

Mechanistic Indicators of Childhood Asthma (MICA): A Systems Biology Approach for the Integration of Multifactorial Environmental Health Data

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Science Question

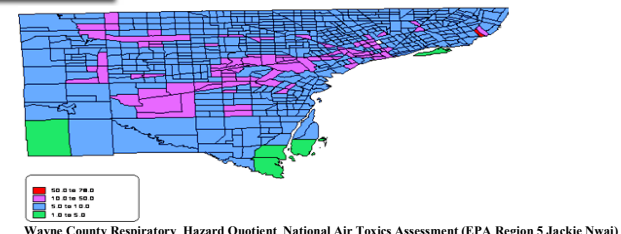
- Can biologically relevant air pollution species and sources (indoor, outdoor, traffic) support the analysis of MICA's genetic, biological and health data?
- Can genomic data viewed together with a spectrum of exposure, effects, clinical and susceptibility markers:
 - Increase the sensitivity needed to define exposure-response-effects relationships?
 - Provide mechanistic explanations as to varied asthmatic responses and susceptibility to environmental exposures?

Research Goals

- Identify biologically relevant air pollution species and sources for MICA through home monitoring, geospatial and exposure-related variables, such as proximity to various stationary and mobile sources
- Relate metrics and surrogates for traffic pollution to clinical markers of asthma allergy and inflammation
- Apply exposure, effect, clinical, and gene biomarkers to evaluate both overarching and specific studies along the source to health outcome paradigm
- Use blood and lung tissues from rodents to inform the blood gene expression analysis from the children's study.
- Identify panels of biomarkers from the array of MICA co variants for future application in large population studies by addressing their reliability, predictive value, sensitivity, specificity, affordability, applicability.
- Apply statistical and methodological approaches to help define asthma phenotypes and identify underlying mechanisms related to perturbation of gene networks related to environmental exposures and/or genetic susceptibility.

Acknowledgments

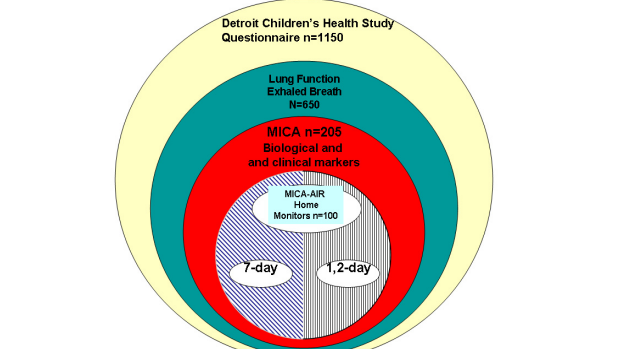
We gratefully acknowledge our many collaborators: US EPA Region 5, NERL (RTP and Cincinnati), NHEERL (EPHD, ECD), Genomic and Chemistry Core, NIEHS, Michigan State University, Rutgers University, Harvard, University of North Carolina, Johns Hopkins, Mercy College, Integrated Laboratory Systems. Expression Analysis, Southwest Research Institute, RTI, Lab Corp, Henry Ford Health Systems and Westat



Why Detroit, Michigan ?

- Other major EPA initiatives DEARS (NHEERL) and DCHS (NHEERL) and NCEA (Near Roadway)
- Wayne county ranks in top 10 % of US counties with respect to criteria pollutants (person days exceeding 8-hr standard for ozone) and several air toxics chemicals

