Andrea A Baccarelli

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4040862/publications.pdf

Version: 2024-02-01

339 papers 22,413 citations

¹⁶⁷⁹¹ 66 h-index

131 g-index

351 all docs

351 docs citations

times ranked

351

24571 citing authors

#	Article	IF	CITATIONS
1	White matter microstructural variability linked to differential attentional skills and impulsive behavior in a pediatric population. Cerebral Cortex, 2023, 33, 1895-1912.	1.6	5
2	A methodological pipeline to generate an epigenetic marker of prenatal exposure to air pollution indicators. Epigenetics, 2022, 17, 32-40.	1.3	8
3	Large-Scale Hypothesis Testing for Causal Mediation Effects with Applications in Genome-wide Epigenetic Studies. Journal of the American Statistical Association, 2022, 117, 67-81.	1.8	35
4	Impact of paternal education on epigenetic ageing in adolescence and mid-adulthood: a multi-cohort study in the USA and Mexico. International Journal of Epidemiology, 2022, 51, 870-884.	0.9	6
5	Associations between infant sex and DNA methylation across umbilical cord blood, artery, and placenta samples. Epigenetics, 2022, 17, 1080-1097.	1.3	14
6	Prenatal maternal phthalate exposures and trajectories of childhood adiposity from four to twelve years. Environmental Research, 2022, 204, 112111.	3.7	8
7	Critical windows of perinatal particulate matter (PM2.5) exposure and preadolescent kidney function. Environmental Research, 2022, 204, 112062.	3.7	5
8	Role of brain extracellular vesicles in air pollution-related cognitive impairment and neurodegeneration. Environmental Research, 2022, 204, 112316.	3.7	13
9	Prenatal particulate matter exposure and mitochondrial mutational load at the maternal-fetal interface: Effect modification by genetic ancestry. Mitochondrion, 2022, 62, 102-110.	1.6	8
10	Marine pollutant exposures and human milk extracellular vesicle-microRNAs in a mother-infant cohort from the Faroe Islands. Environment International, 2022, 158, 106986.	4.8	11
11	Department Chairs Weigh In: Environmental Health Education Is More Essential Than Ever. American Journal of Public Health, 2022, 112, 75-76.	1.5	O
12	The relationship between persistent organic pollutants and Attention Deficit Hyperactivity Disorder phenotypes: Evidence from task-based neural activity in an observational study of a community sample of Canadian mother-child dyads. Environmental Research, 2022, 206, 112593.	3.7	7
13	Prenatal lead exposure, telomere length in cord blood, and DNA methylation age in the PROGRESS prenatal cohort. Environmental Research, 2022, 205, 112577.	3.7	11
14	Associations of Childhood and Perinatal Blood Metals with Children's Gut Microbiomes in a Canadian Gestation Cohort. Environmental Health Perspectives, 2022, 130, 17007.	2.8	13
15	DunedinPACE, a DNA methylation biomarker of the pace of aging. ELife, 2022, 11, .	2.8	214
16	Prenatal phthalates, gestational weight gain, and long-term weight changes among Mexican women. Environmental Research, 2022, 209, 112835.	3.7	4
17	An exposomic framework to uncover environmental drivers of aging. Exposome, 2022, 2, osac002.	1.2	12
18	Mid-life epigenetic age, neuroimaging brain age, and cognitive function: coronary artery risk development in young adults (CARDIA) study. Aging, 2022, 14, 1691-1712.	1.4	16

#	Article	IF	CITATIONS
19	Gestational Perfluoroalkyl Substance Exposure and DNA Methylation at Birth and 12 Years of Age: A Longitudinal Epigenome-Wide Association Study. Environmental Health Perspectives, 2022, 130, 37005.	2.8	24
20	Association of Prenatal Acetaminophen Exposure Measured in Meconium With Adverse Birth Outcomes in a Canadian Birth Cohort. Frontiers in Pediatrics, 2022, 10, 828089.	0.9	0
21	Predicting chemical ecotoxicity by learning latent space chemical representations. Environment International, 2022, 163, 107224.	4.8	5
22	The role of outdoor and indoor air quality in the spread of SARS-CoV-2: Overview and recommendations by the research group on COVID-19 and particulate matter (RESCOP commission). Environmental Research, 2022, 211, 113038.	3.7	42
23	Environmental Exposures and Extracellular Vesicles: Indicators of Systemic Effects and Human Disease. Current Environmental Health Reports, 2022, 9, 465-476.	3.2	18
24	Integrative analysis of clinical and epigenetic biomarkers of mortality. Aging Cell, 2022, 21, e13608.	3.0	8
25	Pulmonary Function and Blood DNA Methylation: A Multiancestry Epigenome-Wide Association Meta-analysis. American Journal of Respiratory and Critical Care Medicine, 2022, 206, 321-336.	2.5	15
26	DNA methylation signature of chronic low-grade inflammation and its role in cardio-respiratory diseases. Nature Communications, 2022, 13, 2408.	5.8	26
27	Gaseous air pollutants and DNA methylation in a methylome-wide association study of an ethnically and environmentally diverse population of U.S. adults. Environmental Research, 2022, 212, 113360.	3.7	7
28	Association of DNA methylation in circulating CD4+T cells with short-term PM2.5 pollution waves: A quasi-experimental study of healthy young adults. Ecotoxicology and Environmental Safety, 2022, 239, 113634.	2.9	3
29	Environmentally Just Futures: A Collection of Community-Driven African Environmental Education and Improvement Initiatives. International Journal of Environmental Research and Public Health, 2022, 19, 6622.	1.2	2
30	Predictors of patterns of weight change 1 year after delivery in a cohort of Mexican women. Public Health Nutrition, 2021, 24, 4113-4123.	1.1	4
31	Prenatal gestational diabetes mellitus exposure and accelerated offspring DNA methylation age in early childhood. Epigenetics, 2021, 16, 186-195.	1.3	25
32	Associations between maternal lifetime stressors and negative events in pregnancy and breast milk-derived extracellular vesicle microRNAs in the programming of intergenerational stress mechanisms (PRISM) pregnancy cohort. Epigenetics, 2021, 16, 389-404.	1.3	20
33	Testing cell-type-specific mediation effects in genome-wide epigenetic studies. Briefings in Bioinformatics, 2021, 22, .	3.2	5
34	Prenatal maternal phthalate exposures and child lipid and adipokine levels at age six: A study from the PROGRESS cohort of Mexico City. Environmental Research, 2021, 192, 110341.	3.7	13
35	DNA methylation-based biomarkers of age acceleration and all-cause death, myocardial infarction, stroke, and cancer in two cohorts: The NAS, and KORA F4. EBioMedicine, 2021, 63, 103151.	2.7	42
36	Cord blood androgen levels of females from same sex and opposite sex twins – A pilot study. Clinical Endocrinology, 2021, 94, 85-89.	1.2	2

