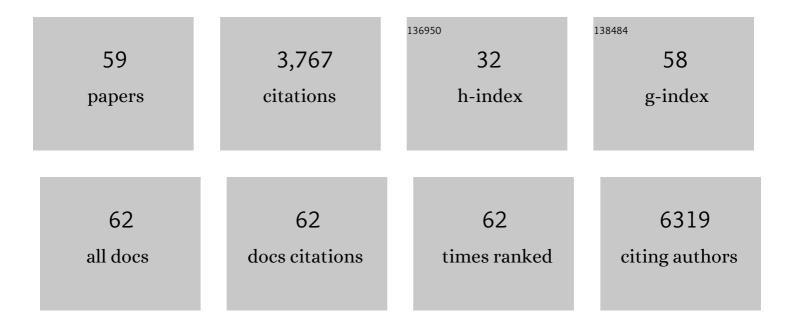
List of Publications by Year in descending order

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KADEN HIJEN

#	Article	IF	CITATIONS
1	DNA Methylation in Newborns and Maternal Smoking in Pregnancy: Genome-wide Consortium Meta-analysis. American Journal of Human Genetics, 2016, 98, 680-696.	6.2	717
2	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.	2.9	211
3	Sex differences in DNA methylation assessed by 450ÂK BeadChip in newborns. BMC Genomics, 2015, 16, 911.	2.8	155
4	Meta-analysis of epigenome-wide association studies in neonates reveals widespread differential DNA methylation associated with birthweight. Nature Communications, 2019, 10, 1893.	12.8	140
5	Genotype–activity relationship for Mn-superoxide dismutase, glutathione peroxidase 1 and catalase in humans. Pharmacogenetics and Genomics, 2006, 16, 279-286.	1.5	133
6	Association of prenatal urinary phthalate metabolite concentrations and childhood BMI and obesity. Pediatric Research, 2017, 82, 405-415.	2.3	129
7	Genome-wide methylation data mirror ancestry information. Epigenetics and Chromatin, 2017, 10, 1.	3.9	120
8	Paraoxonase Polymorphisms, Haplotypes, and Enzyme Activity in Latino Mothersand Newborns. Environmental Health Perspectives, 2006, 114, 985-991.	6.0	113
9	Organophosphate pesticide levels in blood and urine of women and newborns living in an agricultural community. Environmental Research, 2012, 117, 8-16.	7.5	110
10	PON1 and Neurodevelopment in Children from the CHAMACOS Study Exposed to Organophosphate Pesticides <i>in Utero</i> . Environmental Health Perspectives, 2010, 118, 1775-1781.	6.0	107
11	Cohort Profile: Pregnancy And Childhood Epigenetics (PACE) Consortium. International Journal of Epidemiology, 2018, 47, 22-23u.	1.9	105
12	Residential Traffic and Children's Respiratory Health. Environmental Health Perspectives, 2008, 116, 1274-1279.	6.0	91
13	Association of Organophosphate Pesticide Exposure and Paraoxonase with Birth Outcome in Mexican-American Women. PLoS ONE, 2011, 6, e23923.	2.5	86
14	Epigenome-wide meta-analysis of blood DNA methylation in newborns and children identifies numerous loci related to gestational age. Genome Medicine, 2020, 12, 25.	8.2	81
15	<i>In Utero</i> and Childhood Polybrominated Diphenyl Ether Exposures and Body Mass at Age 7 Years: The CHAMACOS Study. Environmental Health Perspectives, 2015, 123, 636-642.	6.0	79
16	Effects of age, sex, and persistent organic pollutants on DNA methylation in children. Environmental and Molecular Mutagenesis, 2014, 55, 209-222.	2.2	74
17	Prenatal phthalate exposure and altered patterns of DNA methylation in cord blood. Environmental and Molecular Mutagenesis, 2017, 58, 398-410.	2.2	71
18	Comparison of DNA methylation measured by Illumina 450K and EPIC BeadChips in blood of newborns and 14-year-old children. Epigenetics, 2018, 13, 655-664.	2.7	65

