Epigenetics as a Biomarker in Environmental Health

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Epigenetics in perinatal programming





Boekelheide et al. Environ Health Persp. 2012

DNA methylation

- 5-methylcytosine residues; usually found in CpG dinucleotides
- Often associated (negatively) with gene expression
- Well studied





Beyond candidate gene approaches: Epigenome-wide platforms







High Density Microarrays

Illumina 450k and 850k

Cost-effective - measures hundreds of thousands of pre-specified sites



flickr.com/photos/jgr/2771104231/

Early exposures leave a fingerprint on the epigenome: results from a meta-analysis of prenatal maternal smoking and cord blood methylation



Joubert et al. The American Journal of Human Genetics 2016 98, 680-696



Swimming in data: a new challenge for environmental health rgctategetaaat ctcagaattact Catctttaggta

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Releaset

Need for new diagnostic tools

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Reliant on multi-step pre-processing pipelines



Mount Sinai

Wilhelm-Benartzi et al. Br J Cancer 2013

Quality Control is Essential

Check your dataset carefully!

- Look for sample mix-ups, failures, and contamination
 R package *ewastools* on github
- Thoughtful detection p-values (fluorescence signal quality) slashes rate of outliers
 Heiss and Just. BioArxiv preprint: https://doi.org/10.1101/245217

Even low numbers of misidentified samples or big outliers can have a devastating impact on power and robustness of findings Technical replicates can give estimate of Signal versus Noise



Chen and Just et al. CpGFilter: Model-based CpG probe filtering with replicates for epigenome-wide association studies. Bioinformatics 2015

Improved analytic approaches

Restrict the search:

- Prioritize best measured features
- Prioritize biologically relevant features

Borrow information:

- Pathway analyses
- Regional analyses



Larger sample sizes

Epigenomic Meta-analysis Consortia:

Prenatal And Childhood Epigenetics (PACE)

Felix JF et al. 2017. Cohort profile: Pregnancy and childhood epigenetics (PACE) consortium. *Int J Epidemiol*.



Pregnancy And Childhood Epigenetics

Cohorts for Heart & Aging Research in Genetic Epidemiology (CHARGE)



Note: correction for multiple comparisons doesn't protect against confounding: high type I error in epigenomic epi



Summary: epigenomics in environmental health

1. Exciting prospects to discover links with early

environmental exposures and later outcomes

- 2. Measurements are standardized; analyses are complicated
- 3. Need care when n<<p; borrow information using biology and statistics



Thank you!

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