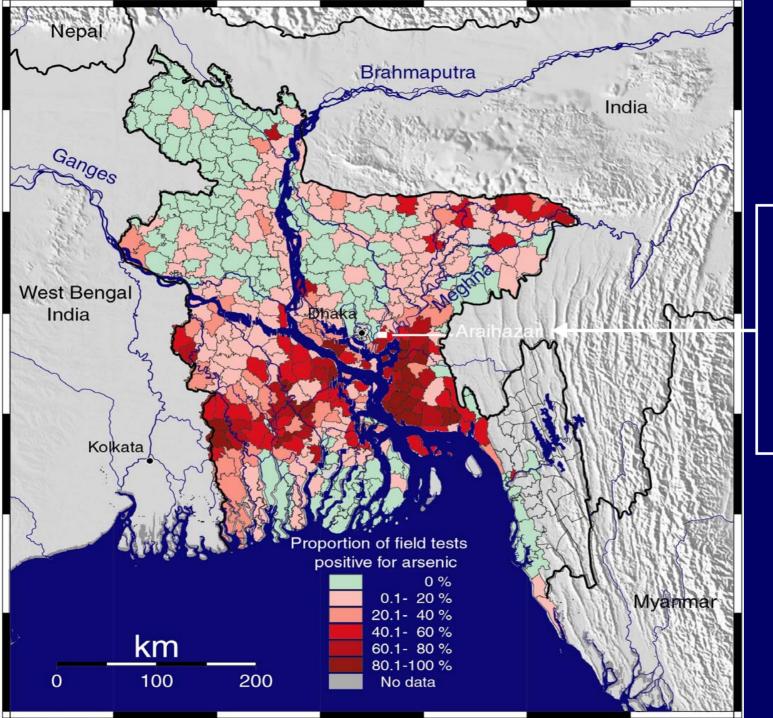


Molecular Epidemiology and Prevention of Health Effects of Arsenic: A Multidisciplinary Approach

