53
    0.177912
54
55
56
              Asymptotic Correlation Matrix of Parameter Estimates
57
```

```
1
                                  rho
                     alpha
                                             beta_0
                                                        beta_1
                     1
2
                                 -0.99
                                                          -0.17
        alpha
                                             0.06
3
                                 1
                                                          0.18
          rho
                     -0.99
                                             -0.065
4
       beta_0
                     0.06
                                -0.065
                                             1
                                                          -0.39
                                             -0.39
5
       beta 1
                     -0.17
                                 0.18
                                                            1
6
7
8
         Table of Data and Estimated Values of Interest
9
10
    Dose
              N
                    Obs Mean
                               Obs Std Dev Est Mean
                                                       Est Std Dev
                                                                   Chi^2
11
    Res.
12
                    _____
                                _____
                                             _____
                                                       _____
                                                                     _____
13
14
                                  8
15
       0
            15
                      27
                                             28.6
                                                         11.6
                                                                      -0.549
                                                         14.1
16
            15
                       37
                                 17.2
                                             34.3
                                                                       0.739
       40
17
      200
            15
                      59
                                  28
                                              57
                                                          24.3
                                                                       0.321
18
     1000
          15
                      159
                                   75
                                               170
                                                          79
                                                                      -0.556
19
20
21
22
     Model Descriptions for likelihoods calculated
23
24
25
     Model A1:
                    Yij = Mu(i) + e(ij)
26
              Var\{e(ij)\} = Sigma^2
27
28
     Model A2:
                   Yij = Mu(i) + e(ij)
29
              Var\{e(ij)\} = Sigma(i)^2
30
31
                     Yij = Mu(i) + e(ij)
     Model A3:
32
              Var\{e(ij)\} = alpha*(Mu(i))^rho
33
34
     Model R:
                      Yi = Mu + e(i)
35
               Var\{e(i)\} = Sigma^2
36
37
38
                          Likelihoods of Interest
39
40
               Model
                         Log(likelihood)
                                          DF
                                                   AIC
                                          5 511.887866
41
                A1
                          -250.943933
42
                A2
                          -216.540867
                                          8
                                                449.081734
                                          6
43
                A3
                          -217.537818
                                                447.075636
                                           4
44
              fitted
                           -219.838125
                                                 447.676251
                                                 567.326624
45
                 R
                           -281.663312
46
47
48
                      Explanation of Tests
49
50
     Test 1: Does response and/or variances differ among Dose
51
    levels?
52
              (A2 vs. R)
53
     Test 2: Are Variances Homogeneous? (A1 vs A2)
54
    Test 3: Are variances adequately modeled? (A2 vs. A3)
55
     Test 4: Does the Model for the Mean Fit? (A3 vs. fitted)
56
57
                        Tests of Interest
```

```
1
2
                -2*log(Likelihood Ratio) Test df
       Test
                                                         p-value
3
       Test 1
4
                            130.245
                                              6
                                                          <.0001
       Test 2
                            68.8061
                                                          <.0001
5
                                              3
6
       Test 3
                             1.9939
                                              2
                                                          0.369
7
       Test 4
                            4.60062
                                                          0.1002
8
9
    The p-value for Test 1 is less than .05. There appears
10
11
    difference between response and/or variances among the
12
    dose levels
13
    It seems appropriate to model the data
14
15
    The p-value for Test 2 is less than .05. A
16
    non-homogeneous variance
17
    model appears to be appropriate
18
19
    The p-value for Test 3 is greater than .05. The
20
    modeled variance appears
21
     to be appropriate here
22
23
    The p-value for Test 4 is greater than .05. The model
24
    chosen seems
25
    to adequately describe the data
26
27
28
     Benchmark Dose Computation
29
    Specified effect =
                                     1
30
31
    Risk Type
                            Estimated standard deviations from the control mean
32
33
34
    Confidence level =
                                  0.95
35
36
                  BMD =
                              81.8426
37
38
39
                 BMDL =
                              58.3727
```



