Patrik L Ståhl

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3452492/publications.pdf

Version: 2024-02-01

25 papers 4,644 citations

430874 18 h-index 610901 24 g-index

30 all docs

30 docs citations

30 times ranked

5591 citing authors

#	Article	IF	CITATIONS
1	Three-dimensional spatial transcriptomics uncovers cell type localizations in the human rheumatoid arthritis synovium. Communications Biology, 2022, 5, 129.	4.4	35
2	Single cell and spatial transcriptomics in human tendon disease indicate dysregulated immune homeostasis. Annals of the Rheumatic Diseases, 2021, 80, 1494-1497.	0.9	33
3	Divergent clonal differentiation trajectories establish CD8+ memory TÂcell heterogeneity during acute viral infections in humans. Cell Reports, 2021, 35, 109174.	6.4	9
4	Spatial mapping reveals human adipocyte subpopulations with distinct sensitivities to insulin. Cell Metabolism, 2021, 33, 1869-1882.e6.	16.2	92
5	Identification and transfer of spatial transcriptomics signatures for cancer diagnosis. Breast Cancer Research, 2020, 22, 6.	5.0	54
6	ST viewer: a tool for analysis and visualization of spatial transcriptomics datasets. Bioinformatics, 2019, 35, 1058-1060.	4.1	30
7	Conbase: a software for unsupervised discovery of clonal somatic mutations in single cells through read phasing. Genome Biology, 2019, 20, 68.	8.8	21
8	High-definition spatial transcriptomics for in situ tissue profiling. Nature Methods, 2019, 16, 987-990.	19.0	708
9	Exploring inflammatory signatures in arthritic joint biopsies with Spatial Transcriptomics. Scientific Reports, 2019, 9, 18975.	3.3	55
10	A Spatiotemporal Organ-Wide Gene Expression and Cell Atlas of the Developing Human Heart. Cell, 2019, 179, 1647-1660.e19.	28.9	470
11	ST Spot Detector: a web-based application for automatic spot and tissue detection for spatial Transcriptomics image datasets. Bioinformatics, 2018, 34, 1966-1968.	4.1	30
12	Barcoded solid-phase RNA capture for Spatial Transcriptomics profiling in mammalian tissue sections. Nature Protocols, 2018, 13, 2501-2534.	12.0	144
13	Spatial maps of prostate cancer transcriptomes reveal an unexplored landscape of heterogeneity. Nature Communications, 2018, 9, 2419.	12.8	374
14	Spatially resolved transcriptome profiling in model plant species. Nature Plants, 2017, 3, 17061.	9.3	135
15	ST Pipeline: an automated pipeline for spatial mapping of unique transcripts. Bioinformatics, 2017, 33, 2591-2593.	4.1	81
16	Spatial detection of fetal marker genes expressed at low level in adult human heart tissue. Scientific Reports, 2017, 7, 12941.	3.3	62
17	05.16â€Transcriptome visualisation of the inflamed rheumatoid arthritis joint. , 2017, , .		0
18	Massive and parallel expression profiling using microarrayed single-cell sequencing. Nature Communications, 2016, 7, 13182.	12.8	44

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#	Article	IF	CITATIONS
19	An automated approach to prepare tissue-derived spatially barcoded RNA-sequencing libraries. Scientific Reports, 2016, 6, 37137.	3.3	52
20	Visualization and analysis of gene expression in tissue sections by spatial transcriptomics. Science, 2016, 353, 78-82.	12.6	1,983
21	The age and genomic integrity of neurons after cortical stroke in humans. Nature Neuroscience, 2014, 17, 801-803.	14.8	108
22	Toward the Single-Hour High-Quality Genome. Annual Review of Biochemistry, 2012, 81, 359-378.	11.1	29
23	Translational Database Selection and Multiplexed Sequence Capture for Up Front Filtering of Reliable Breast Cancer Biomarker Candidates. PLoS ONE, 2011, 6, e20794.	2.5	2
24	Sun-Induced Nonsynonymous p53 Mutations Are Extensively Accumulated and Tolerated in Normal Appearing Human Skin. Journal of Investigative Dermatology, 2011, 131, 504-508.	0.7	49
25	Visual DNA - Identification of DNA sequence variations by bead trapping. Genomics, 2007, 90, 741-745.	2.9	14