

## Supplementary Information

**Table S1.** Confirmed and predicted Imprinted Genes.

Gene ID	Full Gene Name	Location	Status	Expressed Allele
<i>ABCC9</i>	ATP-binding cassette, sub-family C, member 9	12p12.1 AS	Predicted	Maternal
<i>ABCG8</i>	ATP-binding cassette, sub-family G, member 8	2p21	Predicted	Maternal
<i>ACD</i>	Adrenocortical dysplasia homolog	16q22.1 AS	Predicted	Maternal
<i>ADAMTS16</i>	ADAM metalloproteinase with thrombospondin type 1 motif, 16	5p15	Predicted	Maternal
<i>AIM1</i>	Absent in melanoma 1	6q21	Imprinted	Paternal
<i>ALDH1L1</i>	Aldehyde dehydrogenase 1 family, member L1	3q21.3 AS	Predicted	Maternal
<i>ANO1</i>	Anoctamin 1, calcium activated chloride channel	11q13.3	Imprinted	Maternal
<i>APBA1</i>	Amyloid beta precursor protein-binding, family A, member 1	9q13-q21.1 AS	Predicted	Paternal
<i>ATP10A</i>	ATPase, class V, type 10A	15q11.2 AS	Imprinted	Maternal
<i>B4GALNT4</i>	Beta-1,4-N-acetyl-galactosaminyl transferase 4	11p15.5	Predicted	Maternal
<i>BLCAP</i>	Bladder cancer associated protein	20q11.2-q12 AS	Imprinted	Isoform Dependent
<i>BMP8B</i>	Bone morphogenetic protein 8b	1p35-p32 AS	Predicted	Paternal
<i>BRP44L</i>	Brain protein 44-like	6q27 AS	Predicted	Paternal
<i>BRUNOLA</i>	Bruno-like 4, RNA binding protein	18q12 AS	Predicted	Maternal
<i>BTNL2</i>	Butyrophilin-like 2	6p21.3 AS	Predicted	Maternal
<i>C10orf91</i>	Chromosome 10 open reading frame 91	10q26.3	Predicted	Maternal
<i>C10orf93</i>	Chromosome 10 open reading frame 93	10q26.3 AS	Predicted	Maternal
<i>C16orf57</i>	Chromosome 16 open reading frame 57	16q21	Predicted	Maternal
<i>C20orf20</i>	Chromosome 20 open reading frame 20	20q13.33	Predicted	Maternal
<i>C20orf82</i>	Chromosome 20 open reading frame 82	20p12.1	Predicted	Paternal
<i>C6orf117</i>	Chromosome 6 open reading frame 117	6q14.3	Predicted	Paternal
<i>C9orf116</i>	Chromosome 9 open reading frame 116	9q34.3 AS	Predicted	Paternal
<i>C9orf85</i>	Chromosome 9 open reading frame 85	9q21.13	Predicted	Paternal
<i>CCDC85A</i>	Coiled-coil domain containing 85A	2p16.1	Predicted	Paternal
<i>CDH18</i>	Cadherin 18, type 2	5p15.2-p15.1 AS	Predicted	Paternal
<i>CDK4</i>	Cyclin-dependent kinase 4	12q14 AS	Predicted	Maternal
<i>CDKN1C</i>	Cyclin-dependent kinase inhibitor 1C	11p15.5 AS	Imprinted	Maternal
<i>CHMP2A</i>	Chromatin modifying protein 2A	19q AS	Predicted	Maternal
<i>CHST8</i>	Carbohydrate sulfotransferase 8	19q13.1	Predicted	Maternal
<i>COL9A3</i>	Collagen, type IX, alpha 3	20q13.3	Predicted	Maternal
<i>COPG2IT1</i>	COPG2 imprinted transcript 1	7q32	Imprinted	Paternal
<i>CPA4</i>	Carboxypeptidase A4	7q32	Imprinted	Maternal
<i>CSF2</i>	Colony stimulating factor 2	5q31.1	Predicted	Maternal
<i>CYP1B1</i>	Cytochrome P450, family 1, subfamily B, polypeptide 1	2p21 AS	Predicted	Paternal
<i>DDC</i>	Dopa decarboxylase	7p12.2 AS	Imprinted	Isoform Dependent
<i>DGCR6</i>	DiGeorge syndrome critical region gene 6	22q11.21	Imprinted	Unknown
<i>DGCR6L</i>	DiGeorge syndrome critical region gene 6-like	22q11 AS	Imprinted	Unknown
<i>DIRAS3</i>	DIRAS family, GTP-binding RAS-like 3	1p31 AS	Imprinted	Paternal
<i>DLGAP2</i>	Discs, large homolog-associated protein 2	8p23	Imprinted	Paternal

Table S1. Cont.

