## Placental DNA Methylation Linking Exposures and Newborn Health

Carmen J. Marsit, PhD

Pharmacology & Toxicology

Epidemiology

Geisel School of Medicine at Dartmouth



THE CHILDREN'S ENVIRONMENTAL HEALTH & DISEASE PREVENTION RESEARCH CENTER AT DARTMOUTH

## Where we study methylation matters

### DNA Methylation is highly tissue specific

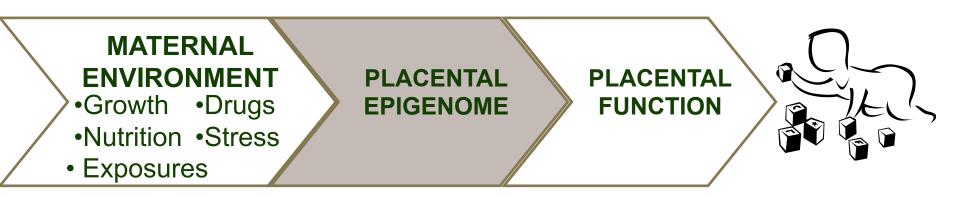
Represents functional alteration

#### Placenta

- First complex organ to form
- Regulates intrauterine environment
- Transport
  - Nutrients
  - Water
  - Gas
  - Waste products
- Immuno-endocrine
  - Hormones
  - Growth factors



# Role of Placental Epigenome



- Demonstrate placental molecular features
   integrate environmental signals
- Link variability in molecular features to Infant Outcomes (and beyond!)



## Study Population: Rhode Island Child Health Study

- Hospital-based Birth Cohort
  - Moms enrolled following delivery at Women & Infant Hospital
  - WIH sees 75% of deliveries in RI
  - 2009-2014
  - Total Enrollment n=899
- Mothers
  - 18-40 years old
  - No history of psychological disorders
  - In good physical health
- Infants
  - Viable
  - No known genetic disorders
  - No life threatening illness
  - Term (≥ 37 weeks)
- Oversampled for SGA (small) and LGA (large) infants. Matched to AGA (appropriate)
  - Gestation time, infant sex, maternal age (±3 yrs)



#### Mom • Medical

- Medical Chart Review
- Questionnaire
- Maternal blood
- Toenail Samples

#### Delivery



#### Infant Clinical Characteristics

- NNNS Assessment
- Placenta
- Cord Blood
- Toenails

Linking Molecular Character with Exposures and Outcomes



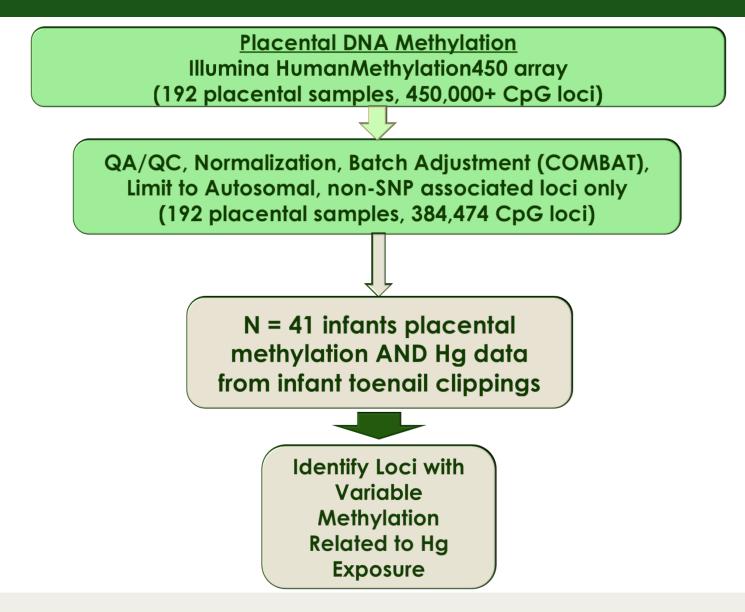
# Mercury exposures in utero

- Sources of exposure
  - Dietary (fish, seafood) i.e. methylmercury (MeHg)
    - Biomethylated, biomagnified
  - Dental amalgams
  - Industrial
    - Minimata disease (Japan) (Harada et al. 1968)
    - Iraqi fungicide contamination incident (Bakir et al. 1973)
  - Air Pollution
  - Can be assessed with finger/toenail measurements (He et al. 2013; Wickre et al. 2004; Xun et al. 2013; Hinners et al. 2012)
- Crosses placenta (National Research Council 2000; Yang et al. 1997; Ilbäck et al. 1991)
  - Interferes with placental function (Boadi et al. 1992)
- Neurobehavioral effects associated with prenatal and childhood exposure
  - Infants: Cerebellum size, CNS damage, poor psychomotor development (Cace et al. 2011; Choi 1989; Llop et al. 2012)





