

Jennifer Brody

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11301876/publications.pdf>

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12
papers

2,741
citations

759233

12
h-index

1199594

12
g-index

14
all docs

14
docs citations

14
times ranked

11454
citing authors

#	ARTICLE	IF	CITATIONS
1	Systematic identification of trans eQTLs as putative drivers of known disease associations. <i>Nature Genetics</i> , 2013, 45, 1238-1243.	21.4	1,544
2	The transcriptional landscape of age in human peripheral blood. <i>Nature Communications</i> , 2015, 6, 8570.	12.8	533
3	Whole-Exome Sequencing Identifies Rare and Low-Frequency Coding Variants Associated with LDL Cholesterol. <i>American Journal of Human Genetics</i> , 2014, 94, 233-245.	6.2	193
4	Dynamic incorporation of multiple in silico functional annotations empowers rare variant association analysis of large whole-genome sequencing studies at scale. <i>Nature Genetics</i> , 2020, 52, 969-983.	21.4	146
5	DNA Methylation Signatures of Depressive Symptoms in Middle-aged and Elderly Persons. <i>JAMA Psychiatry</i> , 2018, 75, 949.	11.0	78
6	Sequence Kernel Association Test for Survival Traits. <i>Genetic Epidemiology</i> , 2014, 38, 191-197.	1.3	58
7	Assessing the causal relationship between obesity and venous thromboembolism through a Mendelian Randomization study. <i>Human Genetics</i> , 2017, 136, 897-902.	3.8	46
8	Association of dietary folate and vitamin B-12 intake with genome-wide DNA methylation in blood: a large-scale epigenome-wide association analysis in 5841 individuals. <i>American Journal of Clinical Nutrition</i> , 2019, 110, 437-450.	4.7	46
9	Association of exome sequences with plasma C-reactive protein levels in >9000 participants. <i>Human Molecular Genetics</i> , 2015, 24, 559-571.	2.9	36
10	FastSKAT: Sequence kernel association tests for very large sets of markers. <i>Genetic Epidemiology</i> , 2018, 42, 516-527.	1.3	26
11	Sequencing of 2 Subclinical Atherosclerosis Candidate Regions in 3669 Individuals. <i>Circulation: Cardiovascular Genetics</i> , 2014, 7, 359-364.	5.1	18
12	Common variation in fatty acid metabolic genes and risk of incident sudden cardiac arrest. <i>Heart Rhythm</i> , 2014, 11, 471-477.	0.7	16