

Integrative Exposomic, Transcriptomic, Epigenomic Analyses of Human Placental Tissues Links Understudied Chemicals to Preeclampsia

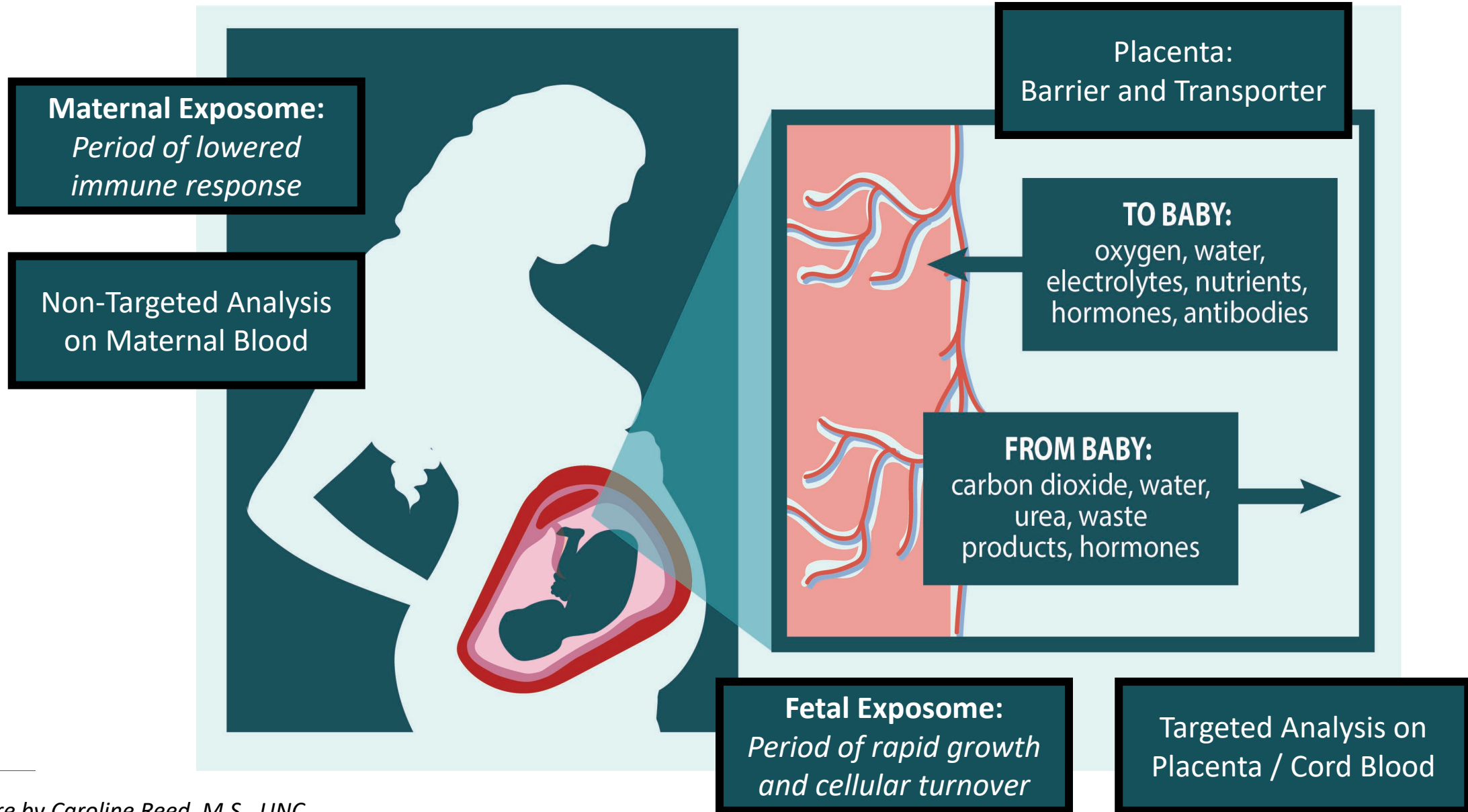
Alex Chao

Jarod N. Grossman, Celeste Carberry, Yunjia Lai, Antony J. Williams, Jeffrey M. Minucci, S. Thomas Purucker, John T. Szilagyi, Kun Lu, Kim Boggess, Rebecca C. Fry, Jon R. Sobus, Julia E. Rager



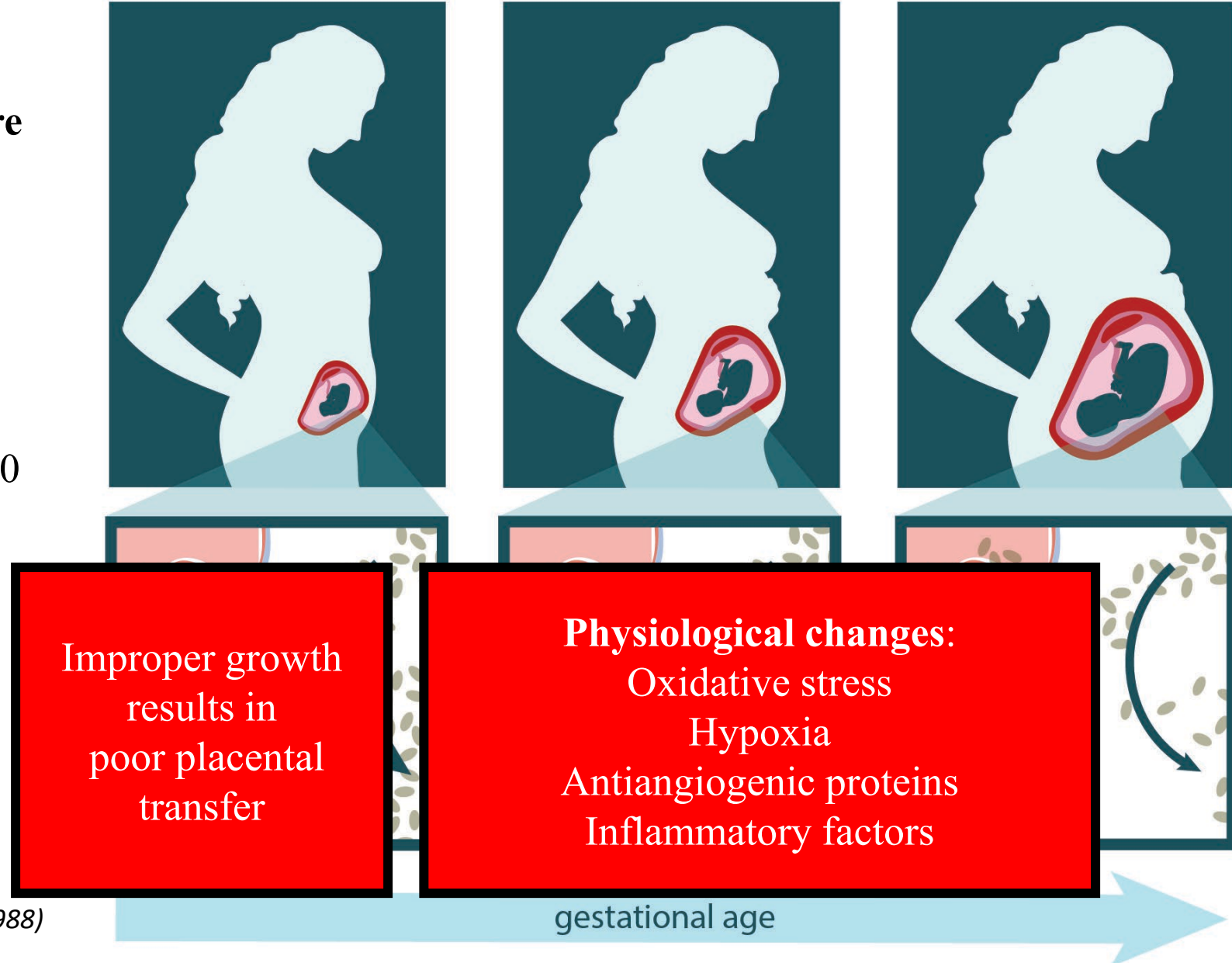
The views expressed in this presentation are those of the author and do not necessarily reflect the views or policies of the U.S. EPA

