Genetic Polymorphisms to Improve Interpretation of Contaminant Exposure and Risk in Inuit – Year #1 Summary

INTRODUCTION

- Inuit communities are exposed to some of the highest levels of mercury and other contaminants worldwide (1)
- The ability of humans to process such contaminants is reliant upon environmentally responsive genes (2)
- Environmentally responsive genes are polymorphic and thus their activity varies tremendously across individuals and ethnicities (2)
- Genetic variability may affect the susceptibility of individuals or populations to the effects of pollutants in the environment (2)
- Little is known about gene-contaminant interactions amongst Inuit communities

OBJECTIVES

•In collaboration with the Inuvialuit Settlement Region (ISR) and using archived samples from the 2007-2008 International Polar Year Inuit Health Survey (IHS), specific aims were to:

- 1. characterize polymorphisms in environmental responsive genes
- 2. Relate polymorphisms with existing data on blood contaminant levels

•long-term objective of our research program is to better understand how Inuit process contaminants so that dietary exposure assessments and linkages to health can be improved

METHODS

•Leverage biospecimens and information from n=288 ISR community members who participated in the 2007-2008 International Polar Year Inuit Health Survey (IHS)

•Select and then genotype single nucleotide polymorphisms (SNPs) in n=360 environmentally responsive genes (e.g., metallothionein, glutathione) at McGill-Genome Quebec Innovation Center using the Sequenom iPlex platform

•Descriptive methods to compare genetic variation of ISR with other ethnicities (via 1000 Genomes Project database)

•ANOVAs and regressions to relate polymorphisms with existing data on blood Hg, Pb, Cd, and Se

RESULTS

1) Demographic Characteristics of ISR Study Population

Demographic	(n) mean (SD) or		% range	
Age (years)	281	44.73 (16.02)	18.00 to 90.00	
Male (%)	281	33.8		
Nutritional				
Body mass index (BMI)	235	30.40 (6.65)	17.3 to 58.3	
Hemoglobin (g/L)	234	13.70 (1.51)	7.6 to 17.0	
Blood Hg (µg/L)#**	249	4.64 (9.31)	0.05 to 55.00	
Blood Cd (µg/L)#**	249	1.31 (1.72)	0.02 to 7.30	
Blood Se (Hg) $(\mu g/g)$ #**	249	304.36 (32.18)	150 to 1300	
Blood Pb (µg/L)#**	249	32.18 (33.36)	4.50 to 210.00	
Estimated average MeHg intake from fish μ g/kg/day	236	4.53 (7.67)	0.01 to 68.23	
Estimated average Se intake from fish μ g/kg/ day	236	10.68 (17.19)	0.02 to 109.82	

GM (SD) reported instead due to distribution of the variable. **Geometric means are reported, but SDs are arithmetic SDs

2) Example of Genetic Variation Across Ethnicities

- 146 SNPs (of 360) successfully studied from 281 participants
- Below is representative data on SNP rs1056836 (CYP1B1)



3) Genetic Polymorphisms and Blood Metals

		Blood Hg		Blood Cd		
Gene symbol	SNP	Main Effect	Genotype * Ln Fish Hg Intake	Genotype	Genotype * Ln packs year	
TXNRD2	rs1139793					
TXNRD2	rs5748469					
SEPHS2	rs1133238					
SELS	rs7178239					
GPX3	rs8177412					
ABCB1	rs1128503					
ABCC1	rs212090					
ABCG2	rs2231142					
ABCB1	rs3842					
NAT2	rs1041983					
CYP2D6	rs1080985					
CYP2D6	rs16947					
AHR	rs2066853					
PRDX2	rs33942654					
CYP24A1	rs6022987					
		Legend:		Positive Beta	Negetive Beta	
		Not Significant	p>0.05			
		Nominal	p<0.05			
		More Significa	p<0.01			
		Bonferonni Co	p<0.0003			
	Gene symbol TXNRD2 TXNRD2 SEPHS2 SELS GPX3 ABCB1 ABCC1 ABCC1 ABCC2 ABCB1 NAT2 CYP2D6 CYP2D6 AHR PRDX2 CYP24A1	Gene symbol SNP TXNRD2 rs1139793 TXNRD2 rs5748469 SEPHS2 rs1133238 SEPHS2 rs1133238 SELS rs7178239 GPX3 rs8177412 ABCB1 rs1128503 ABCC1 rs212090 ABCG2 rs2231142 ABCB1 rs1041983 CYP2D6 rs1080985 CYP2D6 rs16947 AHR rs2066853 PRDX2 rs33942654 CYP24A1 rs6022987	Gene symbolSNPMain EffectTXNRD2rs1139793TXNRD2rs5748469SEPHS2rs113238SEPHS2rs113238SELSrs7178239GPX3rs8177412ABCB1rs1128503ABCC1rs212090ABCG2rs2231142ABCB1rs3842NAT2rs1041983CYP2D6rs1080985CYP2D6rs16947AHRrs2066853PRDX2rs33942654CYP24A1rs6022987Image: Second Stress StressMore SignificantMore SignificantMore SignificantStress Stress StressStress Stress StressStress Stress StressStress Stress Stress Stress StressStress Stress Stress Stress Stress Stress StressStress Stress	Gene symbolSNPMain EffectGenotype * Ln Fish Hg Intake $TXNRD2$ rs1139793	Gene symbolSNPMain EffectGenotype * Ln Fish Hg IntakeGenotype * Cenotype * Ln Fish Hg IntakeGenotype * Genotype * Cenotype * Ln Fish Hg IntakeGenotype * Genotype * Cenotype * Cenotype * Intake $TXNRD2$ rs1139793 $TXNRD2$ rs5748469 $SEPHS2$ rs1133238 $SEPHS2$ rs113238 $GPX3$ rs8177412 $ABCB1$ rs1128503 $ABCC1$ rs212090 $ABCG2$ rs2231142 $ABCG1$ rs3842 $NAT2$ rs1041983 $CYP2D6$ rs1080985 $CYP2D6$ rs1080985 $PRDX2$ rs33942654 $PRDX2$ rs33942654-Positive Beta $Not Significan p>0.05IntakNominalp<0.05$	









DISCUSSION

- Many environmentally responsive genes are polymorphic amongst study participants
- Frequency of several polymorphisms are different than in other ethnic groups
- Carriers of certain genetic variants have different levels of blood metals including mercury, cadmium and lead
- These type of results are similar to what is being found in other populations worldwide
- Results such as these are expected to improve interpretation of dietary exposure and biomarker relationships, exposure-disease relationships, and ultimately improved risk assessments and guidance for Inuit communities
- Additional work is underway to: A) conduct more detailed epidemiological analyses; and B) perform a similar study with Inuit from Nunavik who participated in the 2004 Qanuippitaa Survey

REFERENCES

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- 2.Basu, N., Goodrich, J., Head, J. 2014. Ecogenetics of mercury: From genetic polymorphisms and epigenetics to risk assessment and decision-making. Environmental Toxicology and Chemistry. 33: 1248-1258.

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