

Dana C Dolinoy

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

133
papers

7,750
citations

41
h-index

87
g-index

137
ext. papers

8,985
ext. citations

5.1
avg. IF

6.24
L-index

#	Paper	IF	Citations
133	Maternal and Neonatal One-Carbon Metabolites and the Epigenome-wide Infant Response.. <i>Journal of Nutritional Biochemistry</i> , 2022 , 101, 108938	6.3	1
132	Gestational exposure to high fat diets and bisphenol A alters metabolic outcomes in dams and offspring, but produces hepatic steatosis only in dams. <i>Chemosphere</i> , 2022 , 286, 131645	8.4	0
131	Prenatal Exposures to Common Phthalates and Prevalent Phthalate Alternatives and Infant DNA Methylation at Birth.. <i>Frontiers in Genetics</i> , 2022 , 13, 793278	4.5	0
130	Adolescent sleep timing and dietary patterns in relation to DNA methylation of core circadian genes: a pilot study of Mexican youth. <i>Epigenetics</i> , 2021 , 16, 894-907	5.7	2
129	Tissue and sex-specific programming of DNA methylation by perinatal lead exposure: implications for environmental epigenetics studies. <i>Epigenetics</i> , 2021 , 16, 1102-1122	5.7	4
128	U.S.-China Collaboration is Vital to Global Plans for a Healthy Environment and Sustainable Development. <i>Environmental Science & Technology</i> , 2021 , 55, 9622-9626	10.3	2
127	Prenatal Lead (Pb) Exposure and Peripheral Blood DNA Methylation (5mC) and Hydroxymethylation (5hmC) in Mexican Adolescents from the ELEMENT Birth Cohort. <i>Environmental Health Perspectives</i> , 2021 , 129, 67002	8.4	4
126	Paradoxical whole genome DNA methylation dynamics of 5mza-deoxycytidine in chronic low-dose exposure in mice. <i>Epigenetics</i> , 2021 , 16, 209-227	5.7	0
125	Association of Maternal-Neonatal Steroids With Early Pregnancy Endocrine Disrupting Chemicals and Pregnancy Outcomes. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2021 , 106, 665-687	5.6	4
124	Genome-Wide Epigenetic Signatures of Adaptive Developmental Plasticity in the Andes. <i>Genome Biology and Evolution</i> , 2021 , 13,	3.9	4
123	DNA methylation at birth potentially mediates the association between prenatal lead (Pb) exposure and infant neurodevelopmental outcomes. <i>Environmental Epigenetics</i> , 2021 , 7, dvab005	2.4	4
122	Perinatal DEHP exposure induces sex- and tissue-specific DNA methylation changes in both juvenile and adult mice. <i>Environmental Epigenetics</i> , 2021 , 7, dvab004	2.4	1
121	Blood lead levels in Peruvian adults are associated with proximity to mining and DNA methylation. <i>Environment International</i> , 2021 , 155, 106587	12.9	3
120	Sex-Specific Alterations in Cardiac DNA Methylation in Adult Mice by Perinatal Lead Exposure. <i>International Journal of Environmental Research and Public Health</i> , 2021 , 18,	4.6	2
119	Maternal lipodome across pregnancy is associated with the neonatal DNA methylome. <i>Epigenomics</i> , 2020 , 12, 2077-2092	4.4	2
118	Single-Cell Analysis of the Gene Expression Effects of Developmental Lead (Pb) Exposure on the Mouse Hippocampus. <i>Toxicological Sciences</i> , 2020 , 176, 396-409	4.4	3
117	Short- and long-term effects of perinatal phthalate exposures on metabolic pathways in the mouse liver. <i>Environmental Epigenetics</i> , 2020 , 6, dvaa017	2.4	1