Gene ID	Full Gene Name	Location	Status	Expressed Allele
<i>DLK1</i>	Delta-like 1 homolog	14q32	Imprinted	Paternal
<i>DLX5</i>	Distal-less homeobox 5	7q22 AS	Imprinted	Maternal
<i>DNMT1</i>	DNA (cytosine-5-)-methyltransferase 1	19p13.2 AS	Imprinted	Paternal
<i>DVL1</i>	Dishevelled, dsh homolog 1	1p36 AS	Predicted	Maternal
<i>E2F7</i>	E2F transcription factor 7	12q21.2 AS	Predicted	Maternal
<i>EGFL7</i>	EGF-like-domain, multiple 7	9q34.3	Predicted	Paternal
<i>EVX1</i>	Even-skipped homeobox 1	7p15-p14	Predicted	Paternal
<i>FAM50B</i>	Family with sequence similarity 50, member B	6p25.2	Imprinted	Paternal
<i>FAM59A</i>	Family with sequence similarity 59, member A	18q12.1 AS	Predicted	Paternal
<i>FAM70B</i>	Family with sequence similarity 70, member B	13q34	Predicted	Maternal
<i>FASTK</i>	Fas-activated serine/threonine kinase	7q35 AS	Predicted	Maternal
<i>FBRSL1</i>	Fibrosin-like 1	12q24.33	Predicted	Maternal
<i>FERMT2</i>	Fermitin family homolog 2	14q22.1 AS	Predicted	Paternal
<i>FGFRL1</i>	Fibroblast growth factor receptor-like 1	4p16	Predicted	Maternal
<i>FLJ20464</i>	Hypothetical protein FLJ20464	22q12.2	Predicted	Paternal
<i>FLJ40296</i>	FLJ40296 protein	13q21.1	Predicted	Maternal
<i>FLJ46321</i>	FAM75-like protein FLJ46321	9q21.32	Predicted	Maternal
<i>FOXF1</i>	Forkhead box F1	16q24	Predicted	Maternal
<i>FOXG1</i>	Forkhead box G1	14q13	Predicted	Paternal
<i>FUCA1</i>	Fucosidase, alpha-L-1, tissue	1p34 AS	Predicted	Paternal
<i>GATA3</i>	GATA binding protein 3	10p15	Predicted	Paternal
<i>GFI1</i>	Growth factor independent 1 transcription repressor	1p22 AS	Predicted	Paternal
<i>GLI3</i>	GLI family zinc finger 3	7p13 AS	Predicted	Maternal
<i>GLIS3</i>	GLIS family zinc finger 3	9p24.2 AS	Imprinted	Paternal
<i>GNAS</i>	GNAS complex locus	20q13.3	Imprinted	Isoform Dependent
<i>GNASAS</i>	GNAS antisense RNA	20q13.32 AS	Imprinted	Paternal
<i>GPR1</i>	G protein-coupled receptor 1	2q33.3 AS	Imprinted	Paternal
<i>GPT</i>	Glutamic-pyruvate transaminase	8q24.3	Predicted	Maternal
<i>GRB10</i>	Growth factor receptor-bound protein 10	7p12-p11.2 AS	Imprinted	Isoform Dependent
<i>H19</i>	H19, imprinted maternally expressed untranslated mRNA	11p15.5 AS	Imprinted	Maternal
<i>HES1</i>	Hairy and enhancer of split 1,	3q28-q29	Predicted	Paternal
<i>HIST3H2BB</i>	Histone cluster 3, H2bb	1q42.13	Predicted	Maternal
<i>HOXA11</i>	Homeobox A11	7p15-p14 AS	Predicted	Maternal
<i>HOXA2</i>	Homeobox A2	7p15-p14 AS	Predicted	Maternal
<i>HOXA3</i>	Homeobox A3	7p15-p14 AS	Predicted	Maternal
<i>HOXA4</i>	Homeobox A4	7p15-p14 AS	Predicted	Maternal
<i>HOXA5</i>	Homeobox A5	7p15-p14 AS	Predicted	Maternal
<i>HOXB2</i>	Homeobox B2	17q21-q22 AS	Predicted	Maternal
<i>HOXB3</i>	Homeobox B3	17q21.3 AS	Predicted	Maternal
<i>HOXC4</i>	Homeobox C4	12q13.3	Predicted	Maternal
<i>HOXC9</i>	Homeobox C9	12q13.3	Predicted	Maternal