#	Article	IF	CITATIONS
37	Blood DNA methylation biomarkers of cumulative lead exposure in adults. Journal of Exposure Science and Environmental Epidemiology, 2021, 31, 108-116.	1.8	21
38	Maternal anxiety during pregnancy and newborn epigenome-wide DNA methylation. Molecular Psychiatry, 2021, 26, 1832-1845.	4.1	24
39	Hallmarks of environmental insults. Cell, 2021, 184, 1455-1468.	13.5	177
40	Mental Health of High-Risk Urban Youth: The Housing Subsidies Paradox. Race and Social Problems, 2021, 13, 22-33.	1.2	0
41	Long-term PM2.5 exposure before diagnosis is associated with worse outcome in breast cancer. Breast Cancer Research and Treatment, 2021, 188, 525-533.	1.1	8
42	DNA methylation architecture of the ACE2 gene in nasal cells of children. Scientific Reports, 2021, 11, 7107.	1.6	21
43	Associations Between Maternal Lifetime Stress and Placental Mitochondrial DNA Mutations in an Urban Multiethnic Cohort. Biological Psychiatry, 2021, 89, 570-578.	0.7	11
44	Human milk extracellular vesicle miRNA expression and associations with maternal characteristics in a population-based cohort from the Faroe Islands. Scientific Reports, 2021, 11, 5840.	1.6	34
45	Nanoparticle Tracking Analysis for the Quantification and Size Determination of Extracellular Vesicles. Journal of Visualized Experiments, 2021, , .	0.2	21
46	Epigenome-wide association study of kidney function identifies trans-ethnic and ethnic-specific loci. Genome Medicine, 2021, 13, 74.	3.6	20
47	Exosomal miRNAs in urine associated with children's cardiorenal parameters: a cross-sectional study. Epigenomics, 2021, 13, 499-512.	1.0	3
48	Short-term air pollution, cognitive performance and nonsteroidal anti-inflammatory drug use in the Veterans Affairs Normative Aging Study. Nature Aging, 2021, 1, 430-437.	5.3	33
49	The association between prenatal concentrations of polybrominated diphenyl ether and child cognitive and psychomotor function. Environmental Epidemiology, 2021, 5, e156.	1.4	3
50	A multi-ethnic epigenome-wide association study of leukocyte DNA methylation and blood lipids. Nature Communications, 2021, 12, 3987.	5.8	18
51	Epigenetic Aging Biomarkers Associated With Cognitive Impairment in Older African American Adults With Human Immunodeficiency Virus (HIV). Clinical Infectious Diseases, 2021, 73, 1982-1991.	2.9	14
52	DNAm-based signatures of accelerated aging and mortality in blood are associated with low renal function. Clinical Epigenetics, 2021, 13, 121.	1.8	13
53	Dietary fat intake during early pregnancy is associated with cord blood DNA methylation at <i>IGF2</i> and <i>H19</i> genes in newborns. Environmental and Molecular Mutagenesis, 2021, 62, 388-398.	0.9	9
54	Placental mitochondrial DNA mutational load and perinatal outcomes: Findings from a multi-ethnic pregnancy cohort. Mitochondrion, 2021, 59, 267-275.	1.6	8

#	Article	IF	CITATIONS
55	Telomere dynamics across the early life course: Findings from a longitudinal study in children. Psychoneuroendocrinology, 2021, 129, 105270.	1.3	10
56	Risks of Macrosomia Associated with Catechol-O-Methyltransferase Genotypes and Genetic–Epigenetic Interactions among Children with and without Gestational Diabetes Exposure. Childhood Obesity, 2021, 17, 365-370.	0.8	0
57	Causal mediation analysis with latent subgroups. Statistics in Medicine, 2021, 40, 5628-5641.	0.8	1
58	Prenatal Household Air Pollution Exposure, Cord Blood Mononuclear Cell Telomere Length and Age Four Blood Pressure: Evidence from a Ghanaian Pregnancy Cohort. Toxics, 2021, 9, 169.	1.6	12
59	Epigenetically mediated electrocardiographic manifestations of sub-chronic exposures to ambient particulate matter air pollution in the Women's Health Initiative and Atherosclerosis Risk in Communities Study. Environmental Research, 2021, 198, 111211.	3.7	4
60	Assessing the Effects of Metal Mixtures in Urine and Blood on Kidney Function. ISEE Conference Abstracts, 2021, 2021, .	0.0	0
61	Associations between air pollution and temperature on glycated hemoglobin levels in women of child bearing age. ISEE Conference Abstracts, 2021, 2021, .	0.0	0
62	Critical windows of perinatal particulate matter (PM2.5) exposure and preadolescent kidney function. ISEE Conference Abstracts, 2021, 2021, .	0.0	0
63	Epigenetic Age in Young African American Adults With Perinatally Acquired HIV. Journal of Acquired Immune Deficiency Syndromes (1999), 2021, 87, 1102-1109.	0.9	9
64	Blood DNA Methylation and Incident Coronary Heart Disease. JAMA Cardiology, 2021, 6, 1237.	3.0	24
65	The Hallmarks of Aging and Environmental Insults: Molecular Mechanisms. ISEE Conference Abstracts, 2021, 2021, .	0.0	0
66	Metal exposure and bone remodeling during pregnancy: Results from the PROGRESS cohort study. Environmental Pollution, 2021, 282, 116962.	3.7	11
67	Prenatal lead exposure, telomere length in cord blood and DNA methylation age in the PROGRESS cohort. ISEE Conference Abstracts, 2021, 2021, .	0.0	0
68	Residential PM2.5 exposure and the nasal methylome in children. Environment International, 2021, 153, 106505.	4.8	10
69	Long-term particulate matter exposure and bone mineral density in the Women's Health Initiative. ISEE Conference Abstracts, 2021, 2021, .	0.0	0
70	Associations between liver PFAS concentrations and plasma extracellular miRNAs in a cohort of adolescents undergoing bariatric surgery. ISEE Conference Abstracts, 2021, 2021, .	0.0	0
71	Nasal DNA Methylation Architecture of the {ACE2} gene and Epigenetic Aging. ISEE Conference Abstracts, 2021, 2021, .	0.0	0
72	Nephrotoxic Metal Mixtures and Preadolescent Kidney Function. Children, 2021, 8, 673.	0.6	5

#	Article	IF	CITATIONS
73	Associations between maternal asthma and atopy and breast milk-derived extracellular vesicle microRNA profiles in the PRISM pregnancy cohort. ISEE Conference Abstracts, 2021, 2021, .	0.0	O
74	Associations between indoor temperature and noise and semen parameters among participants in the US-based general population Growing Up Today Study. ISEE Conference Abstracts, 2021, 2021, .	0.0	0
75	Infant sex and DNA methylation: differentially methylated regions and positions across umbilical cord blood, artery, and placenta samples. ISEE Conference Abstracts, 2021, 2021, .	0.0	1
76	Prenatal metal exposure, cord blood DNA methylation and persistence in childhood: epigenome-wide association study of twelve metals. ISEE Conference Abstracts, 2021, 2021, .	0.0	0
77	Novel epigenetic link between gestational diabetes mellitus and macrosomia. Epigenomics, 2021, 13, 1221-1230.	1.0	4
78	In Utero Exposure to Persistent Organic Pollutants and Childhood Lipid Levels. Metabolites, 2021, 11, 657.	1.3	10
79	Prenatal blood lead levels and reduced preadolescent glomerular filtration rate: Modification by body mass index. Environment International, 2021, 154, 106414.	4.8	10
80	Effect of School Integrated Pest Management or Classroom Air Filter Purifiers on Asthma Symptoms in Students With Active Asthma. JAMA - Journal of the American Medical Association, 2021, 326, 839.	3.8	45
81	Short-Term Ambient Particulate Air Pollution and Hospitalization Expenditures of Cause-Specific Cardiorespiratory Diseases in China: A Multicity Analysis. The Lancet Regional Health - Western Pacific, 2021, 15, 100232.	1.3	15
82	Prenatal urinary concentrations of phthalate metabolites and behavioral problems in Mexican children: The Programming Research in Obesity, Growth Environment and Social Stress (PROGRESS) study. Environmental Research, 2021, 201, 111338.	3.7	6
83	Short-term exposure to PM2.5 components and renal health: Findings from the Veterans Affairs Normative Aging Study. Journal of Hazardous Materials, 2021, 420, 126557.	6.5	20
84	The associations of phthalate biomarkers during pregnancy with later glycemia and lipid profiles. Environment International, 2021, 155, 106612.	4.8	14
85	L-arginine supplementation to mitigate cardiovascular effects of walking outside in the context of traffic-related air pollution in participants with elevated blood pressure: A randomized, double-blind, placebo-controlled trial. Environment International, 2021, 156, 106631.	4.8	5
86	Isolation and characterization of extracellular vesicles in saliva of children with asthma. , 2021, 2, 29-48.		11
87	Extracellular vesicles and female reproduction. Journal of Assisted Reproduction and Genetics, 2021, 38, 549-557.	1.2	17
88	Special focus issue on epigenomics and health disparities: foreword. Epigenomics, 2021, 13, 1673-1676.	1.0	0
89	Stress and spirituality in relation to HPA axis gene methylation among US Black women: results from the Black Women's Health Study and the Study on Stress, Spirituality and Health. Epigenomics, 2021, 13, 1711-1734.	1.0	3
90	Is your environment making you older? Molecular biomarkers and new approaches to investigate the influences of environmental chemicals through aging. Medicina Del Lavoro, 2021, 112, 8-14.	0.3	3