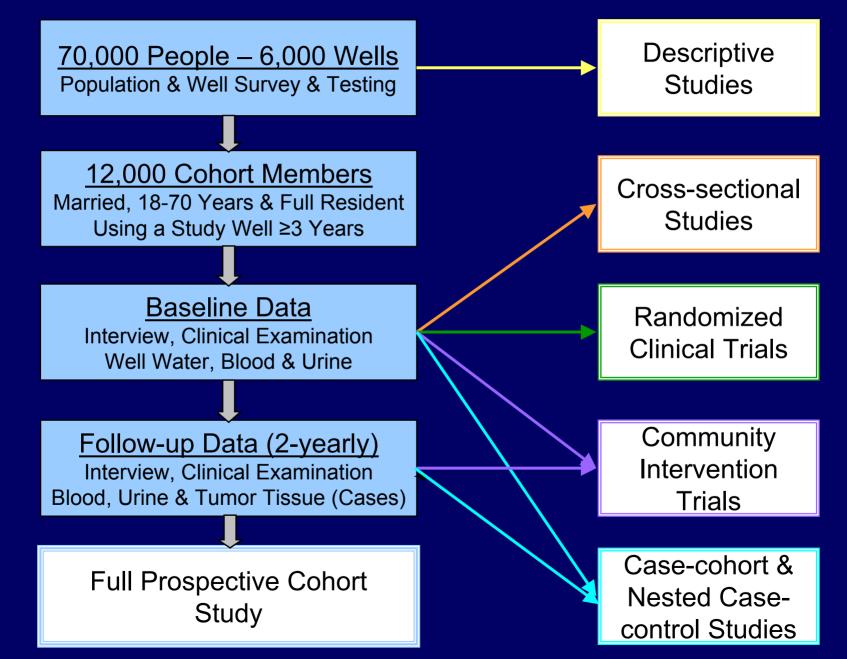
Habibul Ahsan, MD

SBRP Annual Meeting, New York January 13, 2006



Columbia University Superfund Program Study Area in Araihazar, Bangladesh

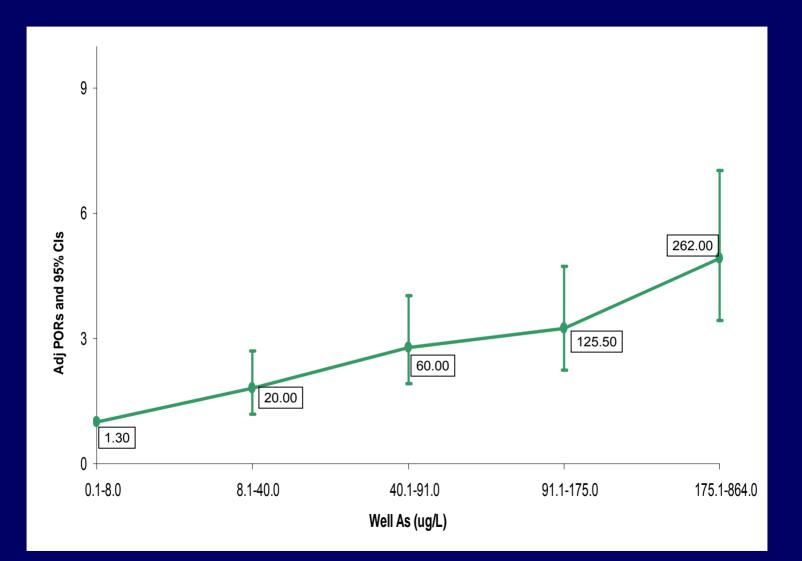
Health Effects of Arsenic Longitudinal Study [HEALS]



Measures of Arsenic Exposure in HEALS

- Well water total As full cohort
- Time-weighted & Cumulative As indices full cohort
- Urinary total As (every 2-year) full cohort
- Blood total As sub-cohort
- Urinary As species sub-cohort
- Toe-nail total As & Blood As species sub-cohort (future)

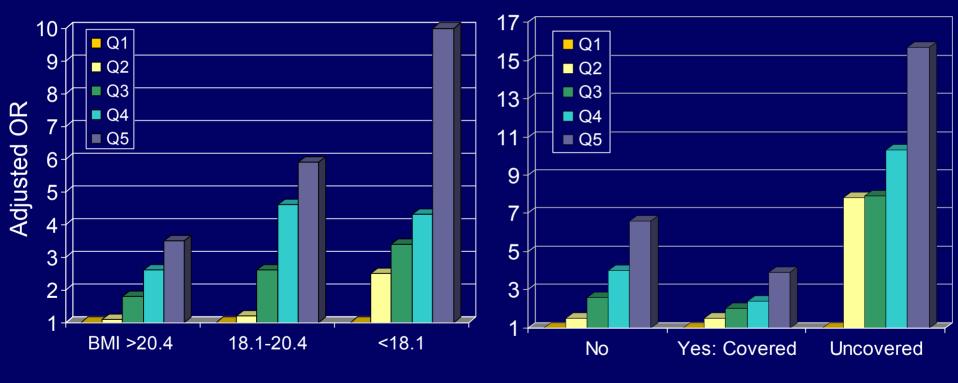
Risk of Skin Lesions by Well Water Arsenic