Table S1. Cont.

Gene ID	Full Gene Name	Location	Status	Expressed Allele
<i>HSPA6</i>	Heat shock 70 kDa protein 6	1q23	Predicted	Maternal
<i>HYMAI</i>	Hydatidiform mole associated and imprinted	6q24.2 AS	Imprinted	Paternal
<i>IFITM1</i>	Interferon induced transmembrane protein 1	11p15.5	Predicted	Maternal
<i>IGF2</i>	Insulin-like growth factor 2	11p15.5 AS	Imprinted	Paternal
<i>IGF2AS</i>	Insulin-like growth factor 2 antisense	11p15.5	Imprinted	Paternal
<i>INPP5F</i>	inositol polyphosphate-5-phosphatase F	10q26.11	Imprinted	Paternal
<i>INS</i>	Insulin	11p15.5 AS	Imprinted	Paternal
<i>ISM1</i>	Isthmin 1 homolog	20p12.1	Predicted	Paternal
<i>KBTD3</i>	Kelch repeat and BTB domain containing 3	11q22.3 AS	Predicted	Paternal
<i>KCNK9</i>	Potassium channel, subfamily K, member 9	8q24.3 AS	Imprinted	Maternal
<i>KCNQ1</i>	Potassium voltage-gated channel, KQT-like subfamily, member 1	11p15.5	Imprinted	Maternal
<i>KCNQ1DN</i>	KCNQ1 downstream neighbor	11p15.4	Imprinted	Maternal
<i>KCNQ1OT1</i>	KCNQ1 overlapping transcript 1	11p15	Imprinted	Paternal
<i>KIAA1530</i>	UV-Stimulated Scaffold Protein A	4p16.3	Predicted	Maternal
<i>KIAA1545</i>	KIAA1545 protein	12q24.33	Predicted	Maternal
<i>KLF14</i>	Kruppel-like factor 14	7q32.3 AS	Imprinted	Maternal
<i>L3MBTL</i>	L(3)mbt-like	20q13.12	Imprinted	Paternal
<i>LDB1</i>	LIM domain binding 1	10q24-q25 AS	Predicted	Maternal
<i>LILRB4</i>	Leukocyte immunoglobulin-like receptor, subfamily B, member 4	19q13.4	Predicted	Maternal
<i>LIN28B</i>	Lin-28 homolog B	6q21	Imprinted	Paternal
<i>LMX1B</i>	LIM homeobox transcription factor 1, beta	9q34	Predicted	Maternal
<i>LOC100131170</i>	Hypothetical LOC100131170	17q25.3 AS	Predicted	Paternal
<i>LRRTM1</i>	Leucine rich repeat transmembrane neuronal 1	2p12 AS	Imprinted	Paternal
<i>LY6D</i>	Lymphocyte antigen 6 complex, locus D	8q24-qter AS	Predicted	Paternal
<i>MAGEL2</i>	MAGE-like 2	15q11-q12 AS	Imprinted	Paternal
<i>MAGI2</i>	Membrane associated guanylate kinase, WW and PDZ domain containing 2	7q21 AS	Imprinted	Maternal
<i>MCTS2</i>	20q11.21	20q11.21	Imprinted	Paternal
<i>MEG3</i>	Maternally expressed 3	14q32	Imprinted	Maternal
<i>MEST</i>	Mesoderm specific transcript homolog	7q32	Imprinted	Paternal
<i>MESTIT1</i>	MEST intronic transcript 1, antisense RNA	7q32.2 AS	Imprinted	Paternal
<i>MIMT1</i>	MER1 repeat containing imprinted transcript 1	19q13.4	Imprinted	Paternal
<i>MIR296</i>	MicroRNA 296	20q13.32 AS	Imprinted	Paternal
<i>MIR298</i>	MicroRNA 298	20q13.32 AS	Imprinted	Paternal
<i>MIR371A</i>	MicroRNA 371a	19q13.42	Imprinted	Paternal
<i>MKRN3</i>	Makorin ring finger protein 3	15q11-q13	Imprinted	Paternal
<i>MRAP2</i>	Melanocortin 2 receptor accessory protein 2	6q14.2	Predicted	Paternal
<i>MYEOV2</i>	Myeloma overexpressed 2	2q37.3 AS	Predicted	Paternal
<i>MZF1</i>	Myeloid zinc finger 1	19q13.4 AS	Predicted	Maternal
<i>NAA60</i>	N(alpha)-acetyltransferase 60, NatF catalytic subunit	16p13.3	Imprinted	Maternal
<i>NAP1L5</i>	Nucleosome assembly protein 1-like 5	4q22.1 AS	Imprinted	Paternal