Pregnancy: Two Susceptible Populations

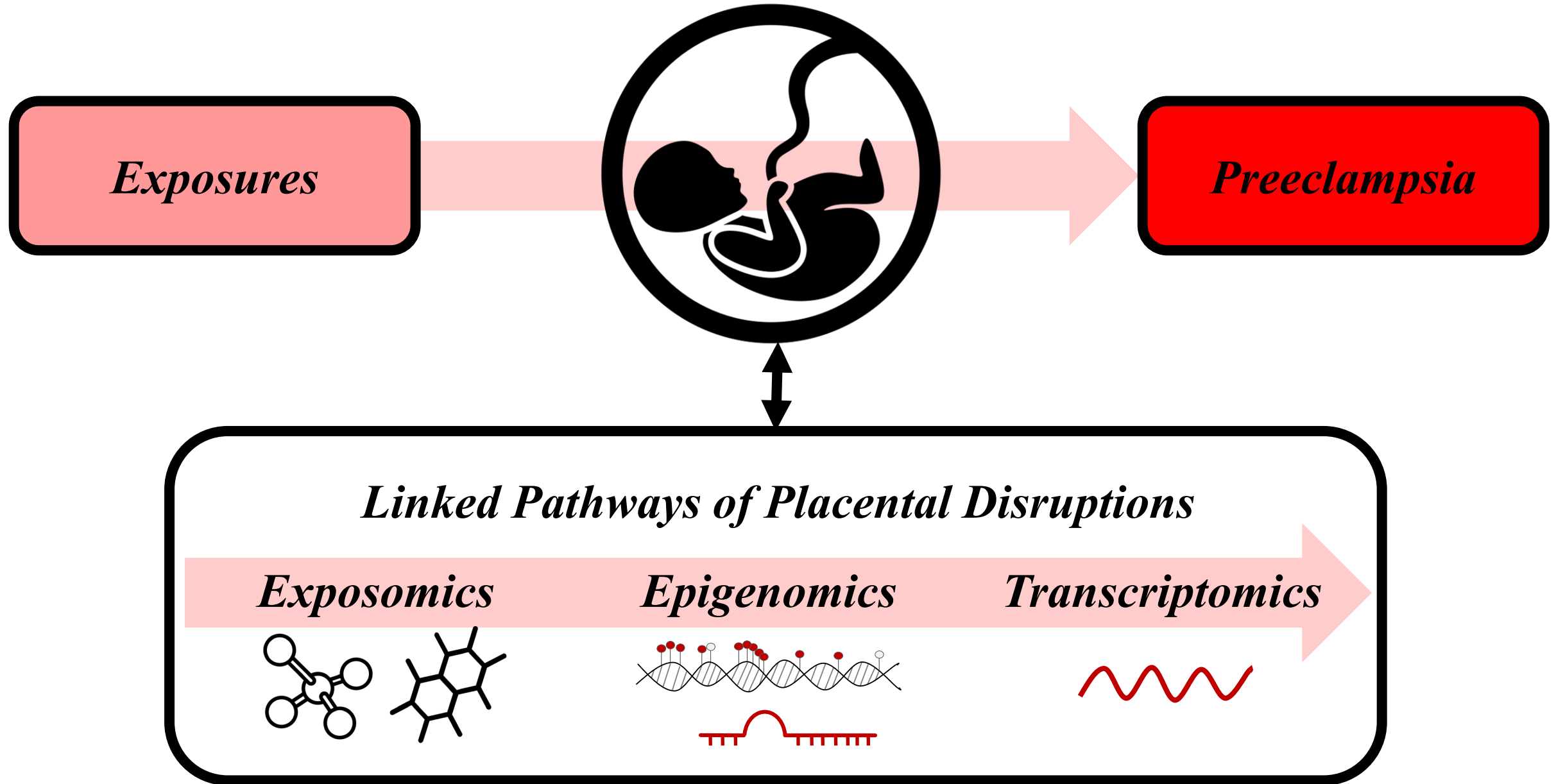


Preeclampsia: A Serious Pregnancy Disorder

- Preeclampsia is a prenatal disease characterized by **high blood pressure** and elevated urine protein levels (indicative of **organ damage**)
- Preeclampsia affects up to 8% of pregnancies worldwide
- Deaths of ~70,000 women / ~500,000 fetuses yearly
- Exact causes are still unknown
- Associated with abnormal formation and growth of the placenta



Multi-Omic Analyses of Disease



Primary Study Cohort

- Cross-sectional study organized by Dr. Kim Boggess (UNC Maternal and Child Health) and Dr. Rebecca C. Fry (UNC Institute for Environmental Health Solutions)
- Cohort of patients with normotensive (N=17) and preeclamptic (N=18) pregnancies



Characteristics	All N (%) / Mean (range)	Controls N (%) / Mean (range)	Cases N (%) / Mean (range)
Subjects	35 (100%)	17 (100%)	18 (100%)
Race			
African American	13 (37%)	5 (29%)	8 (44%)
Asian	2 (6%)	2 (12%)	0 (0%)
Caucasian	10 (29%)	5 (29%)	5 (28%)
Hispanic	7 (20%)	5 (29%)	2 (11%)
Other	3 (9%)	0 (0%)	3 (17%)
Parity			
Primipara	20 (57%)	9 (53%)	11 (61%)
Multipara	15 (43%)	8 (47%)	7 (39%)
Smoking Status			
Smoker	3 (9%)	1 (6%)	2 (11%)
Non-Smoker	32 (91%)	16 (94%)	16 (89%)
Maternal Age (years)	28.2 (19-38)	27.5 (19-38)	28.6 (19-37)
Gestational age (weeks)	35.3 (22-41)	38.9 (36-41)	31.9 (22-38)