#	Article	IF	CITATIONS
91	Prospective Associations of Early Pregnancy Metal Mixtures with Mitochondria DNA Copy Number and Telomere Length in Maternal and Cord Blood. Environmental Health Perspectives, 2021, 129, 117007.	2.8	28
92	Mitochondrial DNA Copy Number Adaptation as a Biological Response Derived from an Earthquake at Intrauterine Stage. International Journal of Environmental Research and Public Health, 2021, 18, 11771.	1.2	3
93	Prenatal metal exposure, cord blood DNA methylation and persistence in childhood: an epigenome-wide association study of 12 metals. Clinical Epigenetics, 2021, 13, 208.	1.8	20
94	Epigenome-wide association study of serum urate reveals insights into urate co-regulation and the SLC2A9 locus. Nature Communications, 2021, 12, 7173.	5.8	8
95	Meta-analyses identify DNA methylation associated with kidney function and damage. Nature Communications, 2021, 12, 7174.	5.8	30
96	Maternal Phthalates Exposure and Blood Pressure during and after Pregnancy in the PROGRESS Study. Environmental Health Perspectives, 2021, 129, 127007.	2.8	11
97	Nonsteroidal Antiinflammatory Drugs Modify the Effect of Short-Term Air Pollution on Lung Function. American Journal of Respiratory and Critical Care Medicine, 2020, 201, 374-378.	2.5	12
98	Maternal Gestational Diabetes Mellitus and Newborn DNA Methylation: Findings From the Pregnancy and Childhood Epigenetics Consortium. Diabetes Care, 2020, 43, 98-105.	4.3	145
99	Age-related DNA hydroxymethylation is enriched for gene expression and immune system processes in human peripheral blood. Epigenetics, 2020, 15, 294-306.	1.3	8
100	Battle of epigenetic proportions: comparing Illumina's EPIC methylation microarrays and TruSeq targeted bisulfite sequencing. Epigenetics, 2020, 15, 174-182.	1.3	26
101	Physical activity, sedentary time and cardiometabolic health indicators among Mexican children. Clinical Obesity, 2020, 10, e12346.	1.1	3
102	Leukocyte Traits and Exposure to Ambient Particulate Matter Air Pollution in the Women's Health Initiative and Atherosclerosis Risk in Communities Study. Environmental Health Perspectives, 2020, 128, 17004.	2.8	17
103	Identifying critical windows of prenatal particulate matter (PM2.5) exposure and early childhood blood pressure. Environmental Research, 2020, 182, 109073.	3.7	36
104	Accelerating the Search for Interventions Aimed at Expanding the Health Span in Humans: The Role of Epidemiology. Journals of Gerontology - Series A Biological Sciences and Medical Sciences, 2020, 75, 77-86.	1.7	7
105	Association of Prenatal Acetaminophen Exposure Measured in Meconium With Risk of Attention-Deficit/Hyperactivity Disorder Mediated by Frontoparietal Network Brain Connectivity. JAMA Pediatrics, 2020, 174, 1073.	3.3	31
106	Association of Neutrophil to Lymphocyte Ratio With Pulmonary Function in a 30-Year Longitudinal Study of US Veterans. JAMA Network Open, 2020, 3, e2010350.	2.8	18
107	DNA methylation and body mass index from birth to adolescence: meta-analyses of epigenome-wide association studies. Genome Medicine, 2020, 12, 105.	3.6	41
108	Association of ambient PM2·5 exposure with maternal bone strength in pregnant women from Mexico City: a longitudinal cohort study. Lancet Planetary Health, The, 2020, 4, e530-e537.	5.1	12

#	Article	IF	CITATIONS
109	Age and mitochondrial DNA copy number influence the association between outdoor temperature and cognitive function. Environmental Epidemiology, 2020, 4, e0108.	1.4	8
110	Outdoor air pollution and cancer: An overview of the current evidence and public health recommendations. Ca-A Cancer Journal for Clinicians, 2020, 70, 460-479.	157.7	348
111	Methylome-wide association study of central adiposity implicates genes involved in immune and endocrine systems. Epigenomics, 2020, 12, 1483-1499.	1.0	6
112	Pregnancy-associated changes in cervical noncoding RNA. Epigenomics, 2020, 12, 1013-1025.	1.0	3
113	Prenatal lead exposure and cord blood DNA methylation in PROGRESS: an epigenome-wide association study. Environmental Epigenetics, 2020, 6, dvaa014.	0.9	14
114	Prenatal particulate air pollution and newborn telomere length: Effect modification by maternal antioxidant intakes and infant sex. Environmental Research, 2020, 187, 109707.	3.7	39
115	Glucose metabolism among obese and non-obese children of mothers with gestational diabetes. BMJ Open Diabetes Research and Care, 2020, 8, e000822.	1.2	4
116	Epigenome-wide meta-analysis of blood DNA methylation in newborns and children identifies numerous loci related to gestational age. Genome Medicine, 2020, 12, 25.	3.6	81
117	Phthalate Exposures and MicroRNA Expression in Uterine Fibroids: The FORGE Study. Epigenetics Insights, 2020, 13, 251686572090405.	0.6	32
118	Trends and Patterns of Phthalates and Phthalate Alternatives Exposure in Pregnant Women from Mexico City during 2007–2010. Environmental Science & Exposure in Pregnant Women from Mexico City during 2007–2010. Environmental Science & Exposure in Pregnant Women from Mexico City during 2007–2010. Environmental Science & Exposure in Pregnant Women from Mexico City during 2007–2010. Environmental Science & Exposure in Pregnant Women from Mexico City during 2007–2010. Environmental Science & Exposure in Pregnant Women from Mexico City during 2007–2010. Environmental Science & Exposure in Pregnant Women from Mexico City during 2007–2010. Environmental Science & Exposure in Pregnant Women from Mexico City during 2007—2010. Environmental Science & Exposure in Pregnant Women from Mexico City during 2007—2010. Environmental Science & Exposure in Pregnant Women from Mexico City during 2007—2010. Environmental Science & Exposure in Pregnant Women from Mexico City during 2007—2010. Environmental Science & Exposure in Pregnant Women from Mexico City during 2007—2010. Environmental Science & Exposure in Pregnant Women from Mexico City during 2007†in Pregnant Women from Mexico City du	4.6	33
119	Patterns of Weight Change One Year after Delivery Are Associated with Cardiometabolic Risk Factors at Six Years Postpartum in Mexican Women. Nutrients, 2020, 12, 170.	1.7	16
120	Modification of the effects of prenatal manganese exposure on child neurodevelopment by maternal anemia and iron deficiency. Pediatric Research, 2020, 88, 325-333.	1.1	15
121	Fine particulate matter exposure and lipid levels among children in Mexico city. Environmental Epidemiology, 2020, 4, e088.	1.4	14
122	Molecular and cellular mechanisms linking air pollution and bone damage. Environmental Research, 2020, 185, 109465.	3.7	47
123	Methylparaben in meconium and risk of maternal thyroid dysfunction, adverse birth outcomes, and Attention-Deficit Hyperactivity Disorder (ADHD). Environment International, 2020, 139, 105716.	4.8	42
124	Epigenetic Intergenerational Transmission: Mothers' Adverse Childhood Experiences and DNA Methylation. Journal of the American Academy of Child and Adolescent Psychiatry, 2020, 59, 900-901.	0.3	13
125	Associations between daily ambient temperature and sedentary time among children 4–6 years old in Mexico City. PLoS ONE, 2020, 15, e0241446.	1.1	4
126	Mitochondria and aging in older individuals: an analysis of DNA methylation age metrics, leukocyte telomere length, and mitochondrial DNA copy number in the VA normative aging study. Aging, 2020, 12, 2070-2083.	1.4	26