Table S1. Cont.

Gene ID	Full Gene Name	Location	Status	Expressed Allele
<i>NDN</i>	Necdin homolog	15q11.2-q12 AS	Imprinted	Paternal
<i>NDUFA4P1</i>	NADH dehydrogenase 1 alpha subcomplex, 4, 9 kDa, pseudogene 1	1p13.3	Predicted	Paternal
<i>NKAIN3</i>	Na <sup>+</sup> /K <sup>+</sup> transporting ATPase interacting 3	8q12.3	Predicted	Paternal
<i>NKX6-2</i>	NK6 homeobox 2	10q26 AS	Predicted	Maternal
<i>NLRP2</i>	NLR family, pyrin domain containing 2	19q13.42	Imprinted	Maternal
<i>NNAT</i>	Neuronatin	20q11.2-q12	Imprinted	Paternal
<i>NPAP1</i>	Nuclear pore associated protein 1	15q11-q13	Imprinted	Unknown
<i>NTM</i>	Neurotrimin	11q25	Imprinted	Maternal
<i>OBSCN</i>	Obscurin, cytoskeletal calmodulin and titin-interacting RhoGEF	1q42.13	Predicted	Paternal
<i>OR11L1</i>	Olfactory receptor, family 11, subfamily L, member 1	1q44 AS	Predicted	Paternal
<i>OSBPL5</i>	Oxysterol binding protein-like 5	11p15.4 AS	Imprinted	Maternal
<i>OTX1</i>	Orthodenticle homeobox 1	2p13	Predicted	Maternal
<i>PAOX</i>	Polyamine oxidase	10q26.3	Predicted	Maternal
<i>PEG10</i>	Paternally expressed 10	7q21	Imprinted	Paternal
<i>PEG3</i>	Paternally expressed gene 3	19q13.4 AS	Imprinted	Paternal
<i>PEX10</i>	Peroxisomal biogenesis factor 10	1p36.32 AS	Predicted	Maternal
<i>PHLDA2</i>	Pleckstrin homology-like domain, family A, member 2	11p15.5 AS	Imprinted	Maternal
<i>PHPT1</i>	Phosphohistidine phosphatase 1	9q34.3	Predicted	Maternal
<i>PKP3</i>	Plakophilin 3	11p15	Predicted	Maternal
<i>PLAGL1</i>	Pleiomorphic adenoma gene-like 1	6q24-q25 AS	Imprinted	Paternal
<i>PPAP2C</i>	Phosphatidic acid phosphatase type 2C	19p13 AS	Predicted	Maternal
<i>PPP1R9A</i>	Protein phosphatase 1, regulatory subunit 9A	7q21.3	Imprinted	Maternal
<i>PRDM16</i>	PR domain containing 16	1p36.23-p33	Predicted	Paternal
<i>PSIMCT-1</i>	Malignant T cell amplified sequence 1 pseudogene	20q11.2	Imprinted	Paternal
<i>PTPN14</i>	Protein tyrosine phosphatase, non-receptor type 14	1q32.2 AS	Predicted	Maternal
<i>PURG</i>	Purine-rich element binding protein G	8p11 AS	Predicted	Paternal
<i>PWCR1</i>	Prader-Willi syndrome chromosome region 1	15q11.2	Imprinted	Paternal
<i>PYY2</i>	Peptide YY, 2	17q11	Predicted	Paternal
<i>RAB1B</i>	RAB1B, member RAS oncogene family	11q12	Predicted	Maternal
<i>RB1</i>	Retinoblastoma 1	13q14.2	Imprinted	Maternal
<i>RBP5</i>	Retinol binding protein 5, cellular	12p13.31 AS	Imprinted	Maternal
<i>RNU5D-1</i>	RNA, U5D small nuclear 1	1p34.1 AS	Imprinted	Paternal
<i>RPL22</i>	Ribosomal protein L22	1p36.3-p36.2 AS	Predicted	Paternal
<i>SALL1</i>	Sal-like 1	16q12.1 AS	Predicted	Maternal
<i>SANG</i>	GNAS1 antisense	20q13.32	Imprinted	Paternal
<i>SGCE</i>	Sarcoglycan, epsilon	7q21-q22 AS	Imprinted	Paternal
<i>SIM2</i>	Single-minded homolog 2	21q22.2	Predicted	Paternal
<i>SLC22A18</i>	Solute carrier family 22, member 18	11p15.5	Imprinted	Maternal
<i>SLC22A2*</i>	Solute carrier family 22, member 2	6q26 AS	Imprinted	Maternal
<i>SLC22A3*</i>	Solute carrier family 22, member 3	6q26-q27	Imprinted	Maternal
<i>SLC26A10</i>	Solute carrier family 26, member 10	12q13	Predicted	Maternal