Non-Targeted Analysis Method

- Mobile Phases: 0.1% Formic Acid in DI water / 0.1% Formic Acid in acetonitrile (w/ 0.5 mM ammonium fluoride in negative mode)
- Zorbax Eclipse Plus C8 1.8 μm 2.1 \times 100 mm
- Column temperature: 40°C
- 2.0 μL sample injection
- Samples acquired in triplicate for MS1 data
- Data-dependent acquisition run for MS2 data
- Run in positive and negative mode



Agilent 6546 Q-TOF

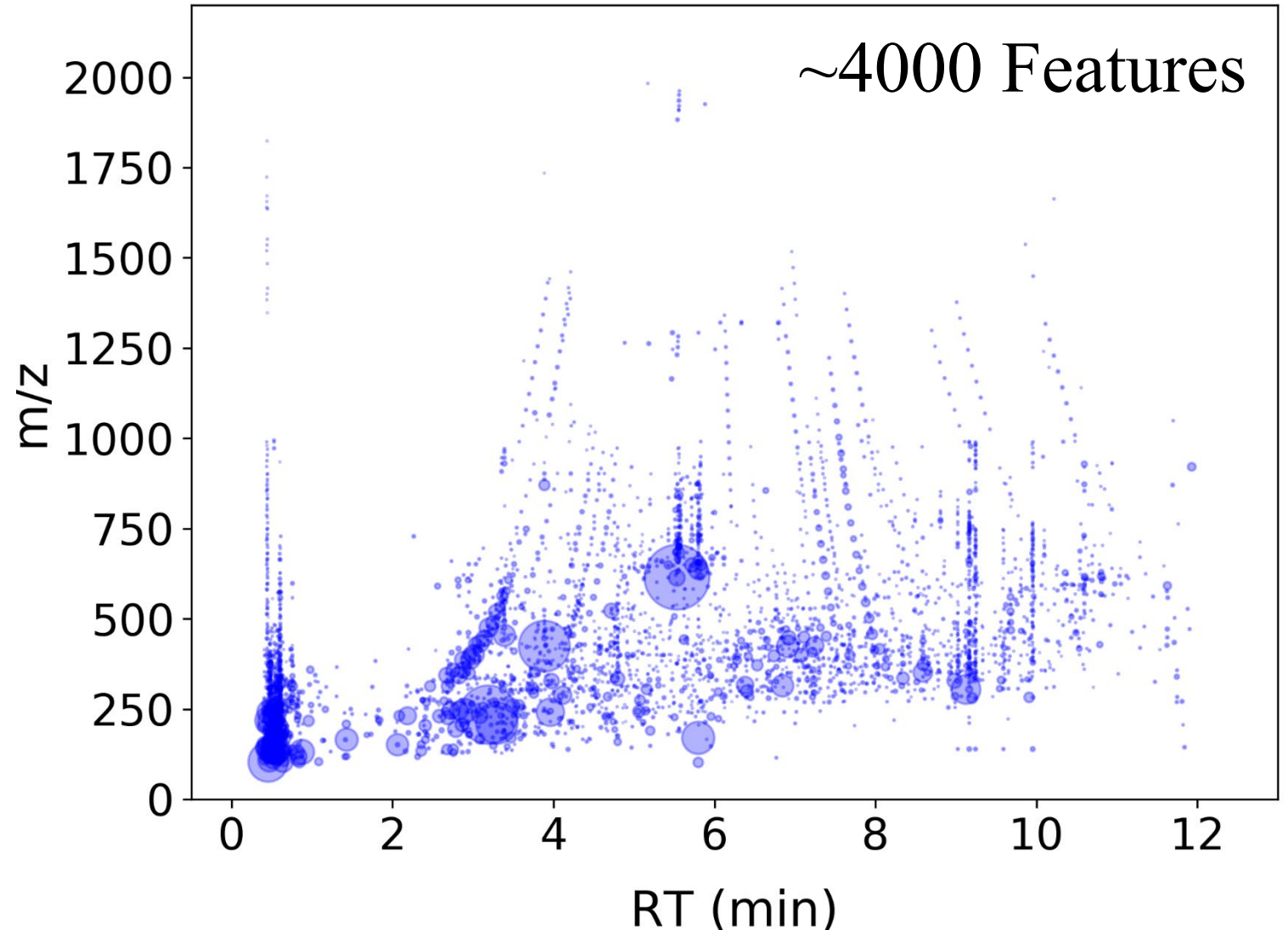
Analytical Filtering of Data

Which features observed are real?

Agilent Profinder / Mass Profiler Professional:
Feature generation and retention time alignment

EPA NTA WebApp: *Feature filtering*

- Adducts
- Duplicate features
- Reproducibility
- Blank subtraction
- Sample/blank thresholds