```
13:00 05/09 2006
1
2
3
    ______
4
           Polynomial Model. Revision: 2.2 Date: 9/12/2002
5
           Input Data File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY
6
    DOCUMENTS\_CYANO TOX REV\MODELING\FAWELL_MALE_ALT.(d)
7
           Gnuplot Plotting File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY
8
    DOCUMENTS\_CYANO TOX REV\MODELING\FAWELL_MALE_ALT.plt
9
                                          Tue May 09 13:01:37 2006
10
     ______
11
12
     BMDS MODEL RUN
13
14
15
       The form of the response function is:
16
17
       Y[dose] = beta_0 + beta_1*dose + beta_2*dose^2 + ...
18
19
20
       Dependent variable = MEAN
21
       Independent variable = COLUMN1
22
       The polynomial coefficients are restricted to be positive
23
       The variance is to be modeled as Var(i) = alpha*mean(i)^rho
24
25
       Total number of dose groups = 4
26
       Total number of records with missing values = 0
27
       Maximum number of iterations = 250
28
       Relative Function Convergence has been set to: 1e-008
29
       Parameter Convergence has been set to: 1e-008
```

A-90

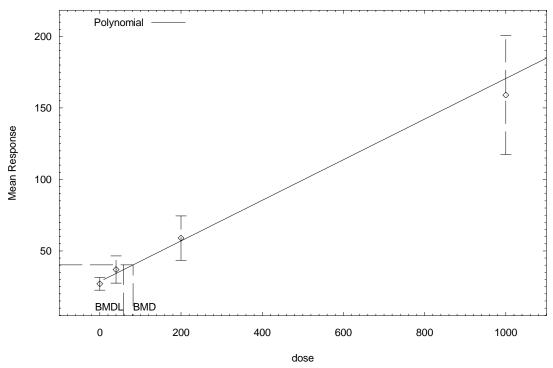
```
1
2
3
                       Default Initial Parameter Values
4
                               alpha =
                                                  1
                                 rho =
5
                                                  0
6
                              beta 0 =
                                           28.5738
7
                              beta 1 =
                                           0.160635
8
                              beta 2 =
9
10
11
12
                                      Parameter Estimates
13
                                                              95.0% Wald
15
    Confidence Interval
16
           Variable
                            Estimate
                                             Std. Err.
                                                          Lower Conf. Limit
17
    Upper Conf. Limit
18
              alpha
                           0.0983241
                                             0.116485
                                                                 -0.129982
19
    0.32663
20
                rho
                            2.15231
                                             0.291636
                                                                   1.58072
21
    2.72391
22
                            28.6439
                                              2.33865
                                                                   24.0602
             beta_0
23
    33.2276
24
             beta 1
                            0.141692
                                           0.0184799
                                                                  0.105472
25
    0.177912
26
                                    Ω
             beta_2
                                                    NA
27
28
    NA - Indicates that this parameter has hit a bound
29
         implied by some inequality constraint and thus
30
         has no standard error.
31
32
33
               Asymptotic Correlation Matrix of Parameter Estimates
34
35
                      alpha
                                                beta_0
                                                             beta_1
                                     rho
36
                                                 0.06
                                                              -0.17
         alpha
                                    -0.99
                        1
                                                               0.18
37
           rho
                       -0.99
                                     1
                                                -0.065
38
        beta 0
                       0.06
                                   -0.065
                                                   1
                                                              -0.39
39
                      -0.17
                                     0.18
                                                 -0.39
        beta_1
40
41
    The following parameter(s) have been estimated at a
42
43
    point or have been specified. Correlations are not
44
    computed:
45
46
    beta_2
47
48
49
         Table of Data and Estimated Values of Interest
50
51
     Dose
                N
                     Obs Mean
                                  Obs Std Dev
                                              Est Mean
                                                          Est Std Dev
52
    Res.
53
54
55
                                                 28.6
56
        0
             15
                        27
                                      8
                                                              11.6
                                                                           -0.549
57
       40
             15
                        37
                                    17.2
                                                34.3
                                                              14.1
                                                                            0.739
```