Table S1. Cont.

Gene ID	Full Gene Name	Location	Status	Expressed Allele
<i>SLC4A2</i>	Solute carrier family 4, anion exchanger, member 2	7q35-q36	Predicted	Maternal
<i>SNORD107</i>	Small nucleolar RNA, C/D box 107	15q11.2	Imprinted	Paternal
<i>SNORD108</i>	Small nucleolar RNA, C/D box 108	15q11.2	Imprinted	Paternal
<i>SNORD109A</i>	Small nucleolar RNA, C/D box 109A	15q11.2	Imprinted	Paternal
<i>SNORD109B</i>	Small nucleolar RNA, C/D box 109B	15q11.2	Imprinted	Paternal
<i>SNORD115@</i>	Small nucleolar RNA, C/D box 115 cluster	15q11.2	Imprinted	Paternal
<i>SNORD115-48</i>	Small nucleolar RNA, C/D box 115-48	15q11.2	Imprinted	Paternal
<i>SNORD116@</i>	Small nucleolar RNA, C/D box 116 cluster	15q11.2	Imprinted	Paternal
<i>SNORD64</i>	Small nucleolar RNA, C/D box 64	15q12	Imprinted	Paternal
<i>SNRPN</i>	Small nuclear ribonucleoprotein polypeptide N	15q11.2	Imprinted	Paternal
<i>SNURF</i>	SNRPN upstream reading frame	15q12	Imprinted	Paternal
<i>SOX8</i>	SRY-box 8	16p13.3	Predicted	Paternal
<i>SPON2</i>	Spondin 2, extracellular matrix protein	4p16.3 AS	Predicted	Paternal
<i>TCEB3C</i>	Transcription elongation factor B polypeptide 3C	18q21.1 AS	Imprinted	Maternal
<i>TFPI2</i>	Tissue factor pathway inhibitor 2	7q22 AS	Imprinted	Maternal
<i>TIGD1</i>	Tigger transposable element derived 1	2q37.1 AS	Predicted	Paternal
<i>TMEM52</i>	Transmembrane protein 52	1p36.33 AS	Predicted	Paternal
<i>TMEM60</i>	Transmembrane protein 60	7q11.23 AS	Predicted	Paternal
<i>TMEM88</i>	Transmembrane protein 88	17p13.1	Predicted	Maternal
<i>TP73</i>	Tumor protein p73	1p36.3	Imprinted	Maternal
<i>TSHZ3</i>	Teashirt zinc finger homeobox 3	19q12 AS	Predicted	Paternal
<i>UBE3A</i>	Ubiquitin protein ligase E3A	15q11-q13 AS	Imprinted	Maternal
<i>VAX2</i>	Ventral anterior homeobox 2	2p13	Predicted	Maternal
<i>VENTX</i>	VENT homeobox homolog	10q26.3	Predicted	Maternal
<i>WDR8</i>	WD repeat domain 8	1p36.3 AS	Predicted	Maternal
<i>WT1</i>	Wilms tumor 1	11p13 AS	Imprinted	Paternal
<i>ZC3H12C</i>	Zinc finger CCCH-type containing 12C	11q22.3	Imprinted	Paternal
<i>ZDBF2</i>	Zinc finger, DBF-type containing 2	2q33.3	Imprinted	Paternal
<i>ZFAT</i>	Zinc finger and AT hook domain containing	8q24.22 AS	Imprinted	Paternal
<i>ZFAT-AS1</i>	Zinc finger and AT hook domain containing antisense RNA 1	8q24.22	Imprinted	Paternal
<i>ZFP36L2</i>	Zinc finger protein 36, C3H type-like 2	2p22.3-p21 AS	Predicted	Maternal
<i>ZIC1</i>	Zic family member 1	3q24	Predicted	Maternal
<i>ZIM2</i>	Zinc finger, imprinted 2	19q13.4 AS	Imprinted	Paternal
<i>ZNF225</i>	Zinc finger protein 225	19q13.2	Predicted	Paternal
<i>ZNF229</i>	Zinc finger protein 229	19q13.31 AS	Predicted	Maternal
<i>ZNF597</i>	Zinc finger protein 597	16p13.3 AS	Imprinted	Maternal