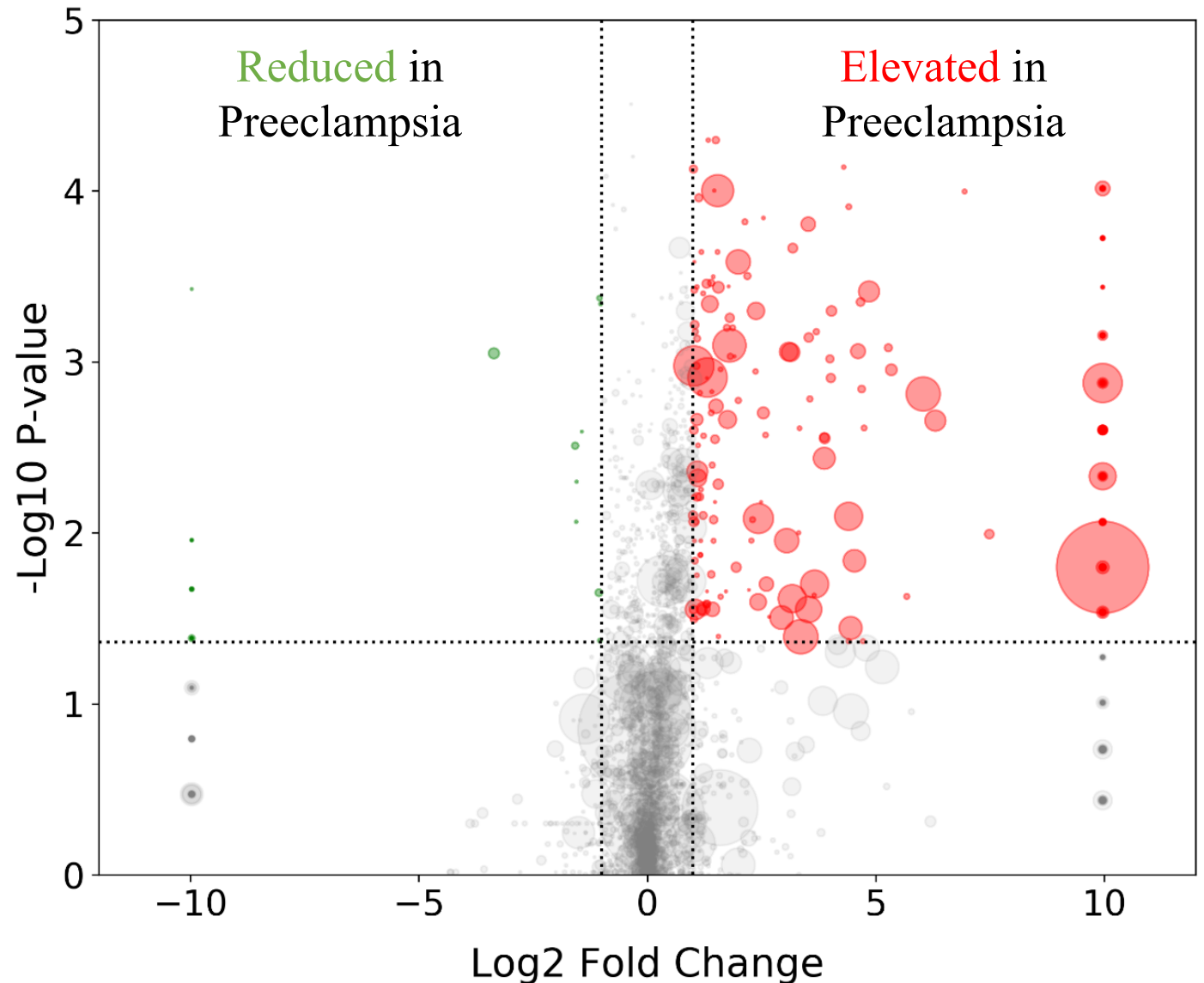


Statistical Filtering of Data

Which features change in correlation with Preeclampsia observations?

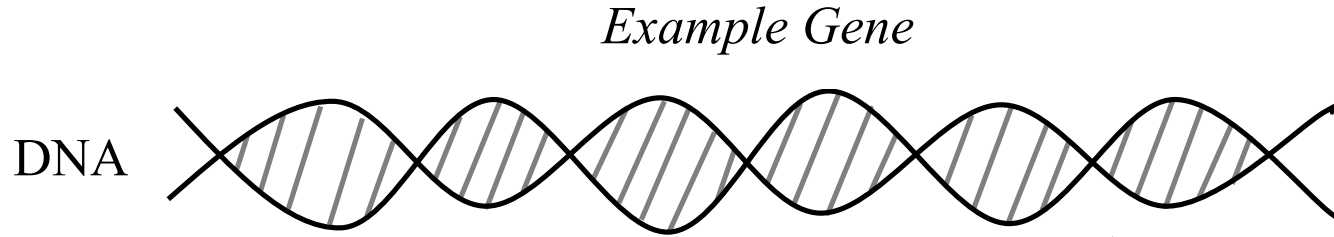
- Normotensive (N=17) and preeclamptic (N=18) pregnancies
- Significance threshold (\log_{10} -space) = 1.36
- Fold change threshold (\log_2 -space) = 1

183 Molecular
Features



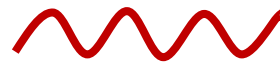
The “Central Dogma” of Biology

**Genomics /
Epigenomics**



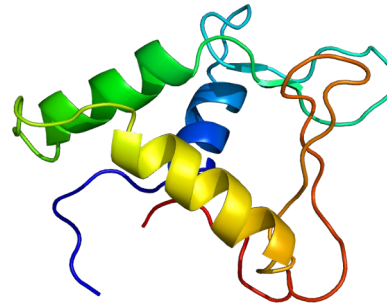
Transcriptomics

mRNA



Proteomics

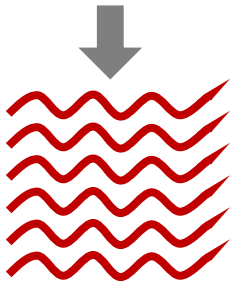
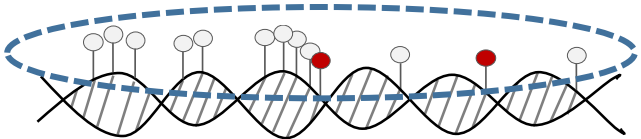
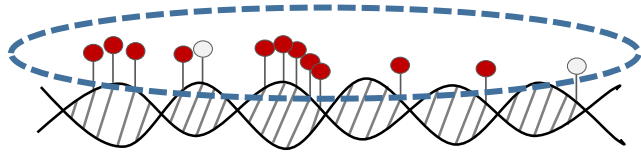
Protein



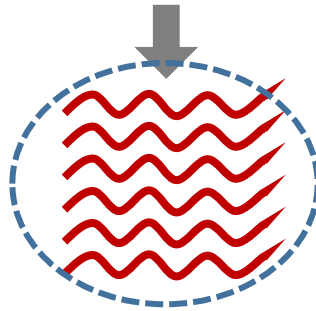
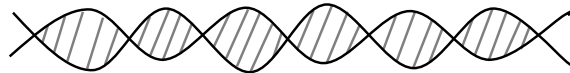
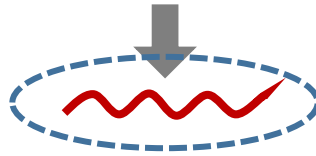
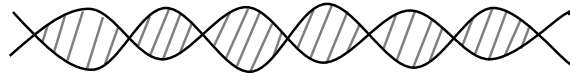
*Regulation occurs at
all levels*

Epigenomic / Transcriptomic Regulation

1. DNA Methylation Levels

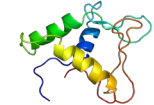
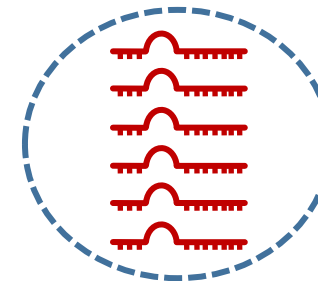
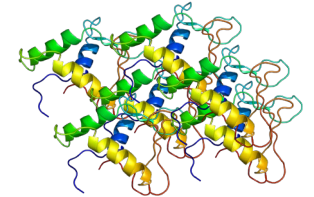
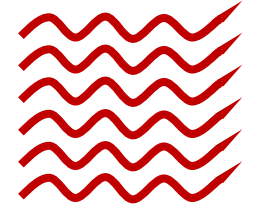
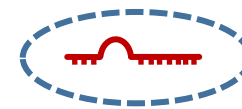


2. Gene Expression Levels



3. MicroRNA Levels

miRNA



Epigenomic / Transcriptomic Associations

 PLOS ONE

RESEARCH ARTICLE

Epigenetics and Preeclampsia: Defining Functional Epimutations in the Preeclamptic Placenta Related to the TGF- β Pathway

Elizabeth Martin¹, Paul D. Bay^{1,2}, Lisa Smeester¹, Matthew R. Grace³, Kim Boggess³,

h, 135
America,
Carolina,
School of

Which molecular features are highly correlated to preeclamptic epigenomic and transcriptomic signatures?



