Jamie L Marshall

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3527577/publications.pdf

Version: 2024-02-01

20 papers 3,636 citations

567281 15 h-index 752698 20 g-index

26 all docs

26 docs citations

26 times ranked

7476 citing authors

| # | Article | IF | Citations |
|----|---|-------------|-----------|
| 1 | Single-Cell Transcriptomics Reveal Disrupted Kidney Filter Cell-Cell Interactions after Early and Selective Podocyte Injury. American Journal of Pathology, 2022, 192, 281-294. | 3.8 | 7 |
| 2 | High-resolution Slide-seqV2 spatial transcriptomics enables discovery of disease-specific cell neighborhoods and pathways. IScience, 2022, 25, 104097. | 4.1 | 32 |
| 3 | Single-nucleus cross-tissue molecular reference maps toward understanding disease gene function. Science, 2022, 376, eabl4290. | 12.6 | 180 |
| 4 | Highly sensitive spatial transcriptomics at near-cellular resolution with Slide-seqV2. Nature Biotechnology, 2021, 39, 313-319. | 17.5 | 569 |
| 5 | Targeting a Braf/Mapk pathway rescues podocyte lipid peroxidation in CoQ-deficiency kidney disease. Journal of Clinical Investigation, 2021, 131, . | 8.2 | 25 |
| 6 | Compressed sensing for highly efficient imaging transcriptomics. Nature Biotechnology, 2021, 39, 936-942. | 17.5 | 33 |
| 7 | Principles of Spatial Transcriptomics Analysis: A Practical Walk-Through in Kidney Tissue. Frontiers in Physiology, 2021, 12, 809346. | 2.8 | 14 |
| 8 | Efficient, continuous mutagenesis in human cells using a pseudo-random DNA editor. Nature Biotechnology, 2020, 38, 165-168. | 17.5 | 59 |
| 9 | The effect of LRRK2 loss-of-function variants in humans. Nature Medicine, 2020, 26, 869-877. | 30.7 | 79 |
| 10 | Disease-associated astrocytes in Alzheimer's disease and aging. Nature Neuroscience, 2020, 23, 701-706. | 14.8 | 525 |
| 11 | HyPR-seq: Single-cell quantification of chosen RNAs via hybridization and sequencing of DNA probes. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 33404-33413. | 7.1 | 21 |
| 12 | Identification of a Novel Deep Intronic Mutation in CAPN3 Presenting a Promising Target for Therapeutic Splice Modulation. Journal of Neuromuscular Diseases, 2019, 6, 475-483. | 2.6 | 6 |
| 13 | Resolving the full spectrum of human genome variation using Linked-Reads. Genome Research, 2019, 29, 635-645. | 5.5 | 182 |
| 14 | Single cell census of human kidney organoids shows reproducibility and diminished off-target cells after transplantation. Nature Communications, 2019, 10, 5462. | 12.8 | 133 |
| 15 | A recurrent COL6A1 pseudoexon insertion causes muscular dystrophy and is effectively targeted by splice-correction therapies. JCl Insight, 2019, 4, . | 5.0 | 33 |
| 16 | Congenital Titinopathy: Comprehensive characterization and pathogenic insights. Annals of Neurology, 2018, 83, 1105-1124. | 5. 3 | 93 |
| 17 | Improving genetic diagnosis in Mendelian disease with transcriptome sequencing. Science Translational Medicine, 2017, 9, . | 12.4 | 516 |
| 18 | Landscape of X chromosome inactivation across human tissues. Nature, 2017, 550, 244-248. | 27.8 | 764 |

| # | Article | IF | CITATIONS |
|----|--|------|-----------|
| 19 | Differentiation-related glycan epitopes identify discrete domains of the muscle glycocalyx. Glycobiology, 2016, 26, 1120-1132. | 2.5 | 10 |
| 20 | Quantifying prion disease penetrance using large population control cohorts. Science Translational Medicine, 2016, 8, 322ra9. | 12.4 | 289 |