

Varun Kilaru

List of Publications by Year in descending order

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Version: 2024-02-01

32
papers

2,185
citations

471509

17
h-index

454955

30
g-index

32
all docs

32
docs citations

32
times ranked

3866
citing authors

#	ARTICLE	IF	CITATIONS
1	Methylation quantitative trait loci are largely consistent across disease states in Crohn's disease. <i>Genes, Genomes, Genetics</i> , 2022, 12, .	1.8	2
2	Cell-Type Specific Methylation Analysis Reveals Multiple Loci Associated With PTSD in African Americans. <i>Biological Psychiatry</i> , 2022, 91, S2.	1.3	0
3	PTSD is associated with increased DNA methylation across regions of HLA-DPB1 and SPATC1L. <i>Brain, Behavior, and Immunity</i> , 2021, 91, 429-436.	4.1	17
4	Genome-wide DNA methylation differences and polychlorinated biphenyl (PCB) exposure in a US population. <i>Epigenetics</i> , 2021, 16, 338-352.	2.7	12
5	Epigenetic prediction of 17 β -estradiol and relationship to trauma-related outcomes in women. <i>Comprehensive Psychoneuroendocrinology</i> , 2021, 6, 100045.	1.7	2
6	Critical evaluation of copy number variant calling methods using DNA methylation. <i>Genetic Epidemiology</i> , 2020, 44, 148-158.	1.3	12
7	Epigenome-wide meta-analysis of PTSD across 10 military and civilian cohorts identifies methylation changes in AHR. <i>Nature Communications</i> , 2020, 11, 5965.	12.8	84
8	Evaluating the impact of trauma and PTSD on epigenetic prediction of lifespan and neural integrity. <i>Neuropsychopharmacology</i> , 2020, 45, 1609-1616.	5.4	63
9	Sex-specific DNA methylation differences in people exposed to polybrominated biphenyl. <i>Epigenomics</i> , 2020, 12, 757-770.	2.1	13
10	Association of HLA locus alleles with posttraumatic stress disorder. <i>Brain, Behavior, and Immunity</i> , 2019, 81, 655-658.	4.1	30
11	Exposure to polybrominated biphenyl (PBB) associates with genome-wide DNA methylation differences in peripheral blood. <i>Epigenetics</i> , 2019, 14, 52-66.	2.7	38
12	25. Longitudinal Epigenome-Wide Changes From Trauma to PTSD Diagnosis. <i>Biological Psychiatry</i> , 2019, 85, S10-S11.	1.3	0
13	Exposure to polybrominated biphenyl and stochastic epigenetic mutations: application of a novel epigenetic approach to environmental exposure in the Michigan polybrominated biphenyl registry. <i>Epigenetics</i> , 2019, 14, 1003-1018.	2.7	14
14	Blood-Derived DNA Methylation Signatures of Crohn's Disease and Severity of Intestinal Inflammation. <i>Gastroenterology</i> , 2019, 156, 2254-2265.e3.	1.3	91
15	Environmental exposure to polybrominated biphenyl (PBB) associates with an increased rate of biological aging. <i>Aging</i> , 2019, 11, 5498-5517.	3.1	14
16	<i>SLC9B1</i> methylation predicts fetal intolerance of labor. <i>Epigenetics</i> , 2018, 13, 33-39.	2.7	16
17	Methylation differences reveal heterogeneity in preterm pathophysiology: results from bipartite network analyses. <i>Journal of Perinatal Medicine</i> , 2018, 46, 509-521.	1.4	13
18	Characterization of gene expression changes over healthy term pregnancies. <i>PLoS ONE</i> , 2018, 13, e0204228.	2.5	13

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19	Epigenome-wide association of PTSD from heterogeneous cohorts with a common multi-site analysis pipeline. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2017, 174, 619-630.	1.7	69
20	Exposure to Violence Accelerates Epigenetic Aging in Children. <i>Scientific Reports</i> , 2017, 7, 8962.	3.3	131
21	A Gene-Based Analysis of Acoustic Startle Latency. <i>Frontiers in Psychiatry</i> , 2017, 8, 117.	2.6	7
22	Resilience and biomarkers of health risk in Black smokers and nonsmokers.. <i>Health Psychology</i> , 2017, 36, 1047-1058.	1.6	12
23	An epigenetic clock for gestational age at birth based on blood methylation data. <i>Genome Biology</i> , 2016, 17, 206.	8.8	193
24	DNA methylation provides insight into intergenerational risk for preterm birth in African Americans. <i>Epigenetics</i> , 2015, 10, 784-792.	2.7	47
25	Accounting for Population Stratification in DNA Methylation Studies. <i>Genetic Epidemiology</i> , 2014, 38, 231-241.	1.3	207
26	Epigenetic changes associated with inflammation in breast cancer patients treated with chemotherapy. <i>Brain, Behavior, and Immunity</i> , 2014, 38, 227-236.	4.1	59
27	Fetal DNA Methylation Associates with Early Spontaneous Preterm Birth and Gestational Age. <i>PLoS ONE</i> , 2013, 8, e67489.	2.5	84
28	DNA methylation in neonates born to women receiving psychiatric care. <i>Epigenetics</i> , 2012, 7, 409-414.	2.7	68
29	Prenatal antiepileptic exposure associates with neonatal DNA methylation differences. <i>Epigenetics</i> , 2012, 7, 458-463.	2.7	46
30	MethLAB: A graphical user interface package for the analysis of array-based DNA methylation data. <i>Epigenetics</i> , 2012, 7, 225-229.	2.7	38
31	Post-traumatic stress disorder is associated with PACAP and the PAC1 receptor. <i>Nature</i> , 2011, 470, 492-497.	27.8	695
32	Neonatal DNA methylation patterns associate with gestational age. <i>Epigenetics</i> , 2011, 6, 1498-1504.	2.7	95