ELSEVIER

journal homepage: www.elsevier.com/locate/foodchemtox

miRNAs as common regulators of the transforming growth factor (TGF)- β pathway in the preeclamptic placenta and cadmium-treated trophoblasts: Links between the environment, the epigenome and preeclampsia

Samira A. Brooks^a, Elizabeth Martin^a, Lisa Smeester^a, Matthew R. Grace^b, Kim Boggess^b, Rebecca C. Fry^{a, c, *}

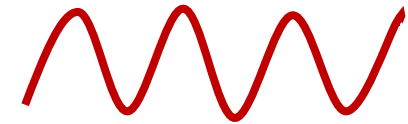
^a Department of Environmental Sciences and Engineering, Gillings School of Global Public Health, 135 Dauer Drive, CB 7431, University of North Carolina, Chapel Hill, NC, USA

^b Department of Obstetrics & Gynecology, University of North Carolina School of Medicine, University of North Carolina, Chapel Hill, NC, USA

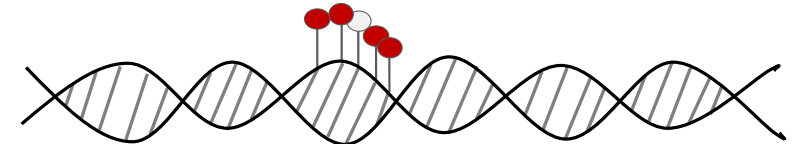
^c Curriculum in Toxicology, School of Medicine, University of North Carolina, Chapel Hill, NC, USA



730 differentially expressed genes



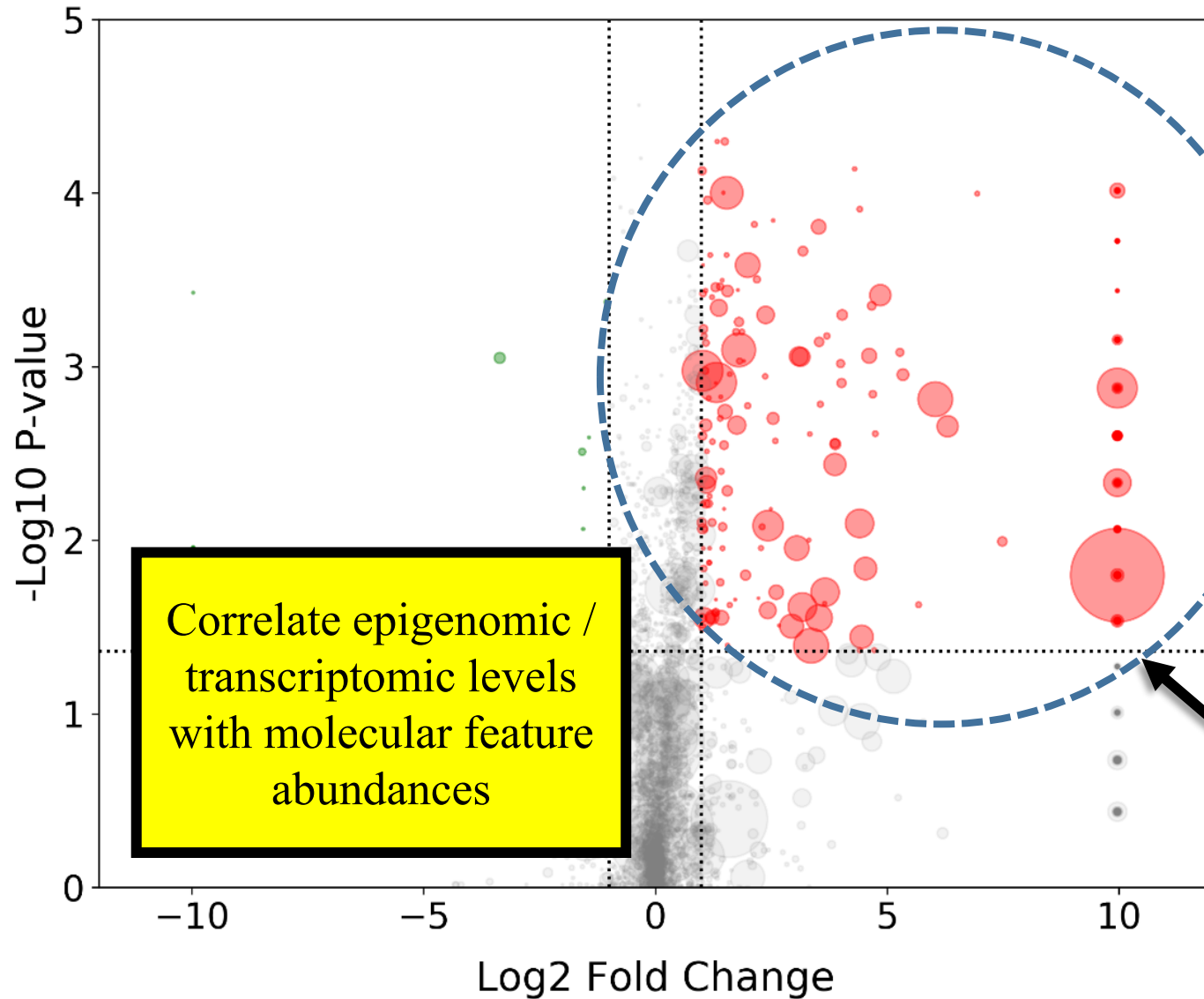
617 differentially methylated genes



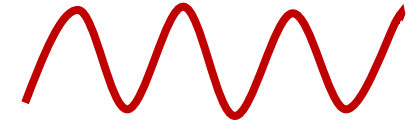
631 differentially expressed miRNAs



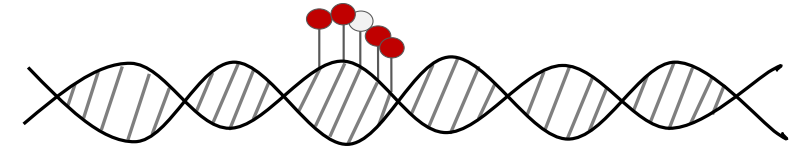
Epigenomic / Transcriptomic Associations