116	Maternal environmental exposure to bisphenols and epigenome-wide DNA methylation in infant cord blood. <i>Environmental Epigenetics</i> , 2020 , 6, dvaa021	2.4	7
115	Accelerometer-measured Physical Activity, Reproductive Hormones, and DNA Methylation. <i>Medicine and Science in Sports and Exercise</i> , 2020 , 52, 598-607	1.2	5
114	Trimester-Specific Associations of Prenatal Lead Exposure With Infant Cord Blood DNA Methylation at Birth. <i>Epigenetics Insights</i> , 2020 , 13, 2516865720938669	3	10
113	Sex-Specific Programming of Cardiac DNA Methylation by Developmental Phthalate Exposure. <i>Epigenetics Insights</i> , 2020 , 13, 2516865720939971	3	5
112	Neonatal Lead (Pb) Exposure and DNA Methylation Profiles in Dried Bloodspots. <i>International Journal of Environmental Research and Public Health</i> , 2020 , 17,	4.6	9
111	Tissue- and Sex-Specific DNA Methylation Changes in Mice Perinatally Exposed to Lead (Pb). <i>Frontiers in Genetics</i> , 2020 , 11, 840	4.5	4
110	Maternal lipid levels across pregnancy impact the umbilical cord blood lipidome and infant birth weight. <i>Scientific Reports</i> , 2020 , 10, 14209	4.9	13
109	Integrative Analysis of Gene-Specific DNA Methylation and Untargeted Metabolomics Data from the ELEMENT Cohort. <i>Epigenetics Insights</i> , 2020 , 13, 2516865720977888	3	1
108	The role of environmental exposures and the epigenome in health and disease. <i>Environmental and Molecular Mutagenesis</i> , 2020 , 61, 176-192	3.2	25
107	Maternal Exposure to Environmental Disruptors and Sexually Dimorphic Changes in Maternal and Neonatal Oxidative Stress. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2020 , 105,	5.6	14
106	Characterization of the mouse white adipose tissue redox environment and associations with perinatal environmental exposures to bisphenol A and high-fat diets. <i>Journal of Nutritional Biochemistry</i> , 2019 , 66, 86-97	6.3	2
105	First trimester maternal exposures to endocrine disrupting chemicals and metals and fetal size in the Michigan Mother-Infant Pairs study. <i>Journal of Developmental Origins of Health and Disease</i> , 2019 , 10, 447-458	2.4	30
104	Phthalate Exposures, DNA Methylation and Adiposity in Mexican Children Through Adolescence. <i>Frontiers in Public Health</i> , 2019 , 7, 162	6	21
103	Sleep duration and fragmentation in relation to leukocyte DNA methylation in adolescents. <i>Sleep</i> , 2019 , 42,	1.1	3
102	Perinatal Lead (Pb) Exposure and Cortical Neuron-Specific DNA Methylation in Male Mice. <i>Genes</i> , 2019 , 10,	4.2	16
101	Dietary exposures, epigenetics and pubertal tempo. <i>Environmental Epigenetics</i> , 2019 , 5, dvz002	2.4	2
100	Genomic Tools for Environmental Epigenetics and Implications for Public Health. <i>Current Opinion in Toxicology</i> , 2019 , 18, 27-33	4.4	5
99	Somatic expression of piRNA and associated machinery in the mouse identifies short, tissue-specific piRNA. <i>Epigenetics</i> , 2019 , 14, 504-521	5.7	31