Modifiers of As Effect on Skin Lesions

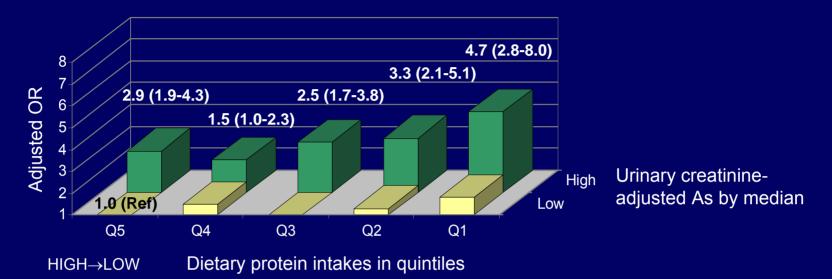
Influence of BMI

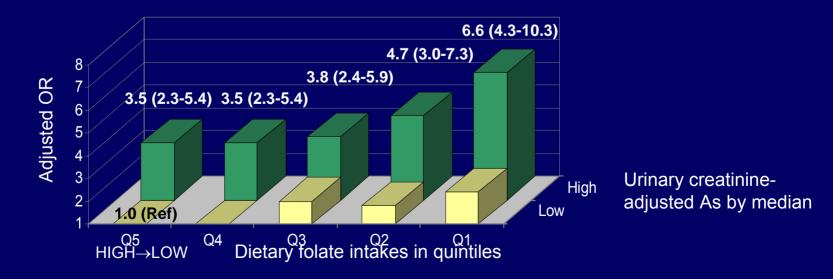
Influence of Sun Exposure

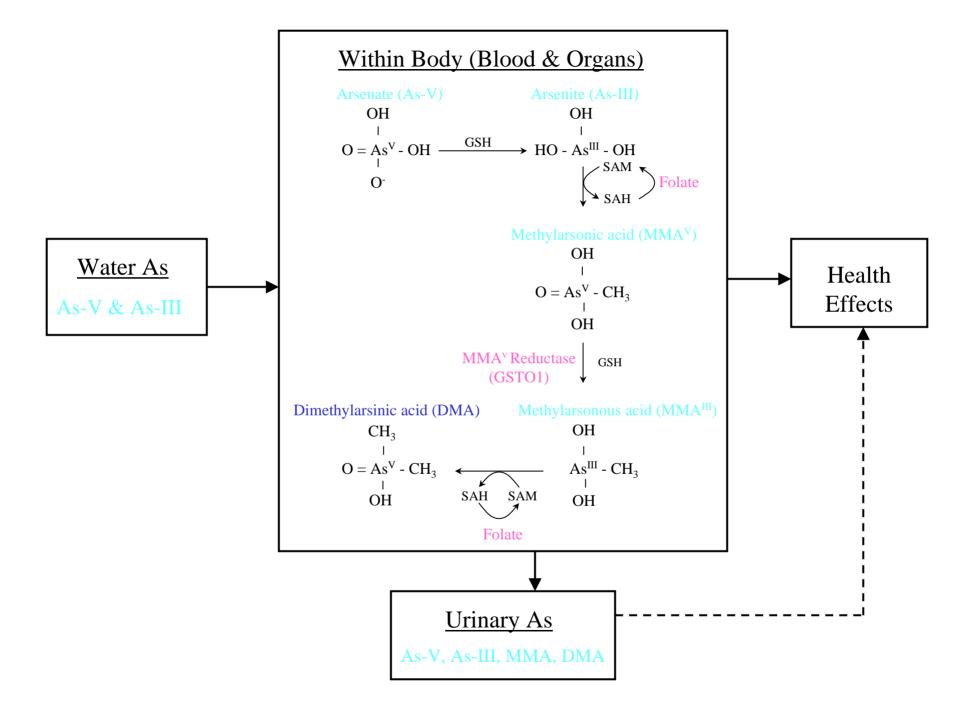


Work outdoor

Influence of Dietary Protein & Folate Intakes on Risk of Arsenic-induced Skin Lesion







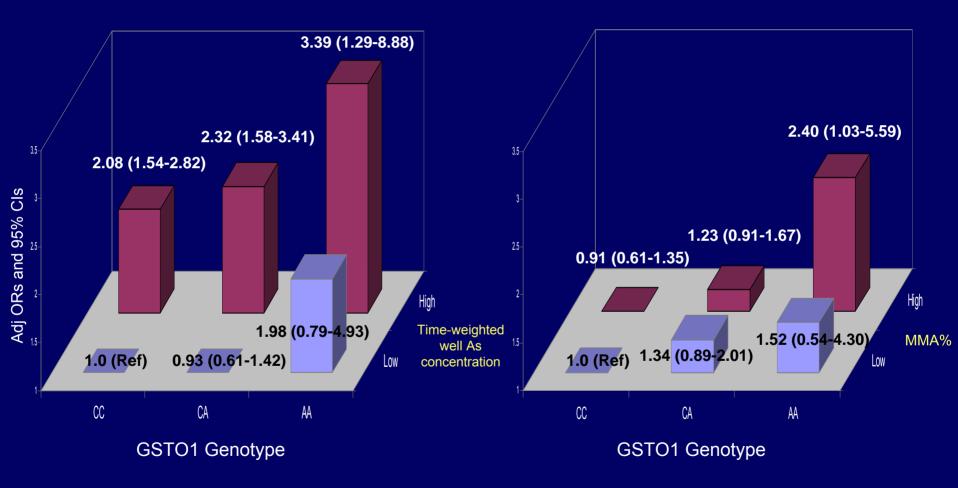
Urinary As Species & Risk of Skin Lesions