#	Article	IF	Citations
127	Biomarkers of aging and lung function in the normative aging study. Aging, 2020, 12, 11942-11966.	1.4	15
128	Blood DNA methylation sites predict death risk in a longitudinal study of 12, 300 individuals. Aging, 2020, 12, 14092-14124.	1.4	15
129	Accelerated epigenetic aging as a risk factor for chronic obstructive pulmonary disease and decreased lung function in two prospective cohort studies. Aging, 2020, 12, 16539-16554.	1.4	13
130	Quantification of the pace of biological aging in humans through a blood test, the DunedinPoAm DNA methylation algorithm. ELife, 2020, 9 , .	2.8	268
131	Blood Leukocyte DNA Methylation Predicts Risk of Future Myocardial Infarction and Coronary Heart Disease. Circulation, 2019, 140, 645-657.	1.6	151
132	Effects of Physical Exercise on Endothelial Function and DNA Methylation. International Journal of Environmental Research and Public Health, 2019, 16, 2530.	1.2	30
133	Smoking-Related DNA Methylation is Associated with DNA Methylation Phenotypic Age Acceleration: The Veterans Affairs Normative Aging Study. International Journal of Environmental Research and Public Health, 2019, 16, 2356.	1.2	22
134	The nasal methylome as a biomarker of asthma and airway inflammation in children. Nature Communications, 2019, 10, 3095.	5.8	129
135	Short-term ambient particle radioactivity level and renal function in older men: Insight from the Normative Aging Study. Environment International, 2019, 131, 105018.	4.8	13
136	Effect of particulate matter-bound metals exposure on prothrombotic biomarkers: A systematic review. Environmental Research, 2019, 177, 108573.	3.7	58
137	Joint Associations of Maternal Gestational Diabetes and Hypertensive Disorders of Pregnancy With Overweight in Offspring. Frontiers in Endocrinology, 2019, 10, 645.	1.5	15
138	The Human Exposome. Journal of the American College of Cardiology, 2019, 74, 1329-1331.	1.2	2
139	miRNA Profiles in Extracellular Vesicles From Serum Early in Pregnancies Complicated by Gestational Diabetes Mellitus. Journal of Clinical Endocrinology and Metabolism, 2019, 104, 5157-5169.	1.8	95
140	Comparison of smoking-related DNA methylation between newborns from prenatal exposure and adults from personal smoking. Epigenomics, 2019, 11, 1487-1500.	1.0	64
141	Supraphysiological Concentrations of Bisphenol A Alter the Expression of Extracellular Vesicle-Enriched miRNAs From Human Primary Granulosa Cells. Toxicological Sciences, 2019, 169, 5-13.	1.4	18
142	Methylome-wide association study provides evidence of particulate matter air pollution-associated DNA methylation. Environment International, 2019, 132, 104723.	4.8	58
143	Hypertensive Disorders of Pregnancy and DNA Methylation in Newborns. Hypertension, 2019, 74, 375-383.	1.3	73
144	Prenatal Particulate Air Pollution and DNA Methylation in Newborns: An Epigenome-Wide Meta-Analysis. Environmental Health Perspectives, 2019, 127, 57012.	2.8	111

#	Article	IF	Citations
145	An integrative cross-omics analysis of DNA methylation sites of glucose and insulin homeostasis. Nature Communications, 2019, 10, 2581.	5.8	62
146	Prenatal salivary sex hormone levels and birth-weight-for-gestational age. Journal of Perinatology, 2019, 39, 941-948.	0.9	11
147	Ambient particulate air pollution and circulating C-reactive protein level: A systematic review and meta-analysis. International Journal of Hygiene and Environmental Health, 2019, 222, 756-764.	2.1	70
148	DNA methylation GrimAge strongly predicts lifespan and healthspan. Aging, 2019, 11, 303-327.	1.4	1,128
149	Comparative validation of an epigenetic mortality risk score with three aging biomarkers for predicting mortality risks among older adult males. International Journal of Epidemiology, 2019, 48, 1958-1971.	0.9	25
150	Meta-analysis of epigenome-wide association studies in neonates reveals widespread differential DNA methylation associated with birthweight. Nature Communications, 2019, 10, 1893.	5.8	140
151	Epigenome-wide association study reveals methylation pathways associated with childhood allergic sensitization. Epigenetics, 2019, 14, 445-466.	1.3	43
152	Prenatal manganese and cord blood mitochondrial DNA copy number: Effect modification by maternal anemic status. Environment International, 2019, 126, 484-493.	4.8	28
153	Impacts of air pollution, temperature, and relative humidity on leukocyte distribution: An epigenetic perspective. Environment International, 2019, 126, 395-405.	4.8	52
154	Association between prenatal particulate air pollution exposure and telomere length in cord blood: Effect modification by fetal sex. Environmental Research, 2019, 172, 495-501.	3.7	51
155	Epigenetic age acceleration is associated with allergy and asthma in children in Project Viva. Journal of Allergy and Clinical Immunology, 2019, 143, 2263-2270.e14.	1.5	43
156	Cognition level and change in cognition during adolescence are associated with cognition in midlife. Annals of Epidemiology, 2019, 35, 48-52.e2.	0.9	4
157	Prenatal maternal antidepressants, anxiety, and depression and offspring DNA methylation: epigenome-wide associations at birth and persistence into early childhood. Clinical Epigenetics, 2019, 11, 56.	1.8	46
158	Prenatal Metal Concentrations and Childhood Cardiometabolic Risk Using Bayesian Kernel Machine Regression to Assess Mixture and Interaction Effects. Epidemiology, 2019, 30, 263-273.	1.2	62
159	Altered cord blood mitochondrial DNA content and pregnancy lead exposure in the PROGRESS cohort. Environment International, 2019, 125, 437-444.	4.8	27
160	ENVIRONMENTAL EPIGENETICS AND AGING. Innovation in Aging, 2019, 3, S735-S735.	0.0	0
161	Association of Prenatal and Perinatal Exposures to Particulate Matter With Changes in Hemoglobin A _{1c} Levels in Children Aged 4 to 6 Years. JAMA Network Open, 2019, 2, e1917643.	2.8	18
162	Length of gestation and birth weight are associated with indices of combined kidney biomarkers in early childhood. PLoS ONE, 2019, 14, e0227219.	1.1	0