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19	Developmental Changes in PON1 Enzyme Activity in Young Children and Effects of PON1 Polymorphisms. Environmental Health Perspectives, 2009, 117, 1632-1638.	6.0	64
20	Organophosphate pesticide exposure, PON1, and neurodevelopment in school-age children from the CHAMACOS study. Environmental Research, 2014, 134, 149-157.	7.5	63
21	Validation of PON1 enzyme activity assays for longitudinal studies. Clinica Chimica Acta, 2009, 402, 67-74.	1.1	62
22	Considerations for normalization of DNA methylation data by Illumina 450K BeadChip assay in population studies. Epigenetics, 2013, 8, 1141-1152.	2.7	60
23	In utero and childhood DDT, DDE, PBDE and PCBs exposure and sex hormones in adolescent boys: The CHAMACOS study. International Journal of Hygiene and Environmental Health, 2017, 220, 364-372.	4.3	58
24	Urinary Phthalate Metabolites and Biomarkers of Oxidative Stress in a Mexican-American Cohort: Variability in Early and Late Pregnancy. Toxics, 2016, 4, 7.	3.7	57
25	Maternal phthalate exposure during pregnancy is associated with DNA methylation of LINE-1 and Alu repetitive elements in Mexican-American children. Environmental Research, 2016, 148, 55-62.	7.5	49
26	Adiponectin and Leptin Trajectories in Mexican-American Children from Birth to 9 Years of Age. PLoS ONE, 2013, 8, e77964.	2.5	46
27	Longitudinal changes in PON1 enzymatic activities in Mexican–American mothers and children with different genotypes and haplotypes. Toxicology and Applied Pharmacology, 2010, 244, 181-189.	2.8	43
28	Estimation of blood cellular heterogeneity in newborns and children for epigenomeâ€wide association studies. Environmental and Molecular Mutagenesis, 2015, 56, 751-758.	2.2	43
29	DNA methylation and body mass index from birth to adolescence: meta-analyses of epigenome-wide association studies. Genome Medicine, 2020, 12, 105.	8.2	41
30	Cytogenetic Damage in Blood Lymphocytes and Exfoliated Epithelial Cells of Children With Inflammatory Bowel Disease. Pediatric Research, 2007, 61, 209-214.	2.3	38
31	Application of a geographic information system to explore associations between air pollution and micronucleus frequencies in African American children and adults. Environmental and Molecular Mutagenesis, 2006, 47, 236-246.	2.2	36
32	DNA methylation of imprinted genes in Mexican–American newborn children with prenatal phthalate exposure. Epigenomics, 2018, 10, 1011-1026.	2.1	33
33	PON1 as a model for integration of genetic, epigenetic, and expression data on candidate susceptibility genes. Environmental Epigenetics, 2015, 1, .	1.8	32
34	Prenatal Exposure to Mixtures of Phthalates, Parabens, and Other Phenols and Obesity in Five-Year-Olds in the CHAMACOS Cohort. International Journal of Environmental Research and Public Health, 2021, 18, 1796.	2.6	30
35	Metabolomic Markers of Phthalate Exposure in Plasma and Urine of Pregnant Women. Frontiers in Public Health, 2018, 6, 298.	2.7	29
36	DNA methylation and socioeconomic status in a Mexican-American birth cohort. Clinical Epigenetics, 2018, 10, 61.	4.1	26