98	Early pregnancy exposure to endocrine disrupting chemical mixtures are associated with inflammatory changes in maternal and neonatal circulation. <i>Scientific Reports</i> , 2019 , 9, 5422	4.9	48
97	Early life social and ecological determinants of global DNA methylation in wild spotted hyenas. <i>Molecular Ecology</i> , 2019 , 28, 3799-3812	5.7	8
96	Longitudinal Metabolic Impacts of Perinatal Exposure to Phthalates and Phthalate Mixtures in Mice. <i>Endocrinology</i> , 2019 , 160, 1613-1630	4.8	11
95	Perinatal Bisphenol A Exposure and Reprogramming of Imprinted Gene Expression in the Adult Mouse Brain. <i>Frontiers in Genetics</i> , 2019 , 10, 951	4.5	7
94	DNA Methylation Changes Are Associated With an Incremental Ascent to High Altitude. <i>Frontiers in Genetics</i> , 2019 , 10, 1062	4.5	12
93	Early Life Exposure in Mexico to ENvironmental Toxicants (ELEMENT) Project. <i>BMJ Open</i> , 2019 , 9, e030437	4.7	39
92	LINE-1 and EPAS1 DNA methylation associations with high-altitude exposure. <i>Epigenetics</i> , 2019 , 14, 1-15	5.7	26
91	Neonatal bloodspot DNA methylation patterns are associated with childhood weight status in the Healthy Families Project. <i>Pediatric Research</i> , 2019 , 85, 848-855	3.2	6
90	The NIEHS TaRGET II Consortium and environmental epigenomics. <i>Nature Biotechnology</i> , 2018 , 36, 225-227	4.5	44
89	Maternal levels of endocrine disrupting chemicals in the first trimester of pregnancy are associated with infant cord blood DNA methylation. <i>Epigenetics</i> , 2018 , 13, 301-309	5.7	44
88	Epigenetics and the maintenance of developmental plasticity: extending the signalling theory framework. <i>Biological Reviews</i> , 2018 , 93, 1323-1338	13.5	28
87	Age-related epigenome-wide DNA methylation and hydroxymethylation in longitudinal mouse blood. <i>Epigenetics</i> , 2018 , 13, 779-792	5.7	18
86	Longitudinal effects of developmental bisphenol A, variable diet, and physical activity on age-related methylation in blood. <i>Environmental Epigenetics</i> , 2018 , 4, dvy017	2.4	7
85	Association of blood leukocyte DNA methylation at LINE-1 and growth-related candidate genes with pubertal onset and progression. <i>Epigenetics</i> , 2018 , 13, 1222-1233	5.7	9
84	Longitudinal Effects of Developmental Bisphenol A Exposure on Epigenome-Wide DNA Hydroxymethylation at Imprinted Loci in Mouse Blood. <i>Environmental Health Perspectives</i> , 2018 , 126, 077006	8.4	19
83	Environmental Deflection: The Impact of Toxicant Exposures on the Aging Epigenome. <i>Toxicological Sciences</i> , 2017 , 156, 325-335	4.4	22
82	Sexually Dimorphic Effects of Early-Life Exposures to Endocrine Disruptors: Sex-Specific Epigenetic Reprogramming as a Potential Mechanism. <i>Current Environmental Health Reports</i> , 2017 , 4, 426-438	6.5	30
81	Hepatic Lipid Accumulation and Nrf2 Expression following Perinatal and Peripubertal Exposure to Bisphenol A in a Mouse Model of Nonalcoholic Liver Disease. <i>Environmental Health Perspectives</i> , 2017 , 125, 087005	8.4	45

80	Small-Magnitude Effect Sizes in Epigenetic End Points are Important in Children's Environmental Health Studies: The Children's Environmental Health and Disease Prevention Research Center's Epigenetics Working Group. <i>Environmental Health Perspectives</i> , 2017 , 125, 511-526	8.4	158
79	Perinatal lead (Pb) exposure results in sex and tissue-dependent adult DNA methylation alterations in murine IAP transposons. <i>Environmental and Molecular Mutagenesis</i> , 2017 , 58, 540-550	3.2	28
78	Novel Epigenetic Biomarkers Mediating Bisphenol A Exposure and Metabolic Phenotypes in Female Mice. <i>Endocrinology</i> , 2017 , 158, 31-40	4.8	28
77	Longitudinal effects of developmental bisphenol A and variable diet exposures on epigenetic drift in mice. <i>Reproductive Toxicology</i> , 2017 , 68, 154-163	3.4	33
76	Gene-specific DNA methylation may mediate atypical antipsychotic-induced insulin resistance. <i>Bipolar Disorders</i> , 2016 , 18, 423-32	3.8	18
75	Environmental Contaminants and Child Development. <i>Child Development Perspectives</i> , 2016 , 10, 228-233	5.5	13
74	Lipid metabolism is associated with developmental epigenetic programming. <i>Scientific Reports</i> , 2016 , 6, 34857	4.9	25
73	Detection of differential DNA methylation in repetitive DNA of mice and humans perinatally exposed to bisphenol A. <i>Epigenetics</i> , 2016 , 11, 489-500	5.7	37
72	Mono-2-ethylhexyl phthalate disrupts neurulation and modifies the embryonic redox environment and gene expression. <i>Reproductive Toxicology</i> , 2016 , 63, 32-48	3.4	22
71	Correlation between Conjugated Bisphenol A Concentrations and Efflux Transporter Expression in Human Fetal Livers. <i>Drug Metabolism and Disposition</i> , 2016 , 44, 1061-5	4	8
70	Perinatal Lead Exposure Alters Gut Microbiota Composition and Results in Sex-specific Bodyweight Increases in Adult Mice. <i>Toxicological Sciences</i> , 2016 , 151, 324-33	4.4	81
69	Genetic polymorphisms are associated with hair, blood, and urine mercury levels in the American Dental Association (ADA) study participants. <i>Environmental Research</i> , 2016 , 149, 247-258	7.9	22
68	Adolescent epigenetic profiles and environmental exposures from early life through peri-adolescence. <i>Environmental Epigenetics</i> , 2016 , 2, dvw018	2.4	30
67	Epigenome-wide DNA methylation analysis implicates neuronal and inflammatory signaling pathways in adult murine hepatic tumorigenesis following perinatal exposure to bisphenol A. <i>Environmental and Molecular Mutagenesis</i> , 2016 , 57, 435-46	3.2	8
66	Prenatal exposures and DNA methylation in newborns: a pilot study in Durban, South Africa. <i>Environmental Sciences: Processes and Impacts</i> , 2016 , 18, 908-17	4.3	17
65	Mono-2-ethylhexyl phthalate (MEHP) alters histiotrophic nutrition pathways and epigenetic processes in the developing conceptus. <i>Journal of Nutritional Biochemistry</i> , 2016 , 27, 211-8	6.3	17
64	Impact of gestational bisphenol A on oxidative stress and free fatty acids: Human association and interspecies animal testing studies. <i>Endocrinology</i> , 2015 , 156, 911-22	4.8	44
63	DNA methylation, insulin resistance and second-generation antipsychotics in bipolar disorder. <i>Epigenomics</i> , 2015 , 7, 343-52	4.4	25

