

***Epigenetic alterations induced by genotoxic
occupational and environmental human
chemical carcinogens:
A systematic literature review***

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Disclaimers:

- Research is funded by grants from NIH and US EPA
- Currently serve on several National Research Council committees that consider issues relevant to this workshop
- Health Assessments Workspace Collaborative (hawcproject.org) online tool was developed and supported, in part, by funding from Texas A&M University and NIH grants

MECHANISTIC DATA:

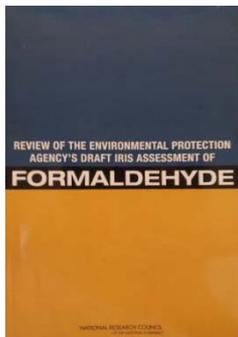
The Third (Rail?) Data Stream in Risk Assessment

A typical statement found in many human health assessment documents:
“The mode of action of [insert your chemical’s name here] is complex and has not been fully elucidated yet”



- “Decisions to protect public health and the environment cannot await “perfection” in scientific knowledge [...]. It is important that risk assessments incorporate **the best available scientific information in scientifically rigorous ways and that they capture and describe the uncertainties** in the information in ways that are useful for decision-makers”

NRC (2009): “Science and Decisions: Advancing Risk Assessment”



- “Clear concise statements of **criteria used to exclude, include, and advance studies** [as part of IRIS process]”
- “**Standardized evidence tables** that provide the methods and results of each study are needed for all health outcomes; if appropriate tables were used, long descriptions of the studies could be moved to an appendix or deleted”

NRC (2011): “Review of the Environmental Protection Agency’s Draft IRIS Assessment of Formaldehyde”

Opportunity: Improved evidence capture and reporting for risk assessments through systematic literature review

Identifying and Presenting “Mechanism Data”: DEHP Case Study [pre-historic years: 3 B.K.C.]

Deliverables:

IRIS Toxic Review

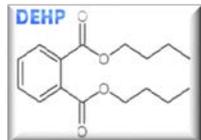
- Evidence tables
- Short narrative summaries

Appendix to the IRIS Toxic Review

- Search terms (*e.g.*, MeSH)
- Literature trees
- Inclusion/exclusion criteria
- Literature tags (for HERO)

HERO Database

- Primary literature (PDF+tags)
- EndNote reference libraries



1. Select a chemical and target tissue



2. Identify relevant mechanisms of action



3. Devise overall search strategy



4. Define assessment factors (i.e., exclusion criteria)



5. Conduct literature searches



6. Construct literature trees for each mechanism



7. Construct evidence tables

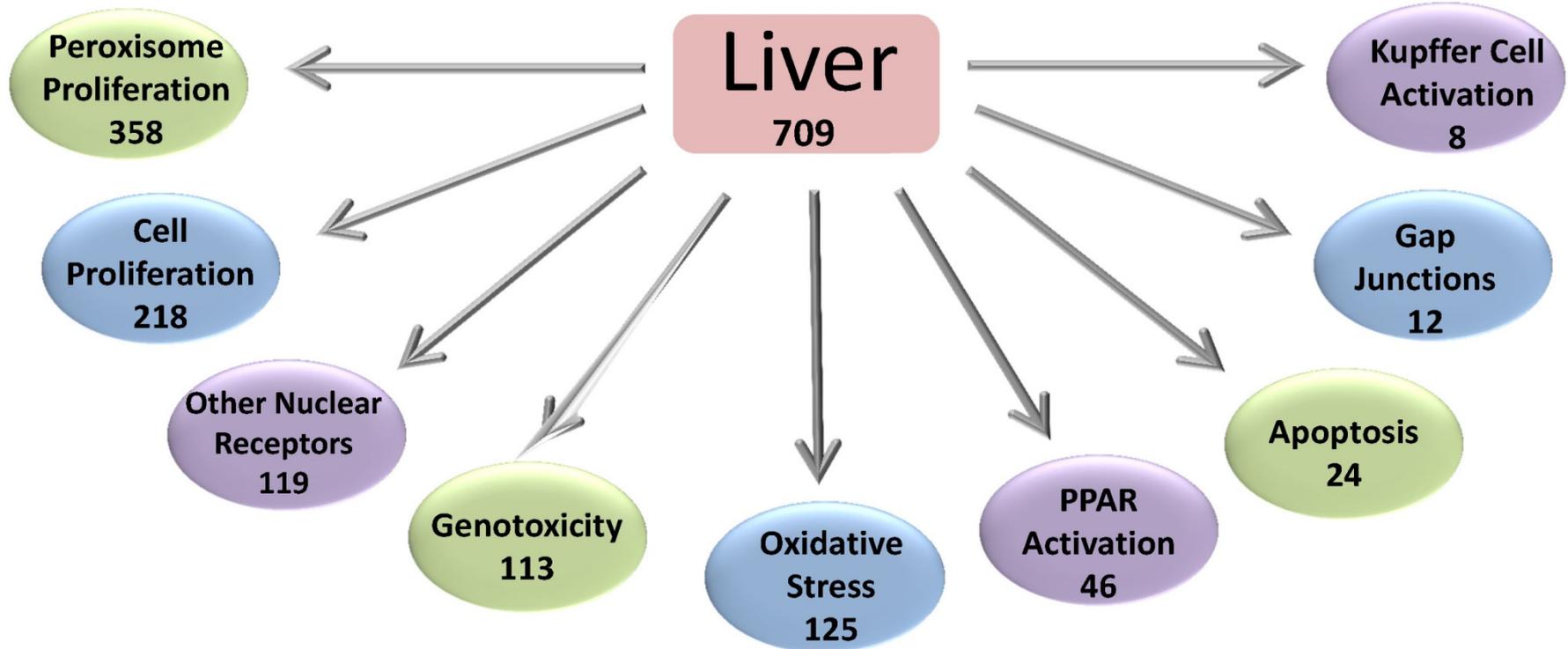


8. Organize and tag evidence

Specify study question:

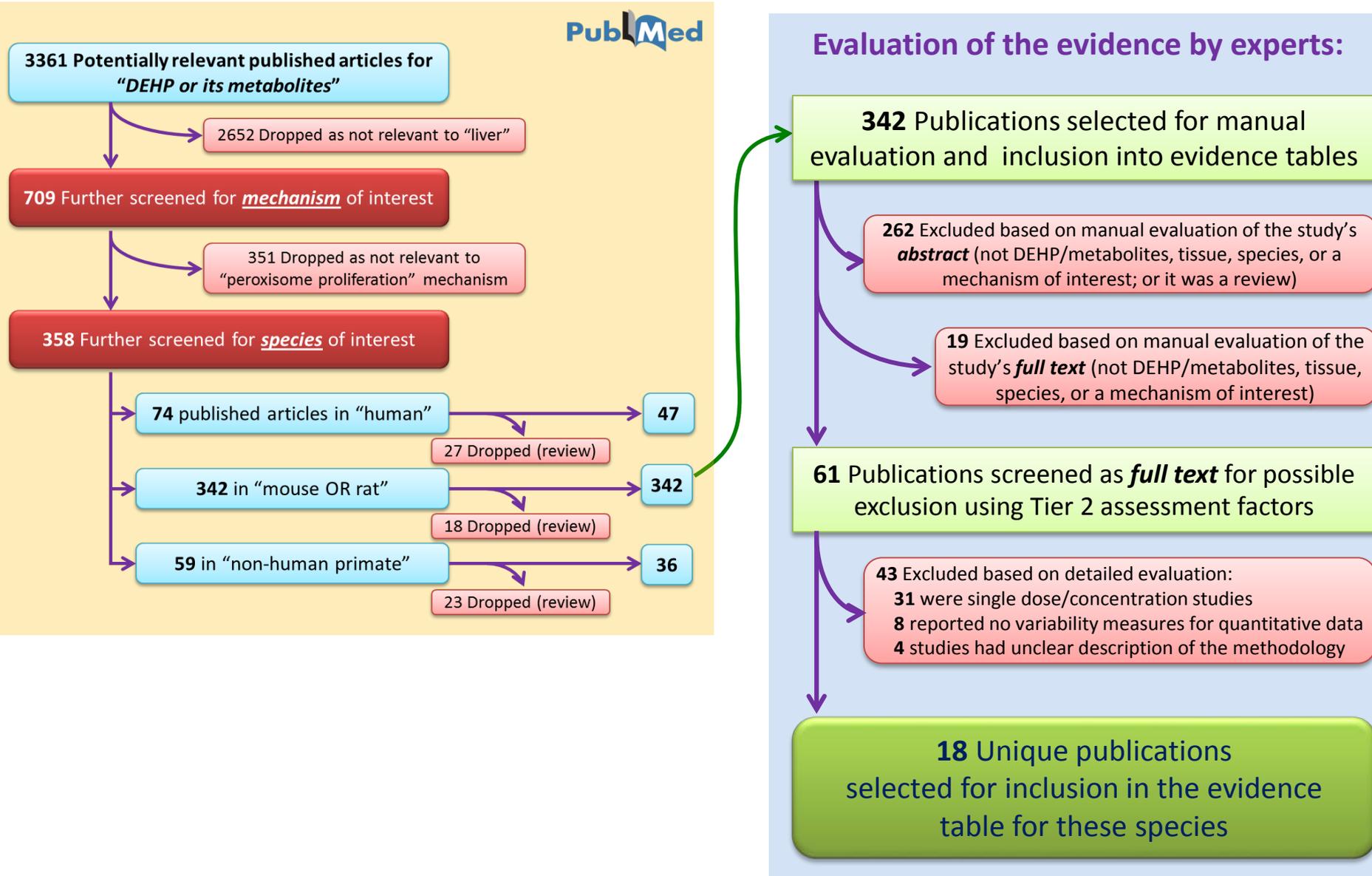
Effects of **DEHP** and its **metabolites** on the **liver**

Identify relevant mechanistic events (**in liver**) from diverse and authoritative literature reviews



Construct Literature Trees for Each Mechanism

Large database example: “*Peroxisome Proliferation*”



Constructing evidence tables for each mechanistic event

Provide sufficient details about:

- the study,
- study design,
- model system,
- endpoint,
- dose & concentration,
- fold change & significance



Study Design and Reference	Endpoint and Assay	Results (% Change from Control)						
Humans								
<i>In Vivo</i>								
<i>No primary literature evidence identified</i>								
<i>In Vitro</i>								
(Kamendulis et al., 2002)		<u>mM MEHP</u>						
Human 1° hepatocytes (3 males)	GJIC disruption, <i>in situ</i> dye transfer (ISDT)	1° Heps, 4 hrs	0	0.05	0.1	0.2	0.3	0.5
HLEC-04 cells (SV-40 transf. hum.heps)		24 hrs	0	-1	-3	nd	-2	nd
Duration: 4 or 24 hrs		HLEC-04, 4 hrs	0	-8	-6	nd	-7	nd
N = 4 cultures/group		24 hrs	0	-8	-6	nd	-7	nd
		24 hrs	0	-7	-6	nd	-7	-2
Non-human primates								
<i>In Vivo</i>								
(Pugh et al., 2000)		<u>mg/kg/day DEHP</u>						
Cynomolgus monkeys (2 yrs of age)	GJIC disruption, ISDT		0	500				
Duration: 2 wks			0	8				
N = 4 males/group								
<i>In Vitro</i>								
(Kamendulis et al., 2002)		<u>mM MEHP</u>						
Cynomolgus monkey 1° heps (5 males)	GJIC disruption, ISDT		0	0.05	0.1	0.2	0.3	0.5
Duration: 4 or 24 hours		4 hrs	0	-1	0	nd	-11	nd
N = 4 cultures/group		24 hrs	0	-3	-2	nd	-5	nd

Group studies by species:

- humans,
- primates,
- rodents

Group studies by model system:

in vivo
in vitro

Highlight studies that have challenged the hypothesized mode of action using knockout and transgenic (e.g., humanized) mouse strains



Study Design and Reference	Endpoint and Assay	Results (% Change from Control ¹)				
<i>In Vivo Chronic Cancer Bioassays</i>						
(Ito et al., 2007)		<u>DEHP (mg/kg/day)¹</u>				
Mice	Neoplastic changes in liver	Adenoma	Wild type	0	13	65
Sv/129 (wild type) and PPARa-null Males (6 wks old)		PPARa-null	0	2	2	
Dietary feeding: 0, 0.01, 0.05%		Carcinoma	Wild type	0	1	6
Duration: 22 months		PPARa-null	0	0	0	
N = 20-31		Cholangiocarcinoma	Wild type	1	0	1
		PPARa-null	0	0	0	
		Total liver tumors	Wild type	0 (0%)	2 (8.7%)	2 (10%)
		PPARa-null	1 (4%)	1 (4%)	8* (26%)	

Systematic Review of “Mechanism Data”: Styrene genotoxicity/mutagenicity (NRC 2014)

BOX D-4 Exclusion Criteria and Search Strategy for Studies of Genotoxicity and Related Mechanisms of Styrene

Exclusion Criteria

- The publication did not evaluate health effects of styrene or its metabolites known to be formed in humans.
- The study evaluated cellular, biochemical, or molecular effects not relevant to the carcinogenesis or the mechanistic event under consideration.
- The publication did not contain primary data.
- The publication did not include information sufficient to determine what species were studied or what experimental methods were used.

Search Strategy

PubMed: [(“Styrene”[Title/Abstract]) AND (“Mutation”[Mesh] OR “Cell Transformation, Neoplastic”[Mesh] OR “Cytogenetic Analysis”[Mesh] OR “Mutagens”[Mesh] OR “Oncogenes”[Mesh] OR “Genetic Processes”[Mesh] OR chromosom* OR clastogen* OR “genetic toxicology” OR “strand break” OR “unscheduled DNA synthesis” OR “DNA damage” OR “DNA adducts”)]. Search run on 05-28-2013; updated on 11-13-2013; and limited to 01-01-2008 to 11-13-2013.