# Preliminary - Discovery Study





## Methylation Impacting Neuro-related Genes

#### 339 CpG loci associated with infant toenail Hg

Within genes or gene promoters associated with neurologic outcomes

Neurodevelopment (DIXDC1, NRBP2, KIF26B, FEZF1, DMRTA2, ACTN1, MYO10, LRFN1) (Singh et al. 2010; Kivimae et al. 2011; Larsson et al. 2008; Heinrich et al. 2012; Eckler et al. 2011; Shimizu and Hibi 2009; Watanabe et al. 2009; Shimizu et al. 2010; Yoshizawa et al. 2011; Konno et al. 2012; Kremerskothen et al. 2002; Silver et al. 2012; Ju et al. 2013; Raines et al. 2012; Yu et al. 2012; Morimura et al. 2006)

Neurobehavior (CPLX1, LMX1B, ADD2) (Drew et al. 2007; Glynn et al. 2007; Barreto-Valer et al. 2013; Porro et al. 2010)

Schizophrenia (DIXDC1, ARVCF, MAGI2, ZIC2) (Bradshaw and Porteous 2012; Sim et al. 2012; Mas et al. 2010; Mas et al. 2009; Chen et al. 2005)

ADHD (TCERG1L) (Neale et al. 2010; Karlsson et al. 2012; Hatayama et al. 2011)

Movement disorders (NOL3, TP53INP2) (Russell et al. 2012; Bennetts et al. 2007)

Autism (PLXNA4, WNT2) (Suda et al. 2011; Lin et al. 2012; Kalkman 2012)



J.Z.J. Maccani, et al. EHP 2015 (epub ahead of print)

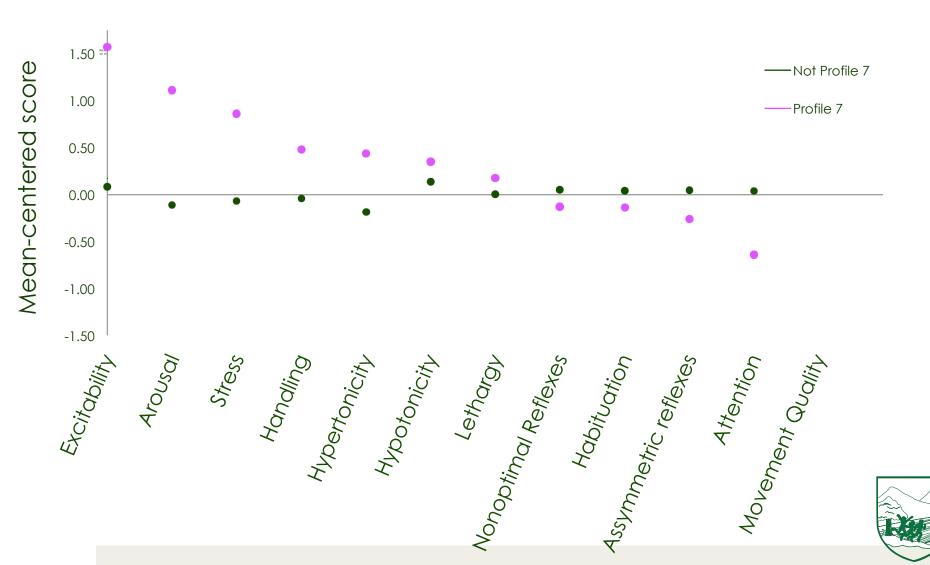
## Looking at Neurobehavioral Effect

### NICU Network Neurobehavioral Scales (NNNS)

- Developed by Lester and Tronick (2004)
  - Built on backbone of Neonatal Behavioral Assessment Scales (NBAS) developed by Brazelton (1973)
- Developed for use in at-risk infant
  - Specifically substance exposed used in the Maternal Lifestyle Study
  - Designed for broad applicability
    - Generalizable
    - Reproducible
    - Sensitive to variety of infant risk factors
    - Infants from 30 weeks gestation to ~2months post-partum
- Approx. 30 minute exam performed after 24 hours of life but prior to discharge
- Examine three major area of neurodevelopment
  - Neurological
  - Behavioral
  - Stress/Abstinence
- Summarized into 13 Summary Scales reflecting various aspects of neurodevelopment
  - Can examine individually
  - Used latent profiling strategy to create profiles of neurobehavior (Liu et al Pediatrics 2010)