#	Article	IF	Citations
163	DNA methylation aging clocks: challenges and recommendations. Genome Biology, 2019, 20, 249.	3.8	552
164	Fried and Baccarelli Comment. American Journal of Public Health, 2019, 109, 1188-1188.	1.5	1
165	Phthalates exposure and uterine fibroid burden among women undergoing surgical treatment for fibroids: a preliminary study. Fertility and Sterility, 2019, 111, 112-121.	0.5	51
166	Maternal corticotropin-releasing hormone is associated with LEP DNA methylation at birth and in childhood: an epigenome-wide study in Project Viva. International Journal of Obesity, 2019, 43, 1244-1255.	1.6	6
167	Monitoring of prenatal exposure to organic and inorganic contaminants using meconium from an Eastern Canada cohort. Environmental Research, 2019, 171, 44-51.	3.7	17
168	Maternal Phthalate and Personal Care Products Exposure Alters Extracellular Placental miRNA Profile in Twin Pregnancies. Reproductive Sciences, 2019, 26, 289-294.	1.1	28
169	Socioeconomic position, lifestyle habits and biomarkers of epigenetic aging: a multi-cohort analysis. Aging, 2019, 11, 2045-2070.	1.4	137
170	DNA methylation-based estimator of telomere length. Aging, 2019, 11, 5895-5923.	1.4	198
171	New epigenomic and genomic frontiers in personalized medicine: direct customer testing, are we ready?. Epidemiologia E Prevenzione, 2019, 43, 295-299.	1.1	0
172	miRNA-Processing Gene Methylation and Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2018, 27, 550-557.	1.1	19
173	Association of Methylation Signals With Incident Coronary Heart Disease in an Epigenome-Wide Assessment of Circulating Tumor Necrosis Factor α. JAMA Cardiology, 2018, 3, 463.	3.0	33
174	GWAS of epigenetic aging rates in blood reveals a critical role for TERT. Nature Communications, 2018, 9, 387.	5.8	151
175	Placental IncRNA Expression Is Associated With Prenatal Phthalate Exposure. Toxicological Sciences, 2018, 163, 116-122.	1.4	31
176	Prenatal particulate matter exposure and mitochondrial dysfunction at the maternal-fetal interface: Effect modification by maternal lifetime trauma and child sex. Environment International, 2018, 112, 49-58.	4.8	70
177	Meta-analysis of epigenome-wide association studies of cognitive abilities. Molecular Psychiatry, 2018, 23, 2133-2144.	4.1	68
178	Acute particulate matter affects cardiovascular autonomic modulation and IFN- \hat{I}^3 methylation in healthy volunteers. Environmental Research, 2018, 161, 97-103.	3.7	38
179	Prenatal fine particulate exposure associated with reduced childhood lung function and nasal epithelia GSTP1 hypermethylation: Sex-specific effects. Respiratory Research, 2018, 19, 76.	1.4	32
180	Epigenome-wide association study of total serum immunoglobulin E in children: a life course approach. Clinical Epigenetics, 2018, 10, 55.	1.8	36

#	Article	IF	Citations
181	Cohort Profile: Pregnancy And Childhood Epigenetics (PACE) Consortium. International Journal of Epidemiology, 2018, 47, 22-23u.	0.9	105
182	The Inflammatory Potential of Dietary Manganese in a Cohort of Elderly Men. Biological Trace Element Research, 2018, 183, 49-57.	1.9	19
183	Aberrant promoter methylation in genes related to hematopoietic malignancy in workers exposed to a VOC mixture. Toxicology and Applied Pharmacology, 2018, 339, 65-72.	1.3	30
184	Prenatal arsenic exposure, child marriage, and pregnancy weight gain: Associations with preterm birth in Bangladesh. Environment International, 2018, 112, 23-32.	4.8	36
185	Maternal alcohol consumption and offspring DNA methylation: findings from six general population-based birth cohorts. Epigenomics, 2018, 10, 27-42.	1.0	58
186	Urinary concentrations of biomarkers of phthalates and phthalate alternatives and IVF outcomes. Environment International, 2018, 111, 23-31.	4.8	85
187	Accelerated DNA methylation age and the use of antihypertensive medication among older adults. Aging, 2018, 10, 3210-3228.	1.4	21
188	An epigenetic biomarker of aging for lifespan and healthspan. Aging, 2018, 10, 573-591.	1.4	1,552
189	Analysis of repeated leukocyte DNA methylation assessments reveals persistent epigenetic alterations after an incident myocardial infarction. Clinical Epigenetics, 2018, 10, 161.	1.8	20
190	Maternal antenatal stress has little impact on child sleep: results from a prebirth cohort in Mexico City. Sleep Health, 2018, 4, 397-404.	1.3	8
191	Exposure to childhood abuse is associated with human sperm DNA methylation. Translational Psychiatry, 2018, 8, 194.	2.4	56
192	Residential Proximity to Major Roadways at Birth, DNA Methylation at Birth and Midchildhood, and Childhood Cognitive Test Scores: Project Viva (Massachusetts, USA). Environmental Health Perspectives, 2018, 126, 97006.	2.8	15
193	Maternal exposure to gasoline and exhaust increases the risk of childhood leukaemia in offspring $\hat{a}\in$ a prospective study in the Norwegian Mother and Child Cohort Study. British Journal of Cancer, 2018, 119, 1028-1035.	2.9	7
194	Extracellular vesicle-enriched microRNAs interact in the association between long-term particulate matter and blood pressure in elderly men. Environmental Research, 2018, 167, 640-649.	3.7	43
195	Diurnal Cortisol Concentrations and Growth Indexes of 12- to 48-Month-Old Children From Mexico City. Journal of Clinical Endocrinology and Metabolism, 2018, 103, 3386-3393.	1.8	0
196	Long noncoding RNA expression in the cervix mid-pregnancy is associated with the length of gestation at delivery. Epigenetics, 2018, 13, 742-750.	1.3	14
197	DNA Methylation Signatures of Depressive Symptoms in Middle-aged and Elderly Persons. JAMA Psychiatry, 2018, 75, 949.	6.0	78
198	Cumulative lifetime maternal stress and epigenome-wide placental DNA methylation in the PRISM cohort. Epigenetics, 2018, 13, 665-681.	1.3	37

