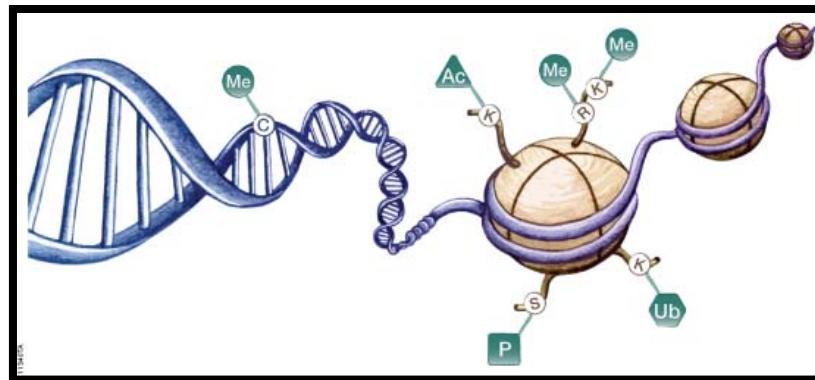


The King of Poisons: Arsenic - Latent Life Health Impacts

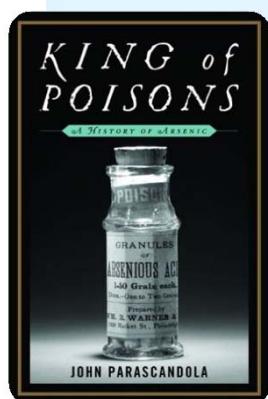
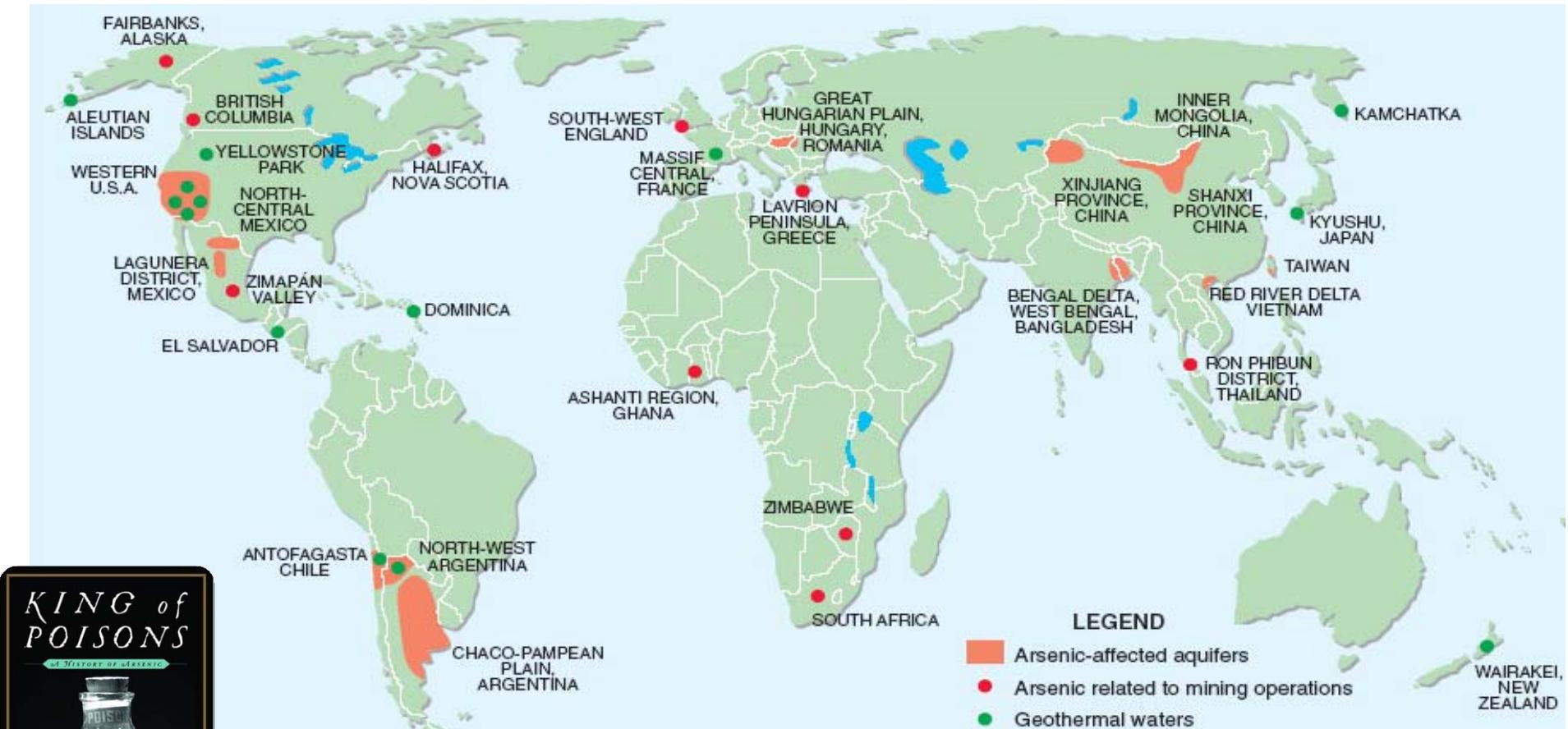


REBECCA FRY, PH.D.