Medline and Embase: [(styrene.ab. or styrene.ti.) and (1 or 2 or 3 or 4 or 5 or 6 or chromosom*.mp. or clastogen*.mp. or genetic toxicology.mp. or strand break.mp. or unscheduled DNA synthesis.mp. or DNA damage.mp. or DNA adducts.mp.), where the following keywords are: 1) Mutation, 2) Cell Transformation, Neoplastic, 3) Cytogenetic Analysis, 4) Mutagens, 5) Oncogenes, 6) Genetic Processes]. Search run on 05-28-2013; updated on 11-13-2013; and limited to 01-01-2008 to 11-13-2013.

Scopus: [(“Styrene”) AND (“mutation” OR “cell transformation, neoplastic” OR “cytogenetic analysis” OR “mutagens” OR “oncogenes” OR “genetic processes” OR “chromosom*” OR “clastogen*” OR “genetic toxicology” OR “strand break” OR “unscheduled DNA synthesis” OR “DNA damage” OR “DNA adducts”)]. Search run on 05-28-2013; updated on 11-13-2013; and limited to 01-01-2008 to 11-13-2013.

Web of Science: [(“Styrene”) AND (“mutation” OR “cell transformation, neoplastic” OR “cytogenetic analysis” OR “mutagens” OR “oncogenes” OR “genetic processes” OR “chromosom*” OR “clastogen*” OR “genetic toxicology” OR “strand break” OR “unscheduled DNA synthesis” OR “DNA damage” OR “DNA adducts”)]. Search run on 05-28-2013; updated on 11-13-2013; and limited to 01-01-2008 to 11-13-2013.

Systematic
Review

144 Publications identified in the literature search for genotoxicity and mutagenicity studies

114 Publications removed according to exclusion criteria

30 Publications were further evaluated using full text

16 Publications removed as not relevant according to exclusion criteria

14

+4 Publications added from IARC (1994a,b; 2002)

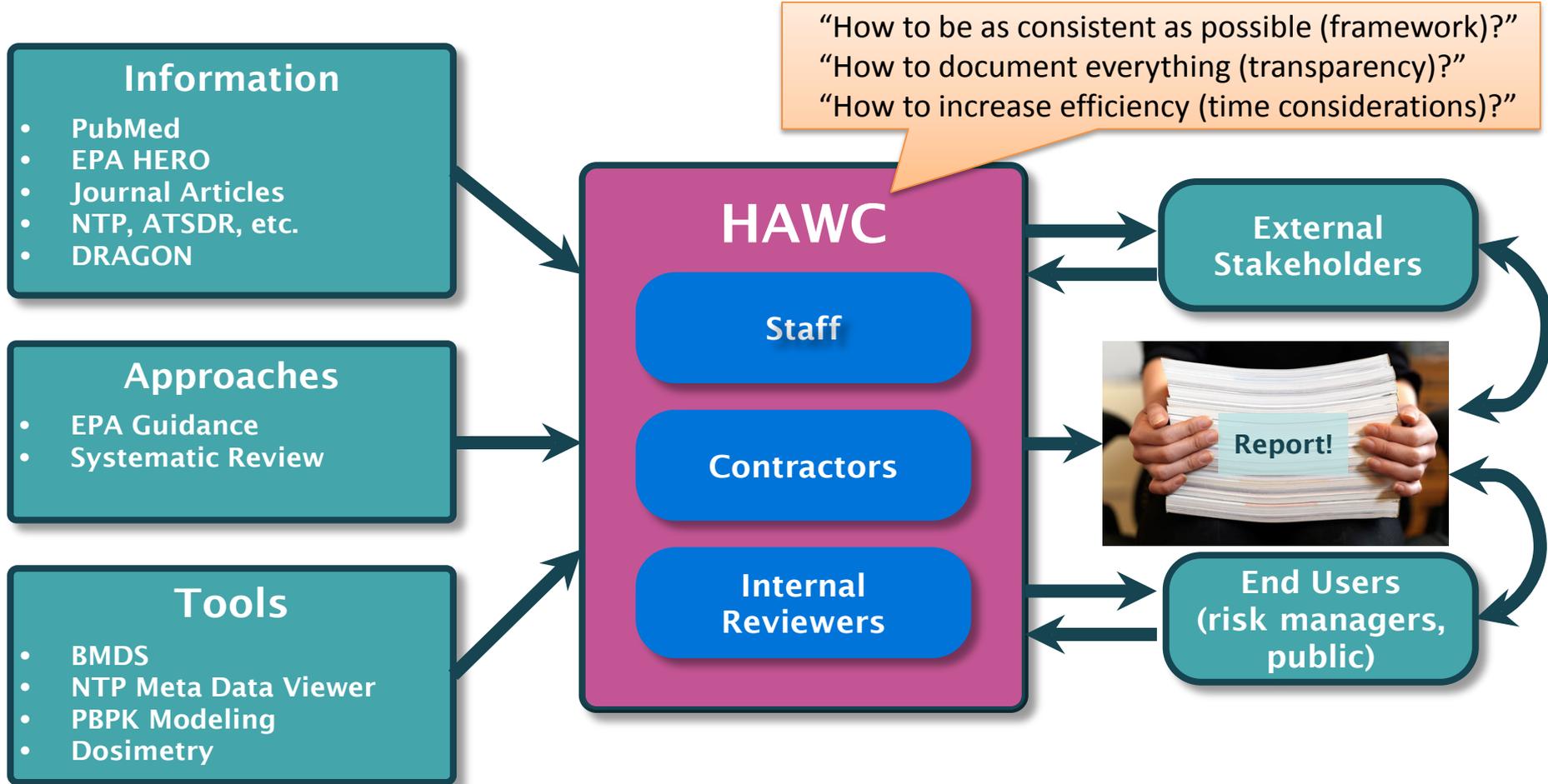
+20 Publications added from Scott and Preston (1994)

38 Publications identified as relevant and evaluated in Chapter 3

integration

Overall, the observations in various studies performed over the last 3 decades have been consistent. Temporal and exposure–response relationships have been established. Not only is the experimental evidence extensive, it is likely to be relevant to all target tissues that have been associated with cancer after exposure to styrene. Causality is strengthened by the large amount of evidence obtained **from studies of exposed humans.**

HAWC: Web-based content management system



Create a HAWC account or view public-assessments:

<https://hawcproject.org>

Actively under development; feedback is appreciated.

