Fan Bai

List of Publications by Year in descending order

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87	7,732	36	80
papers	citations	h-index	g-index
93	93	93	11965 citing authors
all docs	docs citations	times ranked	

#	Article	IF	Citations
1	Spatiotemporal Immune Landscape of Colorectal Cancer Liver Metastasis at Single-Cell Level. Cancer Discovery, 2022, 12, 134-153.	9.4	286
2	Cancer biology deciphered by single-cell transcriptomic sequencing. Protein and Cell, 2022, 13, 167-179.	11.0	17
3	A spatial and cellular distribution of rabies virus infection in the mouse brain revealed by fMOST and singleâ€cell RNA sequencing. Clinical and Translational Medicine, 2022, 12, e700.	4.0	6
4	Single Cell RNA Sequencing Identifies a Unique Inflammatory Macrophage Subset as a Druggable Target for Alleviating Acute Kidney Injury. Advanced Science, 2022, 9, e2103675.	11.2	37
5	Lung cancer scRNA-seq and lipidomics reveal aberrant lipid metabolism for early-stage diagnosis. Science Translational Medicine, 2022, 14, eabk2756.	12.4	57
6	Singleâ€cell RNA sequencing reveals the multiâ€cellular ecosystem in different radiological components of pulmonary partâ€solid nodules. Clinical and Translational Medicine, 2022, 12, e723.	4.0	7
7	Single-cell transcriptomics links malignant T cells to the tumor immune landscape in cutaneous T cell lymphoma. Nature Communications, 2022, 13, 1158.	12.8	29
8	Global transcriptomic characterization of T cells in individuals with chronic HIV-1 infection. Cell Discovery, 2022, 8, 29.	6.7	18
9	Expression Analysis of Same-Patient Metachronous and Synchronous Upper Tract and Bladder Urothelial Carcinoma. Letter Journal of Urology, 2022, , 101097JU0000000000002698.	0.4	0
10	Excessive deubiquitination of NLRP3-R779C variant contributes to very-early-onset inflammatory bowel disease development. Journal of Allergy and Clinical Immunology, 2021, 147, 267-279.	2.9	38
11	Metabolic-Pathway-Based Subtyping of Triple-Negative Breast Cancer Reveals Potential Therapeutic Targets. Cell Metabolism, 2021, 33, 51-64.e9.	16.2	211
12	Decoding the multicellular ecosystem of lung adenocarcinoma manifested as pulmonary subsolid nodules by single-cell RNA sequencing. Science Advances, 2021, 7, .	10.3	88
13	Single-cell DNA methylome analysis of circulating tumor cells. Chinese Journal of Cancer Research: Official Journal of China Anti-Cancer Association, Beijing Institute for Cancer Research, 2021, 33, 391-404.	2.2	6
14	The Effect of Face Mask Use on COVID-19 Models. Epidemiologia, 2021, 2, 75-83.	2.2	10
15	COVID-19 outbreak in Wuhan demonstrates the limitations of publicly available case numbers for epidemiological modeling. Epidemics, 2021, 34, 100439.	3.0	16
16	Flagella and Their Properties Affect the Transport and Deposition Behaviors of <i>Escherichia coli</i> in Quartz Sand. Environmental Science & Environm	10.0	26
17	COVID-19 immune features revealed by a large-scale single-cell transcriptome atlas. Cell, 2021, 184, 1895-1913.e19.	28.9	512
18	Probing bacterial cell wall growth by tracing wall-anchored protein complexes. Nature Communications, 2021, 12, 2160.	12.8	6