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Curriculum in Toxicology and Environmental Medicine



Arsenic, the king of poisons, contaminates the water of millions around the globe

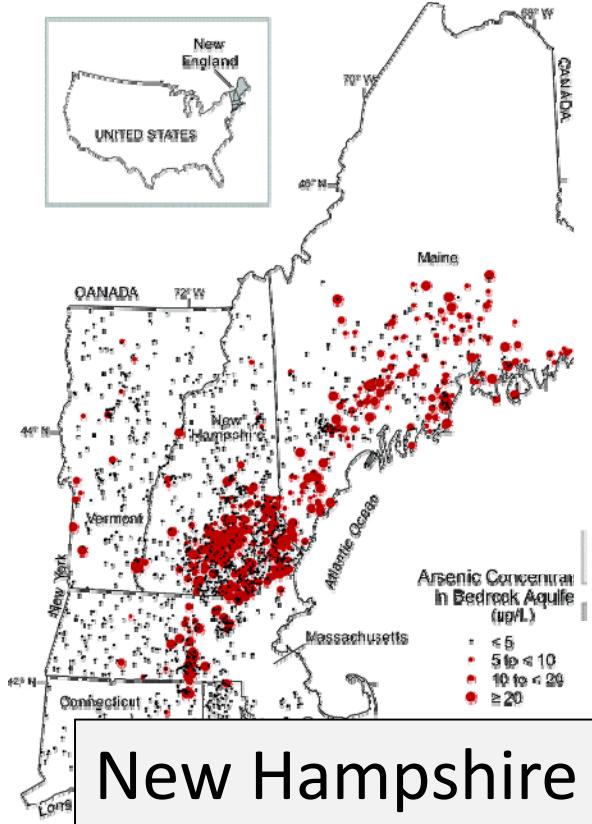


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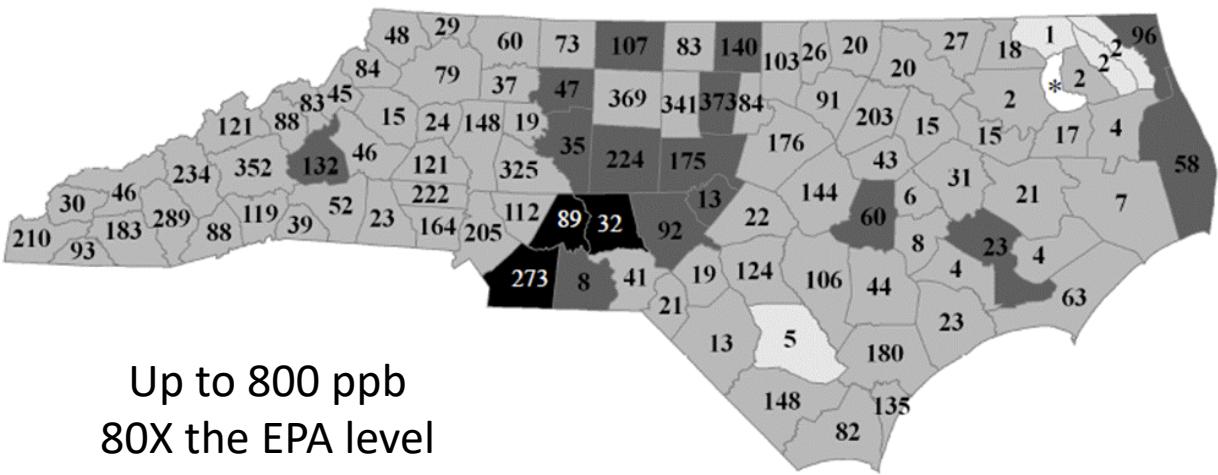