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37	Cholinesterase and paraoxonase (PON1) enzyme activities in Mexican–American mothers and children from an agricultural community. Journal of Exposure Science and Environmental Epidemiology, 2012, 22, 641-648.	3.9	25
38	Associations of PON1 and Genetic Ancestry with Obesity in Early Childhood. PLoS ONE, 2013, 8, e62565.	2.5	25
39	Risk Factors Associated With SARS-CoV-2 Infection Among Farmworkers in Monterey County, California. JAMA Network Open, 2021, 4, e2124116.	5.9	25
40	Meta-analysis of epigenome-wide association studies in newborns and children show widespread sex differences in blood DNA methylation. Mutation Research - Reviews in Mutation Research, 2022, 789, 108415.	5.5	24
41	Folate concentrations in pediatric patients with newly diagnosed inflammatory bowel disease. American Journal of Clinical Nutrition, 2009, 89, 545-550.	4.7	23
42	Relationship between expression and methylation of obesity-related genes in children. Mutagenesis, 2015, 30, 411-420.	2.6	23
43	Epigenome-wide association study and epigenetic age acceleration associated with cigarette smoking among Costa Rican adults. Scientific Reports, 2022, 12, 4277.	3.3	22
44	PON1 DNA methylation and neurobehavior in Mexican-American children with prenatal organophosphate exposure. Environment International, 2018, 121, 31-40.	10.0	21
45	Effects of <i>PON</i> polymorphisms and haplotypes on molecular phenotype in Mexicanâ€American mothers and children. Environmental and Molecular Mutagenesis, 2011, 52, 105-116.	2.2	18
46	Recent progress in the genetics and epigenetics of paraoxonase. Current Opinion in Pediatrics, 2015, 27, 240-247.	2.0	18
47	Reduced Intracellular T-Helper 1 Interferon-Gamma in Blood of Newly Diagnosed Children With Crohn's Disease and Age-Related Changes in Th1/Th2 Cytokine Profiles. Pediatric Research, 2008, 63, 257-262.	2.3	16
48	miRNAs differentially expressed by next-generation sequencing in cord blood buffy coat samples of boys and girls. Epigenomics, 2016, 8, 1619-1635.	2.1	16
49	Associations between perinatal factors and adiponectin and leptin in 9â€yearâ€old <scp>M</scp> exican– <scp>A</scp> merican children. Pediatric Obesity, 2013, 8, 454-463.	2.8	15
50	Vitamin C intervention may lower the levels of persistent organic pollutants in blood of healthy women – A pilot study. Food and Chemical Toxicology, 2016, 92, 197-204.	3.6	15
51	DNA methylation of LINE-1 and Alu repetitive elements in relation to sex hormones and pubertal timing in Mexican-American children. Pediatric Research, 2016, 79, 855-862.	2.3	15
52	Discovery of tetrahydroisoquinoline (THIQ) derivatives as potent and orally bioavailable LFA-1/ICAM-1 antagonists. Bioorganic and Medicinal Chemistry Letters, 2010, 20, 5269-5273.	2.2	14
53	CpG Methylation across the adipogenic PPARÎ ³ gene and its relationship with birthweight and child BMI at 9Âyears. BMC Medical Genetics, 2017, 18, 7.	2.1	13
54	Meta-analysis of epigenome-wide associations between DNA methylation at birth and childhood cognitive skills. Molecular Psychiatry, 2022, 27, 2126-2135.	7.9	13

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55	AHR gene-dioxin interactions and birthweight in the Seveso Second Generation Health Study. International Journal of Epidemiology, 2018, 47, 1992-2004.	1.9	8
56	Age-Related Differences in miRNA Expression in Mexican-American Newborns and Children. International Journal of Environmental Research and Public Health, 2019, 16, 524.	2.6	8
57	Pregnancy lipidomic profiles and DNA methylation in newborns from the CHAMACOS cohort. Environmental Epigenetics, 2019, 5, dvz004.	1.8	7
58	Early-Life Home Environment and Obesity in a Mexican American Birth Cohort: The CHAMACOS Study. Psychosomatic Medicine, 2019, 81, 209-219.	2.0	2
59	Comparison of DNA methylation measurements from EPIC BeadChip and SeqCap targeted bisulphite sequencing in PON1 and nine additional candidate genes. Epigenetics, 2022, 17, 1944-1955.	2.7	1