

Jamie L Marshall

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

3,636
citations

643344

15
h-index

843174

20
g-index

26
all docs

26
docs citations

26
times ranked

8130
citing authors

#	ARTICLE	IF	CITATIONS
1	Single-Cell Transcriptomics Reveal Disrupted Kidney Filter Cell-Cell Interactions after Early and Selective Podocyte Injury. <i>American Journal of Pathology</i> , 2022, 192, 281-294.	1.9	7
2	High-resolution Slide-seqV2 spatial transcriptomics enables discovery of disease-specific cell neighborhoods and pathways. <i>iScience</i> , 2022, 25, 104097.	1.9	32
3	Single-nucleus cross-tissue molecular reference maps toward understanding disease gene function. <i>Science</i> , 2022, 376, eabl4290.	6.0	180
4	Highly sensitive spatial transcriptomics at near-cellular resolution with Slide-seqV2. <i>Nature Biotechnology</i> , 2021, 39, 313-319.	9.4	569
5	Targeting a Braf/Mapk pathway rescues podocyte lipid peroxidation in CoQ-deficiency kidney disease. <i>Journal of Clinical Investigation</i> , 2021, 131, .	3.9	25
6	Compressed sensing for highly efficient imaging transcriptomics. <i>Nature Biotechnology</i> , 2021, 39, 936-942.	9.4	33
7	Principles of Spatial Transcriptomics Analysis: A Practical Walk-Through in Kidney Tissue. <i>Frontiers in Physiology</i> , 2021, 12, 809346.	1.3	14
8	Efficient, continuous mutagenesis in human cells using a pseudo-random DNA editor. <i>Nature Biotechnology</i> , 2020, 38, 165-168.	9.4	59
9	The effect of LRRK2 loss-of-function variants in humans. <i>Nature Medicine</i> , 2020, 26, 869-877.	15.2	79
10	Disease-associated astrocytes in Alzheimer's disease and aging. <i>Nature Neuroscience</i> , 2020, 23, 701-706.	7.1	525
11	HyPR-seq: Single-cell quantification of chosen RNAs via hybridization and sequencing of DNA probes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 33404-33413.	3.3	21
12	Identification of a Novel Deep Intronic Mutation in CAPN3 Presenting a Promising Target for Therapeutic Splice Modulation. <i>Journal of Neuromuscular Diseases</i> , 2019, 6, 475-483.	1.1	6
13	Resolving the full spectrum of human genome variation using Linked-Reads. <i>Genome Research</i> , 2019, 29, 635-645.	2.4	182
14	Single cell census of human kidney organoids shows reproducibility and diminished off-target cells after transplantation. <i>Nature Communications</i> , 2019, 10, 5462.	5.8	133
15	A recurrent COL6A1 pseudoexon insertion causes muscular dystrophy and is effectively targeted by splice-correction therapies. <i>JCI Insight</i> , 2019, 4, .	2.3	33
16	Congenital Titinopathy: Comprehensive characterization and pathogenic insights. <i>Annals of Neurology</i> , 2018, 83, 1105-1124.	2.8	93
17	Improving genetic diagnosis in Mendelian disease with transcriptome sequencing. <i>Science Translational Medicine</i> , 2017, 9, .	5.8	516
18	Landscape of X chromosome inactivation across human tissues. <i>Nature</i> , 2017, 550, 244-248.	13.7	764

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19	Differentiation-related glycan epitopes identify discrete domains of the muscle glycocalyx. <i>Glycobiology</i> , 2016, 26, 1120-1132.	1.3	10
20	Quantifying prion disease penetrance using large population control cohorts. <i>Science Translational Medicine</i> , 2016, 8, 322ra9.	5.8	289