Elevated levels of arsenic in private US drinking wells

50,000 on private wells

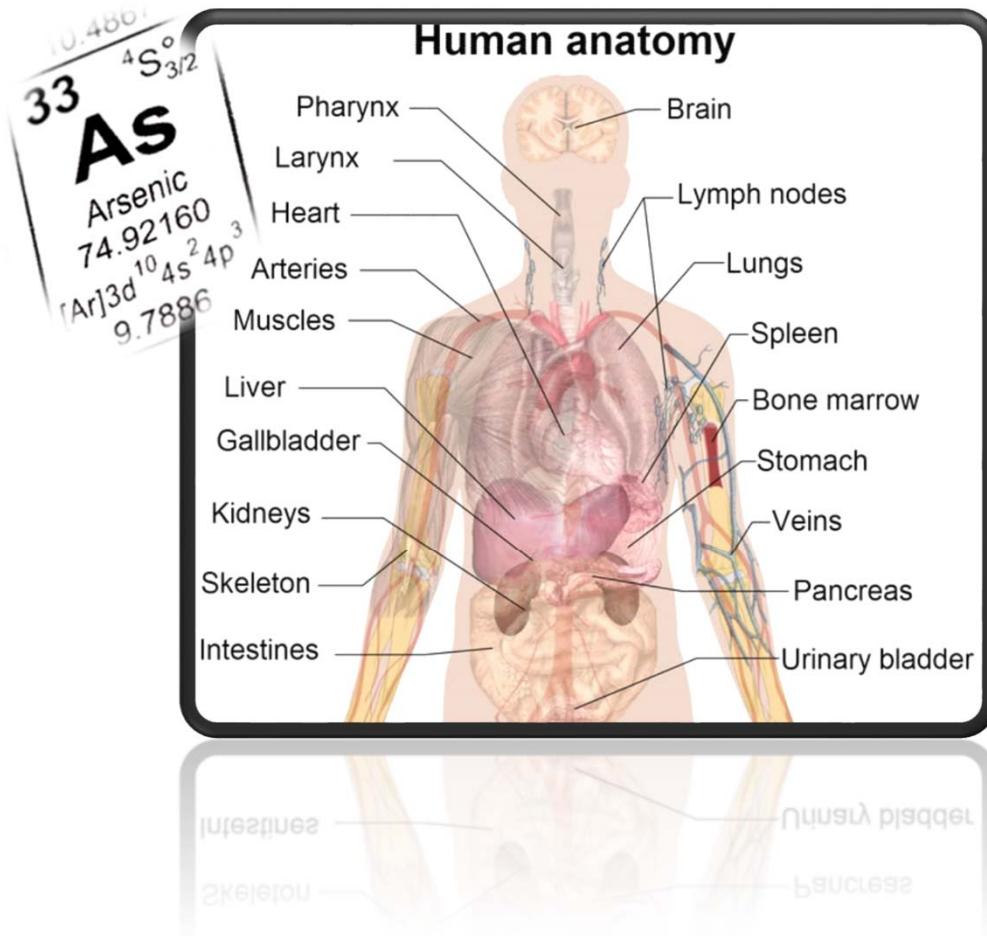


2 million on private wells



>13 million in US on private drinking wells

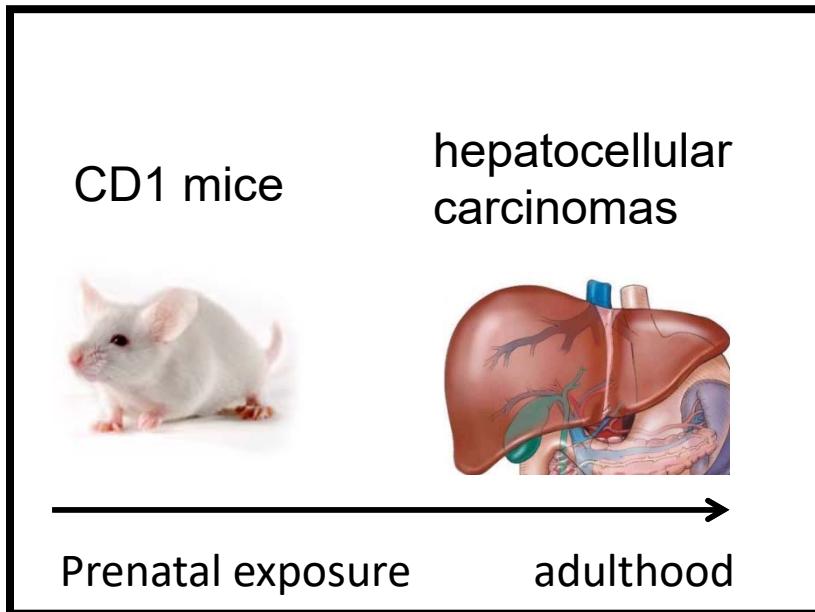
Arsenic is associated with both cancer and non-cancer endpoints



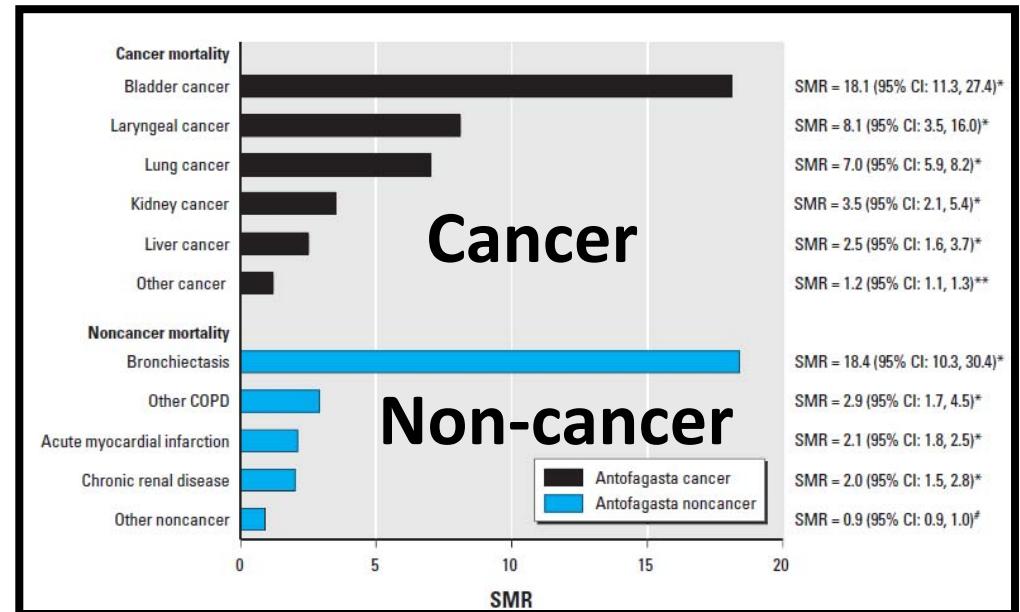
- & Classified as Group 1 Carcinogen by the International Agency for Research on Cancer (IARC): Chronic exposure results in many cancers: **skin, bladder, lung, liver, prostate and kidney**
- & Exposure is associated with non-cancer endpoints: neurological disorders, reproductive effects, cardiovascular disease, diabetes