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19	Genomic and transcriptomic profiling of hepatoid adenocarcinoma of the stomach. Oncogene, 2021, 40, 5705-5717.	5.9	20
20	<i>IDH</i> Mutation Subgroup Status Associates with Intratumor Heterogeneity and the Tumor Microenvironment in Intrahepatic Cholangiocarcinoma. Advanced Science, 2021, 8, e2101230.	11.2	26
21	Single-cell sequencing of immune cells from anticitrullinated peptide antibody positive and negative rheumatoid arthritis. Nature Communications, 2021, 12, 4977.	12.8	73
22	A body map of somatic mutagenesis in morphologically normal human tissues. Nature, 2021, 597, 398-403.	27.8	107
23	The genomic architecture of EBV and infected gastric tissue from precursor lesions to carcinoma. Genome Medicine, 2021, 13, 146.	8.2	9
24	Membraneless organelles formed by liquid-liquid phase separation increase bacterial fitness. Science Advances, 2021, 7, eabh2929.	10.3	55
25	Genomic characterisation of pulmonary subsolid nodules: mutational landscape and radiological features. European Respiratory Journal, 2020, 55, 1901409.	6.7	42
26	A General Workflow for Characterization ofÂNernstian Dyes and Their Effects on Bacterial Physiology. Biophysical Journal, 2020, 118, 4-14.	0.5	28
27	Buckling behavior of cold-formed C/Z-section purlins incorporating the effects of diaphragm and the screw location. Advances in Structural Engineering, 2020, 23, 1114-1128.	2.4	0
28	Macroscopic somatic clonal expansion in morphologically normal human urothelium. Science, 2020, 370, 82-89.	12.6	115
29	Liver Immune Profiling Reveals Pathogenesis and Therapeutics for Biliary Atresia. Cell, 2020, 183, 1867-1883.e26.	28.9	70
30	Single-cell landscape of immunological responses in patients with COVID-19. Nature Immunology, 2020, 21, 1107-1118.	14.5	508
31	Single-cell transcriptomic analysis defines the interplay between tumor cells, viral infection, and the microenvironment in nasopharyngeal carcinoma. Cell Research, 2020, 30, 950-965.	12.0	111
32	Integrative genomic study of Chinese clear cell renal cell carcinoma reveals features associated with thrombus. Nature Communications, 2020, 11, 739.	12.8	39
33	Liveâ€cell fluorescence imaging reveals dynamic production and loss of bacterial flagella. Molecular Microbiology, 2020, 114, 279-291.	2.5	23
34	Mapping the spreading routes of lymphatic metastases in human colorectal cancer. Nature Communications, 2020, 11, 1993.	12.8	68
35	Potential therapeutic effects of dipyridamole in the severely ill patients with COVID-19. Acta Pharmaceutica Sinica B, 2020, 10, 1205-1215.	12.0	193
36	Somatic mutation landscape reveals differential variability of cell-of-origin for primary liver cancer. Heliyon, 2020, 6, e03350.	3.2	7

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37	Mucosal Profiling of Pediatric-Onset Colitis and IBD Reveals Common Pathogenics and Therapeutic Pathways. Cell, 2019, 179, 1160-1176.e24.	28.9	163
38	Graded regulation of cellular quiescence depth between proliferation and senescence by a lysosomal dimmer switch. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 22624-22634.	7.1	84
39	Indole Reverses Intrinsic Antibiotic Resistance by Activating a Novel Dual-Function Importer. MBio, 2019, 10, .	4.1	31
40	Inferring the Evolution and Progression of Small-Cell Lung Cancer by Single-Cell Sequencing of Circulating Tumor Cells. Clinical Cancer Research, 2019, 25, 5049-5060.	7.0	66
41	Genomic and Transcriptomic Profiling of Combined Hepatocellular and Intrahepatic Cholangiocarcinoma Reveals Distinct Molecular Subtypes. Cancer Cell, 2019, 35, 932-947.e8.	16.8	182
42	Genomic and Transcriptomic Landscape of Triple-Negative Breast Cancers: Subtypes and Treatment Strategies. Cancer Cell, 2019, 35, 428-440.e5.	16.8	571
43	Spatial clustering and common regulatory elements correlate with coordinated gene expression. PLoS Computational Biology, 2019, 15, e1006786.	3.2	26
44	A Streptococcus aquaporin acts as peroxiporin for efflux of cellular hydrogen peroxide and alleviation of oxidative stress. Journal of Biological Chemistry, 2019, 294, 4583-4595.	3.4	20
45	Comparison of Escherichia coli surface attachment methods for single-cell microscopy. Scientific Reports, 2019, 9, 19418.	3.3	11
46	ATP-Dependent Dynamic Protein Aggregation Regulates Bacterial Dormancy Depth Critical for Antibiotic Tolerance. Molecular Cell, 2019, 73, 143-156.e4.	9.7	221
47	Transcriptional profiles of different states of cancer stem cells in triple-negative breast cancer. Molecular Cancer, 2018, 17, 65.	19.2	48
48	Early metastasis detected in patients with multifocal pulmonary ground-glass opacities (GGOs). Thorax, 2018, 73, 290-292.	5.6	43
49	Molecular characterization of circulating tumor cells—from bench to bedside. Seminars in Cell and Developmental Biology, 2018, 75, 88-97.	5.0	38
50	Apj+ Vessels Drive Tumor Growth and Represent a Tractable Therapeutic Target. Cell Reports, 2018, 25, 1241-1254.e5.	6.4	26
51	Frequent pauses in Escherichia coli flagella elongation revealed by single cell real-time fluorescence imaging. Nature Communications, 2018, 9, 1885.	12.8	31
52	Leukaemic alterations of IKZF1 prime stemness and malignancy programs in human lymphocytes. Cell Death and Disease, 2018, 9, 526.	6.3	6
53	Photosynthetic Accumulation of Lutein in Auxenochlorella protothecoides after Heterotrophic Growth. Marine Drugs, 2018, 16, 283.	4.6	32
54	Active efflux in dormant bacterial cells – New insights into antibiotic persistence. Drug Resistance Updates, 2017, 30, 7-14.	14.4	38

