## **Bob Chen**

## List of Publications by Year in descending order

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759055 839398 18 903 12 18 citations h-index g-index papers 25 25 25 1448 docs citations all docs times ranked citing authors

#	Article	IF	Citations
1	Matrix stiffness enhances cancer-macrophage interactions and M2-like macrophage accumulation in the breast tumor microenvironment. Acta Biomaterialia, 2023, 163, 365-377.	4.1	17
2	Single-Cell Transcriptomics Reveals a Conserved Metaplasia Program in Pancreatic Injury. Gastroenterology, 2022, 162, 604-620.e20.	0.6	43
3	MTG16 regulates colonic epithelial differentiation, colitis, and tumorigenesis by repressing E protein transcription factors. JCI Insight, 2022, 7, .	2.3	9
4	Mining tasks and task characteristics from electronic health record audit logs with unsupervised machine learning. Journal of the American Medical Informatics Association: JAMIA, 2021, 28, 1168-1177.	2.2	19
5	Automated quality control and cell identification of droplet-based single-cell data using dropkick. Genome Research, 2021, 31, 1742-1752.	2.4	25
6	Processing single-cell RNA-seq data for dimension reduction-based analyses using open-source tools. STAR Protocols, 2021, 2, 100450.	0.5	8
7	Measuring Collaboration Through Concurrent Electronic Health Record Usage: Network Analysis Study. JMIR Medical Informatics, 2021, 9, e28998.	1.3	3
8	Deciphering the cancer microenvironment from bulk data with EcoTyper. Cell, 2021, 184, 5306-5308.	13.5	7
9	Differential pre-malignant programs and microenvironment chart distinct paths to malignancy in human colorectal polyps. Cell, 2021, 184, 6262-6280.e26.	13.5	125
10	Succinate Produced by Intestinal Microbes Promotes Specification of Tuft Cells to Suppress Ileal Inflammation. Gastroenterology, 2020, 159, 2101-2115.e5.	0.6	123
11	Coregulator Sin3a Promotes Postnatal Murine $\hat{l}^2$ -Cell Fitness by Regulating Genes in Ca2+ Homeostasis, Cell Survival, Vesicle Biosynthesis, Glucose Metabolism, and Stress Response. Diabetes, 2020, 69, 1219-1231.	0.3	9
12	Use of Single-Cell -Omic Technologies to Study the Gastrointestinal Tract and Diseases, From Single Cell Identities to Patient Features. Gastroenterology, 2020, 159, 453-466.e1.	0.6	17
13	Dual indexed library design enables compatibility of in-Drop single-cell RNA-sequencing with exAMP chemistry sequencing platforms. BMC Genomics, 2020, 21, 456.	1.2	22
14	ATAC-Me Captures Prolonged DNA Methylation of Dynamic Chromatin Accessibility Loci during Cell Fate Transitions. Molecular Cell, 2020, 77, 1350-1364.e6.	4.5	47
15	The Human Tumor Atlas Network: Charting Tumor Transitions across Space and Time at Single-Cell Resolution. Cell, 2020, 181, 236-249.	13.5	334
16	pyNVR: investigating factors affecting feature selection from scRNA-seq data for lineage reconstruction. Bioinformatics, 2019, 35, 2335-2337.	1.8	15
17	Single-Cell Computational Strategies for Lineage Reconstruction in Tissue Systems. Cellular and Molecular Gastroenterology and Hepatology, 2018, 5, 539-548.	2.3	33
18	Quantitative assessment of cell population diversity in single-cell landscapes. PLoS Biology, 2018, 16, e2006687.	2.6	40