Early life exposures associated with both short and long term health effects

Mice



Humans



Smith, A et al 20012

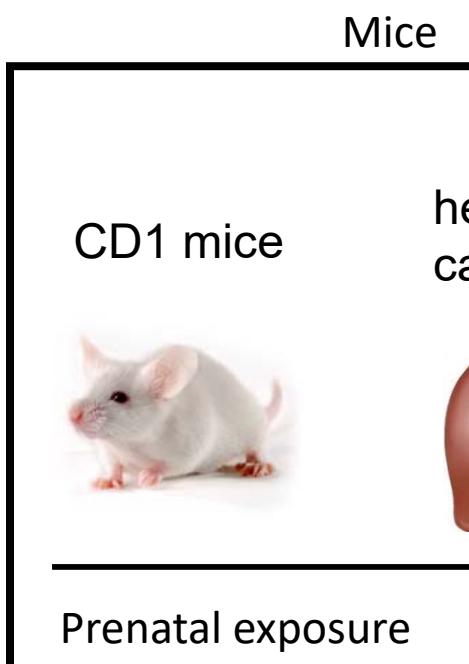
Permanent changes in gene expression



Waalkes, M. et al. Carcinogenesis 2004;
Xie, Y. et al TAP, 2007



Early life exposure and long-term health



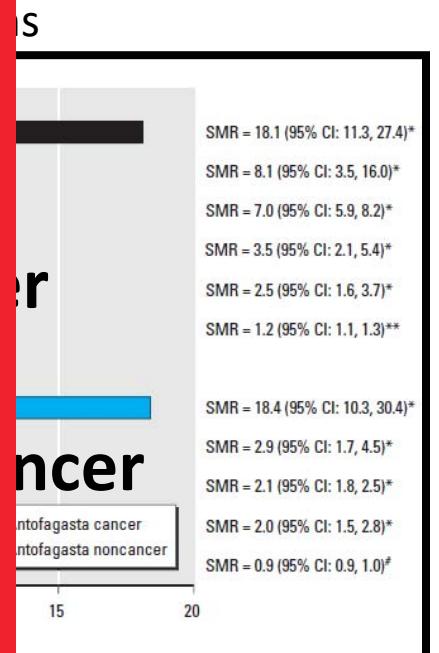
Permanent changes in



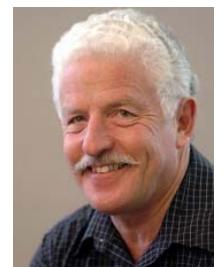
Waalkes, M. et al.



short and long



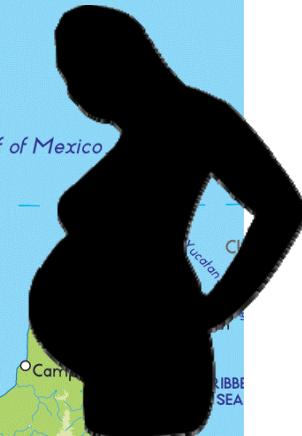
Smith, A et al 20012



What are the health effects of prenatal arsenic exposure and links to the epigenome?



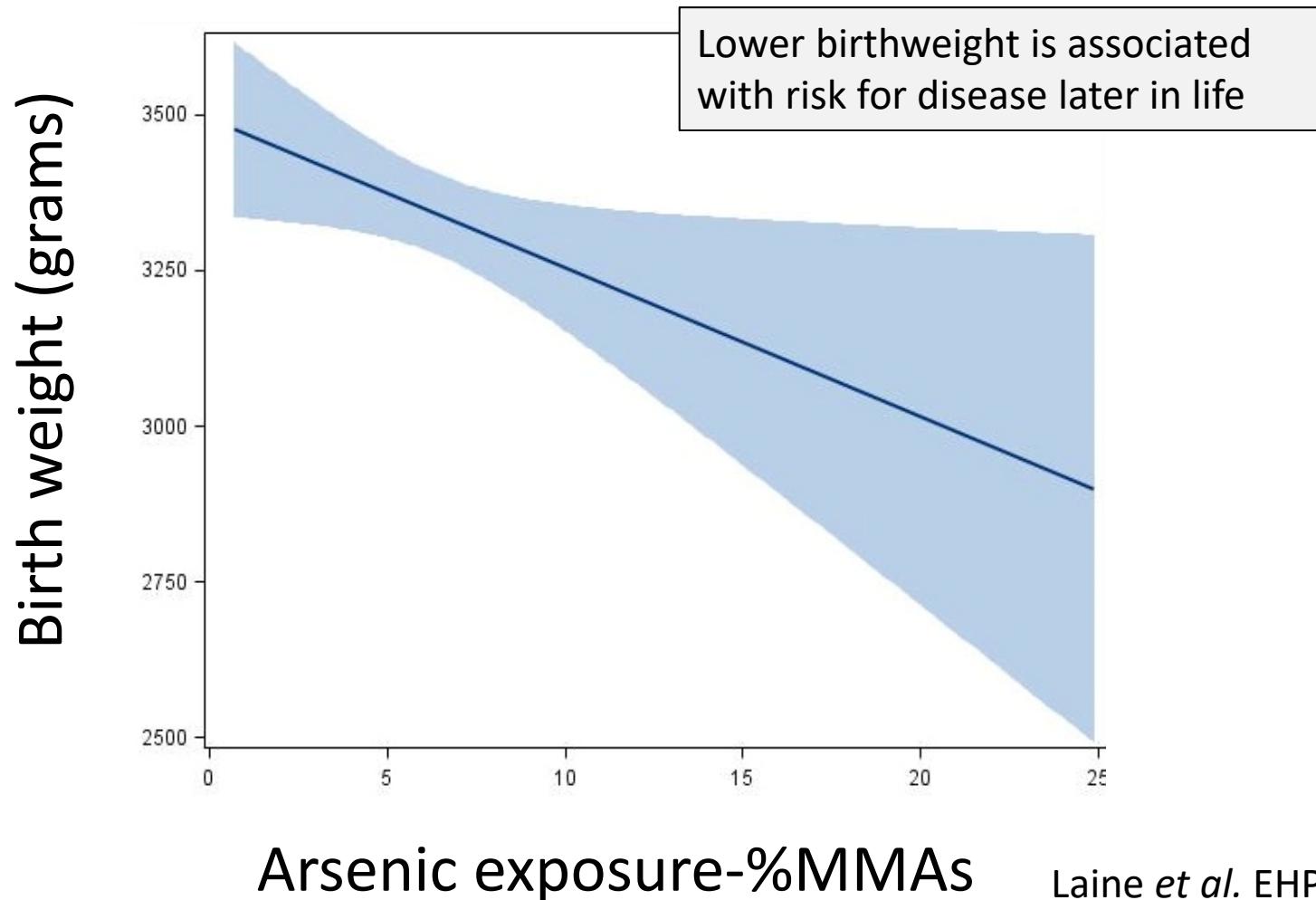
- Recruited 200 pregnant women
- Measured arsenic in drinking water and urine
- Collected cord blood for fetal DNA, RNA, protein isolation
- Measured a series of birth outcomes