```
1
       200
              15
                         59
                                       28
                                                    57
                                                                 24.3
                                                                              0.321
2
      1000
              15
                                       75
                                                    170
                                                                  79
                                                                              -0.556
                        159
3
4
5
6
     Model Descriptions for likelihoods calculated
7
8
9
      Model A1:
                       Yij = Mu(i) + e(ij)
10
                Var\{e(ij)\} = Sigma^2
11
12
     Model A2:
                       Yij = Mu(i) + e(ij)
13
                Var\{e(ij)\} = Sigma(i)^2
14
15
                       Yij = Mu(i) + e(ij)
     Model A3:
16
                Var\{e(ij)\} = alpha*(Mu(i))^rho
17
18
     Model R:
                        Yi = Mu + e(i)
19
                 Var\{e(i)\} = Sigma^2
20
21
22
                             Likelihoods of Interest
23
24
                 Model
                             Log(likelihood)
                                                DF
                                                          AIC
25
                  A1
                              -250.943933
                                                5
                                                       511.887866
26
                  A2
                              -216.540867
                                                 8
                                                       449.081734
27
                  A3
                              -217.537818
                                                 6
                                                       447.075636
28
                fitted
                              -219.838125
                                                       447.676251
                                                 4
29
                   R
                              -281.663312
                                                 2
                                                       567.326624
30
31
32
                         Explanation of Tests
33
34
     Test 1: Does response and/or variances differ among Dose
35
     levels?
36
               (A2 vs. R)
37
      Test 2:
               Are Variances Homogeneous? (A1 vs A2)
38
               Are variances adequately modeled? (A2 vs. A3)
39
     Test 4: Does the Model for the Mean Fit? (A3 vs. fitted)
40
41
                           Tests of Interest
42
43
                -2*log(Likelihood Ratio) Test df
       Test
                                                          p-value
44
45
       Test 1
                             130.245
                                               6
                                                          <.0001
46
       Test 2
                             68.8061
                                               3
                                                          <.0001
47
       Test 3
                              1.9939
                                               2
                                                           0.369
48
       Test 4
                             4.60062
                                               1
                                                         0.03196
49
50
    The p-value for Test 1 is less than .05. There appears
51
     to be a
52
    difference between response and/or variances among the
53
    dose levels
54
     It seems appropriate to model the data
55
56
     The p-value for Test 2 is less than .05. A
57
    non-homogeneous variance
```

A-92

```
1
    model appears to be appropriate
2
3
    The p-value for Test 3 is greater than .05. The
4
    modeled variance appears
5
     to be appropriate here
6
7
    The p-value for Test 4 is less than .05. You may want
8
    to try a different
9
    model
10
11
12
     Benchmark Dose Computation
13
     Specified effect =
                                     1
14
15
    Risk Type
                             Estimated standard deviations from the control mean
16
17
18
    Confidence level =
                                  0.95
19
20
                               81.8426
                  BMD =
21
22
23
                               58.3727
                 BMDL =
```

Polynomial Model with 0.95 Confidence Level



13:01 05/09 2006

```
1
2
     ______
3
           Power Model. $Revision: 2.1 $ $Date: 2000/10/11 20:57:36 $
4
           Input Data File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY
5
    DOCUMENTS\ CYANO TOX REV\MODELING\FAWELL MALE ALT.(d)
6
           Gnuplot Plotting File: C:\DOCUMENTS AND SETTINGS\HCLYNCH\MY
7
    DOCUMENTS\ CYANO TOX REV\MODELING\FAWELL MALE ALT.plt
8
                                           Tue May 09 13:03:23 2006
9
     ______
10
11
    BMDS MODEL RUN
12
13
14
      The form of the response function is:
15
16
       Y[dose] = control + slope * dose^power
17
18
19
       Dependent variable = MEAN
20
       Independent variable = COLUMN1
21
       The power is restricted to be greater than or equal to 1
22
       The variance is to be modeled as Var(i) = alpha*mean(i)^rho
23
24
       Total number of dose groups = 4
25
       Total number of records with missing values = 0
26
      Maximum number of iterations = 250
27
      Relative Function Convergence has been set to: 1e-008
28
       Parameter Convergence has been set to: 1e-008
29
30
31
32
                     Default Initial Parameter Values
33
                            alpha = 1692.21
34
                              rho =
35
                          control =
                                             27
36
                            slope =
                                       0.519763
37
                            power =
                                       0.801589
38
39
40
              Asymptotic Correlation Matrix of Parameter Estimates
41
42
                     alpha
                                         control
                                                         slope
                                 rho
                                                                    power
43
44
                               -0.99
        alpha
                        1
                                             0.13
                                                         -0.46
                                                                     0.48
45
46
          rho
                    -0.99
                                    1
                                            -0.12
                                                         0.41
                                                                    -0.43
47
48
                    0.13
                                -0.12
                                               1
                                                         -0.76
                                                                     0.74
       control
49
50
                                 0.41
                                            -0.76
                                                           1
         slope
                    -0.46
                                                                       -1
51
52
                    0.48
                                -0.43
                                             0.74
                                                           -1
        power
53
54
```

Parameter Estimates

55 56