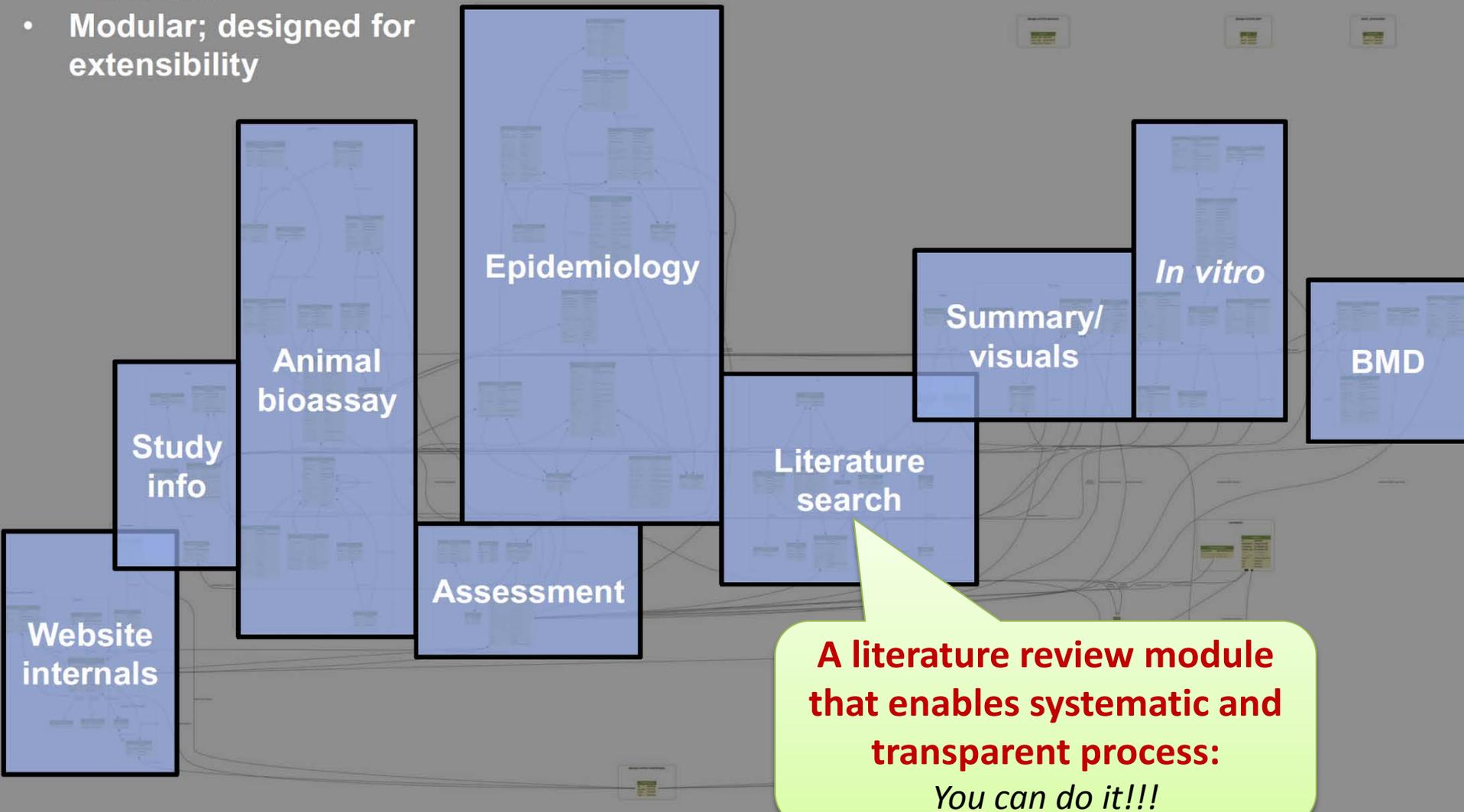
Compatible browsers:



*Recommended browser

HAWC: Web-based content management system

- 102 tables
- 9 modules
- Modular; designed for extensibility



Benzene (2014): Literature Tree

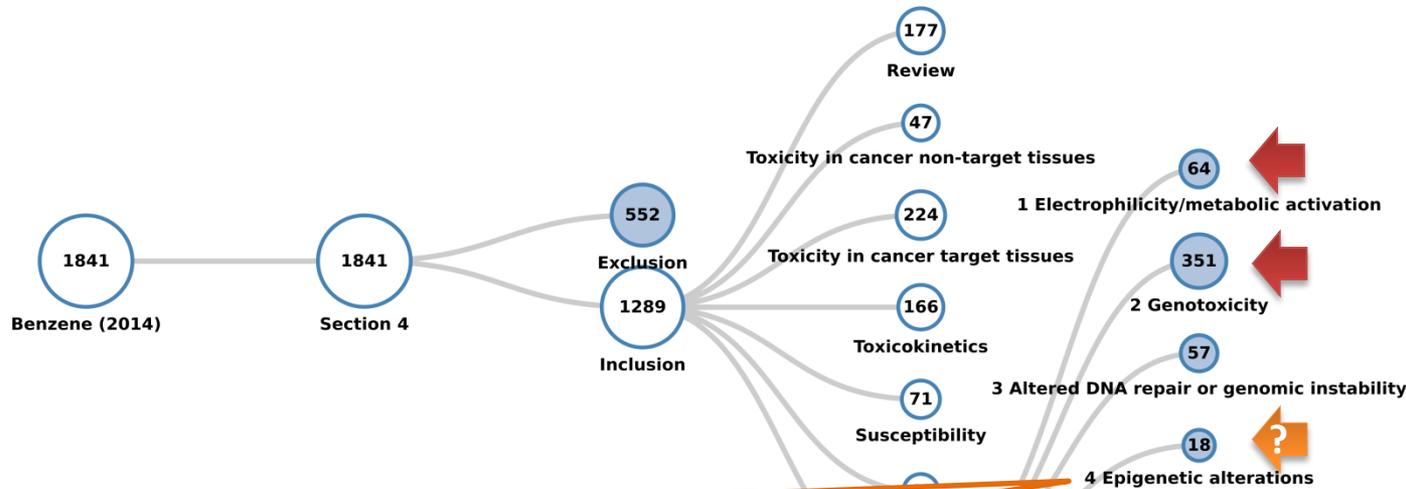
Key Characteristics of Carcinogens as a Basis for Organizing Data on Mechanisms of Carcinogenesis

Martyn T. Smith, Kathryn Z. Guyton, Catherine F. Gibbons, Jason M. Fritz, Christopher J. Portier, Ivan Rusyn, David M. DeMarini, Jane C. Caldwell, Robert J. Kavlock, Paul Lambert, Stephen S. Hecht, John R. Bucher, Bernard W. Stewart, Robert Baan, Vincent J. Cogliano, and Kurt Straif

<http://dx.doi.org/10.1289/ehp.1509912>

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(New) Epigenetics (#4)

Actions ▾

Description	Epigenetics
Search Type	Search
Search Database	PubMed
Search Text	Benzene[Mesh] AND ("rna"[MeSH] OR "epigenesis, genetic"[MeSH] OR rna OR "rna, messenger"[MeSH] OR "rna" OR "messenger rna" OR mrna OR "histones"[MeSH] OR histones OR epigenetic OR miRNA OR methylation)
Created	Dec. 1, 2014, 1:38 p.m.
Last Updated	Dec. 1, 2014, 1:38 p.m.

