

Sarah J Marzi

List of Publications by Citations

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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.

19
papers

297
citations

7
h-index

17
g-index

22
ext. papers

447
ext. citations

9.5
avg, IF

3.31
L-index

#	Paper	IF	Citations
19	Analysis of DNA Methylation in Young People: Limited Evidence for an Association Between Victimization Stress and Epigenetic Variation in Blood. <i>American Journal of Psychiatry</i> , 2018 , 175, 517-529	11.9	83
18	A histone acetylome-wide association study of Alzheimers disease identifies disease-associated H3K27ac differences in the entorhinal cortex. <i>Nature Neuroscience</i> , 2018 , 21, 1618-1627	25.5	72
17	Severe psychosocial deprivation in early childhood is associated with increased DNA methylation across a region spanning the transcription start site of CYP2E1. <i>Translational Psychiatry</i> , 2016 , 6, e830	8.6	41
16	Epigenetics in eating disorders: a systematic review. <i>Molecular Psychiatry</i> , 2019 , 24, 901-915	15.1	32
15	Tissue-specific patterns of allelically-skewed DNA methylation. <i>Epigenetics</i> , 2016 , 11, 24-35	5.7	18
14	DNA methylation in human sperm: a systematic review. <i>Human Reproduction Update</i> , 2020 , 26, 841-873	15.8	15
13	Genetic risk variants for brain disorders are enriched in cortical H3K27ac domains. <i>Molecular Brain</i> , 2019 , 12, 7	4.5	11
12	Early life diet conditions the molecular response to post-weaning protein restriction in the mouse. <i>BMC Biology</i> , 2018 , 16, 51	7.3	7
11	The histone modification H3K4me3 is altered at the locus in Alzheimers disease brain. <i>Future Science OA</i> , 2021 , 7, FSO665	2.7	5
10	The DNA methylome of human sperm is distinct from blood with little evidence for tissue-consistent obesity associations. <i>PLoS Genetics</i> , 2020 , 16, e1009035	6	4
9	Iterative Fragmentation Improves the Detection of ChIP-seq Peaks for Inactive Histone Marks. <i>Bioinformatics and Biology Insights</i> , 2016 , 10, 209-224	5.3	3
8	Genetic variation at mouse and human ribosomal DNA influences associated epigenetic states.. <i>Genome Biology</i> , 2022 , 23, 54	18.3	3
7	DNA methylation covariation in human whole blood and sperm: implications for studies of intergenerational epigenetic effects		2
6	Genetic polymorphisms and their association with brain and behavioural measures in heterogeneous stock mice. <i>Scientific Reports</i> , 2017 , 7, 41204	4.9	1
5	Chd8 regulates X chromosome inactivation in mouse through fine-tuning control of Xist expression. <i>Communications Biology</i> , 2021 , 4, 485	6.7	0
4	The DNA methylome of human sperm is distinct from blood with little evidence for tissue-consistent obesity associations 2020 , 16, e1009035		
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2 The DNA methylome of human sperm is distinct from blood with little evidence for tissue-consistent obesity associations **2020**, 16, e1009035

1 The DNA methylome of human sperm is distinct from blood with little evidence for tissue-consistent obesity associations **2020**, 16, e1009035