

New directions integrating genetic, environment, and possible epigenetic effects to understand causes of ADHD



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DISCLOSURES

- Royalties from Guilford Press
 - Getting Ahead of ADHD (2017)
 - What Causes ADHD (2007)
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OUTLINE

- Plausibility of an epigenetic model or paradigm for ADHD
 - Lessons from complex disease studies
- What exactly is epigenetic change?
- Are environmental correlates of ADHD causal?
 - Using genetically informed studies as one approach to find out
- Preliminary human DNA methylation findings in ADHD
- Conclusions

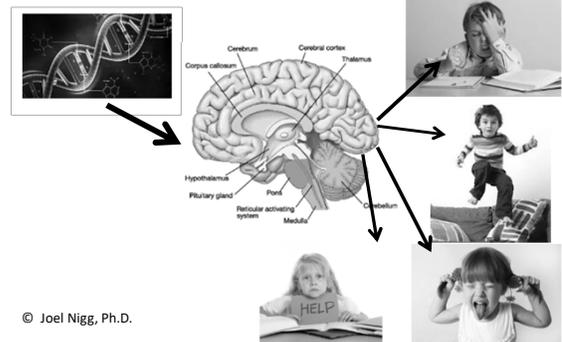
PLAUSIBILITY OF AN EPIGENETIC PARADIGM FOR ADHD

What is the paradigm?

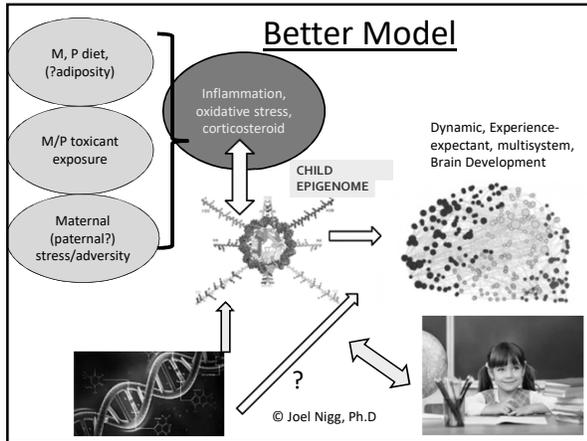
- Paradigm=exemplar (Aristotle, Kuhn)
- Wrong paradigm 1: metabolic disease
 - “find the gene, solve the disease”
- Wrong paradigm 2: Linear causality
 - “like a machine; mass=force x acceleration. Find the causal chain, solve the disease”

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OLD VIEW combined those two ideas into a single paradigms(1980s-2000's)



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3 reasons to reconsider the environment and integrate it with our progress in genetics

- Complex disease model more appropriate
- GxE (heritability of liability) hidden in heritability
- Epigenetic insight— GxE determines phenotype biologically (if not always statistically)

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Simple versus complex disease: Is it “genetic”? What does that mean?

- Single gene disorder
- Complex disease

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Simple versus complex disease: Is it “genetic”? What does that mean?

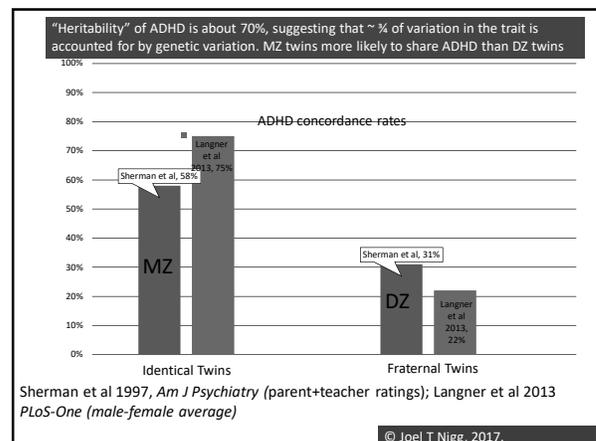
- Single gene disorder
- Deterministic
- Rare (< 1/10,000)
- Large risk increase in relatives
- PKU, Huntingtons’

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Simple versus complex disease: Is it “genetic”? What does that mean?