## Focused on "At-risk" profile



# Discovery Scheme

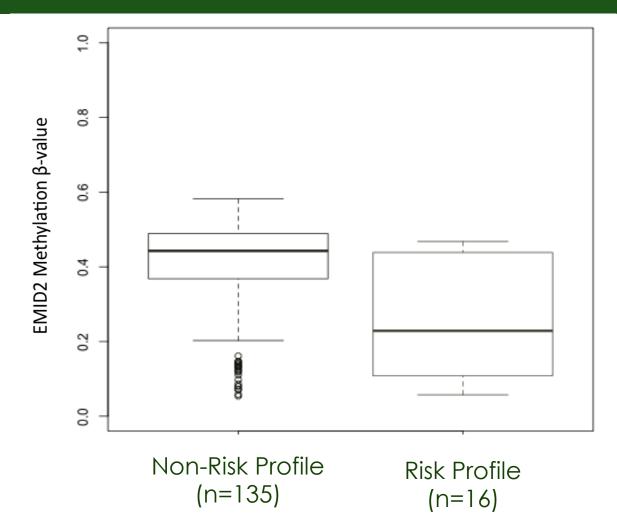
Illumina HumanMethylation450 placental tissue array (192 placental samples, 450,000+ CpG loci) QA/QC, Normalization, Batch Adjustment (COMBAT), Limit to Autosomal, non-SNP associated loci only (192 placental samples, 384,474 CpG loci) Subset 1: Subset 2: N = 41 infants placental N = 151 infants with placental methylation AND Hg data from methylation AND infant infant toenail clippings **Neurobehavioral Assessments Examine Relationship of** 339 Loci Related to Hg-Associated Loci with **Hg Exposure NNNS** Profiles

# 6 CpG Hg-associated loci are associated with NNNS High Excitability Profile

Illumina CpG Designation	Gene Symt	ool P Value	UCSC CpG Island Designation
cg13267931	EMID2	8.25x10⁻ <sup>6</sup>	Island
cg14874750	EMID2	6.06x10 <sup>-5</sup>	Island
cg23424003	EMID2	7.30x10 <sup>-5</sup>	Island
cg27179533	EMID2	5.46x10 <sup>-5</sup>	Island
cg27528510	EMID2	9.00x10 <sup>-5</sup>	Island
cg14048874	EMID2	0.0023	Island
cg14175932		2.84x10 <sup>-5</sup>	
cg17128947	CPLX1	0.0054	Island
cg25385940	ПС23	0.0059	N Shore
cg10470368		0.0075	

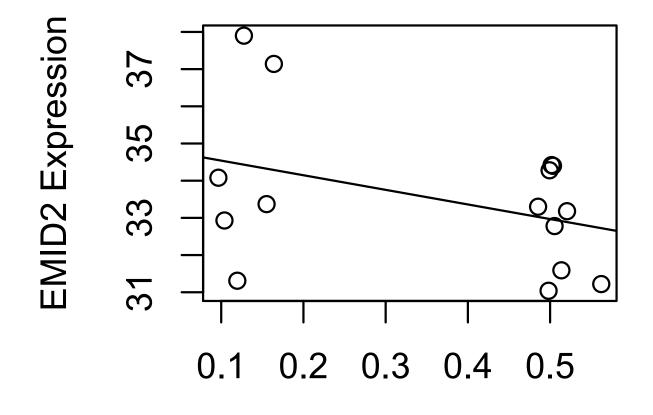


# Placental hypomethylation of *EMID2* associated with Risk Behavioral Profile





# Methylation Correlated with Expression



**EMID2** Mean Methylation



# EMID2

Collagen protein, unknown placental function

- Variant associated with asthma/airway hyperresponsiveness in nasal passages (Pasaje et al. 2011; Pasaje et al. 2012)
- A SNP within EMID2 mediates side effects on vision and hearing in response to an antidepressant (Adkins et al. 2012)
- More work needed to understand the functional role of this gene in placenta

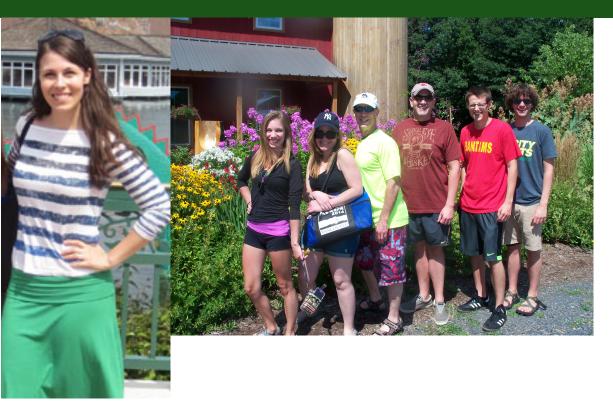


## DNA Methylation as Toxicant Mechanism

- Alterations to genes or pathways can have long-term consequences on development
- DNA methylation is susceptible to environmental signals
  - Toxicant Exposures
  - Maternal Factors/Lifestyle
  - Stress, Psychosocial adversity
- Can then link altered DNA methylation to critical outcomes
- Ongoing work
  - Better defining what environment can do
  - Consequences of these alterations Various Health Outcomes



## Acknowledgments



<u>Funding</u> NIMH R01MH094609 NIEHS R01ES022223 NIEHS P01 ES022832; EPA RD83544201

#### **Dartmouth**

Dave Armstrong Corina Lesseur Perez Allison Paquette Allison Appleton Dylan Guerin Brian Jackson Margaret Karagas & Children's Center Team

#### **Oregon State**

Andy Houseman <u>Women and Infants</u> <u>Barry Lester</u> James Padbury Joyce Lee Erica Oliviera <u>Brown</u> Jennifer Maccani Karl Kelsey



THE CHILDREN'S ENVIRONMENTAL HEALTH & DISEASE PREVENTION RESEARCH CENTER AT DARTMOUTH