What are the health effects of prenatal arsenic exposure and links to the epigenome?

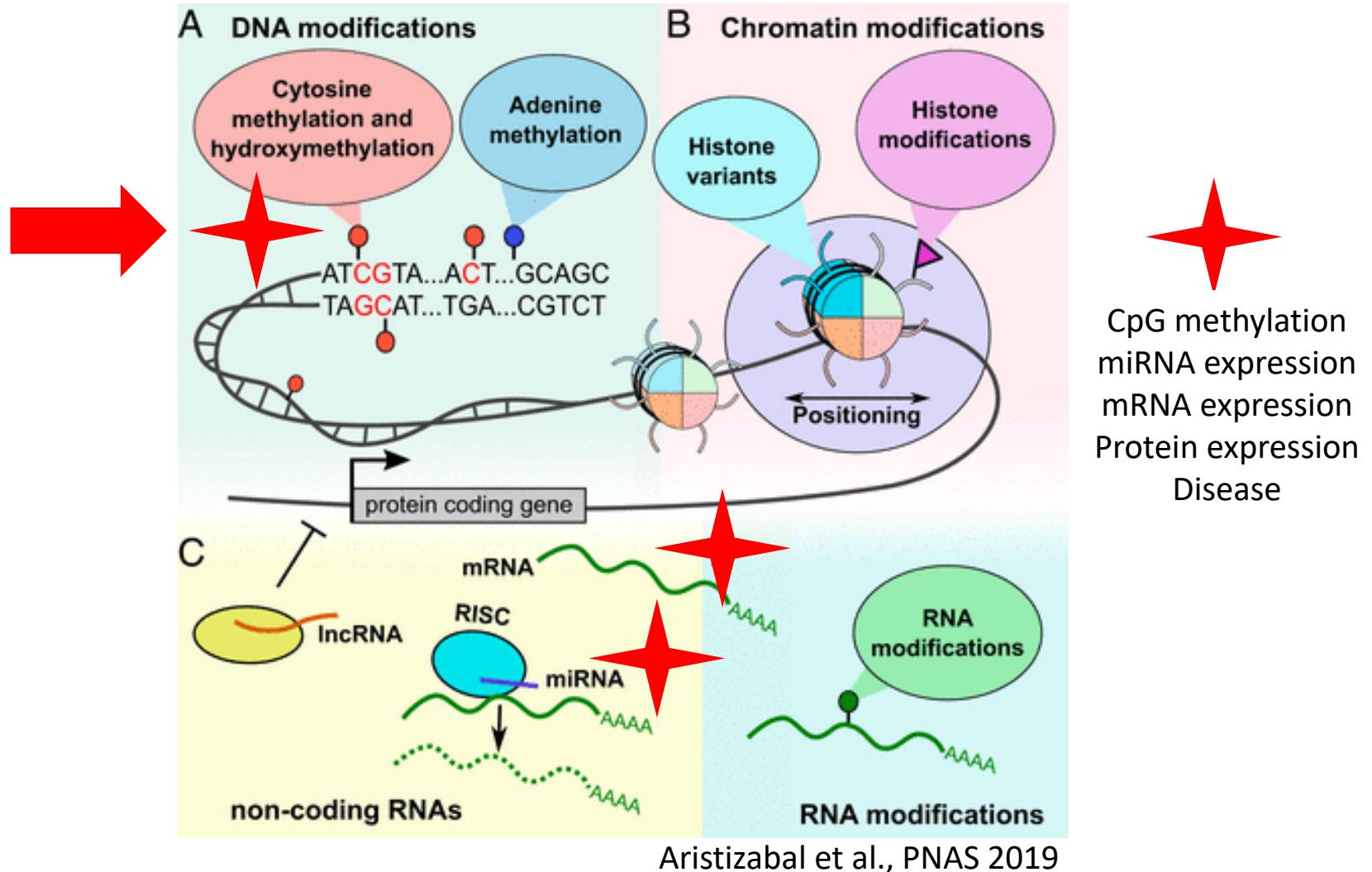


Prenatal arsenic exposure is associated with lower birthweight in infants



Huyck et al., JOEM, 2007 (Bangladesh); Kile et al. Epidemiology, 2016 (Bangladesh);
Fei et al. EH, 2013 (US); Hopenhayn et al. Epidemiology, 2003 (Chile)

What are the health effects of prenatal arsenic exposure and links to the epigenome?