**Table S2.** Associations between the imprinted gene set and Arsenic, Cadmium, Lead, and Mercury in human studies curated through the Comparative Toxicogenomics Database.

Gene ID	Metal	Interaction	Status	PMID
<i>ACD</i>	Lead	decreased expression of ACD mRNA	Predicted	19921347
<i>ANO1</i>	Arsenic	increased methylation of ANO1 promoter region	Imprinted	21291286
<i>APBA1</i>	Cadmium	decreased expression of APBA1 mRNA	Predicted	12634122
	Lead	decreased expression of APBA1 mRNA	Predicted	12634122
<i>C9ORF116</i>	Cadmium	increased expression of C9ORF116 mRNA	Predicted	21457566
<i>CDK4</i>	Arsenic	increased expression of CDK4 mRNA	Predicted	20654705 17530484 19769630 20016248
	Lead	increased expression of CDK4 mRNA	Predicted	19921347
<i>CDKN1C</i>	Arsenic	increased expression of CDKN1C mRNA	Imprinted	18585445
<i>CSF2</i>	Arsenic	decreased expression of RELA, CASP3, and CASP8 proteins	Predicted	16174796
		increased expression of CSF2 protein	Predicted	11312651
	Cadmium	increased expression and secretion of CSF2 protein	Predicted	18082304
<i>CYP1B1</i>	Arsenic	increased expression of CYP1B1 mRNA	Predicted	17530438 15894607
	Cadmium	decreased expression of CYP1B1 mRNA	Predicted	12760830 18560533
<i>DLK1</i>	Arsenic	DLK1 protein affects the susceptibility to Arsenic	Imprinted	18575777
<i>DNMT1</i>	Arsenic	decreased expression of DNMT1 mRNA	Imprinted	23159075 20596618 21176356 16613325
		decreased expression of DNMT1 protein	Imprinted	22558281 21687957
	Cadmium	decreased expression of DNMT1 mRNA	Imprinted	22112500
<i>E2F7</i>	Arsenic	decreased expression of E2F7 mRNA	Predicted	19945496
<i>EGFL7</i>	Lead	decreased expression of EGFL7 mRNA	Predicted	19921347
<i>FOXF1</i>	Arsenic	increased methylation of FOXF1 promoter	Predicted	21291286
<i>FUCA1</i>	Arsenic	increased expression of FUCA1 mRNA	Predicted	20458559
<i>GATA3</i>	Arsenic	inhibits the binding of GATA3 and IL13 promoter and KPNA2 protein	Predicted	19080345
<i>GNAS</i>	Lead	increased expression of GNAS mRNA	Imprinted	19921347
<i>GPT</i>	Cadmium	decreased expression of GPT protein	Predicted	11281253
<i>GRB10</i>	Arsenic	increased expression of GRB10 mRNA	Imprinted	20458559
	Arsenic	GRB10 mRNA affects the susceptibility to arsenic	Predicted	23911876
	Cadmium	decreased expression of GRB10 mRNA	Predicted	12160620
<i>HES1</i>	Arsenic	decreases expression of HES1 mRNA	Predicted	23832679 20458559 23497375 18633435
<i>HOXA5</i>	Arsenic	decreased expression of HOXA5 mRNA	Predicted	21461292
<i>HSPA6</i>	Arsenic	increased expression of HSPA6 mRNA	Predicted	19417148 16132727 15978632