#	Article	IF	Citations
199	Urinary concentrations of phthalate metabolites, bisphenols and personal care product chemical biomarkers in pregnant women in Israel. Environment International, 2018, 116, 319-325.	4.8	53
200	Regulation of birthweight by placenta-derived miRNAs: evidence from an arsenic-exposed birth cohort in Bangladesh. Epigenetics, 2018, 13, 573-590.	1.3	28
201	Prenatal lead exposure modifies the effect of shorter gestation on increased blood pressure in children. Environment International, 2018, 120, 464-471.	4.8	30
202	Promoter methylation of <i>PGC1A</i> and <i>PGC1B</i> predicts cancer incidence in a veteran cohort. Epigenomics, 2018, 10, 733-743.	1.0	12
203	DNA Methylation of Telomere-Related Genes and Cancer Risk. Cancer Prevention Research, 2018, 11, 511-522.	0.7	12
204	Epigenetic clock for skin and blood cells applied to Hutchinson Gilford Progeria Syndrome and ex vivo studies. Aging, 2018, 10, 1758-1775.	1.4	406
205	An epigenome-wide association study of total serum IgE in Hispanic children. Journal of Allergy and Clinical Immunology, 2017, 140, 571-577.	1.5	53
206	Bacterial and cytokine mixtures predict the length of gestation and are associated with miRNA expression in the cervix. Epigenomics, 2017, 9, 33-45.	1.0	11
207	Placental mitochondrial DNA and CYP1A1 gene methylation as molecular signatures for tobacco smoke exposure in pregnant women and the relevance for birth weight. Journal of Translational Medicine, 2017, 15, 5.	1.8	56
208	Air pollution and <i>in utero</i> programming of poor fetal growth. Epigenomics, 2017, 9, 213-216.	1.0	18
209	Epigenome-wide cross-tissue predictive modeling and comparison of cord blood and placental methylation in a birth cohort. Epigenomics, 2017, 9, 231-240.	1.0	23
210	Extracellular microRNAs in follicular fluid and their potential association with oocyte fertilization and embryo quality: an exploratory study. Journal of Assisted Reproduction and Genetics, 2017, 34, 525-533.	1.2	76
211	Reply to Lucock et al.: Significance of interpretation and misinterpretation of a small mechanistic study. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E3880-E3881.	3.3	1
212	Empirical comparison of reduced representation bisulfite sequencing and Infinium BeadChip reproducibility and coverage of DNA methylation in humans. Npj Genomic Medicine, 2017, 2, 13.	1.7	26
213	Maternal Lifetime Stress and Prenatal Psychological Functioning and Decreased Placental Mitochondrial DNA Copy Number in the PRISM Study. American Journal of Epidemiology, 2017, 186, 1227-1236.	1.6	65
214	B-vitamin Supplementation Mitigates Effects of Fine Particles on Cardiac Autonomic Dysfunction and Inflammation: A Pilot Human Intervention Trial. Scientific Reports, 2017, 7, 45322.	1.6	31
215	Epigenetic Pathways in Human Disease: The Impact of DNA Methylation on Stress-Related Pathogenesis and Current Challenges in Biomarker Development. EBioMedicine, 2017, 18, 327-350.	2.7	113
216	Persistent DNA methylation changes associated with prenatal mercury exposure and cognitive performance during childhood. Scientific Reports, 2017, 7, 288.	1.6	95

#	Article	IF	Citations
217	Short-term effects of air temperature and mitochondrial DNA lesions within an older population. Environment International, 2017, 103, 23-29.	4.8	3
218	Ambient temperature and cardiovascular biomarkers in a repeated-measure study in healthy adults: A novel biomarker index approach. Environmental Research, 2017, 156, 231-238.	3.7	21
219	B vitamins attenuate the epigenetic effects of ambient fine particles in a pilot human intervention trial. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 3503-3508.	3.3	121
220	Traffic-derived particulate matter exposure and histone H3 modification: A repeated measures study. Environmental Research, 2017, 153, 112-119.	3.7	52
221	Prenatal lead exposure and fetal growth: Smaller infants have heightened susceptibility. Environment International, 2017, 99, 228-233.	4.8	44
222	Effects of environmental noise exposure on DNA methylation in the brain and metabolic health. Environmental Research, 2017, 153, 73-82.	3.7	39
223	Differential DNA methylation and PM _{2.5} species in a 450K epigenome-wide association study. Epigenetics, 2017, 12, 139-148.	1.3	52
224	High pesticide exposure events and <scp>DNA</scp> methylation among pesticide applicators in the agricultural health study. Environmental and Molecular Mutagenesis, 2017, 58, 19-29.	0.9	48
225	Bisphenol-A exposure and gene expression in human luteinized membrana granulosa cells in vitro. Human Reproduction, 2017, 32, 409-417.	0.4	19
226	Association between preconception maternal beverage intake and inÂvitro fertilization outcomes. Fertility and Sterility, 2017, 108, 1026-1033.	0.5	30
227	Testing for the indirect effect under the null for genomeâ€wide mediation analyses. Genetic Epidemiology, 2017, 41, 824-833.	0.6	60
228	Promoter methylation status in genes related with inflammation, nitrosative stress and xenobiotic metabolism in low-level benzene exposure: Searching for biomarkers of oncogenesis. Food and Chemical Toxicology, 2017, 109, 669-676.	1.8	23
229	Maternal BMI at the start of pregnancy and offspring epigenome-wide DNA methylation: findings from the pregnancy and childhood epigenetics (PACE) consortium. Human Molecular Genetics, 2017, 26, 4067-4085.	1.4	211
230	Prenatal exposure to PM 2.5 and birth weight: A pooled analysis from three North American longitudinal pregnancy cohort studies. Environment International, 2017, 107, 173-180.	4.8	36
231	Prenatal particulate air pollution exposure and body composition in urban preschool children: Examining sensitive windows and sex-specific associations. Environmental Research, 2017, 158, 798-805.	3.7	56
232	Second trimester extracellular microRNAs in maternal blood and fetal growth: An exploratory study. Epigenetics, 2017, 12, 804-810.	1.3	70
233	Investigating causal relation between prenatal arsenic exposure and birthweight: Are smaller infants more susceptible?. Environment International, 2017, 108, 32-40.	4.8	34
234	DNA Methylation Analysis Identifies Loci for Blood Pressure Regulation. American Journal of Human Genetics, 2017, 101, 888-902.	2.6	154

#	Article	IF	Citations
235	A longitudinal study of DNA methylation as a potential mediator of age-related diabetes risk. GeroScience, 2017, 39, 475-489.	2.1	62
236	Association of air particulate pollution with bone loss over time and bone fracture risk: analysis of data from two independent studies. Lancet Planetary Health, The, 2017, 1, e337-e347.	5.1	96
237	The effect of morphine upon DNA methylation in ten regions of the rat brain. Epigenetics, 2017, 12, 1038-1047.	1.3	36
238	Histone 3 modifications and blood pressure in the Beijing Truck Driver Air Pollution Study. Biomarkers, 2017, 22, 584-593.	0.9	16
239	Regulating role of fetal thyroid hormones on placental mitochondrial DNA methylation: epidemiological evidence from the ENVIRONAGE birth cohort study. Clinical Epigenetics, 2017, 9, 66.	1.8	21
240	Identifying sensitive windows for prenatal particulate air pollution exposure and mitochondrial DNA content in cord blood. Environment International, 2017, 98, 198-203.	4.8	56
241	The association of lead exposure during pregnancy and childhood anthropometry in the Mexican PROGRESS cohort. Environmental Research, 2017, 152, 226-232.	3.7	50
242	Increased methylation of repetitive elements and DNA repair genes is associated with higher DNA oxidation in children in an urbanized, industrial environment. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2017, 813, 27-36.	0.9	41
243	Whole blood microRNA markers are associated with acute respiratory distress syndrome. Intensive Care Medicine Experimental, 2017, 5, 38.	0.9	44
244	Pesticide use and LINE-1 methylation among male private pesticide applicators in the Agricultural Health Study. Environmental Epigenetics, 2017, 3, dvx005.	0.9	16
245	0066â \in Maternal occupational exposure to benzene increases the risk of childhood leukaemia in offspring â \in " a prospective study in the norwegian mother and child cohort study. , 2017, , .		0
246	Cord blood DNA methylation and adiposity measures in early and mid-childhood. Clinical Epigenetics, 2017, 9, 86.	1.8	18
247	Exposure to Low Levels of Lead <i>in Utero</i> and Umbilical Cord Blood DNA Methylation in Project Viva: An Epigenome-Wide Association Study. Environmental Health Perspectives, 2017, 125, 087019.	2.8	73
248	Prenatal Exposure to Mercury: Associations with Global DNA Methylation and Hydroxymethylation in Cord Blood and in Childhood. Environmental Health Perspectives, 2017, 125, 087022.	2.8	57
249	Epigenetic clock analysis of diet, exercise, education, and lifestyle factors. Aging, 2017, 9, 419-446.	1.4	521
250	Telomere Length, Long-Term Black Carbon Exposure, and Cognitive Function in a Cohort of Older Men: The VA Normative Aging Study. Environmental Health Perspectives, 2017, 125, 76-81.	2.8	36
251	Leukocyte telomere length and renal cell carcinoma survival in two studies. British Journal of Cancer, 2017, 117, 752-755.	2.9	17
252	Changes in DNA Methylation in Mouse Lungs after a Single Intra-Tracheal Administration of Nanomaterials. PLoS ONE, 2017, 12, e0169886.	1.1	47