730 differentially expressed genes



617 differentially methylated genes



631 differentially expressed miRNAs



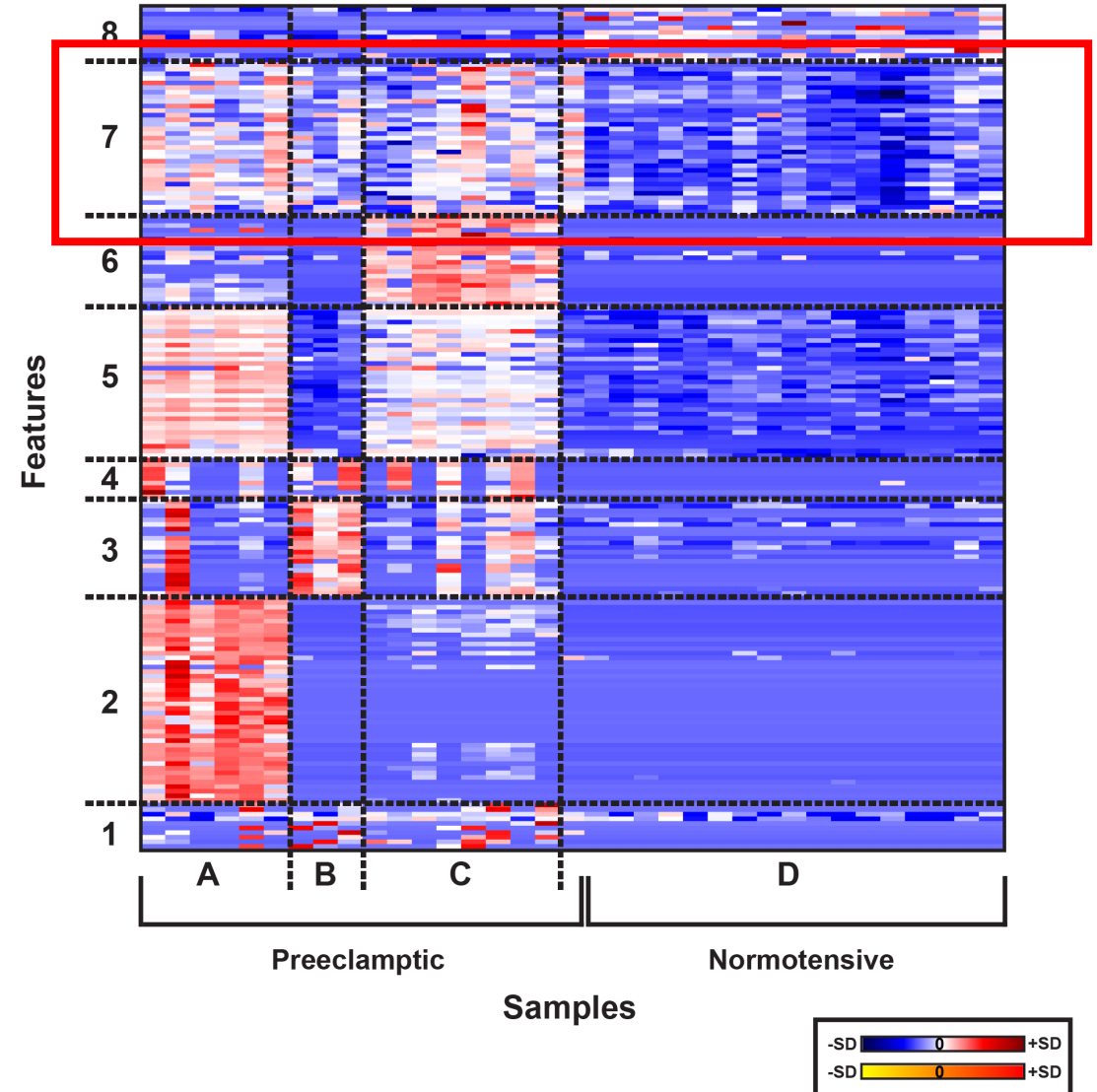
Feature and Sample Cluster Results

How can the remaining features and samples be further grouped/prioritized?

***Unsupervised
Machine Learning
Clustering***

***Preeclampsia
“Omics” data***

- Samples cluster into three groups of preeclamptic placenta samples, and one containing control samples (plus one preeclamptic)
- Features cluster into eight groups
 - Cluster 7 contains highest association with preeclamptic Omics



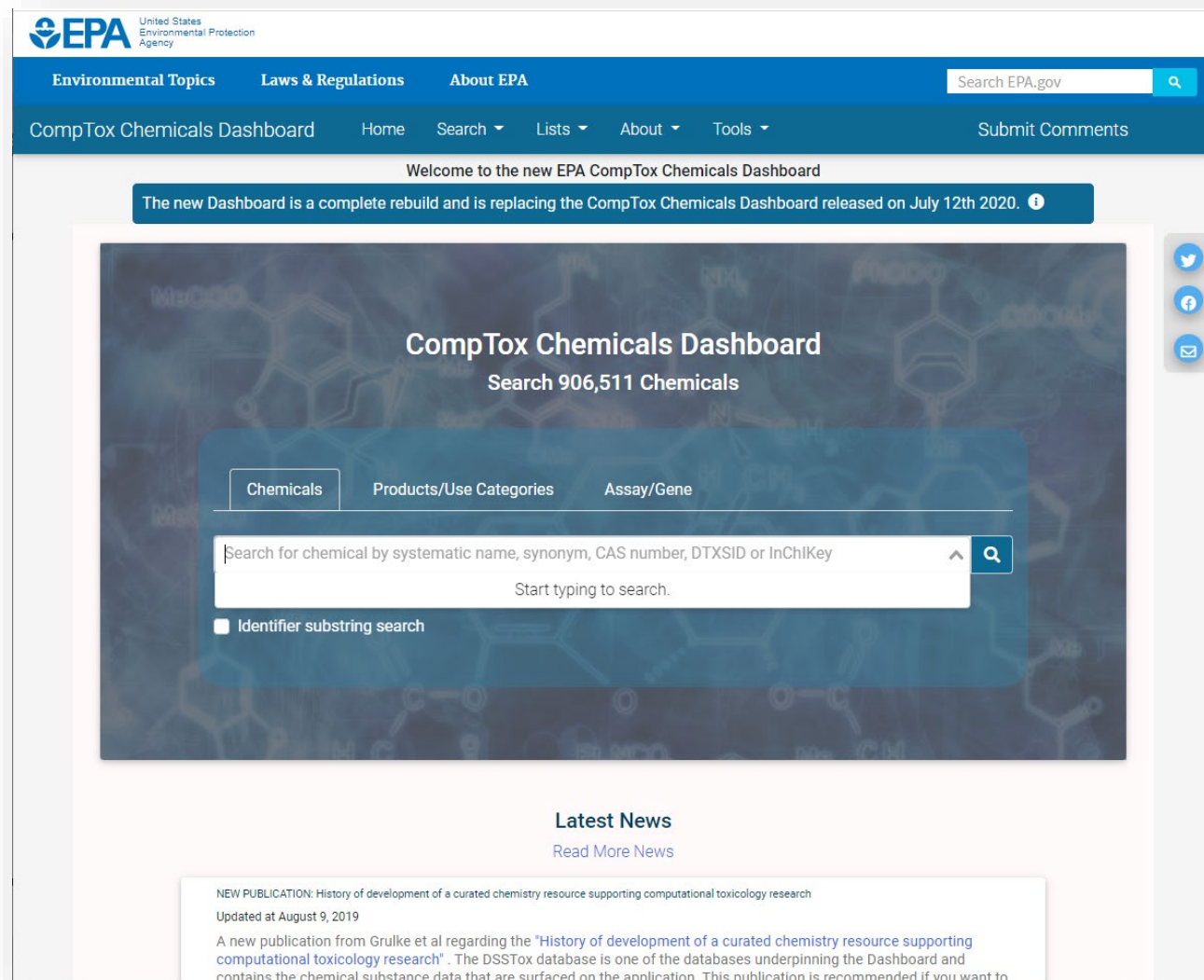
Chemical Candidate Prioritization

For each feature, what are the potential chemicals (candidates)?