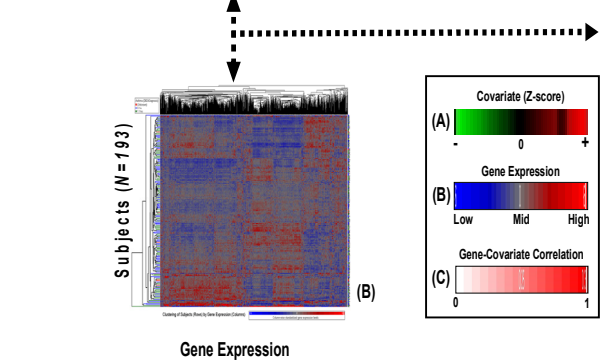
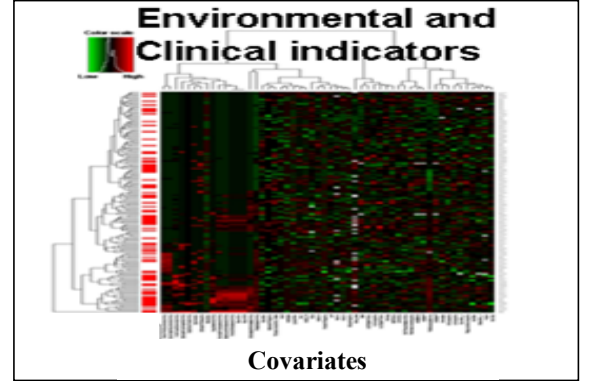
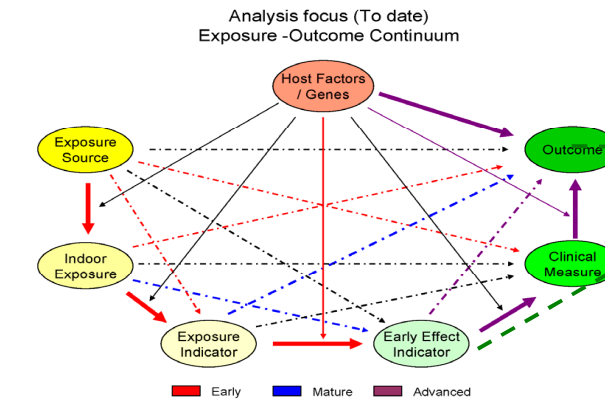
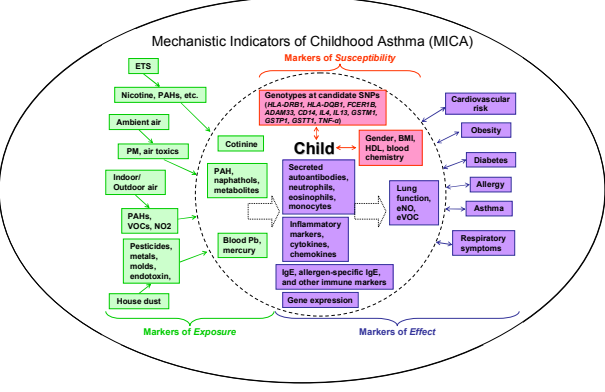
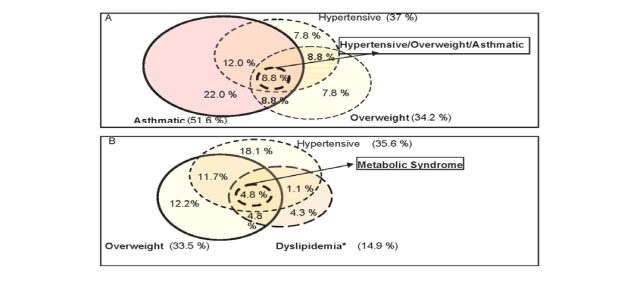
MICA provides biomarker and genomics component For the Detroit Children's Health Study DCHS (L. Neas)



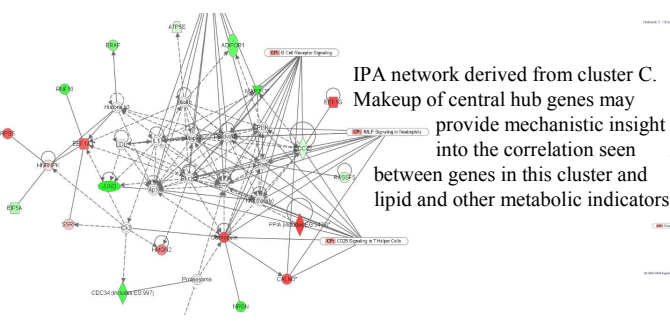
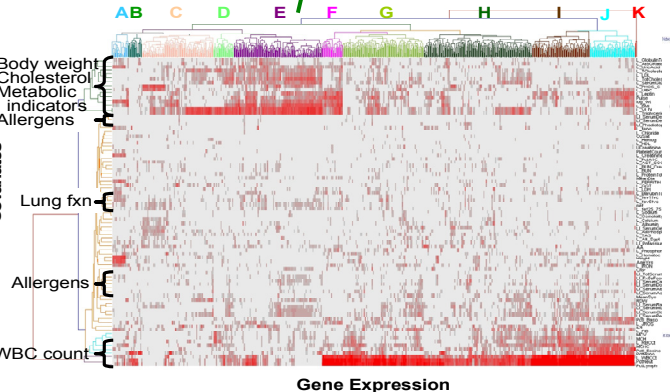
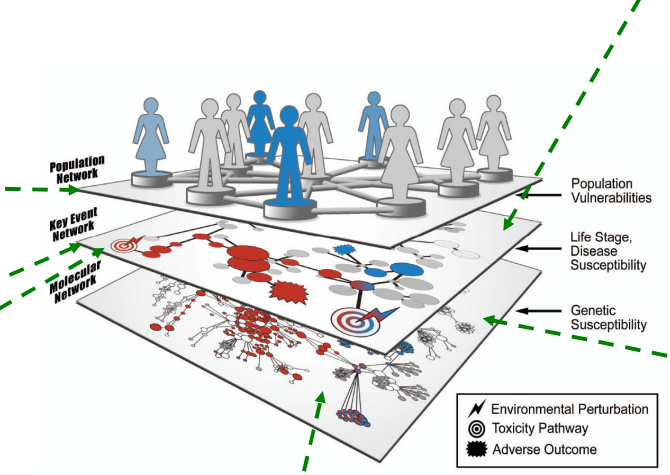
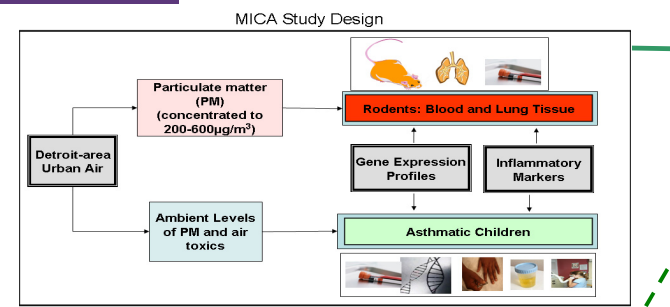
Selected Demographics