- Single gene disorder
- Deterministic
- Rare (< 1/10,000)
- Large risk increase in relatives
- PKU, Huntingtons’
- Complex disease
- Probabilistic
- Common (> 1/500)
- Small risk increase in relatives
- Hypertension

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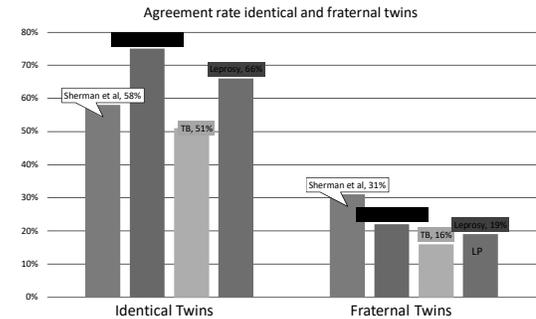


BUT: ADHD does not behave like a single gene metabolic disease

Feature	Single Gene Disorder (PKU, Huntington's)	ADHD
Role of genes	Mostly deterministic	probabilistic
Disease incidence	Rare (< 1/10,000)	Common (>1/100)
Increased sibling risk	Large (~1000x)	Small (~3-10x)

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AND the picture is not much different for major infectious diseases.



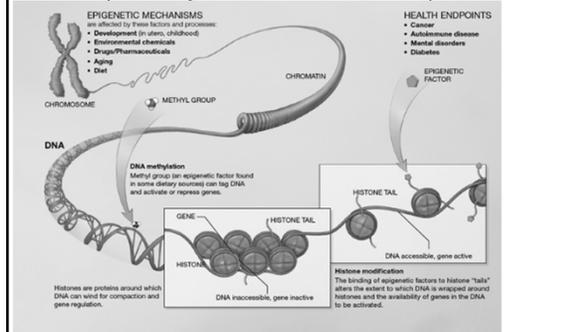
© Joel T Nigg, 2017. TB & Leprosy rates estimated from studies cited in Hill, AVS, 1998; *Ann Rev Immunol*; and in Fine PE, 1981, *Int J Lepr Other Mycobact Dis*, 49, 437-454

Proposed view of ADHD as epigenetic

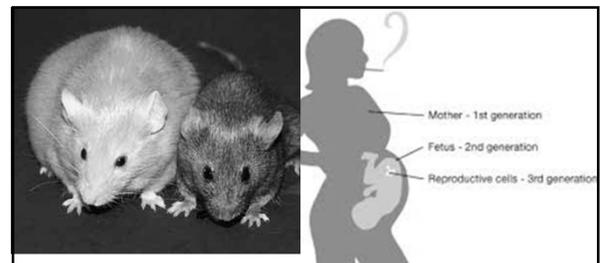
- Does NOT mean “all environmental” and not “2 types”
- ADHD heterogeneous
 - but many routes are potentially GxE or epigenetic.
 - Likely very little “all or none”
- Thus: susceptibility (substantially genetic) + experience (epigenetically mediated effects) = complex syndrome
- With
 - varying manifestations
 - temporal variations,
 - multiple routes to emergence and recovery

SHORT DETOUR: WHAT IS EPIGENETICS ANYWAY?

What is it? Epigenetic markings change gene activity, respond to gene and environmental inputs



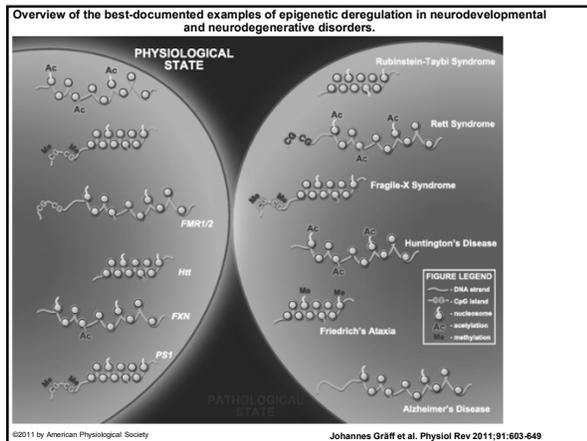
<https://commonfund.nih.gov/epigenomics/figure>. © NIH