Ray et al. Frontiers in Genetics 2014; Rojas *et al.* Tox Sci 2015; Laine *et al.* EHP 2015; Rager *et al.* Tox Sci 2014; Bailey *et al.* Tox Sci 2014

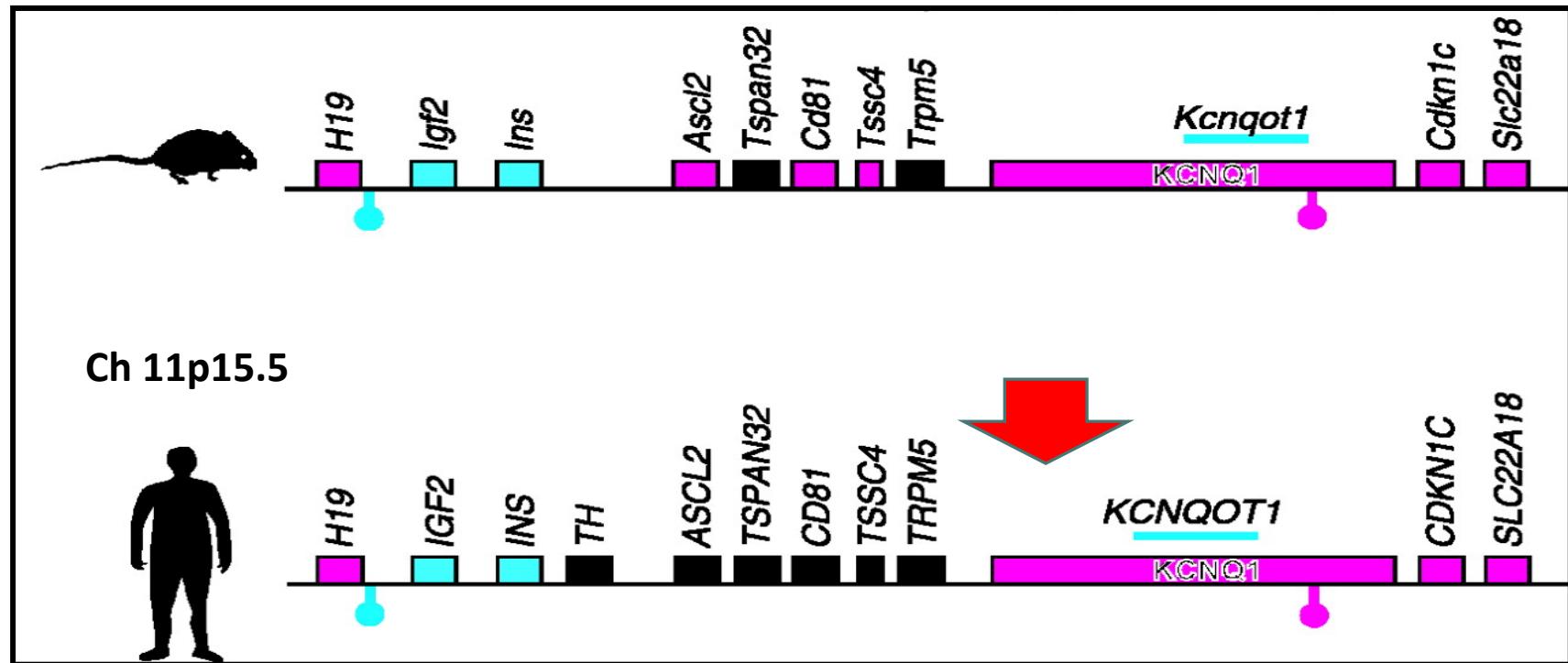
Prenatal Arsenic Exposure and the Epigenome: Identifying Sites of 5-methylcytosine Alterations that Predict Functional Changes in Gene Expression in Newborn Cord Blood and Subsequent Birth Outcomes

Daniel Rojas*, Julia E. Rager[†], Lisa Smeester[†], Kathryn A. Bailey[†],
Zuzana Drobná[‡], Marisela Rubio-Andrade[¶], Miroslav Stýblo^{*,‡},
Gonzalo García-Vargas[¶], and Rebecca C. Fry^{*,†,1}

*Curriculum in Toxicology, [†]Department of Environmental Sciences and Engineering, [‡]Department of Nutrition and [§]Department of Epidemiology, Gillings School of Global Public Health, University of North Carolina, Chapel Hill, North Carolina and [¶]Facultad de Medicina, Universidad Juárez del Estado de Durango, Gómez Palacio, Durango, Mexico