Literature Tagging Statistics

Total References	249
Total Tagged	240
Total Untagged	9
Reference details	View by tag Visualization

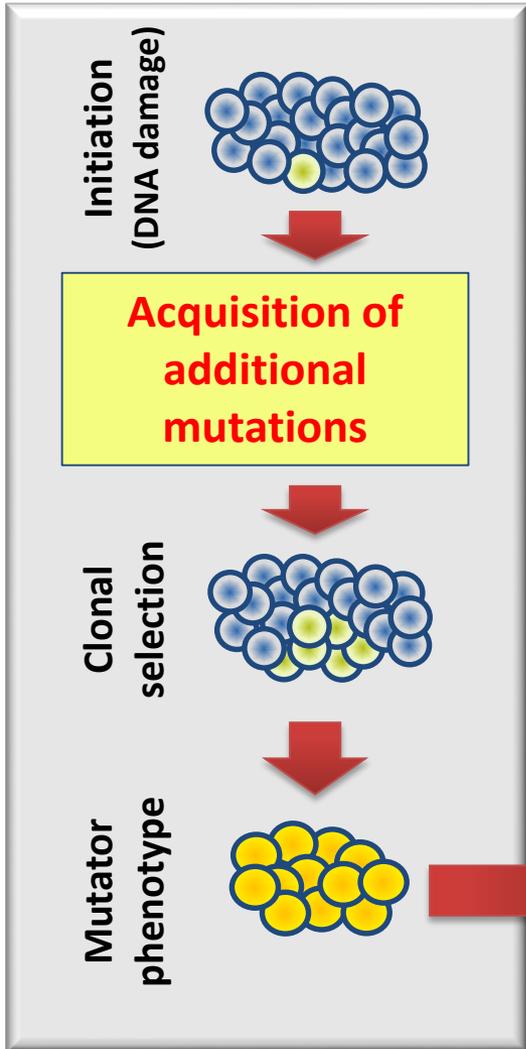
Results from queries

Date last executed	Total references found	References added	References removed
Dec. 12, 2015, 1:43 p.m.	249	9	0
Dec. 1, 2014, 1:38 p.m.	240	240	0

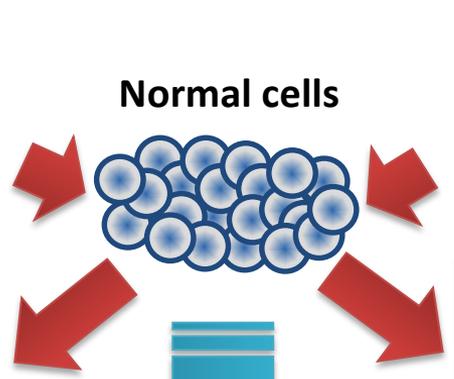
[Tag Literature](#)

[Rerun search](#)

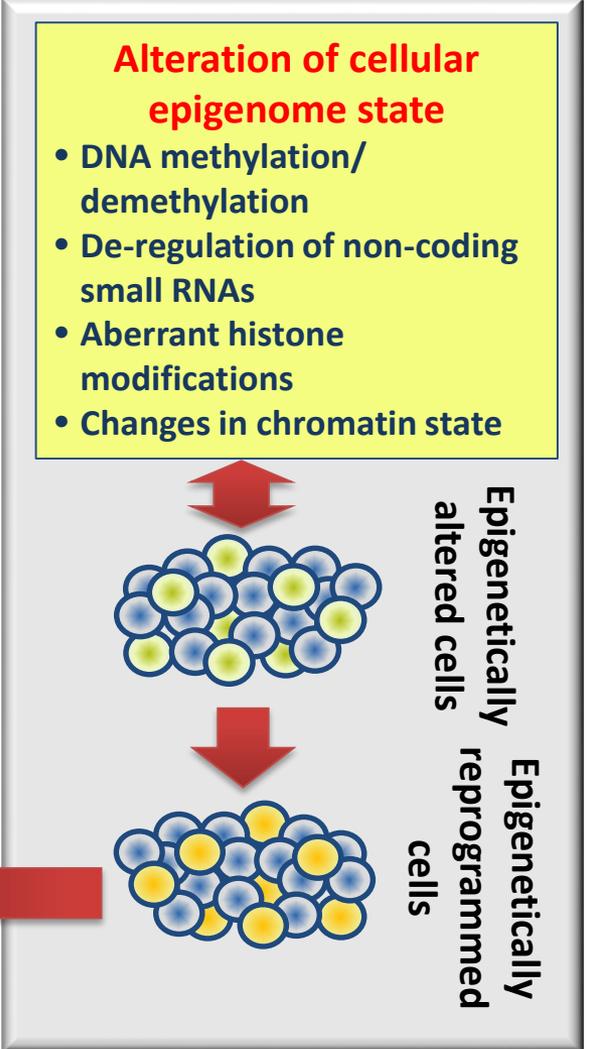
Genotoxicity Pathway to Carcinogenesis



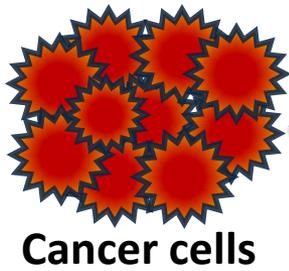
Exogenous damage



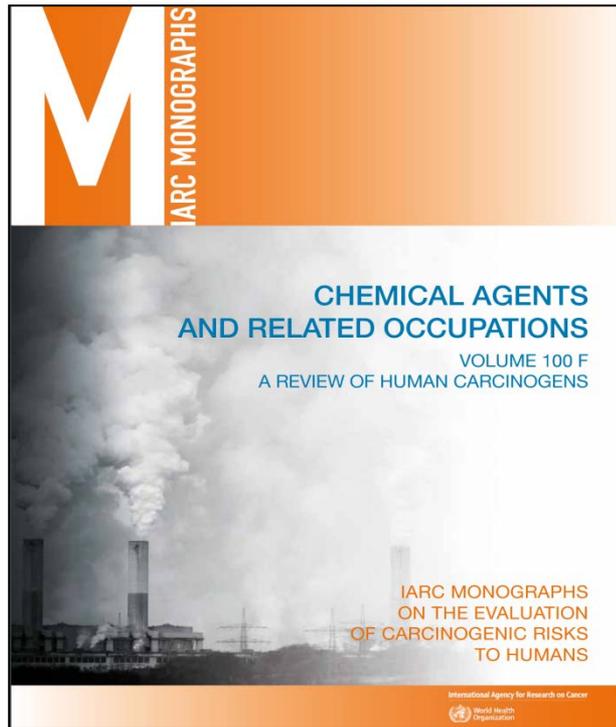
Endogenous damage



Epigenotoxicity Pathway to Carcinogenesis



- Epigenetic alterations play an important role in chemical carcinogenesis
- Although changes to the epigenome may be as important in carcinogenicity, the genotoxicity of chemicals has been studied more thoroughly
- **“Most carcinogens were evaluated by IARC before new data on their epigenetic effects became available”**
(Herceg *et al.*, 2013)



Agents and Related Occupations (Volume 100F)

