Smita Krishnaswamy

List of Publications by Year in descending order

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Version: 2024-02-01

46 papers 6,245 citations

361413 20 h-index 36 g-index

68 all docs

68 docs citations

68 times ranked 11621 citing authors

#	Article	IF	CITATIONS
1	viSNE enables visualization of high dimensional single-cell data and reveals phenotypic heterogeneity of leukemia. Nature Biotechnology, 2013, 31, 545-552.	17.5	1,481
2	Recovering Gene Interactions from Single-Cell Data Using Data Diffusion. Cell, 2018, 174, 716-729.e27.	28.9	1,197
3	Normalization of mass cytometry data with bead standards. Cytometry Part A: the Journal of the International Society for Analytical Cytology, 2013, 83A, 483-494.	1.5	655
4	Visualizing structure and transitions in high-dimensional biological data. Nature Biotechnology, 2019, 37, 1482-1492.	17.5	597
5	Palladium-based mass tag cell barcoding with a doublet-filtering scheme and single-cell deconvolution algorithm. Nature Protocols, 2015, 10, 316-333.	12.0	466
6	Exploring single-cell data with deep multitasking neural networks. Nature Methods, 2019, 16, 1139-1145.	19.0	222
7	Conditional density-based analysis of T cell signaling in single-cell data. Science, 2014, 346, 1250689.	12.6	188
8	PD-1 marks dysfunctional regulatory T cells in malignant gliomas. JCI Insight, 2016, 1, .	5.0	182
9	A reservoir of stem-like CD8 ⁺ T cells in the tumor-draining lymph node preserves the ongoing antitumor immune response. Science Immunology, 2021, 6, eabg7836.	11.9	123
10	Manifold learning-based methods for analyzing single-cell RNA-sequencing data. Current Opinion in Systems Biology, 2018, 7, 36-46.	2.6	103
11	Quantifying the effect of experimental perturbations at single-cell resolution. Nature Biotechnology, 2021, 39, 619-629.	17.5	98
12	Endocrine-Exocrine Signaling Drives Obesity-Associated Pancreatic Ductal Adenocarcinoma. Cell, 2020, 181, 832-847.e18.	28.9	77
13	Transcriptomic and clonal characterization of T cells in the human central nervous system. Science Immunology, 2020, 5, .	11.9	73
14	Structural and developmental principles of neuropil assembly in C. elegans. Nature, 2021, 591, 99-104.	27.8	60
15	Macrophage Migration Inhibitory Factor Regulates U1 Small Nuclear RNP Immune Complex–Mediated Activation of the NLRP3 Inflammasome. Arthritis and Rheumatology, 2019, 71, 109-120.	5.6	59
16	Single-cell mass cytometry of TCR signaling: Amplification of small initial differences results in low ERK activation in NOD mice. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 16466-16471.	7.1	50
17	MLL-AF9 initiates transformation from fast-proliferating myeloid progenitors. Nature Communications, 2019, 10, 5767.	12.8	41
18	Uncovering axes of variation among single-cell cancer specimens. Nature Methods, 2020, 17, 302-310.	19.0	39

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19	Multiscale PHATE identifies multimodal signatures of COVID-19. Nature Biotechnology, 2022, 40, 681-691.	17.5	39
20	The landscape of pioneer factor activity reveals the mechanisms of chromatin reprogramming and genome activation. Molecular Cell, 2022, 82, 986-1002.e9.	9.7	38
21	Mapping Phenotypic Plasticity upon the Cancer Cell State Landscape Using Manifold Learning. Cancer Discovery, 2022, 12, 1847-1859.	9.4	26
22	Single cell immune profiling of dengue virus patients reveals intact immune responses to Zika virus with enrichment of innate immune signatures. PLoS Neglected Tropical Diseases, 2020, 14, e0008112.	3.0	20
23	Multiparameter Single Cell Profiling of Airway Inflammatory Cells. Cytometry Part B - Clinical Cytometry, 2017, 92, 12-20.	1.5	19
24	Identification and Analysis of Islet Antigen–Specific CD8+ T Cells with T Cell Libraries. Journal of Immunology, 2018, 201, 1662-1670.	0.8	19
25	Learning time-varying information flow from single-cell epithelial to mesenchymal transition data. PLoS ONE, 2018, 13, e0203389.	2.5	18
26	Dissecting alterations in human CD8+ T cells with aging by high-dimensional single cell mass cytometry. Clinical Immunology, 2019, 200, 24-30.	3.2	18
27	Modeling uniquely human gene regulatory function via targeted humanization of the mouse genome. Nature Communications, 2022, 13, 304.	12.8	16
28	Multimodal Data Visualization and Denoising with Integrated Diffusion. , 2021, 2021, .		15
29	Neural network predicts need for red blood cell transfusion for patients with acute gastrointestinal bleeding admitted to the intensive care unit. Scientific Reports, 2021, 11, 8827.	3.3	11
30	Visualizing Structure and Transitions for Biological Data Exploration. SSRN Electronic Journal, 0, , .	0.4	11
31	Recovering Gene Interactions from Single-Cell Data Using Data Diffusion. SSRN Electronic Journal, 0, ,	0.4	11
32	Coarse Graining of Data via Inhomogeneous Diffusion Condensation., 2019, 2019, 2624-2633.		9
33	Finding Archetypal Spaces Using Neural Networks. , 2019, , .		9
34	Harmonic Alignment. , 2020, 2020, 316-324.		9
35	IL-7 receptor alpha defines heterogeneity and signature of human effector memory CD8+ T cells in high dimensional analysis. Cellular Immunology, 2020, 355, 104155.	3.0	7
36	Population Genomics Approaches for Genetic Characterization of SARS-CoV-2 Lineages. Frontiers in Medicine, 2022, 9, 826746.	2.6	7

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37	1-deoxysphingolipids bind to COUP-TF to modulate lymphatic and cardiac cell development. Developmental Cell, 2021, 56, 3128-3145.e15.	7.0	6
38	Compressed Diffusion. , 2019, , .		5
39	Generating hard-to-obtain information from easy-to-obtain information: Applications in drug discovery and clinical inference. Patterns, 2021, 2, 100288.	5.9	5
40	Fixing Bias in Reconstruction-based Anomaly Detection with Lipschitz Discriminators. Journal of Signal Processing Systems, 2022, 94, 229-243.	2.1	4
41	Voices of biotech research. Nature Biotechnology, 2021, 39, 281-286.	17.5	3
42	TrajectoryNet: A Dynamic Optimal Transport Network for Modeling Cellular Dynamics. Proceedings of Machine Learning Research, 2020, 119, 9526-9536.	0.3	3
43	Modeling Global Dynamics from Local Snapshots with Deep Generative Neural Networks. , 2019, , .		1
44	MURAL: An Unsupervised Random Forest-Based Embedding for Electronic Health Record Data., 2021,,.		1
45	Learning General Transformations of Data for Out-of-Sample Extensions. , 2020, 2020, .		0
46	Abstract LT013: Endocrine-exocrine signaling is a driver of obesity-associated pancreatic ductal adenocarcinoma., 2021,,.		0