TOXICOLOGICAL SCIENCES, 143(1), 2015, 97–106

KCNQ1 is an imprinted gene on the short arm of Ch11



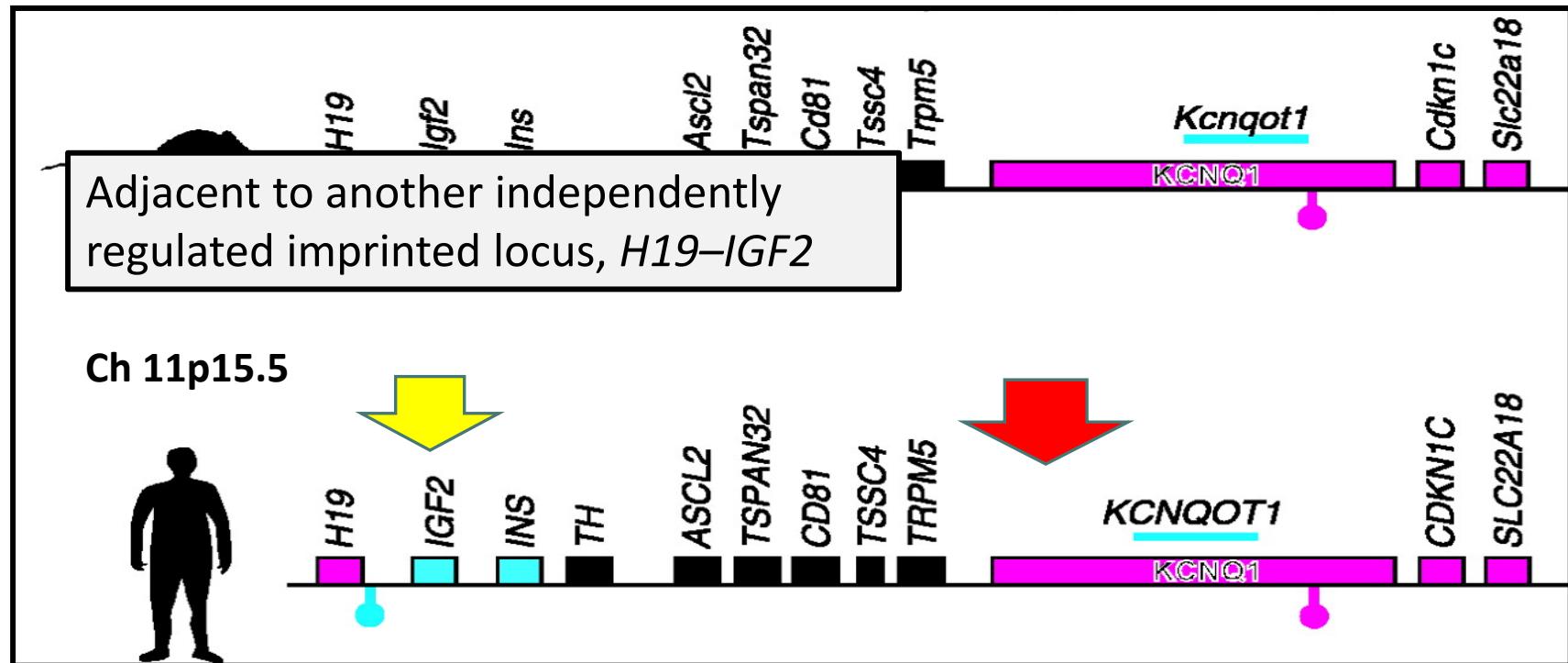
Blue=paternally expressed alleles

Pink=maternally expressed alleles

black indicates non-imprinted genes

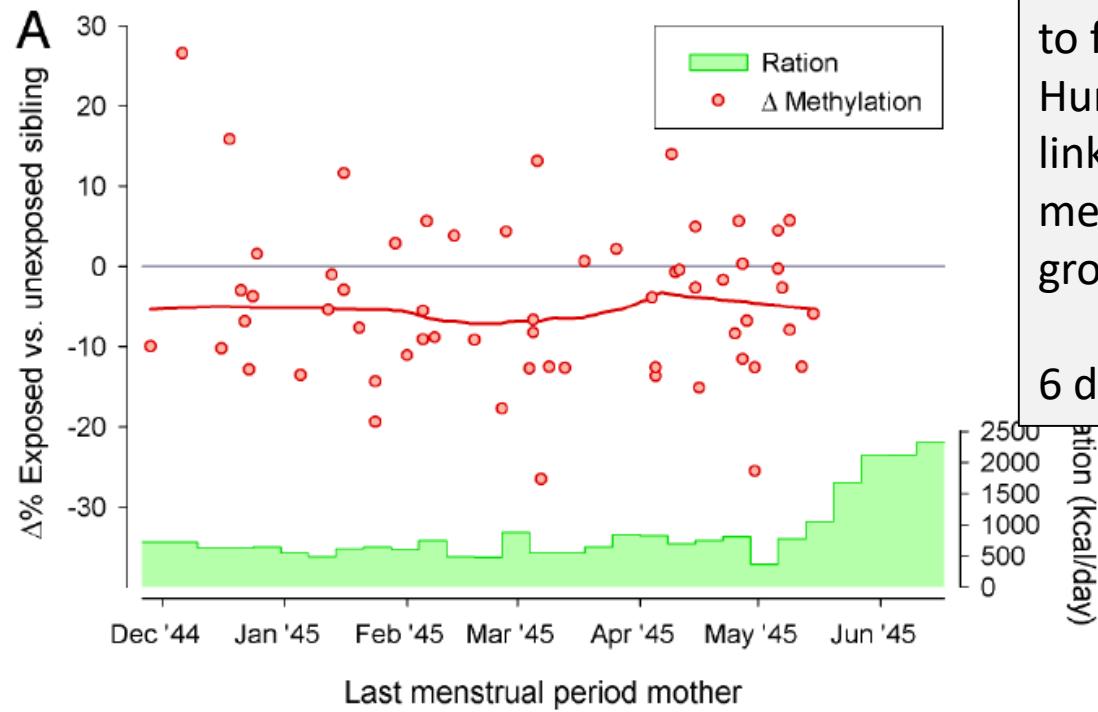
DNA methylation of *KCNQ1* (potassium channel protein) is negatively associated with gene expression and birthweight

KCNQ1 is an imprinted gene on the short arm of Ch11



Blue=paternally expressed alleles
Pink=maternally expressed alleles
black indicates non-imprinted genes

IGF2 is an imprinted gene on Ch 11p15.5, susceptible to prenatal famine-induced changes in DNA methylation



Periconceptual exposure
to famine during the Dutch
Hunger Winter (1944-45)
linked to decreased
methylation of Insulin-like
growth factor 2

6 decades later!