	N		_ Adjusted ORs for Skin	
Urinary As Species (%)	Cases	Controls	Lesions (95% CI)	
InAs%				
0-12.2	190	344	1.0	
12.3-17.1	202	335	1.15 (0.87-1.52)	
17.2-69.3	197	355	1.20 (0.91-1.60)	
MMA%				
0-10.8	130	406	1.0	
10.9-14.9	183	353	1.18 (0.87-1.58)	
15.0-33.7	276	275	1.84 (1.37-2.47)	
DMA%				
27.9-67.0	232	305	1.0	
67.1-74.2	188	346	0.75 (0.58-1.00)	
74.3-1.0	169	383	0.73 (0.55-0.96)	

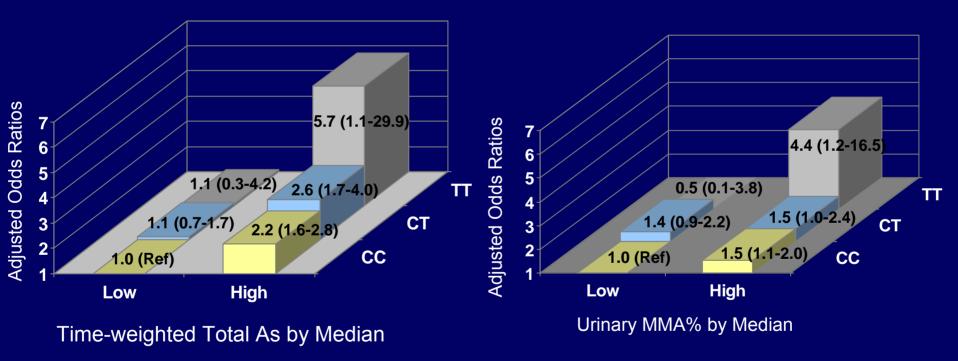
Predictors of Urinary As Species

	DMA%		MMA%		
	Parameter estimates (SE)	P value	Parameter estimates (SE)	P value	
BMI	0.130 (0.077)	0.09	-0.067 (0.044)	0.13	
Age (years)	0.088 (0.031)	<0.01	0.002 (0.017)	0.17	
Gender					
Female	Reference group		Reference group		
Male	-4.554 (0.768)	<0.01	3.625 (0.440)	<0.01	
Well water As (per 100 µg/L)	-0.791 (0.298)	<0.01	0.116 (0.170)	0.49	
Total urinary As (per 100 μg/L)	-0.435 (0.241)	0.07	0.283 (0.138)	0.04	
Urinary creatinine (g)	0.053 (0.006)	<0.01	-0.015 (0.004)	<0.01	

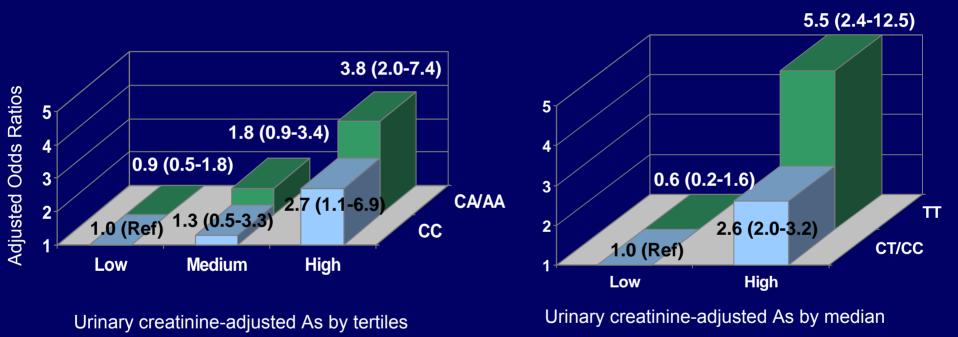
Influence of *GSTO1* Genotype on Risk of Skin Lesion in Relation to Time-weighted As and Urinary MMA%



Influence of MTHFR C677T Genotype on Risk of Skin Lesion in Relation to Time-weighted As and Urinary MMA%