62	Bisphenol A-associated alterations in genome-wide DNA methylation and gene expression patterns reveal sequence-dependent and non-monotonic effects in human fetal liver. <i>Environmental Epigenetics</i> , 2015 , 1,	2.4	45
61	Stat3 is a candidate epigenetic biomarker of perinatal Bisphenol A exposure associated with murine hepatic tumors with implications for human health. <i>Epigenetics</i> , 2015 , 10, 1099-110	5.7	20
60	Assessing Human Health Risk to Endocrine Disrupting Chemicals: a Focus on Prenatal Exposures and Oxidative Stress. <i>Endocrine Disruptors (Austin, Tex)</i> , 2015 , 3,		20
59	In utero bisphenol A concentration, metabolism, and global DNA methylation across matched placenta, kidney, and liver in the human fetus. <i>Chemosphere</i> , 2015 , 124, 54-60	8.4	89
58	New insights and updated guidelines for epigenome-wide association studies. <i>Neuroepigenetics</i> , 2015 , 1, 14-19		20
57	Quality control and statistical modeling for environmental epigenetics: a study on in utero lead exposure and DNA methylation at birth. <i>Epigenetics</i> , 2015 , 10, 19-30	5.7	41
56	Perinatal bisphenol A exposure promotes dose-dependent alterations of the mouse methylome. <i>BMC Genomics</i> , 2014 , 15, 30	4.5	61
55	Emerging issues in public health genomics. <i>Annual Review of Genomics and Human Genetics</i> , 2014 , 15, 461-480	9.7	30
54	Epigenetics: relevance and implications for public health. <i>Annual Review of Public Health</i> , 2014 , 35, 105-220.6	20.6	66
53	Bisphenol A-associated alterations in the expression and epigenetic regulation of genes encoding xenobiotic metabolizing enzymes in human fetal liver. <i>Environmental and Molecular Mutagenesis</i> , 2014 , 55, 184-95	3.2	41
52	Early-Life Exposures and the Epigenome: Interactions between Nutrients and the Environment. <i>Oxidative Stress and Disease</i> , 2014 , 3-52		
51	Perinatal lead (Pb) exposure results in sex-specific effects on food intake, fat, weight, and insulin response across the murine life-course. <i>PLoS ONE</i> , 2014 , 9, e104273	3.7	51
50	Epigenomic Indicators of Age in African Americans. <i>Hereditary Genetics: Current Research</i> , 2014 , 3,		8
49	Longitudinal epigenetic drift in mice perinatally exposed to lead. <i>Epigenetics</i> , 2014 , 9, 934-41	5.7	38
48	Dose-dependent incidence of hepatic tumors in adult mice following perinatal exposure to bisphenol A. <i>Environmental Health Perspectives</i> , 2014 , 122, 485-91	8.4	101
47	The psychology of Vegretable substitutions\Examining consumer judgements of Bisphenol A and its alternatives. <i>Health, Risk and Society</i> , 2014 , 16, 649-666	2	15
46	Bisphenol A at concentrations relevant to human exposure enhances histamine and cysteinyl leukotriene release from bone marrow-derived mast cells. <i>Journal of Immunotoxicology</i> , 2014 , 11, 84-9	3.1	13
45	Perinatal bisphenol A exposures increase production of pro-inflammatory mediators in bone marrow-derived mast cells of adult mice. <i>Journal of Immunotoxicology</i> , 2014 , 11, 205-12	3.1	44