Genetically identical animals with very different phenotype based on epigenetic change caused by feeding different toxicant-diet combination to the mother (R Jirtle et al, Duke University; *Mol Cell Biology*, 23, p. 5293, 2003; © Elsevier Inc.) MCB)

Most events affect only the person exposed, but germ line (transgenerational, 2nd or 3rd generation) effects are known. (Jablonka & Raz 2009, *Quarterly Review of Biology*, 84:131–76; © University of Chicago Press)

Epigenetic Effects based on experiences can, in principle, be as large as genetic effects, although this is unknown in humans



ARE ENVIRONMENTAL CORRELATES OF ADHD CAUSAL?

If we accept a susceptibility model of ADHD: Which Environments do we study and how do we do it?

- Sociological Effects
 - Collapse of civilization
 - Too much pharma marketing
 - Performance pressures on children, starting school too young
- Caregiver Problems
 - Over-indulgent or else hostile/intrusive parenting
 - Under-trained or inexperienced teachers
- Developmental and Biological Context
 - Rare events
 - Perinatal problems, teratogens (alcohol, drugs); micro-ischemias
 - Extreme toxicant exposures, extreme neglect (Romanian orphans)
 - *** • Common but harmful contexts
 - Modern screen Media
 - Moderate psychosocial stress/distress (esp. prenatal)
 - Poor diet
 - Low grade Toxicant/pollutant exposures (pre-natal, post-natal)

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Substantial literature links ADHD to environmental risk factors in development

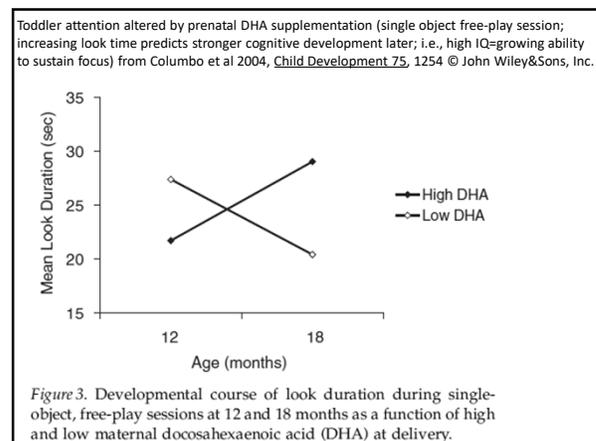
- Toxicants
 - Lead; PCB's; BPA; Pesticides
- Dietary insults
 - Western high-fat diet during gestation
 - Western diet (additives) in development
- Gestational and perinatal risks
 - Parental stress, BMI, smoking, other exposures
 - Infant distress, birthweight, delivery complications