Influence of XPD Codon 751 A-G & hGPX1 Codon 223 C-T Genotypes on Risk of Arsenic-induced Skin Lesion



Association Between Arsenic Levels in Water and Urine and Glycosylated Hemoglobin in Blood (HgbA1c%)

Water As & Blood HgbA1c%

	Parameter Estimates	t Value	P-value
Intercept	4.101	83.180	<.0001
9∼57 µg/l	0.988	-1.700	0.089
58∼138 µg/l	0.994	-0.900	0.366
139~240 µg/l	1.003	0.380	0.703
241~864 µg/l	0.994	-0.930	0.355
BMI	1.005	6.420	<.0001
Education in years	1.000	0.480	0.634
Age in years	1.002	9.200	<.0001
Male	1.018	3.490	0.001
Have TV	1.007	1.480	0.139

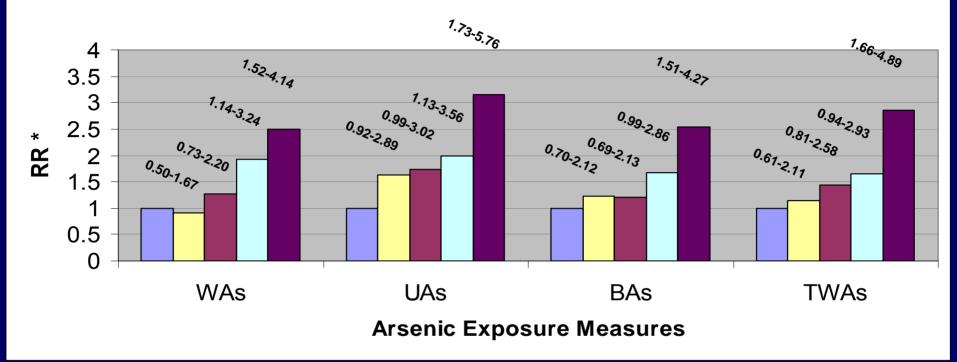
Reference group: 5-8 µg/l. Categories were based on guintiles among the 2102 subjects

Urinary As & Blood HgbA1c%

	Parameter Estimates	t Value	P-value
Intercept	4.082	79.150	<.0001
110~190 µg/g	1.003	0.480	0.631
191~301 µg/g	0.999	-0.170	0.861
302~500 µg/g	0.993	-1.010	0.310
500+ µg/g	1.002	0.290	0.769
BMI	1.005	6.410	<.0001
Education in years	1.000	0.490	0.625
Age in years	1.002	9.210	<.0001
Male	1.017	3.390	0.001
Have TV Reference group: <110	1.007 µg-g creatinine.	1.430 Categories	0.153 were based o

quintiles among the 2102 subjects

Prospective Association between As exposure Measures & Skin Lesion Risk



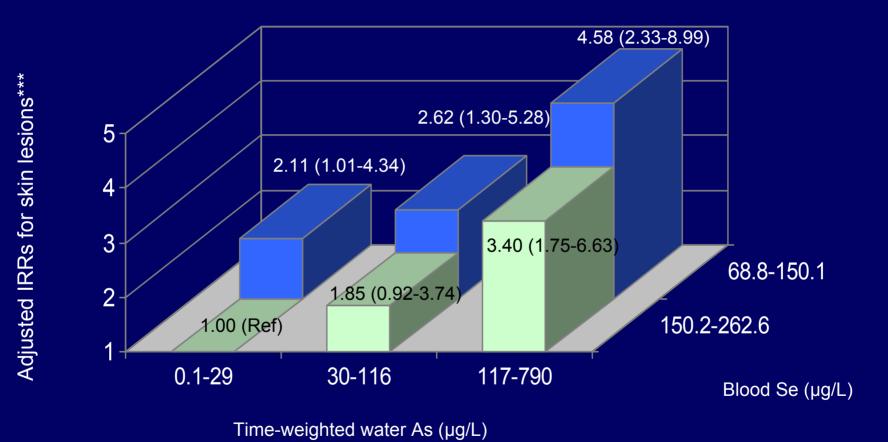
RRs adjusted for age, BMI, gender & smoking status RRs for UAs adjusted for urinary creatinine WAs = water arsenic

