Matt J Silver

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.

31	1,392	18	35
papers	citations	h-index	g-index
35	1,731 ext. citations	8.8	4.2
ext. papers		avg, IF	L-index

#	Paper	IF	Citations
31	DNA methylation signatures associated with cardiometabolic risk factors in children from India and The Gambia: results from the EMPHASIS study <i>Clinical Epigenetics</i> , 2022 , 14, 6	7.7	1
30	Environmentally sensitive hotspots in the methylome of the early human embryo <i>ELife</i> , 2022 , 11,	8.9	3
29	DNA methylation at a nutritionally sensitive region of the gene is associated with thyroid volume and function in Gambian children. <i>Science Advances</i> , 2021 , 7, eabj1561	14.3	2
28	The Role of Nutrition in COVID-19 Susceptibility and Severity of Disease: A Systematic Review. Journal of Nutrition, 2021 , 151, 1854-1878	4.1	24
27	Periconceptional environment predicts leukocyte telomere length in a cross-sectional study of 7-9 year old rural Gambian children. <i>Scientific Reports</i> , 2020 , 10, 9675	4.9	1
26	Intergenerational Influences on Child Development: An Epigenetic Perspective. <i>Nestle Nutrition Institute Workshop Series</i> , 2020 , 93, 145-152	1.9	3
25	Fetal programming and epigenetics. <i>Current Opinion in Endocrine and Metabolic Research</i> , 2020 , 13, 1-6	1.7	9
24	Effect of maternal preconceptional and pregnancy micronutrient interventions on childrens DNA methylation: Findings from the EMPHASIS study. <i>American Journal of Clinical Nutrition</i> , 2020 , 112, 1099-	·7113	5
23	A genomic atlas of systemic interindividual epigenetic variation in humans. <i>Genome Biology</i> , 2019 , 20, 105	18.3	37
22	Meta-analysis of epigenome-wide association studies in neonates reveals widespread differential DNA methylation associated with birthweight. <i>Nature Communications</i> , 2019 , 10, 1893	17.4	79
21	A novel nutritional supplement to reduce plasma homocysteine in nonpregnant women: A randomised controlled trial in The Gambia. <i>PLoS Medicine</i> , 2019 , 16, e1002870	11.6	1
20	Maternal One-Carbon Metabolism and Infant DNA Methylation between Contrasting Seasonal Environments: A Case Study from The Gambia. <i>Current Developments in Nutrition</i> , 2019 , 3,	0.4	9
19	Estimation of a significance threshold for epigenome-wide association studies. <i>Genetic Epidemiology</i> , 2018 , 42, 20-33	2.6	67
18	Establishment of environmentally sensitive DNA methylation states in the very early human embryo. <i>Science Advances</i> , 2018 , 4, eaat2624	14.3	36
17	Epigenetic supersimilarity of monozygotic twin pairs. <i>Genome Biology</i> , 2018 , 19, 2	18.3	52
16	Candidate genes linking maternal nutrient exposure to offspring health via DNA methylation: a review of existing evidence in humans with specific focus on one-carbon metabolism. <i>International Journal of Epidemiology</i> , 2018 , 47, 1910-1937	7.8	33
15	Protocol for the EMPHASIS study; epigenetic mechanisms linking maternal pre-conceptional nutrition and childrens health in India and Sub-Saharan Africa. <i>BMC Nutrition</i> , 2017 , 3,	2.5	7

LIST OF PUBLICATIONS

14	Influence of intergenerational parental energy and nutrient restriction on offspring growth in rural Gambia. <i>FASEB Journal</i> , 2017 , 31, 4928-4934	0.9	13
13	Interindividual Variation in DNA Methylation at a Putative POMC Metastable Epiallele Is Associated with Obesity. <i>Cell Metabolism</i> , 2016 , 24, 502-509	24.6	82
12	Possible relationship between common genetic variation and white matter development in a pilot study of preterm infants. <i>Brain and Behavior</i> , 2016 , 6, e00434	3.4	21
11	Vitamin D binding protein genotype is associated with plasma 25OHD concentration in West African children. <i>Bone</i> , 2015 , 74, 166-70	4.7	27
10	Evidence for negative selection of gene variants that increase dependence on dietary choline in a Gambian cohort. <i>FASEB Journal</i> , 2015 , 29, 3426-35	0.9	14
9	Exposure to aflatoxin B1 in utero is associated with DNA methylation in white blood cells of infants in The Gambia. <i>International Journal of Epidemiology</i> , 2015 , 44, 1238-48	7.8	69
8	Independent genomewide screens identify the tumor suppressor VTRNA2-1 as a human epiallele responsive to periconceptional environment. <i>Genome Biology</i> , 2015 , 16, 118	18.3	119
7	Maternal nutrition at conception modulates DNA methylation of human metastable epialleles. <i>Nature Communications</i> , 2014 , 5, 3746	17.4	362
6	Pathways-driven sparse regression identifies pathways and genes associated with high-density lipoprotein cholesterol in two Asian cohorts. <i>PLoS Genetics</i> , 2013 , 9, e1003939	6	22
5	Fast identification of biological pathways associated with a quantitative trait using group lasso with overlaps. <i>Statistical Applications in Genetics and Molecular Biology</i> , 2012 , 11, Article 7	1.2	36
4	Identification of gene pathways implicated in Alzheimers disease using longitudinal imaging phenotypes with sparse regression. <i>NeuroImage</i> , 2012 , 63, 1681-94	7.9	60
3	False positives in neuroimaging genetics using voxel-based morphometry data. <i>NeuroImage</i> , 2011 , 54, 992-1000	7.9	118
2	Spatial effects favour the evolution of niche construction. <i>Theoretical Population Biology</i> , 2006 , 70, 387	-400	72
1	Could nutrition modulate COVID-19 susceptibility and severity of disease? A systematic review		8