44	Bisphenol A-associated epigenomic changes in prepubescent girls: a cross-sectional study in Gharbiah, Egypt. <i>Environmental Health</i> , 2013 , 12, 33	6	55
43	Complex Phenotypes: Epigenetic Manifestation of Environmental Exposures. <i>Epigenetics and Human Health</i> , 2013 , 77-97		1
42	Early-life lead exposure results in dose- and sex-specific effects on weight and epigenetic gene regulation in weanling mice. <i>Epigenomics</i> , 2013 , 5, 487-500	4.4	78
41	Mercury biomarkers and DNA methylation among Michigan dental professionals. <i>Environmental and Molecular Mutagenesis</i> , 2013 , 54, 195-203	3.2	73
40	Inhibition of proteolysis in histiotrophic nutrition pathways alters DNA methylation and one-carbon metabolism in the organogenesis-stage rat conceptus. <i>Journal of Nutritional Biochemistry</i> , 2013 , 24, 1479-87	6.3	12
39	Phylogenetic and DNA methylation analysis reveal novel regions of variable methylation in the mouse IAP class of transposons. <i>BMC Genomics</i> , 2013 , 14, 48	4.5	23
38	Perinatal bisphenol A exposure promotes hyperactivity, lean body composition, and hormonal responses across the murine life course. <i>FASEB Journal</i> , 2013 , 27, 1784-92	0.9	84
37	Adaptive radiation-induced epigenetic alterations mitigated by antioxidants. <i>FASEB Journal</i> , 2013 , 27, 665-71	0.9	83
36	Fetal liver bisphenol A concentrations and biotransformation gene expression reveal variable exposure and altered capacity for metabolism in humans. <i>Journal of Biochemical and Molecular Toxicology</i> , 2013 , 27, 116-23	3.4	64
35	Comprehensive analysis of DNA methylation in head and neck squamous cell carcinoma indicates differences by survival and clinicopathologic characteristics. <i>PLoS ONE</i> , 2013 , 8, e54742	3.7	41
34	Low dose effects of bisphenol A. <i>Endocrine Disruptors (Austin, Tex)</i> , 2013 , 1, e26490		139
33	Perinatal bisphenol A exposure promotes hyperactivity with corresponding hormonal responses. <i>FASEB Journal</i> , 2013 , 27, 1073.10	0.9	
32	Patterns of cellular and HPV 16 methylation as biomarkers for cervical neoplasia. <i>Journal of Virological Methods</i> , 2012 , 184, 84-92	2.6	21
31	Nutrition and epigenetics: an interplay of dietary methyl donors, one-carbon metabolism and DNA methylation. <i>Journal of Nutritional Biochemistry</i> , 2012 , 23, 853-9	6.3	452
30	Epigenetics for ecotoxicologists. <i>Environmental Toxicology and Chemistry</i> , 2012 , 31, 221-7	3.8	60
29	DNA methylation screening and analysis. <i>Methods in Molecular Biology</i> , 2012 , 889, 385-406	1.4	25
28	Concordance in hippocampal and fecal Nr3c1 methylation is moderated by maternal behavior in the mouse. <i>Ecology and Evolution</i> , 2012 , 2, 3123-31	2.8	18
27	LRpath analysis reveals common pathways dysregulated via DNA methylation across cancer types. <i>BMC Genomics</i> , 2012 , 13, 526	4.5	50