Table S2. Cont.

Gene ID	Metal	Interaction	Status	PMID
		increased expression of HSPA6 protein	Predicted	15978632 19364129
	Cadmium	Increased expression of HSPA6 mRNA	Predicted	21172416 22562489 12064557
	Mercury	Increased expression of HSPA6 mRNA	Predicted	22129749
<i>IFITM1</i>	Arsenic	decreased expression of IFITM1 mRNA	Predicted	12890387
<i>IGF2</i>	Cadmium	promotes binding of IGF2 protein to IGF1R protein	Validated	9648927
<i>INS</i>	Arsenic	decreased methylation of INS promoter	Imprinted	23315758
	Cadmium	decreased expression of INS protein	Predicted	17303580
<i>LILRB4</i>	Lead	Altered expression of LILRB4 in relation to cord blood level concentrations	Predicted	22313790
<i>MRAP2</i>	Mercury	MRAP2 protein increases susceptibility to mercury	Predicted	21139347
<i>NDUFA4</i>	Arsenic	increased expression of NDUFA4 mRNA	Predicted	20458559
<i>OSBPL5</i>	Arsenic	altered expression of OSBPL5 mRNA	Imprinted	18414638
<i>PAOX</i>	Arsenic	increased expression of PAOX mRNA	Predicted	15761015
<i>PHLDA2</i>	Arsenic	decreased expression of PHLDA2 mRNA	Imprinted	15761015
<i>PKP3</i>	Arsenic	PKP3 protein results in decreased susceptibility to arsenic	Predicted	20707922
<i>PPAP2C</i>	Arsenic	PPAP2C protein results in decreased susceptibility to arsenic	Predicted	20707922
<i>PURG</i>	Arsenic	increased expression of PURG mRNA	Predicted	15761015
<i>RB1</i>	Arsenic	decreased expression of RB1 proteins	Imprinted	20953137
		decreased phosphorylation of RB1 proteins	Imprinted	12480548
	Cadmium	promotes phosphorylation of RB1 protein, binding of E2F1 protein	Imprinted	20839231
		decreased expression of RB1 mRNA	Imprinted	20839231
<i>SLC22A18</i>	Arsenic	decreased expression of SLC22A18 mRNA	Imprinted	15725085
<i>SLC22A2</i>	Mercury	decreased activity of SLC22A2 protein	Predicted	17287197
<i>SLC26A10</i>	Arsenic	decreased expression of SLC26A10 mRNA	Predicted	22521957
<i>SLC4A2</i>	Arsenic	increased expression of SLC4A2 protein	Predicted	17018029
<i>SPON2</i>	Arsenic	decreased expression of SPON2 mRNA	Predicted	19128835
	Cadmium	decreased expression of SPON2 mRNA	Predicted	21457566
<i>TIGD1</i>	Arsenic	affects the expression of TIGD1 mRNA	Predicted	18414638
<i>TP73</i>	Arsenic	increased expression of TP73 protein alternative form	Imprinted	16467208 15031205
		increased TP73 protein binding to TP53AIP1 promoter	Predicted	15031205
		increased acetylation of TP73 protein	Predicted	15031205
		increased expression of TP73 mRNA	Predicted	19763917 20458559
<i>UBE3A</i>	Arsenic	increased expression of UBE3A mRNA	Imprinted	20458559
<i>WT1</i>	Arsenic	decreased expression of WT1 mRNA	Predicted	16966277 20471514
		decreased expression of WT1 protein	Predicted	16966277