

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

516710

16
h-index

888059

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. Nature Medicine, 2021, 27, 546-559.	30.7	261
2	COVID-19 tissue atlases reveal SARS-CoV-2 pathology and cellular targets. Nature, 2021, 595, 107-113.	27.8	537
3	A Distinct Transcriptional Program in Human CAR T Cells Bearing the 4-1BB Signaling Domain Revealed by scRNA-Seq. Molecular Therapy, 2020, 28, 2577-2592.	8.2	58
4	Cumulus provides cloud-based data analysis for large-scale single-cell and single-nucleus RNA-seq. Nature Methods, 2020, 17, 793-798.	19.0	134
5	A single-cell and single-nucleus RNA-Seq toolbox for fresh and frozen human tumors. Nature Medicine, 2020, 26, 792-802.	30.7	381
6	High-definition spatial transcriptomics for in situ tissue profiling. Nature Methods, 2019, 16, 987-990.	19.0	708
7	Optimal-Transport Analysis of Single-Cell Gene Expression Identifies Developmental Trajectories in Reprogramming. Cell, 2019, 176, 928-943.e22.	28.9	411
8	The Drug Repurposing Hub: a next-generation drug library and information resource. Nature Medicine, 2017, 23, 405-408.	30.7	689
9	Genetic interrogation of circulating multiple myeloma cells at single-cell resolution. Science Translational Medicine, 2016, 8, 363ra147.	12.4	126
10	Targeting MTHFD2 in acute myeloid leukemia. Journal of Experimental Medicine, 2016, 213, 1285-1306.	8.5	118
11	Comprehensive Genetic Interrogation of Circulating Multiple Myeloma Cells at Single Cell Resolution. Blood, 2016, 128, 800-800.	1.4	0
12	Harnessing Connectivity in a Large-Scale Small-Molecule Sensitivity Dataset. Cancer Discovery, 2015, 5, 1210-1223.	9.4	575
13	Widespread Genetic Heterogeneity in Multiple Myeloma: Implications for Targeted Therapy. Cancer Cell, 2014, 25, 91-101.	16.8	847
14	Discovery and prioritization of somatic mutations in diffuse large B-cell lymphoma (DLBCL) by whole-exome sequencing. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 3879-3884.	7.1	853
15	Tumour micro-environment elicits innate resistance to RAF inhibitors through HGF secretion. Nature, 2012, 487, 500-504.	27.8	1,561
16	Systematic investigation of genetic vulnerabilities across cancer cell lines reveals lineage-specific dependencies in ovarian cancer. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 12372-12377.	7.1	383
17	Multiple Myeloma Sequencing Reveals Subclonality and Timing of Genetic Alterations. Blood, 2011, 118, 2897-2897.	1.4	0
18	The Mutational Landscape of Diffuse Large B Cell Lymphoma. Blood, 2011, 118, 259-259.	1.4	0

#	ARTICLE	IF	CITATIONS
19	GenePattern 2.0. Nature Genetics, 2006, 38, 500-501.	21.4	1,848