26	Urinary bisphenol A concentrations in girls from rural and urban Egypt: a pilot study. <i>Environmental Health</i> , 2012 , 11, 20	6	34
25	Delivery type not associated with global methylation at birth. <i>Clinical Epigenetics</i> , 2012 , 4, 8	7-7	33
24	Bioinformatics for High-Throughput Toxico-Epigenomics Studies 2012 , 569-588		1
23	Epigenetic responses following maternal dietary exposure to physiologically relevant levels of bisphenol A. <i>Environmental and Molecular Mutagenesis</i> , 2012 , 53, 334-42	3-2	121
22	Alzheimer's disease and environmental exposure to lead: the epidemiologic evidence and potential role of epigenetics. <i>Current Alzheimer Research</i> , 2012 , 9, 563-73	3	131
21	Pretreatment dietary intake is associated with tumor suppressor DNA methylation in head and neck squamous cell carcinomas. <i>Epigenetics</i> , 2012 , 7, 883-91	5-7	31
20	Introduction: The use of animals models to advance epigenetic science. <i>ILAR Journal</i> , 2012 , 53, 227-31	1-7	7
19	Genome-wide DNA methylation differences between late-onset Alzheimer's disease and cognitively normal controls in human frontal cortex. <i>Journal of Alzheimer's Disease</i> , 2012 , 29, 571-88	4-3	184
18	Epigenetic Manifestation of Environmental Exposures 2011 , 287-307		1
17	Genome-wide methylation and expression differences in HPV(+) and HPV(-) squamous cell carcinoma cell lines are consistent with divergent mechanisms of carcinogenesis. <i>Epigenetics</i> , 2011 , 6, 777-87	5-7	118
16	An expression microarray approach for the identification of metastable epialleles in the mouse genome. <i>Epigenetics</i> , 2011 , 6, 1105-13	5-7	31
15	Timing is everything: the when and how of environmentally induced changes in the epigenome of animals. <i>Epigenetics</i> , 2011 , 6, 791-7	5-7	224
14	Variable histone modifications at the A(vy) metastable epiallele. <i>Epigenetics</i> , 2010 , 5, 637-44	5-7	75
13	Epigenetic Gene Regulation: Linking Early Development Environment to Adult Disease. Dana Dolinoy, Ph.D.. <i>Biology of Reproduction</i> , 2009 , 81, 113-113	3-9	
12	The agouti mouse model: an epigenetic biosensor for nutritional and environmental alterations on the fetal epigenome. <i>Nutrition Reviews</i> , 2008 , 66 Suppl 1, S7-11	6-4	182
11	Environmental epigenomics in human health and disease. <i>Environmental and Molecular Mutagenesis</i> , 2008 , 49, 4-8	3-2	267
10	Epigenetic gene regulation: linking early developmental environment to adult disease. <i>Reproductive Toxicology</i> , 2007 , 23, 297-307	3-4	394
9	Metastable epialleles, imprinting, and the fetal origins of adult diseases. <i>Pediatric Research</i> , 2007 , 61, 30R-37R	3-2	193

8	Cancer susceptibility: epigenetic manifestation of environmental exposures. <i>Cancer Journal (Sudbury, Mass)</i> , 2007 , 13, 9-16	2.2	78
7	Maternal nutrient supplementation counteracts bisphenol A-induced DNA hypomethylation in early development. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007 , 104, 13056-61	11.5	1080
6	Maternal methyl supplements increase offspring DNA methylation at Axin Fused. <i>Genesis</i> , 2006 , 44, 401-69	4.0	400
5	Maternal genistein alters coat color and protects Avy mouse offspring from obesity by modifying the fetal epigenome. <i>Environmental Health Perspectives</i> , 2006 , 114, 567-72	8.4	758
4	Using GIS-based approaches to support research on neurotoxicants and other children's environmental health threats. <i>NeuroToxicology</i> , 2005 , 26, 223-8	4.4	13
3	GIS modeling of air toxics releases from TRI-reporting and non-TRI-reporting facilities: impacts for environmental justice. <i>Environmental Health Perspectives</i> , 2004 , 112, 1717-24	8.4	52
2	Mapping for prevention: GIS models for directing childhood lead poisoning prevention programs. <i>Environmental Health Perspectives</i> , 2002 , 110, 947-53	8.4	83
1	Toxicopigenetics and Effects on Life Course Disease Susceptibility	439-472	