

Joshua Gould

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2194188/publications.pdf>

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19
papers

9,603
citations

516561

16
h-index

887953

17
g-index

25
all docs

25
docs citations

25
times ranked

22139
citing authors

#	ARTICLE	IF	CITATIONS
1	Single-cell meta-analysis of SARS-CoV-2 entry genes across tissues and demographics. <i>Nature Medicine</i> , 2021, 27, 546-559.	15.2	261
2	COVID-19 tissue atlases reveal SARS-CoV-2 pathology and cellular targets. <i>Nature</i> , 2021, 595, 107-113.	13.7	537
3	A Distinct Transcriptional Program in Human CART T Cells Bearing the 4-1BB Signaling Domain Revealed by scRNA-Seq. <i>Molecular Therapy</i> , 2020, 28, 2577-2592.	3.7	58
4	Cumulus provides cloud-based data analysis for large-scale single-cell and single-nucleus RNA-seq. <i>Nature Methods</i> , 2020, 17, 793-798.	9.0	134
5	A single-cell and single-nucleus RNA-Seq toolbox for fresh and frozen human tumors. <i>Nature Medicine</i> , 2020, 26, 792-802.	15.2	381
6	High-definition spatial transcriptomics for in situ tissue profiling. <i>Nature Methods</i> , 2019, 16, 987-990.	9.0	708
7	Optimal-Transport Analysis of Single-Cell Gene Expression Identifies Developmental Trajectories in Reprogramming. <i>Cell</i> , 2019, 176, 928-943.e22.	13.5	411
8	The Drug Repurposing Hub: a next-generation drug library and information resource. <i>Nature Medicine</i> , 2017, 23, 405-408.	15.2	689
9	Genetic interrogation of circulating multiple myeloma cells at single-cell resolution. <i>Science Translational Medicine</i> , 2016, 8, 363ra147.	5.8	126
10	Targeting MTHFD2 in acute myeloid leukemia. <i>Journal of Experimental Medicine</i> , 2016, 213, 1285-1306.	4.2	118
11	Comprehensive Genetic Interrogation of Circulating Multiple Myeloma Cells at Single Cell Resolution. <i>Blood</i> , 2016, 128, 800-800.	0.6	0
12	Harnessing Connectivity in a Large-Scale Small-Molecule Sensitivity Dataset. <i>Cancer Discovery</i> , 2015, 5, 1210-1223.	7.7	575
13	Widespread Genetic Heterogeneity in Multiple Myeloma: Implications for Targeted Therapy. <i>Cancer Cell</i> , 2014, 25, 91-101.	7.7	847
14	Discovery and prioritization of somatic mutations in diffuse large B-cell lymphoma (DLBCL) by whole-exome sequencing. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 3879-3884.	3.3	853
15	Tumour micro-environment elicits innate resistance to RAF inhibitors through HGF secretion. <i>Nature</i> , 2012, 487, 500-504.	13.7	1,561
16	Systematic investigation of genetic vulnerabilities across cancer cell lines reveals lineage-specific dependencies in ovarian cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 12372-12377.	3.3	383
17	Multiple Myeloma Sequencing Reveals Subclonality and Timing of Genetic Alterations. <i>Blood</i> , 2011, 118, 2897-2897.	0.6	0
18	The Mutational Landscape of Diffuse Large B Cell Lymphoma. <i>Blood</i> , 2011, 118, 259-259.	0.6	0

#	ARTICLE	IF	CITATIONS
19	GenePattern 2.0. Nature Genetics, 2006, 38, 500-501.	9.4	1,848