#	Article	IF	Citations
253	Blood pressure and expression of microRNAs in whole blood. PLoS ONE, 2017, 12, e0173550.	1.1	12
254	Extracellular vesicles: roles in gamete maturation, fertilization and embryo implantation. Human Reproduction Update, 2016, 22, dmv055.	5.2	248
255	Particulate Air Pollution, Exceptional Aging, and Rates of Centenarians: A Nationwide Analysis of the United States, 1980–2010. Environmental Health Perspectives, 2016, 124, 1744-1750.	2.8	18
256	Birth weight-for-gestational age is associated with DNA methylation at birth and in childhood. Clinical Epigenetics, 2016, 8, 118.	1.8	61
257	DNA methylation signatures of chronic low-grade inflammation are associated with complex diseases. Genome Biology, 2016, 17, 255.	3.8	251
258	Countervailing effects of income, air pollution, smoking, and obesity on aging and life expectancy: population-based study of U.S. Counties. Environmental Health, 2016, 15, 86.	1.7	17
259	DNA Methylation in Newborns and Maternal Smoking in Pregnancy: Genome-wide Consortium Meta-analysis. American Journal of Human Genetics, 2016, 98, 680-696.	2.6	717
260	Short-term exposure to high ambient air pollution increases airway inflammation and respiratory symptoms in chronic obstructive pulmonary disease patients in Beijing, China. Environment International, 2016, 94, 76-82.	4.8	131
261	Chemical constituents and sources of ambient particulate air pollution and biomarkers of endothelial function in a panel of healthy adults in Beijing, China. Science of the Total Environment, 2016, 560-561, 141-149.	3.9	48
262	Whole blood microRNAs as a prognostic classifier for acute respiratory distress syndrome 28-day mortality. Intensive Care Medicine, 2016, 42, 1824-1825.	3.9	7
263	APOE ε4 allele modifies the association of lead exposure with age-related cognitive decline in older individuals. Environmental Research, 2016, 151, 101-105.	3.7	10
264	An epigenetic clock for gestational age at birth based on blood methylation data. Genome Biology, 2016, 17, 206.	3.8	193
265	Endocrine Disruptors: A Potential Risk Factor for Gestational Diabetes Mellitus. American Journal of Perinatology, 2016, 33, 1313-1318.	0.6	81
266	Detection of long non-coding RNAs in human breastmilk extracellular vesicles: Implications for early child development. Epigenetics, 2016, 11, 721-729.	1.3	83
267	Long-term ambient particle exposures and blood DNA methylation age: findings from the VA normative aging study. Environmental Epigenetics, 2016, 2, dvw006.	0.9	68
268	Distributional changes in gene-specific methylation associated with temperature. Environmental Research, 2016, 150, 38-46.	3.7	14
269	Childhood abuse, promoter methylation of leukocyte <i>NR3C1</i> and the potential modifying effect of emotional support. Epigenomics, 2016, 8, 1507-1517.	1.0	48
270	CpGFilter: model-based CpG probe filtering with replicates for epigenome-wide association studies. Bioinformatics, 2016, 32, 469-471.	1.8	27

#	Article	IF	Citations
271	Maternal gut and fetal brain connection: Increased anxiety and reduced social interactions in Wistar rat offspring following peri-conceptional antibiotic exposure. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2016, 71, 76-82.	2.5	82
272	Environmental Health and Long Non-coding RNAs. Current Environmental Health Reports, 2016, 3, 178-187.	3.2	82
273	Prospective changes in global DNA methylation and cancer incidence and mortality. British Journal of Cancer, 2016, 115, 465-472.	2.9	41
274	Traffic-Related Air Pollution, Blood Pressure, and Adaptive Response of Mitochondrial Abundance. Circulation, 2016, 133, 378-387.	1.6	77
275	Effects of particulate matter exposure on multiple sclerosis hospital admission in Lombardy region, Italy. Environmental Research, 2016, 145, 68-73.	3.7	68
276	Epigenetics: linking social and environmental exposures to preterm birth. Pediatric Research, 2016, 79, 136-140.	1.1	49
277	Epigenetics—a potential mediator between air pollution and preterm birth. Environmental Epigenetics, 2016, 2, dvv008.	0.9	27
278	Psychological factors and DNA methylation of genes related to immune/inflammatory system markers: the VA Normative Aging Study. BMJ Open, 2016, 6, e009790.	0.8	45
279	Blood Epigenetic Age may Predict Cancer Incidence and Mortality. EBioMedicine, 2016, 5, 68-73.	2.7	162
280	Long-term exposure to black carbon, cognition and single nucleotide polymorphisms in microRNA processing genes in older men. Environment International, 2016, 88, 86-93.	4.8	21
281	The Role of DNA Methylation in Cardiovascular Risk and Disease. Circulation Research, 2016, 118, 119-131.	2.0	167
282	Fetal growth restriction and methylation of growth-related genes in the placenta. Epigenomics, 2016, 8, 33-42.	1.0	50
283	DNA methylation-based measures of biological age: meta-analysis predicting time to death. Aging, 2016, 8, 1844-1865.	1.4	786
284	Long-term exposure to air pollution is associated with biological aging. Oncotarget, 2016, 7, 74510-74525.	0.8	126
285	Ambient particulate matter and microRNAs in extracellular vesicles: a pilot study of older individuals. Particle and Fibre Toxicology, 2015, 13, 13.	2.8	96
286	Characterization of genome-wide H3K27ac profiles reveals a distinct PM2.5-associated histone modification signature. Environmental Health, 2015, 14, 65.	1.7	37
287	A distinct and replicable variant of the squamous cell carcinoma gene inositol polyphosphateâ€5â€phosphatase modifies the susceptibility of arsenicâ€associated skin lesions in Bangladesh. Cancer, 2015, 121, 2222-2229.	2.0	10
288	Maternal Glucose during Pregnancy and after Delivery in Women with Gestational Diabetes Mellitus on Overweight Status of Their Children. BioMed Research International, 2015, 2015, 1-9.	0.9	17