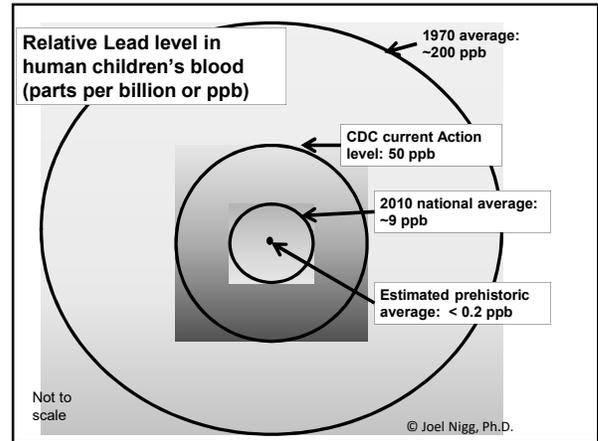
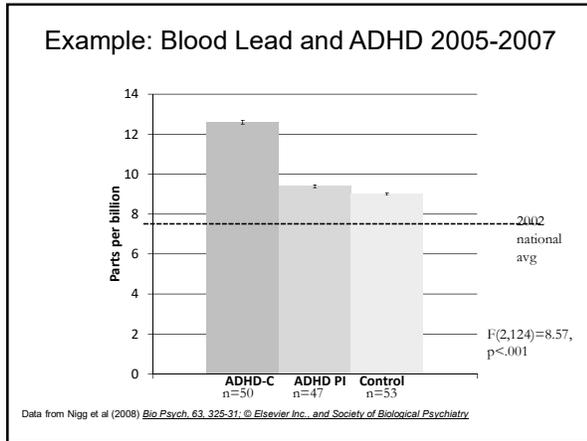
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But are these causal?

- Plausibility (can "low amounts" do harm?)
- rGE and unexamined genetic effects
- Unmeasured confounders

