

Jeffrey A Rosenfeld

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6582723/publications.pdf>

Version: 2024-02-01

13
papers

3,337
citations

840776

11
h-index

1058476

14
g-index

18
all docs

18
docs citations

18
times ranked

6100
citing authors

#	ARTICLE	IF	CITATIONS
1	A complete reference genome improves analysis of human genetic variation. <i>Science</i> , 2022, 376, eabl3533.	12.6	144
2	The complete sequence of a human genome. <i>Science</i> , 2022, 376, 44-53.	12.6	1,222
3	Benchmarking challenging small variants with linked and long reads. <i>Cell Genomics</i> , 2022, 2, 100128.	6.5	77
4	RNA-seq profile of African American men with a clinically localized prostate cancer. <i>Prostate International</i> , 2021, 9, 125-131.	2.3	8
5	A robust benchmark for detection of germline large deletions and insertions. <i>Nature Biotechnology</i> , 2020, 38, 1347-1355.	17.5	233
6	Single-cell sperm transcriptomes and variants from fathers of children with and without autism spectrum disorder. <i>Npj Genomic Medicine</i> , 2020, 5, 14.	3.8	10
7	TCF4 induces enzalutamide resistance via neuroendocrine differentiation in prostate cancer. <i>PLoS ONE</i> , 2019, 14, e0213488.	2.5	20
8	An essential domain of an early-diverged RNA polymerase II functions to accurately decode a primitive chromatin landscape. <i>Nucleic Acids Research</i> , 2017, 45, 7886-7896.	14.5	17
9	Patient-Derived Xenograft Models of Non-Small Cell Lung Cancer and Their Potential Utility in Personalized Medicine. <i>Frontiers in Oncology</i> , 2017, 7, 2.	2.8	63
10	Multi-platform assessment of transcriptome profiling using RNA-seq in the ABRF next-generation sequencing study. <i>Nature Biotechnology</i> , 2014, 32, 915-925.	17.5	217
11	E value cutoff and eukaryotic genome content phylogenetics. <i>Molecular Phylogenetics and Evolution</i> , 2012, 63, 342-350.	2.7	14
12	Random roots and lineage sorting. <i>Molecular Phylogenetics and Evolution</i> , 2012, 64, 12-20.	2.7	38
13	A Systematic Survey of Loss-of-Function Variants in Human Protein-Coding Genes. <i>Science</i> , 2012, 335, 823-828.	12.6	1,095