	N	Percent	Mean	SE	GeoMean	GeoStd
Male	205	58.0				
White/Only	203	9.9				
OtherRace	203	83.9				
Asthma Diagnosis	195	6.9				
AgeYrs	205		11.6	0.08		
School/Fox High School	203	19.7				
+HighSchool	203	3.0				
HighSchool	203	15.8				
College	203	29.1				
Post College	203	32.5				
Height	205		151.6	0.8		
Weight	205		52.7	1.3	21.8	1.3
BMI	205		22.5	6.2		
BMI<25	205	66.3				
BMI=30 (obese)	205	22.9				
BMI>30	205	10.7				
Cardiovascular	203		110.2	0.8		
MeanArts	203		70.4	0.6		
Hypertensive	47	23.1	122.5	0.5		
Hypertensive Diastolic	52	25.6	61.4	0.7		
Selected lipids	189		166.6	31.5	163.6	1.2
Triglycerides	189		153.9	36.2	76.9	1.9
HDL	189		54.4	12.8	52.8	1.4
LDL	189		95.0	28.4	90.7	1.4
TotCholesterol<200	189	14.8				
Triglycerides<150	189	7.9				
HDL<40	189	12.2				
LDL<130	189	11.1				
Immunological Indicators	189		94.6	5.1		
Total serum IgE	189					
Food Screen allergen (+)	196	35.7				
Allergen screen (+)	196	62.2				
Atopic Asthmatic	98	70.4				

MICA more than an asthma study Health status (proportion) asthmatic, overweight, hypertensive, dyslipidemia




Approach



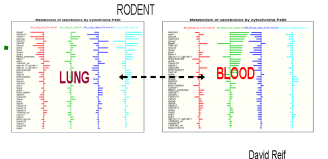
ANIMAL STUDY

✦ Informs Children's Asthma Study

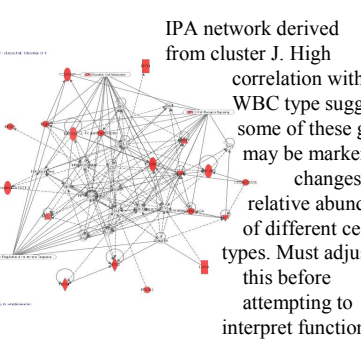
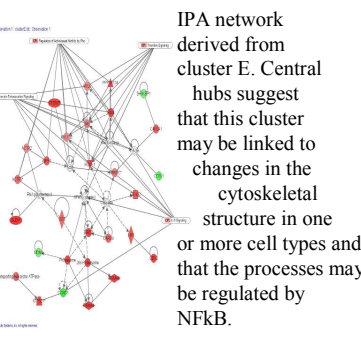


PHASE 1 MICA Rodent Study

- Gene Array data consistent and a more sensitive indicator of exposure/effect compared to more traditional -cell, histology morphometry measures.
- Perturbation of inflammatory and airway remodeling pathways.



David Reif



Results/Conclusions

- Participant-based monitoring is a cost-effective approach to exposure assessment.
- Rodent gene expression data is both consistent and more sensitive than the more traditional morphometric, histological and phenotypic data.
- Clinical measurements and gene expression data provide unique insights into the mechanistic basis for asthma. Combining with exposure measures enhances understanding of the impact of environmental stressors.

Impact and Outcomes

- Environmental/gene markers and MICA's mechanism-based paradigm informs two significant EPA initiatives in the Detroit Area: Detroit Exposure Aerosol Research Study (DEARS) and the Detroit Children's Health Study (DCHS).
- Source to outcome data can inform the National Children's Study and other large epidemiological studies.
- Global gene expression data is unbiased with regard to a priori assumptions on the mechanisms involved. This enhances the interpretation of mechanistic models built from the mechanism (or mode) of action.
- Leveraging both rodent (controlled exposures, blood & lung tissue) and human studies (relevance for risk assessment) help to better understand childhood asthma development.

Future Directions

- Identify impact of exposures on key pathways identified from the gene expression data
- Characterize the genetic variants that increase risk for childhood asthma.
- Estimate combined effects of different environmental exposures and genetic susceptibility in determining the likelihood of developing/exacerbating childhood asthma.
- Evaluate other endpoints of regulatory importance (i.e. hypertension, obesity, metabolic syndrome)

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