

Slide-seq: A scalable technology for measuring genome-resolution

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Questions and (some) answers on reactive astrocytes. <i>Glia</i> , 2019, 67, 2221-2247.	2.5	185
2	RNA sequencing: the teenage years. <i>Nature Reviews Genetics</i> , 2019, 20, 631-656.	7.7	1,192
3	The Spatial and Genomic Hierarchy of Tumor Ecosystems Revealed by Single-Cell Technologies. <i>Trends in Cancer</i> , 2019, 5, 411-425.	3.8	44
4	The Unmixing Problem: A Guide to Applying Single-Cell RNA Sequencing to Bone. <i>Journal of Bone and Mineral Research</i> , 2019, 34, 1207-1219.	3.1	34
6	Computational approaches for characterizing the tumor immune microenvironment. <i>Immunology</i> , 2019, 158, 70-84.	2.0	30
7	Characteristics of a novel cell line ZJU-0430 established from human gallbladder carcinoma. <i>Cancer Cell International</i> , 2019, 19, 190.	1.8	6
8	Highly Multiplexed, Quantitative Tissue Imaging at Cellular Resolution. <i>Current Pathobiology Reports</i> , 2019, 7, 109-118.	1.6	2
9	The Potential of Astrocytes as Immune Modulators in Brain Tumors. <i>Frontiers in Immunology</i> , 2019, 10, 1314.	2.2	36
10	The evolving concept of cell identity in the single cell era. <i>Development (Cambridge)</i> , 2019, 146, .	1.2	115
11	Multi-modal image cytometry approach “ From dynamic to whole organ imaging. <i>Cellular Immunology</i> , 2019, 344, 103946.	1.4	3
12	Recording development with single cell dynamic lineage tracing. <i>Development (Cambridge)</i> , 2019, 146, .	1.2	115
13	Exploring single cells in space and time during tissue development, homeostasis and regeneration. <i>Development (Cambridge)</i> , 2019, 146, .	1.2	51
14	Immunology Driven by Large-Scale Single-Cell Sequencing. <i>Trends in Immunology</i> , 2019, 40, 1011-1021.	2.9	62
15	High-Throughput Mapping of Long-Range Neuronal Projection Using In Situ Sequencing. <i>Cell</i> , 2019, 179, 772-786.e19.	13.5	146
16	Landscape of Noncoding RNA in Prostate Cancer. <i>Trends in Genetics</i> , 2019, 35, 840-851.	2.9	114
17	Revisiting airway epithelial remodeling in type 2 immunity: Beyond goblet cell metaplasia. <i>Journal of Allergy and Clinical Immunology</i> , 2019, 144, 1158-1160.	1.5	6
18	Transcriptome profiling of brain myeloid cells revealed activation of <i>Itgal</i> , <i>Trem1</i> , and <i>Spp1</i> in western diet-induced obesity. <i>Journal of Neuroinflammation</i> , 2019, 16, 169.	3.1	32
19	Implications for Tumor Microenvironment and Epithelial Crosstalk in the Management of Gastrointestinal Cancers. <i>Journal of Oncology</i> , 2019, 2019, 1-11.	0.6	2

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20	SpatialDB: a database for spatially resolved transcriptomes. <i>Nucleic Acids Research</i> , 2020, 48, D233-D237.	6.5	37
21	Studying immune to non-immune cell cross-talk using single-cell technologies. <i>Current Opinion in Systems Biology</i> , 2019, 18, 87-94.	1.3	5
22	A single-cell view of tissue regeneration in plants. <i>Current Opinion in Plant Biology</i> , 2019, 52, 149-154.	3.5	24
23	A Point of Inflection and Reflection on Systems Chemical Biology. <i>ACS Chemical Biology</i> , 2019, 14, 2497-2511.	1.6	8
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25	Network modeling of single-cell omics data: challenges, opportunities, and progresses. <i>Emerging Topics in Life Sciences</i> , 2019, 3, 379-398.	1.1	48
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58	The encyclopedia of maize kernel gene expression. <i>Journal of Integrative Plant Biology</i> , 2020, 62, 879-881.	4.1	6
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