```
1
           Variable
                                                  Std. Err.
                              Estimate
2
                               0.098324
                                                   0.141722
              alpha
3
                rho
                                2.15231
                                                    0.343016
4
                                28.6439
            control
                                                    3.51452
5
                                0.141692
              slope
                                                    0.218496
6
              power
                                                   0.231677
7
8
9
10
         Table of Data and Estimated Values of Interest
11
12
    Dose
               N
                     Obs Mean
                                  Obs Std Dev Est Mean
                                                           Est Std Dev
                                                                        Chi^2
13
    Res.
14
                      _____
15
16
                                  8
17.2
                                                28.6
                                                             11.6
17
        0
             15
                        27
                                                                           -0.142
                                                34.3
                                                                            0.191
18
       40
             15
                        37
                                                              14.1
19
      200 15
                        59
                                   28
                                                 57
                                                             24.3
                                                                            0.083
                                                              79
20
     1000 15
                                     75
                                                 170
                      159
                                                                            -0.144
21
22
23
24
     Model Descriptions for likelihoods calculated
25
26
27
     Model A1: Yij = Mu(i) + e(ij)
28
               Var\{e(ij)\} = Sigma^2
29
30
     Model A2: Yij = Mu(i) + e(ij)
31
               Var\{e(ij)\} = Sigma(i)^2
32
33
     Model A3:
                      Yij = Mu(i) + e(ij)
34
               Var\{e(ij)\} = alpha*(Mu(i))^rho
35
36
     Model R:
                      Yi = Mu + e(i)
37
                 Var\{e(i)\} = Sigma^2
38
39
40
                            Likelihoods of Interest
41
42
                Model
                           Log(likelihood)
                                            DF
                                                       AIC
                            -250.943933 5 511.887866

-216.540867 8 449.081734

-217.537818 6 447.075636

-219.838125 5 449.676251

-281.663312 2 567.326624
43
                 A1
44
                 A2
                 A3
45
46
                fitted
47
                 R
48
49
50
                        Explanation of Tests
51
52
     Test 1: Does response and/or variances differ among Dose levels?
53
              (A2 vs. R)
54
     Test 2: Are Variances Homogeneous? (A1 vs A2)
     Test 3: Are variances adequately modeled? (A2 vs. A3)
     Test 4: Does the Model for the Mean Fit? (A3 vs. fitted)
56
57
```

1	Tests of Interest									
2										
3	Test	-2*log(Lik	elihood Ratio)	d.f	p-value					
4										
5	Test 1		130.245	6	<.00001					
6	Test 2		68.8061	3	<.00001					
7	Test 3		1.9939	2	0.369					
8	Test 4		4.60062	1	0.03196					
9										
10	The p-valu	e for Test	1 is less than	05 The	re appears to be a					
11	The p-value for Test 1 is less than .05. There appears to be a difference between response and/or variances among the dose levels									
12	It seems appropriate to model the data									
13	ic beenib a	ppropriace	co moder che di	aca						
14	The p-valu	e for Test	2 is less than	.05. A no	on-homogeneous variance					
15	The p-value for Test 2 is less than .05. A non-homogeneous variance model appears to be appropriate									
16	moder appe	arb co be a	PPIOPIIACC							
17	The n-valu	e for Test	3 is greater th	han O5 '	The modeled variance appe	arg				
18		ropriate he		.05.	ine modered variance appe	alb				
19	co be app	ropriace ne	10							
20	The n-walu	e for Test	4 is less than	05 VOU	may want to try a differ	ent				
21	model	C IOI ICSC	1 15 1C55 CHAII	.03. 104	may want to try a drifter	CIIC				
22	moder									
23										
24	Ponahmark	Dose Compu	tation							
25	Specified									
25 26	Specified	errect =	1							
27	Dial mass		Tatimated at.		intions from the souther					
28	Risk Type	=	ESCIMALEG SC	andard dev	iations from the control	mean				
28 29	0	11	0 0 5							
	Confidence	rever =	0.95							
30		DIE	01 0406							
31		BMD =	81.8426							
32										
33										
34		BMDL =	58.3727							
35										
36										
37										