- **4-Aminobiphenyl**
- **Benzidine**
- **Dyes Metabolized to Benzidine**
- **4,4'-Methylenebis(2-chloroaniline)**
- **2-Naphthylamine**
- **ortho-Toluidine**
- Auramine and Auramine Production
- Magenta and Magenta Production
- **Benzo[a]pyrene**
- **Coal gasification**
- **Occupational Exposures during Coal-tar Distillation**
- **Coal-tar pitch**
- **Coke Production**
- **Mineral Oils, Untreated or Mildly Treated**
- **Shale Oils**
- **Soot, as found in occupational exposure of chimney-sweeps**
- **Occupational Exposures during Aluminium Production**
- **Aflatoxins**
- **Benzene**
- **Bis(chloromethyl)ether and Chloromethyl Methyl Ether**
- **1,3-Butadiene**
- 2,3,7,8-TCDD, 2,3,4,7,8-PeCDF, and PCB 126
- **Ethylene Oxide**
- **Formaldehyde**
- **Sulfur Mustard**
- **Vinyl Chloride**
- **Isopropyl Alcohol Manufacture by the Strong-acid Process**
- **Mists from Strong Inorganic Acids**
- **Occupational Exposures during Iron and Steel Founding**
- **Occupational Exposure as a Painter**
- **Occupational Exposures in the Rubber Manufacturing Industry**

“Epigenotoxic” Literature Search Terms

IARC (vol 112 and 113) search string:

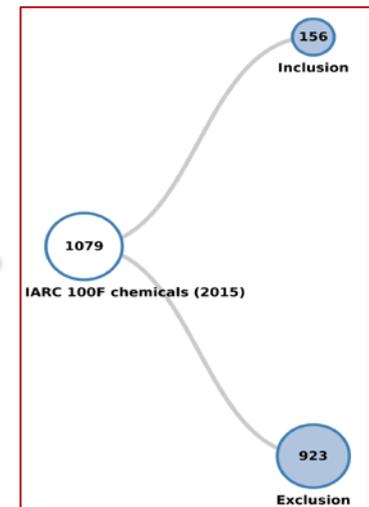
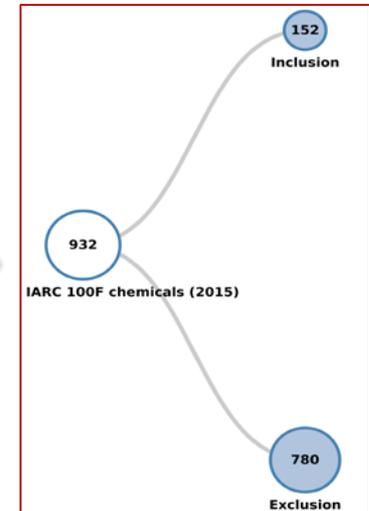
"rna"[MeSH] OR "epigenesis, genetic"[MeSH] OR rna OR "rna, messenger"[MeSH] OR "rna" OR "messenger rna" OR mrna OR "histones"[MeSH] OR histones OR epigenetic OR miRNA OR methylation

TAMU initial search string of [All Fields] terms:

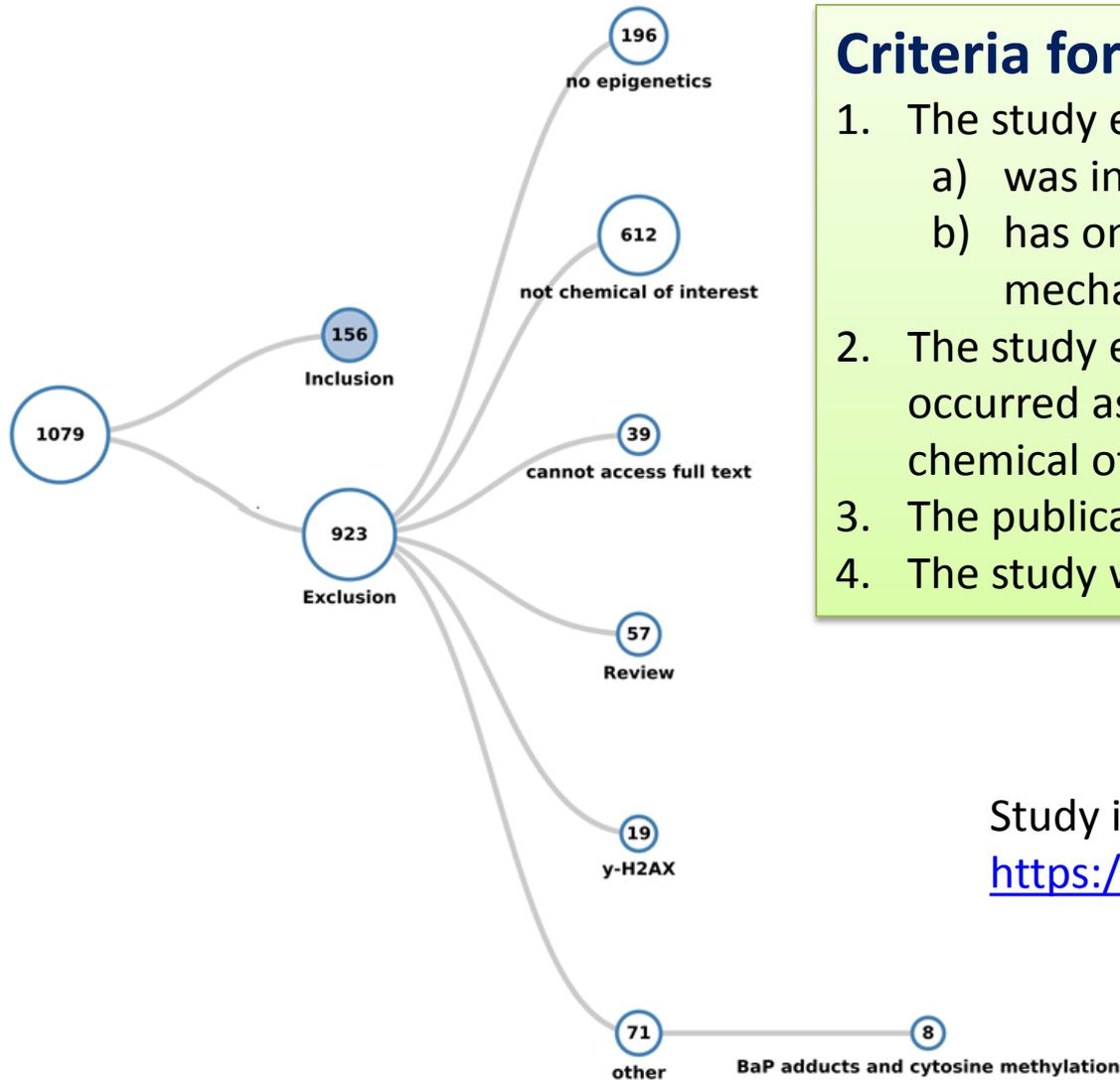
"epigenetic"[All Fields] OR "microRNA"[All Fields] OR "miRNA"[All Fields] OR "lncRNA"[All Fields] OR "non-coding RNA"[All Fields] OR "non coding RNA"[All Fields] OR "ncRNA"[All Fields] OR "small RNA"[All Fields] OR "smallRNA"[All Fields] OR "DNA methylation"[All Fields] OR "methylated DNA"[All Fields] OR "promoter methylation"[All Fields] OR "chromatin modification"[All Fields] OR "open chromatin"[All Fields] OR "histone"[All Fields] OR "histone positioning"[All Fields] OR "histone methylation"[All Fields] OR "histone acetylation"[All Fields] OR "histone mark"[All Fields] OR "histone modification"[All Fields]