UAs = total urinary arsenic

BAs = blood arsenic

TWAs = time-weighted arsenic

Joint Effect of As Exposure and Low Blood Se on Risk of Skin Lesion



***Categories of blood Se and time-weighted water As were determined based on tertile and median values, respectively, in the subcohort. IRRs were adjusted for age, BMI, gender, and smoking status

Differentially Expressed Genes in Response to Arsenic Exposure

Gene symbol	Gene description	GM of:	Skin Lesions	Non-lesions	Ratio of \mathbf{GM}^*
CXCL2	chemokine (C-X-C motif) ligand 2		42.9	1146.1	0.037
CCL20	chemokine (C-C motif) ligand 20		30.6	522.5	0.059
IL1B	interleukin 1, beta		378.8	4894.3	0.077
ZA20D3	Zinc finger, A20 domain containing 3		46.0	378.5	0.122
SOD2	superoxide dismutase 2, mitochondrial		26.7	239.1	0.112
CD44	CD44 antigen (homing function and Indian blood group system)		60.3	539.3	0.112
CCL3	chemokine (C-C motif) ligand 3		411.0	4513.1	0.091
PTX3	pentraxin-related gene, rapidly induced by IL-1 beta		34.4	416.3	0.083
TNF	tumor necrosis factor (TNF superfamily, member 2)		152.5	1971.2	0.077
KCNJ2	potassium inwardly-rectifying channel, subfamily J, member 2		89.6	889.4	0.101
PAPD4	PAP associated domain containing 4		96.9	612.0	0.158
CCRL2	chemokine (C-C motif) receptor-like 2		84.1	655.2	0.128
IER3	immediate early response 3		401.8	2875.0	0.140
SRP68	signal recognition particle 68kDa		70.2	596.7	0.118
PFKFB3	6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 3		100.8	686.9	0.147
PFAAP5	Phosphonoformate immuno-associated protein 5		79.1	466.0	0.170
SOX18	SRY (sex determining region Y)-box 18		73.2	480.2	0.152
HSPA1B	heat shock 70kDa protein 1B		44.4	515.1	0.086
TNFAIP6	tumor necrosis factor, alpha-induced protein 6		55.4	481.6	0.115
SFPQ	Splicing factor proline/glutamine rich		73.1	387.4	0.189
AKAP9	A kinase (PRKA) anchor protein (yotiao) 9		16.1	81.5	0.198
ZNF267	zinc finger protein 267		63.5	351.2	0.181
NR4A2	nuclear receptor subfamily 4, group A, member 2		132.9	783.0	0.170
RFX3	Regulatory factor X, 3 (influences HLA class II expression)		30.3	148.4	0.204

Differentially Expressed Genes in Response to Selenium Treatment

Gene Title	Gene Symbol
immediate early response 3	IER3
interleukin 8	IL8
chemokine (C-C motif) ligand 3 /// chemokine (C-C motif) ligand 3-like 1 /// chemokine (C-C motif) ligand 3-like, centromeric	CCL3 /// CCL3L1 /// MGC12815
zinc finger protein 91 (HPF7, HTF10)	ZNF91
Kruppel-like factor 12	KLF12
tumor necrosis factor (TNF superfamily, member 2)	TNF
chemokine (C-X-C motif) ligand 2	CXCL2
interleukin 8	IL8
putative lymphocyte G0/G1 switch gene	G0S2
Transformer-2 alpha	TRA2A
inhibitor of DNA binding 2, dominant negative helix-loop-helix protein /// inhibitor of DNA binding 2B, dominant negative helix-loop-helix protein	ID2 /// ID2B
superoxide dismutase 2, mitochondrial	SOD2
MRNA; cDNA DKFZp434A202 (from clone DKFZp434A202)	
Development and differentiation enhancing factor 1	DDEF1
cold autoinflammatory syndrome 1	CIAS1
serum/glucocorticoid regulated kinase-like	SGKL
Phosphodiesterase 4B, cAMP-specific (phosphodiesterase E4 dunce homolog, Drosophila)	PDE4B
zinc finger and BTB domain containing 20	ZBTB20
Hypothetical protein FLJ43663	FLJ43663
interleukin 1, beta	IL1B

Changes in Urinary Creatinine-adjusted As (Follow up - Baseline) by Well Switching Status and Baseline Well As



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