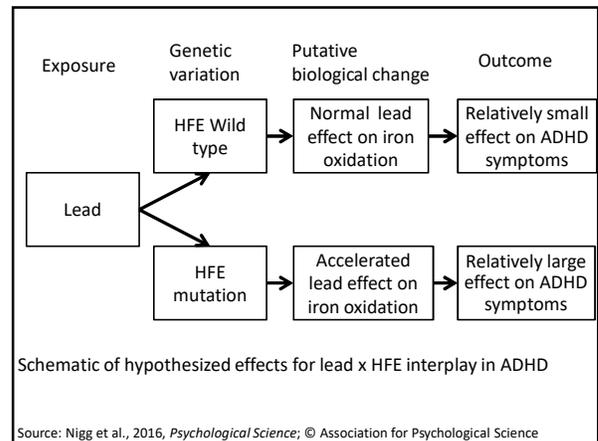
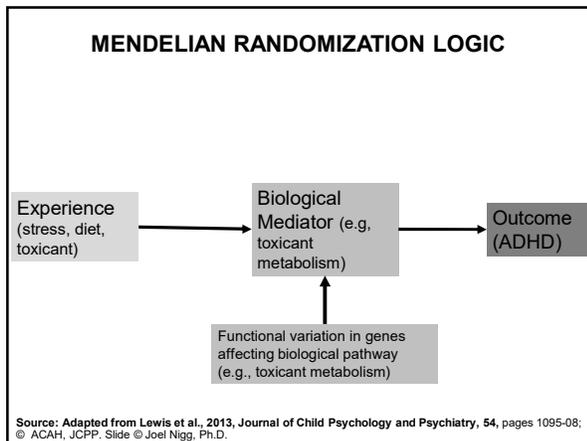
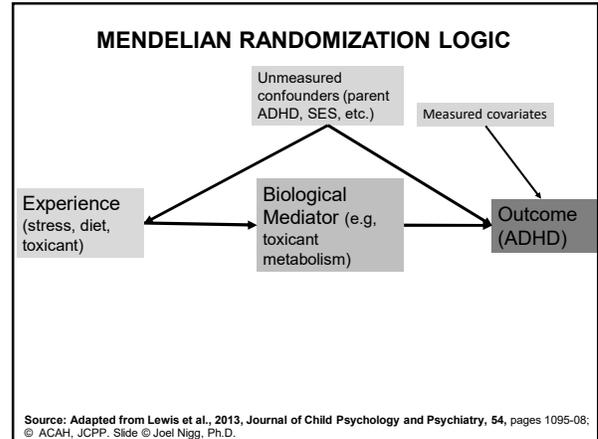
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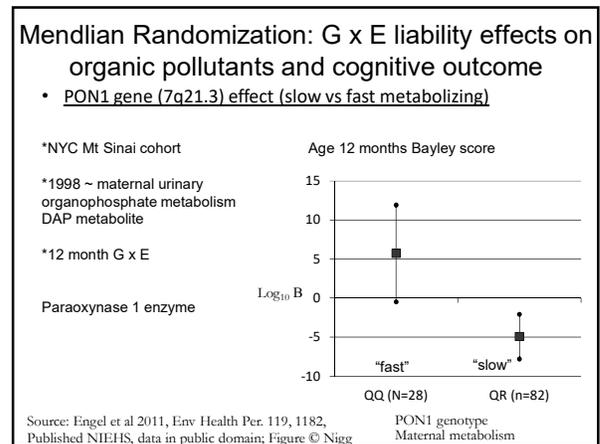
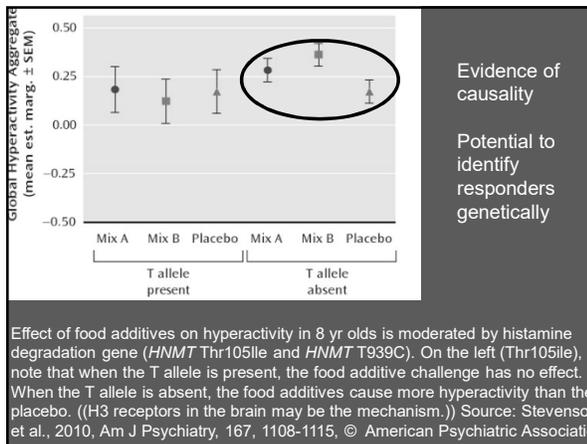
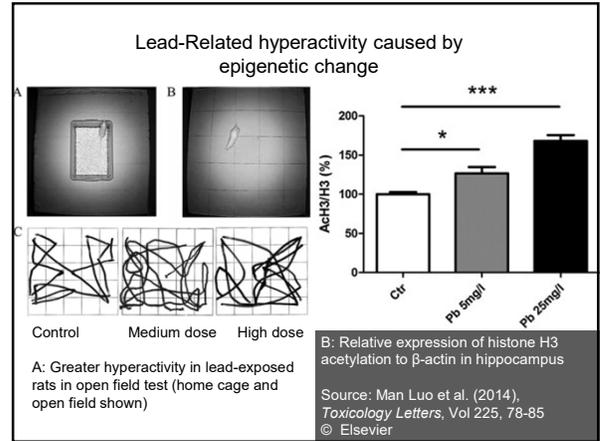
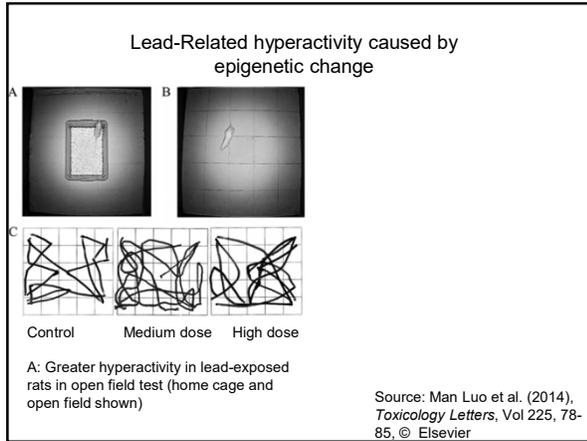
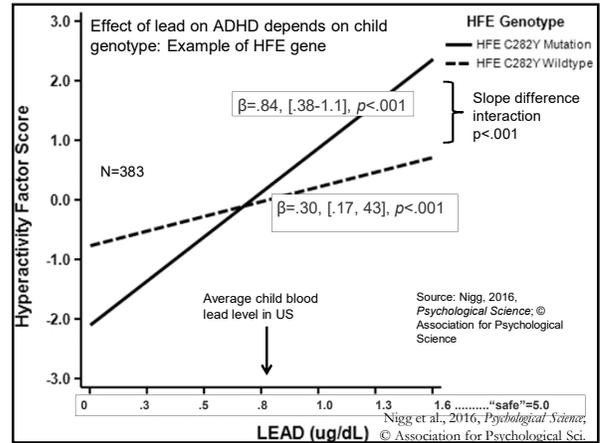
How can we evaluate causality of environmental influences on ADHD in humans when experimental tests are not possible?