TAMU final search string of a refined set of [All Fields] and [MeSH]:

"epigenomics"[MeSH Terms] OR "epigenomics"[All Fields] OR "epigenetic"[All Fields] OR "microrna"[All Fields] OR "micrnas"[All Fields] OR "miRNAs"[All Fields] OR "miRNA"[All Fields] OR "lncRNA"[All Fields] OR "rna, long noncoding"[MeSH Terms] OR "non-coding RNA"[All Fields] OR "non coding RNA"[All Fields] OR "ncRNA"[All Fields] OR "small RNA"[All Fields] OR "smallRNA"[All Fields] OR "dna methylation"[All Fields] OR "promoter methylation"[All Fields] OR "methylated DNA"[All Fields] OR "chromatin modification"[All Fields] OR "open chromatin"[All Fields] OR "histones"[MeSH Terms] OR "histones"[All Fields] OR "histone"[All Fields]



Epigenetic alterations induced by genotoxic occupational and environmental human chemical carcinogens



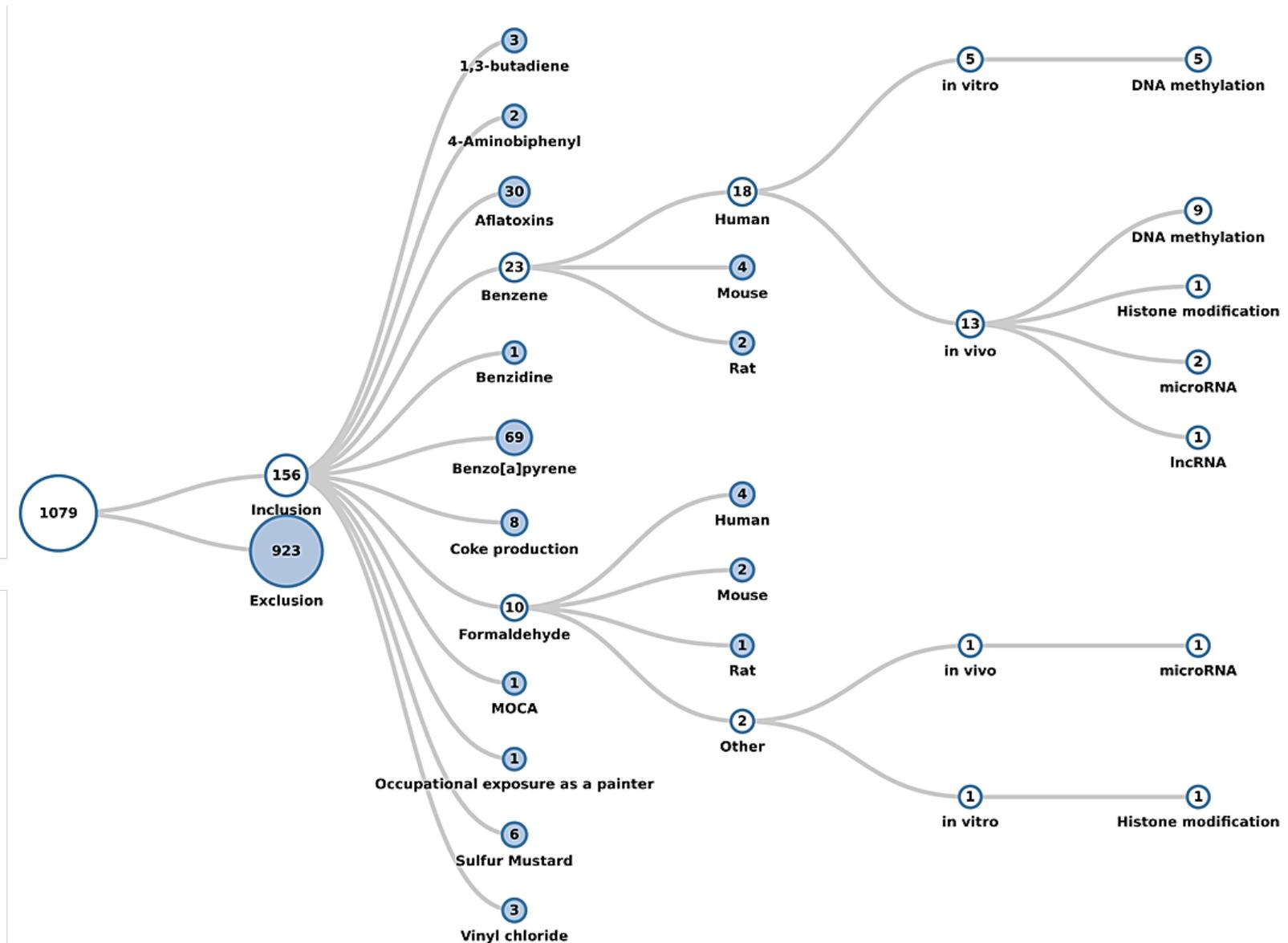
Criteria for inclusion in the review:

1. The study evaluated a chemical that:
 - a) was included in IARC Monograph vol.100F
 - b) has one or more demonstrated genotoxic mechanism(s) of carcinogenesis
2. The study evaluated epigenetic alterations that occurred as a consequence of exposure to the chemical of interest
3. The publication included original data
4. The study was published in English

Study is publicly available in HAWC:

<https://hawcproject.org/assessment/185/>

Epigenetic alterations induced by genotoxic occupational and environmental human chemical carcinogens



Epigenetic alterations induced by genotoxic occupational and environmental human chemical carcinogens

in vitro
in vivo
min
max

		DNA methylation				Histone modification				Non-coding RNA				
		Human	Mouse	Rat	Other	Human	Mouse	Rat	Other	Human	Mouse	Rat	Other	total:
Benzo[a]pyrene	in vitro	11	6		3	6	2		1	16	3			48
	in vivo	5	4		5		1	1		1	7	1		25
Aflatoxins	in vitro	1			2				1	7	1			12
	in vivo	10	2	1	1		1		1	3		2		21
Benzene	in vitro	5	1	1				1						8
	in vivo	9	1	1		1	1			3	2			18
Formaldehyde	in vitro	1				2			1	1				5
	in vivo										2	1	1	4
Coke production	in vitro													0
	in vivo	6								2				8
1,3-butadiene	in vitro													0
	in vivo		3				3							6
Sulfur mustard	in vitro	1								1				2
	in vivo									3	1			4
Vinyl chloride	in vitro													0
	in vivo	3												3
4-aminobiphenyl	in vitro					1				1				2
	in vivo													0
Benzidine	in vitro													0
	in vivo		1											1
MOCA	in vitro							1						1
	in vivo													0
Occupational exp. as a painter	in vitro													0
	in vivo	1												1

85
25
59

“Epigenotoxic” literature search terms: Splitting hairs?