- Search by accurate mass within EPA's **DSSTox** database
- 183 features → ~6000 candidates

Can we prioritize candidates to investigate?

- MS2 spectral matching
- Exposure database metadata
- Multi-Omics scoring
- **46 chemicals** prioritized and acquired



The screenshot shows the EPA CompTox Chemicals Dashboard. At the top is the EPA logo and navigation links: Environmental Topics, Laws & Regulations, and About EPA. Below this is a search bar labeled "Search EPA.gov" and a "Submit Comments" link. The main header reads "Welcome to the new EPA CompTox Chemicals Dashboard" and "The new Dashboard is a complete rebuild and is replacing the CompTox Chemicals Dashboard released on July 12th 2020." The central section is titled "CompTox Chemicals Dashboard" and "Search 906,511 Chemicals". It features three tabs: "Chemicals", "Products/Use Categories", and "Assay/Gene". Below the tabs is a search input field with the placeholder text "Search for chemical by systematic name, synonym, CAS number, DTXSID or InChIKey" and a search button. A checkbox for "Identifier substring search" is also present. At the bottom, there is a "Latest News" section with a link to "Read More News" and a news item titled "NEW PUBLICATION: History of development of a curated chemistry resource supporting computational toxicology research" dated August 9, 2019.

<https://comptox.epa.gov/dashboard>

Targeted Confirmation Analysis

- LC conditions matched NTA LC method
- MS1 and MS2 data collected
- Subset of placental samples re-run for retention time and accurate mass alignment
- **23 unique chemicals** confirmed at level 1 identification via matching of accurate mass, retention time and MS2 fragment ions
- Additional matching on **2 features** which consisted of isomeric pairs (**4 unique chemicals**) unable to be chromatographically resolved



UNC
GILLINGS SCHOOL OF
GLOBAL PUBLIC HEALTH

*Chemistry and
Analytical Core*



Thermo Q-Exactive Orbitrap

Chemical ID and Categorization Results

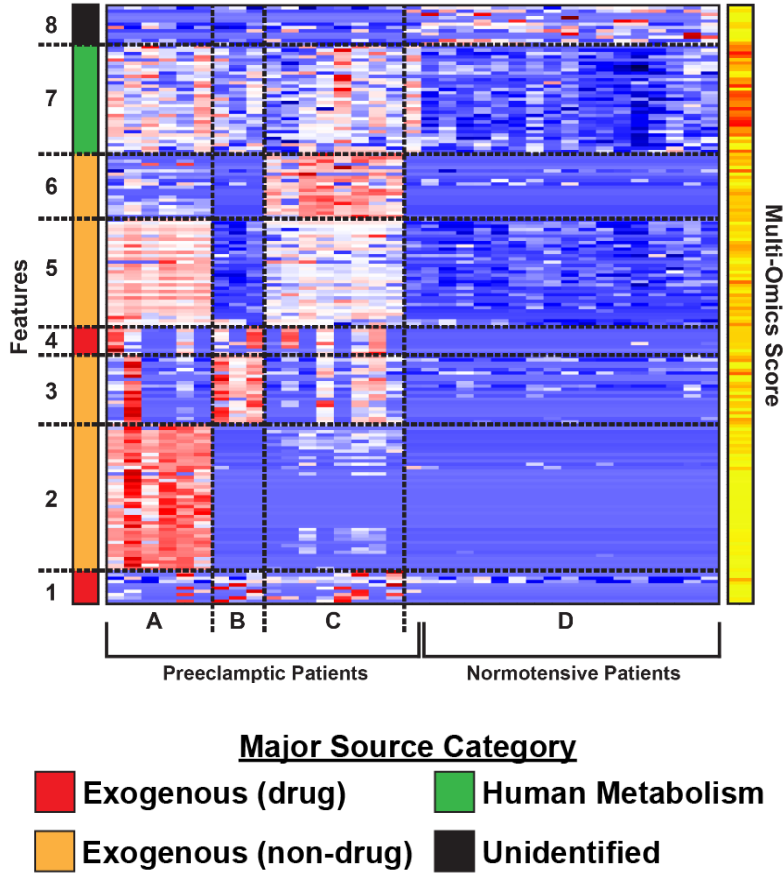
53 total features (corresponding to **40 unique chemicals**) identified at levels 1-3 confidence

- **Endogenous metabolites** previously associated with preeclampsia (nucleotides, acyl carnitines, amino acids)
- **Exogenous drugs** known for administration with pregnancy and/or preeclampsia
 - *Antibiotics*: cefazolin, clindamycin
 - *Nausea*: diphenhydramine, ondansetron
 - *Hypertension*: acetaminophen, labetalol
 - *Steroids*: dexamethasone, betamethasone
- **Exogenous non-drugs** limited in known associations with preeclampsia
 - *Ethanolamides/endocannabinoids*: surfactants in personal care products / neurotransmitters and receptors for biological regulation

