

# Paul D Yousefi

## 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.

27  
papers

1,747  
citations

15  
h-index

29  
g-index

29  
ext. papers

2,364  
ext. citations

7.6  
avg, IF

3.53  
L-index

#	Paper	IF	Citations
27	The EWAS Catalog: a database of epigenome-wide association studies.. <i>Wellcome Open Research</i> , <b>2022</b> , 7, 41	4.8	3
26	DNA methylation-based predictors of health: applications and statistical considerations.. <i>Nature Reviews Genetics</i> , <b>2022</b> ,	30.1	3
25	Epigenetic modelling of former, current and never smokers. <i>Clinical Epigenetics</i> , <b>2021</b> , 13, 206	7.7	3
24	Maternal anxiety during pregnancy and newborn epigenome-wide DNA methylation. <i>Molecular Psychiatry</i> , <b>2021</b> , 26, 1832-1845	15.1	6
23	Do Mass Spectrometry-Derived Metabolomics Improve the Prediction of Pregnancy-Related Disorders? Findings from a UK Birth Cohort with Independent Validation. <i>Metabolites</i> , <b>2021</b> , 11,	5.6	2
22	DNA methylation changes associated with prenatal mercury exposure: A meta-analysis of prospective cohort studies from PACE consortium. <i>Environmental Research</i> , <b>2021</b> , 204, 112093	7.9	1
21	Do nuclear magnetic resonance (NMR)-based metabolomics improve the prediction of pregnancy-related disorders? Findings from a UK birth cohort with independent validation. <i>BMC Medicine</i> , <b>2020</b> , 18, 366	11.4	4
20	DNA methylation and body mass index from birth to adolescence: meta-analyses of epigenome-wide association studies. <i>Genome Medicine</i> , <b>2020</b> , 12, 105	14.4	15
19	Prenatal Particulate Air Pollution and DNA Methylation in Newborns: An Epigenome-Wide Meta-Analysis. <i>Environmental Health Perspectives</i> , <b>2019</b> , 127, 57012	8.4	58
18	Meta-analysis of epigenome-wide association studies in neonates reveals widespread differential DNA methylation associated with birthweight. <i>Nature Communications</i> , <b>2019</b> , 10, 1893	17.4	79
17	Validation and characterisation of a DNA methylation alcohol biomarker across the life course. <i>Clinical Epigenetics</i> , <b>2019</b> , 11, 163	7.7	6
16	Cohort Profile: Pregnancy And Childhood Epigenetics (PACE) Consortium. <i>International Journal of Epidemiology</i> , <b>2018</b> , 47, 22-23u	7.8	62
15	Prenatal phthalate exposure and altered patterns of DNA methylation in cord blood. <i>Environmental and Molecular Mutagenesis</i> , <b>2017</b> , 58, 398-410	3.2	52
14	Maternal BMI at the start of pregnancy and offspring epigenome-wide DNA methylation: findings from the pregnancy and childhood epigenetics (PACE) consortium. <i>Human Molecular Genetics</i> , <b>2017</b> , 26, 4067-4085	5.6	151
13	CpG Methylation across the adipogenic PPAR $\alpha$ gene and its relationship with birthweight and child BMI at 9 years. <i>BMC Medical Genetics</i> , <b>2017</b> , 18, 7	2.1	12
12	Genome-wide methylation data mirror ancestry information. <i>Epigenetics and Chromatin</i> , <b>2017</b> , 10, 1	5.8	48
11	Epigenetic Signatures of Cigarette Smoking. <i>Circulation: Cardiovascular Genetics</i> , <b>2016</b> , 9, 436-447		442

10	DNA Methylation in Newborns and Maternal Smoking in Pregnancy: Genome-wide Consortium Meta-analysis. <i>American Journal of Human Genetics</i> , <b>2016</b> , 98, 680-96	11	489
9	Maternal phthalate exposure during pregnancy is associated with DNA methylation of LINE-1 and Alu repetitive elements in Mexican-American children. <i>Environmental Research</i> , <b>2016</b> , 148, 55-62	7.9	39
8	PON1 as a model for integration of genetic, epigenetic, and expression data on candidate susceptibility genes. <i>Environmental Epigenetics</i> , <b>2015</b> , 1,	2.4	29
7	Sex differences in DNA methylation assessed by 450 K BeadChip in newborns. <i>BMC Genomics</i> , <b>2015</b> , 16, 911	4.5	106
6	Estimation of blood cellular heterogeneity in newborns and children for epigenome-wide association studies. <i>Environmental and Molecular Mutagenesis</i> , <b>2015</b> , 56, 751-8	3.2	39
5	Relationship between expression and methylation of obesity-related genes in children. <i>Mutagenesis</i> , <b>2015</b> , 30, 411-20	2.8	22
4	Genetic modification of the effect of maternal household air pollution exposure on birth weight in Guatemalan newborns. <i>Reproductive Toxicology</i> , <b>2014</b> , 50, 19-26	3.4	5
3	Human urinary mutagenicity after wood smoke exposure during traditional temazcal use. <i>Mutagenesis</i> , <b>2014</b> , 29, 367-77	2.8	11
2	Considerations for normalization of DNA methylation data by Illumina 450K BeadChip assay in population studies. <i>Epigenetics</i> , <b>2013</b> , 8, 1141-52	5.7	52
1	Do nuclear magnetic resonance (NMR)-based metabolomics improve the prediction of pregnancy-related disorders?		2