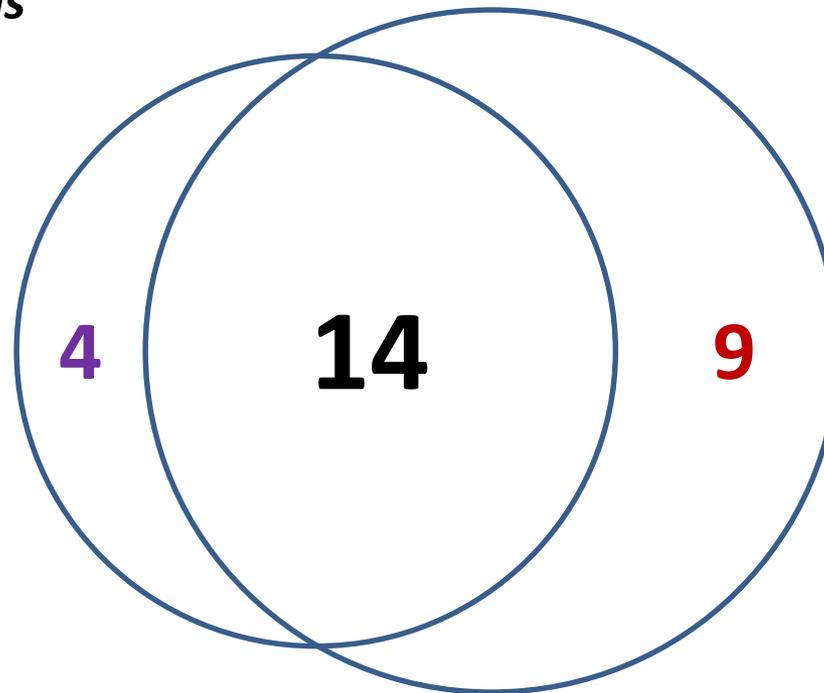
“Benzene” AND “Epigenetic alterations”

IARC search terms (in 2014):

18 publications

TAMU search terms (in 2015):

23 publications



- A review
- Two non-English publications
- One study of chemical reactivity with proteins (histones)

- Five studies published in 2014 or after (more recent than the review)
- Four studies are relevant and published pre-2014 (omitted?)

A refined literature search compendium of terms is more “specific”

“Epigenotoxic” literature search terms: Splitting hairs?

[All Fields] *versus* [MeSH Terms]

[All Fields] terms:

932 total

152 Included

780 Excluded

[All Fields] and [MeSH Terms]:

1083 total

155 Included

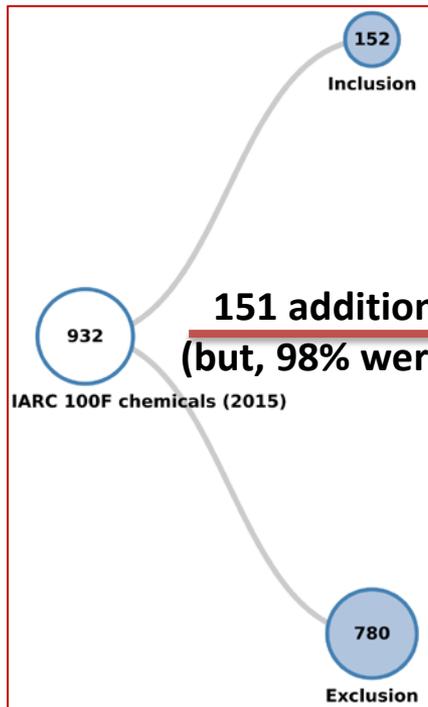
928 Excluded

Removing “redundant” terms:

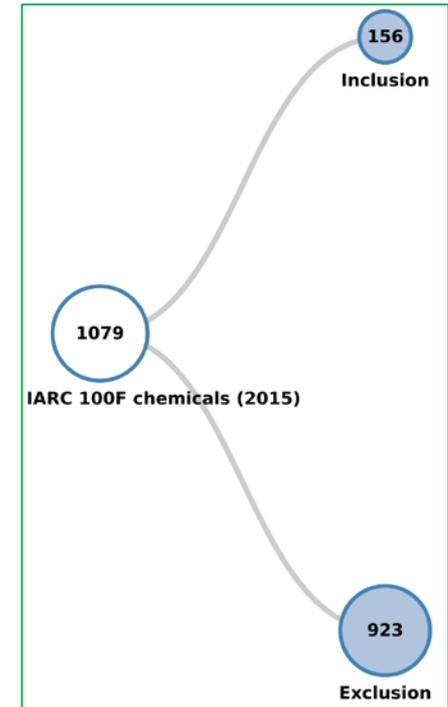
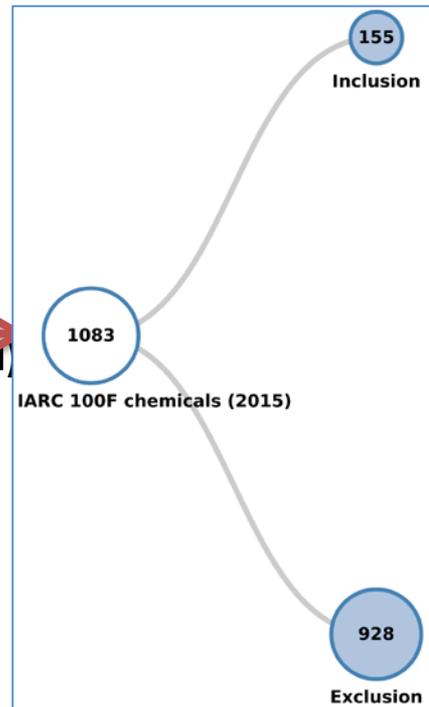
1079 total

156 Included

923 Excluded



151 additional papers
(but, 98% were excluded)



Conclusions

1. While the number of studies devoted to understanding the epigenetic alterations caused by exposure to chemical carcinogens is rapidly increasing, there remains a dearth of well-designed comprehensive studies that identify epigenetic alterations that are associated with the carcinogenesis:

- Only 1/3 (4 out of 12) of the chemicals (and occupational hazard) included in the review had a maximum of two published reports of epigenetic alterations

2. While there is extensive information about the fundamental role of epigenetic alterations in cancer development and progression, the understanding of the mechanistic significance and specificity of carcinogen-induced epigenetic abnormalities in the carcinogenic process is insufficient:

- Delineation between normal epigenetic processes in cells and the epigenetic alterations that have a causal relationship to cancer is needed for all epigenetic marks, not only for DNA methylation

3. To fully understand the importance of epigenetic and epigenomic responses to environmental stressors, studies that investigate and compare both epigenetic data with functional measures (such as gene and protein expression) and within the same study and controlled exposure scenario are needed

4. Epigenetic marks represent a class of biomarkers with great potential in the identification of exposure status, damage response, and/or disease state; however, few studies looked beyond epigenetic effects immediately after exposure

Acknowledgements

Literature reviews:

Mary Kushman
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Weihsueh Chiu
Kate Guyton
Dana Loomis
IARC staff and
Monographers

ToxCast Data:

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David Reif
Weihsueh Chiu
Kate Guyton
Chris Portier
IARC staff and
Monographers