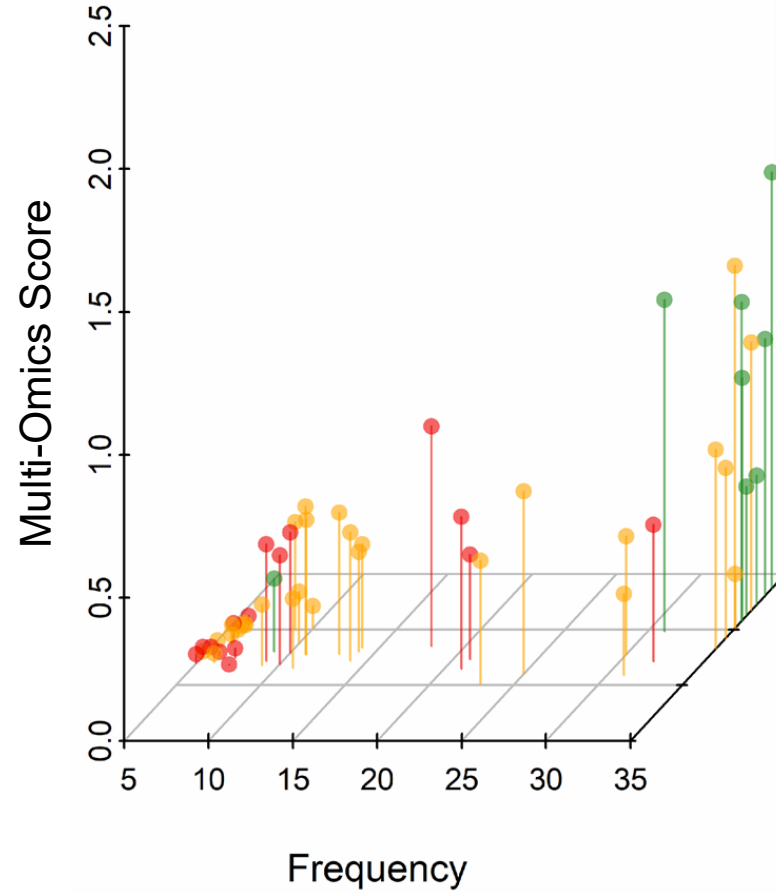
Chemical Name	Cluster	Source Category
Desmethyldiphenhydramine	1	Exogenous (drug)
Diphenhydramine		Exogenous (drug)
Formoterol		Exogenous (drug)
Dexamethasone Betamethasone		Exogenous (drug)
Clindamycin		Exogenous (drug)
Labetalol		Exogenous (drug)
Phentermine		Exogenous (drug)
N-Linoleoylethanolamide	2	Exogenous (non-drug)
3-(1-ethyl-1-methylpropyl)-5-isoxazamine		Exogenous (non-drug)
Methyl 5-(dimethylamino)-2-methyl-5-oxopentanoate		Exogenous (non-drug)
Undecylenoyl monoethanolamide		Exogenous (non-drug)
Methyl hydroxymethyl oleyl oxazoline		Exogenous (non-drug)
Linoleic diethanolamide		Exogenous (non-drug)
Ricinoleoyl diethanolamide		Exogenous (non-drug)
PEG (polyethylene glycol)	3	Exogenous (non-drug)
Ondansetron	4	Exogenous (drug)
Cefazolin		Exogenous (drug)
Dimethyl cyanocarbonodithioimide		Exogenous (non-drug)
Etisomicin		Exogenous (drug)
Hexaminolevulinate	5	Exogenous (drug)
12-Aminododecanoic acid		Exogenous (non-drug)
Icaridin		Exogenous (non-drug)
N,N-Dimethyldodecylamine-N-oxide		Exogenous (non-drug)
Adenosine		Human metabolism
Gabapentin		Exogenous (drug)
Tyrosine		Human metabolism
Stearic acid diethanolamide Isostearic acid diethanolamide	6	Exogenous (non-drug)
N-Dodecanoyl-N-methylglycine		Exogenous (non-drug)
2,4-Bis(1-methyl-1-phenylethyl)phenol		Exogenous (non-drug)
Acetaminophen		Exogenous (drug)
3,3'-[Oxybis(ethane-2,1-diyloxy)]dipropyl-1-amine	7	Exogenous (non-drug)
N-Benzyl-N,N-dimethyldodecan-1-aminium		Exogenous (non-drug)
Lactobionic acid		Exogenous (non-drug)
Acetyl-L-carnitine		Human metabolism
3-Hydroxypentanedioic acid Citramalic acid		Human metabolism
L-Tryptophan		Human metabolism
Valylleucine		Human metabolism
Palmitoylcarnitine		Human metabolism
Acetylcholine		Human metabolism
L-Oleoylcarnitine		Human metabolism

Chemical Feature Trends

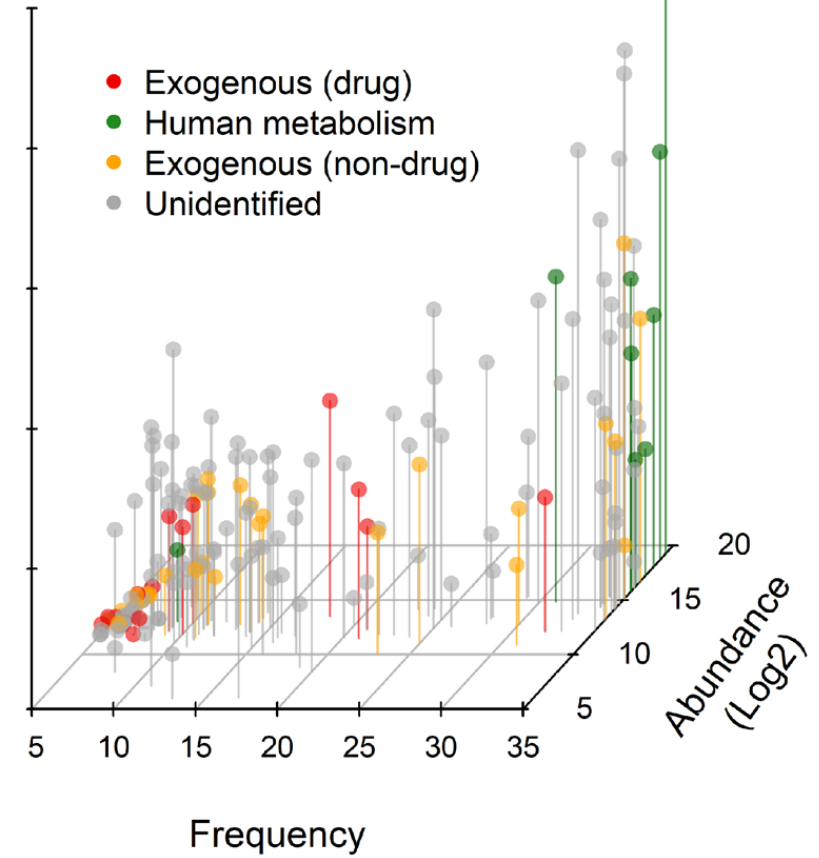
Source Categorization of Clusters



Identified Features (Level 1-3)

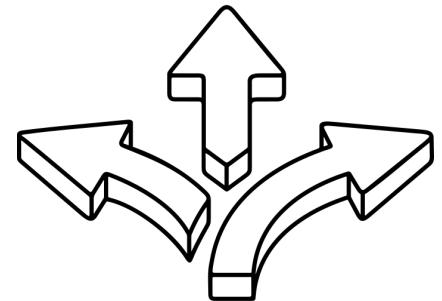
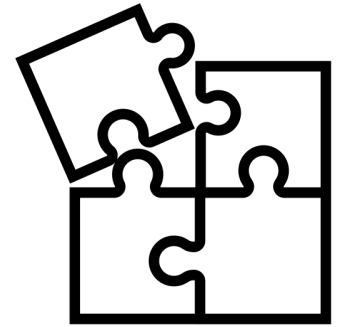


All Features



Summary

- Features corresponding to xenobiotics and endogenous metabolites were found associated with preeclampsia
- Integration of exposomic, epigenomic and transcriptomic analyses supports characterization chemical-biological relationships
- Evaluation of exposomic data trends can inform on exposure source pathways



Acknowledgements



UNC
GILLINGS SCHOOL OF
GLOBAL PUBLIC HEALTH



UNC
SCHOOL OF MEDICINE



Agilent

Jacqueline Bangma

Kim Boggess

Celeste Carberry

Rebecca Fry

Yunjia Lai

Kun Lu

Julia Rager

John Szilagyi

Katherine Coutros

Kathie Dionisio

Louis Groff

Kristin Isaacs

Charles Lowe

Jeff Minucci

Katherine Phillips

Tom Purucker

Jon Sobus

Elin Ulrich

John Wambaugh

Antony Williams

Jarod Grossman