#	Article	IF	Citations
289	Expert position paper on air pollution and cardiovascular disease. European Heart Journal, 2015, 36, 83-93.	1.0	646
290	Platelet mitochondrial DNA methylation: a potential new marker of cardiovascular disease. Clinical Epigenetics, 2015, 7, 44.	1.8	132
291	In <i>utero</i> arsenic exposure and epigenome-wide associations in placenta, umbilical artery, and human umbilical vein endothelial cells. Epigenetics, 2015, 10, 1054-1063.	1.3	56
292	Epigenetic effects of low perinatal doses of flame retardant BDE-47 on mitochondrial and nuclear genes in rat offspring. Toxicology, 2015, 328, 152-159.	2.0	44
293	Cohort Profile: Project Viva. International Journal of Epidemiology, 2015, 44, 37-48.	0.9	275
294	CYP2E1 epigenetic regulation in chronic, low-level toluene exposure: Relationship with oxidative stress and smoking habit. Toxicology and Applied Pharmacology, 2015, 286, 207-215.	1.3	34
295	Using High-Resolution Satellite Aerosol Optical Depth To Estimate Daily PM _{2.5} Geographical Distribution in Mexico City. Environmental Science & Eamp; Technology, 2015, 49, 8576-8584.	4.6	165
296	Endotoxin and \hat{l}^2 -1,3- <scp>d</scp> -Glucan in Concentrated Ambient Particles Induce Rapid Increase in Blood Pressure in Controlled Human Exposures. Hypertension, 2015, 66, 509-516.	1.3	37
297	GDM Women's Pre-Pregnancy Overweight/Obesity and Gestational Weight Gain on Offspring Overweight Status. PLoS ONE, 2015, 10, e0129536.	1.1	50
298	Blood Telomere Length Attrition and Cancer Development in the Normative Aging Study Cohort. EBioMedicine, 2015, 2, 591-596.	2.7	62
299	Effects of particulate matter exposure on blood 5-hydroxymethylation: results from the Beijing truck driver air pollution study. Epigenetics, 2015, 10, 633-642.	1.3	63
300	Cardiac Autonomic Dysfunction: Particulate Air Pollution Effects Are Modulated by Epigenetic Immunoregulation of <i>Tollâ€like Receptor 2</i> and Dietary Flavonoid Intake. Journal of the American Heart Association, 2015, 4, e001423.	1.6	40
301	DNA methylation age of blood predicts all-cause mortality in later life. Genome Biology, 2015, 16, 25.	3.8	928
302	Altered miRNA expression in the cervix during pregnancy associated with lead and mercury exposure. Epigenomics, 2015, 7, 885-896.	1.0	53
303	Offspring DNA methylation of the aryl-hydrocarbon receptor repressor gene is associated with maternal BMI, gestational age, and birth weight. Epigenetics, 2015, 10, 913-921.	1.3	65
304	Longitudinal Study of DNA Methylation of Inflammatory Genes and Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1531-1538.	1.1	26
305	Alternate methods of nasal epithelial cell sampling for airway genomic studies. Journal of Allergy and Clinical Immunology, 2015, 136, 1120-1123.e4.	1.5	16
306	Pesticide Use and Relative Leukocyte Telomere Length in the Agricultural Health Study. PLoS ONE, 2015, 10, e0133382.	1.1	42

#	Article	IF	Citations
307	Association between length of gestation and cervical DNA methylation of <i>PTGER2 </i> li>and LINE 1-HS. Epigenetics, 2014, 9, 1083-1091.	1.3	29
308	Effect of prenatal arsenic exposure on DNA methylation and leukocyte subpopulations in cord blood. Epigenetics, 2014, 9, 774-782.	1.3	140
309	Air pollution exposure and lung function in highly exposed subjects in Beijing, China: a repeated-measure study. Particle and Fibre Toxicology, 2014, 11, 51.	2.8	76
310	Short-term airborne particulate matter exposure alters the epigenetic landscape of human genes associated with the mitogen-activated protein kinase network: a cross-sectional study. Environmental Health, 2014, 13, 94.	1.7	55
311	Blood methylomics in response to arsenic exposure in a low-exposed US population. Journal of Exposure Science and Environmental Epidemiology, 2014, 24, 145-149.	1.8	28
312	Air pollution and gene-specific methylation in the Normative Aging Study. Epigenetics, 2014, 9, 448-458.	1.3	159
313	Effects of shortâ€term exposure to inhalable particulate matter on DNA methylation of tandem repeats. Environmental and Molecular Mutagenesis, 2014, 55, 322-335.	0.9	59
314	Maternal OGTT Glucose Levels at 26–30 Gestational Weeks with Offspring Growth and Development in Early Infancy. BioMed Research International, 2014, 2014, 1-11.	0.9	13
315	Pessimistic orientation in relation to telomere length in older men: The VA Normative Aging Study. Psychoneuroendocrinology, 2014, 42, 68-76.	1.3	26
316	Prenatal exposure to mixtures of xenoestrogens and repetitive element DNA methylation changes in human placenta. Environment International, 2014, 71, 81-87.	4.8	52
317	Influence of multiple APOE genetic variants on cognitive function in a cohort of older men – results from the Normative Aging Study. BMC Psychiatry, 2014, 14, 223.	1.1	20
318	0288†Gene-specific DNA methylation as a valuable tool for risk assessment: the case of occupational exposure to different VOC's in Mexican workers0288†Gene-specific DNA methylation as a valuable tool for risk assessment: the case of occupational exposure to different VOC's in Mexican workers. Occupational and Environmental Medicine, 2014, 71, A36,2-A36.	1.3	3
319	A Novel Genetic Score Approach Using Instruments to Investigate Interactions between Pathways and Environment: Application to Air Pollution. PLoS ONE, 2014, 9, e96000.	1.1	30
320	Maternal Prepregnancy Body Mass Index and Gestational Weight Gain on Pregnancy Outcomes. PLoS ONE, 2013, 8, e82310.	1.1	266
321	Air Pollution and Markers of Coagulation, Inflammation, and Endothelial Function. Epidemiology, 2012, 23, 332-340.	1.2	259
322	Environmental exposures, epigenetics and cardiovascular disease. Current Opinion in Clinical Nutrition and Metabolic Care, 2012, 15, 323-329.	1.3	90
323	Nasal cell DNA methylation, inflammation, lung function and wheezing in children with asthma. Epigenomics, 2012, 4, 91-100.	1.0	66
324	Triggers of MI for the individual and in the community. Lancet, The, 2011, 377, 694-696.	6.3	11

#	Article	IF	CITATIONS
325	Extracellular histones affect blood coagulation in subjects exposed to metal-rich air particles. Occupational and Environmental Medicine, 2011, 68, A93-A93.	1.3	O
326	Effects of particulate air pollution on blood pressure in a highly exposed population in Beijing, China: a repeated-measure study. Environmental Health, 2011, 10, 108.	1.7	76
327	Effects of metal-rich air particles on DNA methylation and on coagulation function among foundry workers in Italy. Occupational and Environmental Medicine, 2011, 68, A52-A53.	1.3	0
328	THE POTENTIAL ROLE OF EPIGENETICS IN MEDIATING AIR POLLUTION EFFECTS. ISEE Conference Abstracts, 2011, 2011, .	0.0	0
329	Prognostic Significance of Telomere Length in Chronic Lymphocytic Leukemia Patients in Early Stage Disease,. Blood, 2011, 118, 3890-3890.	0.6	7
330	Cardiovascular Epigenetics. Circulation: Cardiovascular Genetics, 2010, 3, 567-573.	5.1	186
331	Repetitive element DNA methylation and circulating endothelial and inflammation markers in the VA normative aging study. Epigenetics, 2010, 5, 222-228.	1.3	106
332	Rapid DNA Methylation Changes after Exposure to Traffic Particles. American Journal of Respiratory and Critical Care Medicine, 2009, 179, 572-578.	2.5	608
333	Effects of Particulate Matter on Genomic DNA Methylation Content and <i>iNOS</i> Promoter Methylation. Environmental Health Perspectives, 2009, 117, 217-222.	2.8	310
334	Exposure to Particulate Air Pollution and Risk of Deep Vein Thrombosis. Archives of Internal Medicine, 2008, 168, 920.	4.3	184
335	Cardiac Autonomic Dysfunction. Circulation, 2008, 117, 1802-1809.	1.6	112
336	Neonatal Thyroid Function in Seveso 25 Years after Maternal Exposure to Dioxin. PLoS Medicine, 2008, 5, e161.	3.9	106
337	Air Pollution, Smoking, and Plasma Homocysteine. Environmental Health Perspectives, 2007, 115, 176-181.	2.8	64
338	Reply to the letter to the Editor:Helicobacter pylori infection and MBL2 haplotypes: Lack of association or lack of evidence?. International Journal of Cancer, 2007, 120, 2750-2750.	2.3	0
339	Breast milk-derived extracellular vesicle miRNAs are associated with maternal asthma and atopy. Epigenomics, 0, , .	1.0	5