- Surrogate pregnancy—(e.g., smoking, Thapar et al)
- Sibling, twin, adoption designs
- Natural stratifications (e.g., Dutch famine; or one city stops vaccinations)
- Mendelian randomization** (focus today)



How we proceeded on lead+ADHD

- Replicated the ADHD-low-lead correlation (n=213)
- Then combined both samples, (Total N=363; ADHD+control)
- Mendelian randomization design
- HFE gene (6p22.2)
 - iron uptake in gut, lead x iron interplay
- Weakness: Lacked an independent replication
- Strengths of our study
 - ADHD very well characterized
 - Genotype frequencies matched the regional population
 - Control group blood lead levels matched the population
 - No high blood levels (max=3ug/dL)
 - rGE controlled
 - Race/ethnicity, SES controlled



Examples Linking findings in ND: causally informative designs

- Lead → ADHD (Nigg et al, 2016)
- Lead → epigenetic change → RNA brain → hyperactivity (Luo et al. 2014)
- Prenatal chemical toxicant → ADHD, IQ, autism (e.g., Engle et al 2011)
- Prenatal omega-3 intake → infant IQ (e.g., Columbo et al 2004)
- Food additives → ADHD (Stevenson et al 2010)
- Epigenetic mediation (e.g. Skinner et al 2014)
- We should not be uncritical but should consider these linkages carefully

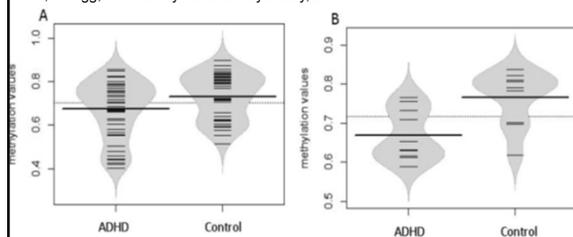
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Epigenetic studies in ADHD in humans

- Strategies
 - Population based vs enriched case-control
 - Candidate gene/probe vs MWAS
- Cautions
 - Tissue specificity and inaccessibility
 - Biomarker vs causal pathway
 - Dynamic
 - Multi-causal (genetic, experience, random)
 - But may help find mechanism of gene action

? Epigenetic biomarkers? First methylome-wide epigenetic study of ADHD—one interesting finding

VIPR2 probe cg 13444538 methylation values of ADHD and non-ADHD boys in a discovery cohort (n=88, panel A, p=.03) and a replication cohort (n=20, panel B, p=.006) showing decreased methylation in ADHD. Source: Wilmot, Fry, Musser, Mill, & Nigg, J Child Psychol and Psychiatry, 2016



VIPR 2 also linked to psychosis, circadian clock, and brain development, a rodent knockout model confirms link to hyperactivity

Key findings in DNA methylation in ADHD in children

- Van Mil et al J Psych Res 2014: 7 candidate genes examined in cord blood → ADHD sx at age 6 (CBCL); DRD4, 5HTT.
- Walton et al (2016), altered DNA methylation at birth associated with future ADHD trajectory (AVON study, n=872)
- Barker et al 2017 (Child Dev)(n=671, ALSPAC), cord blood MWAS related to ODD and overlap with ADHD symptoms
- Wilmot et al 2016 (JCPP), n=92 boys case control MWAS, VIPR2 (later confirmed for boys) and MYT1L (confirmed in mixed male-female sample later)

Conclusions

- ADHD hypothesized as at least in part an epigenetic response to widespread low grade insults in genetically susceptible children
- At least some exposure effects appear to have causal link to ADHD
- Genetically informed studies of environment can clarify causal effects
- Initial epigenetic effects in ADHD show promise, although many cautions are in order



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