Persistent epigenetic differences associated with
prenatal exposure to famine in humans

Bastiaan T. Heijmans^{a,1,2}, Elmar W. Tobi^{a,2}, Aryeh D. Stein^b, Hein Putter^c, Gerard J. Blauw^d, Ezra S. Susser^{e,f},
P. Eline Slagboom^a, and L. H. Lumey^{e,1}

17046–17049 | PNAS | November 4, 2008 | vol. 105 | no. 44

KCNQ1 is associated with weight at birth, later life growth and metabolism

J Assist Reprod Genet (2014) 31:1361–1368
DOI 10.1007/s10815-014-0278-0

EPIGENETICS

Relevance of genomic imprinting in intrauterine human growth expression of *CDKN1C*, *H19*, *IGF2*, *KCNQ1* and *PHLDA2* imprinted genes

Amilcar Cordeiro · Ana Paula Neto · Filipa Carvalho ·
Carla Ramalho · Sofia Dória

Nat Genet. 2013 January ; 45(1): 76–82. doi:10.1038/ng.2477.

New loci associated with birth weight identify genetic links between intrauterine growth and adult height and metabolism

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Mechanisms of action of arsenic:

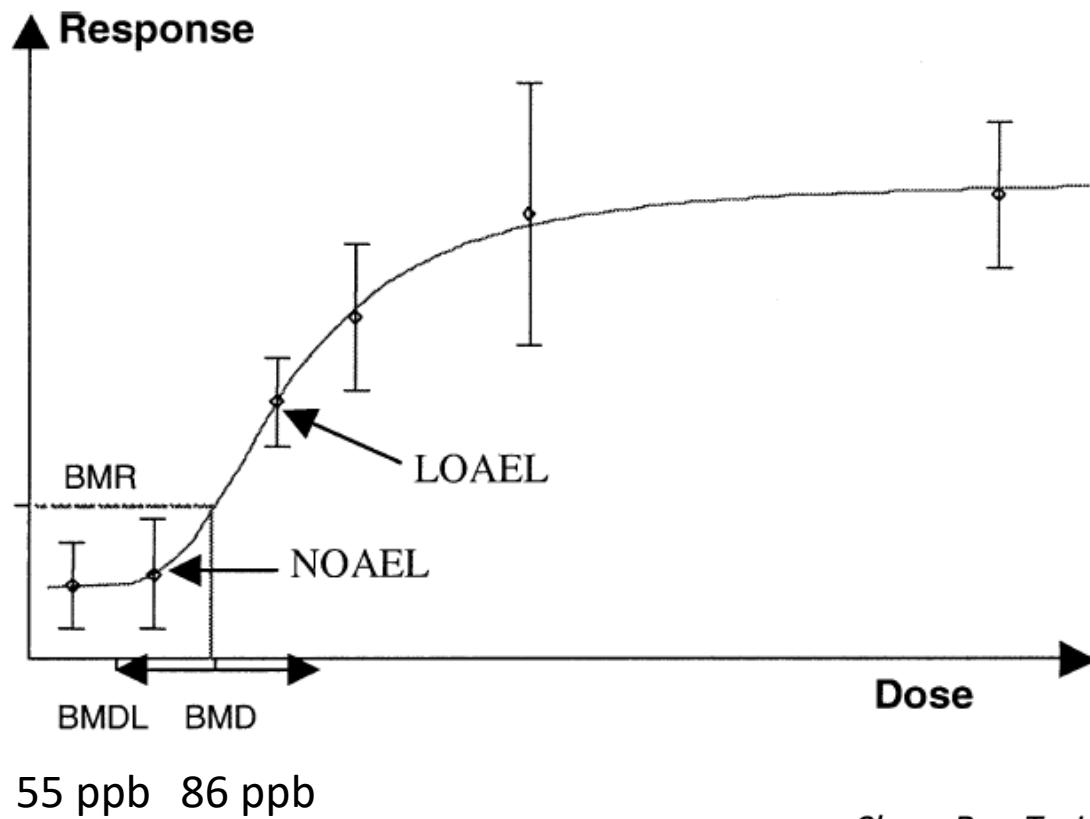
**critical genes that REGULATE
fetal growth and later life**

**health are targeted for arsenic-
associated DNA methylation**

*happens
womb lasts
fetime*

Benchmark Dose Modeling Estimates of the Concentrations of Inorganic Arsenic That Induce Changes to the Neonatal Transcriptome, Proteome, and Epigenome in a Pregnancy Cohort

Julia E. Rager^{*†} , Scott S. Auerbach[‡], Grace A. Chappell[†], Elizabeth Martin[§], Chad M. Thompson[¶] and Rebecca C. Fry[§] 



Chem. Res. Toxicol. 2017, 30, 10, 1911-

- Inorganic arsenic continues to poison individuals around the globe
- Increasing evidence for the role of the epigenome in arsenic-induced disease
- Developing strategies to integrate epigenetic data into a risk assessment framework

Acknowledgements

Funding

NIEHS Superfund:

P42 ES005948; P42ES031007

NIEHS (R01ES028721; R01ES029925;
R01ES029531; R01ES019315; P30ES010126)

NICHD: R01HD092374

