

Paul J Hoffman

List of Publications by Citations

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Version: 2024-04-23

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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.

14
papers

8,974
citations

11
h-index

19
g-index

19
ext. papers

17,929
ext. citations

27.9
avg, IF

6.84
L-index

#	Paper	IF	Citations
14	Integrating single-cell transcriptomic data across different conditions, technologies, and species. <i>Nature Biotechnology</i> , 2018 , 36, 411-420	44.5	4181
13	Comprehensive Integration of Single-Cell Data. <i>Cell</i> , 2019 , 177, 1888-1902.e21	56.2	3858
12	Integrated analysis of multimodal single-cell data. <i>Cell</i> , 2021 , 184, 3573-3587.e29	56.2	558
11	Integrated analysis of multimodal single-cell data		91
10	A Quantitative Proteome Map of the Human Body. <i>Cell</i> , 2020 , 183, 269-283.e19	56.2	73
9	Comprehensive integration of single cell data		58
8	The Role of Deleterious Substitutions in Crop Genomes. <i>Molecular Biology and Evolution</i> , 2016 , 33, 2307-8.7		54
7	Genetic regulatory variation in populations informs transcriptome analysis in rare disease. <i>Science</i> , 2019 , 366, 351-356	33.3	42
6	A vast resource of allelic expression data spanning human tissues. <i>Genome Biology</i> , 2020 , 21, 234	18.3	18
5	Comparative Genomics Approaches Accurately Predict Deleterious Variants in Plants. <i>G3: Genes, Genomes, Genetics</i> , 2018 , 8, 3321-3329	3.2	16
4	angsd-wrapper: utilities for analysing next-generation sequencing data. <i>Molecular Ecology Resources</i> , 2016 , 16, 1449-1454	8.4	11
3	Environmental Association Identifies Candidates for Tolerance to Low Temperature and Drought. <i>G3: Genes, Genomes, Genetics</i> , 2019 , 9, 3423-3438	3.2	6
2	Transcriptome variation in human tissues revealed by long-read sequencing		6
1	Comparative genomics approaches accurately predict deleterious variants in plants		2