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55	Single-cell sequencing deciphers a convergent evolution of copy number alterations from primary to circulating tumor cells. Genome Research, 2017, 27, 1312-1322.	5.5	81
56	Single-Cell Real-Time Visualization and Quantification of Perylene Bioaccumulation in Microorganisms. Environmental Science &	10.0	3
57	Genomic comparison of esophageal squamous cell carcinoma and its precursor lesions by multi-region whole-exome sequencing. Nature Communications, 2017, 8, 524.	12.8	103
58	Genomic signatures of pancreatic adenosquamous carcinoma (PASC). Journal of Pathology, 2017, 243, 155-159.	4.5	43
59	Mutagenic Factors and Complex Clonal Relationship of Multifocal Urothelial Cell Carcinoma. European Urology, 2017, 71, 841-843.	1.9	20
60	Multi-region and single-cell sequencing reveal variable genomic heterogeneity in rectal cancer. BMC Cancer, 2017, 17, 787.	2.6	30
61	A race to uncover a panoramic view of primary liver cancer. Cancer Biology and Medicine, 2017, 14, 335.	3.0	7
62	Length-dependent flagellar growth of Vibrio alginolyticus revealed by real time fluorescent imaging. ELife, 2017, 6, .	6.0	27
63	Enhanced Efflux Activity Facilitates Drug Tolerance in Dormant Bacterial Cells. Molecular Cell, 2016, 62, 284-294.	9.7	317
64	Dynamics of <i>Escherichia coli</i> 's passive response to a sudden decrease in external osmolarity. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E5838-E5846.	7.1	57
65	Bacterial Flagellar Motor Switch in Response to CheY-P Regulation and Motor Structural Alterations. Biophysical Journal, 2016, 110, 1411-1420.	0.5	12
66	Variable Intra-Tumor Genomic Heterogeneity of Multiple Lesions in Patients With Hepatocellular Carcinoma. Gastroenterology, 2016, 150, 998-1008.	1.3	178
67	A Delicate Nanoscale Motor Made by Natureâ€"The Bacterial Flagellar Motor. Advanced Science, 2015, 2, 1500129.	11.2	23
68	Real-Time Visualization of Perylene Nanoclusters in Water and Their Partitioning to Graphene Surface and Macrophage Cells. Environmental Science & Env	10.0	17
69	Single-cell analyses of circulating tumor cells. Cancer Biology and Medicine, 2015, 12, 184-92.	3.0	24
70	TGF-β–induced epithelial-to-mesenchymal transition proceeds through stepwise activation of multiple feedback loops. Science Signaling, 2014, 7, ra91.	3.6	395
71	Assembly and stoichiometry of <scp>FliF</scp> and <scp>FlhA</scp> in <scp><i>S</i></scp> <i>almonella</i> flagellar basal body. Molecular Microbiology, 2014, 91, 1214-1226.	2.5	86
72	Bacterial persistence. Science China Chemistry, 2014, 57, 1625-1633.	8.2	2

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73	Assembly dynamics and the roles of Flil ATPase of the bacterial flagellar export apparatus. Scientific Reports, 2014, 4, 6528.	3.3	72
74	Thermophoretic Manipulation of DNA Translocation through Nanopores. ACS Nano, 2013, 7, 538-546.	14.6	77
75	Populational Heterogeneity vs. Temporal Fluctuation in Escherichia coli Flagellar Motor Switching. Biophysical Journal, 2013, 105, 2123-2129.	0.5	11
76	Reproducible copy number variation patterns among single circulating tumor cells of lung cancer patients. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 21083-21088.	7.1	396
77	Conformational Spread in the Flagellar Motor Switch: A Model Study. PLoS Computational Biology, 2012, 8, e1002523.	3.2	13
78	Coupling between Switching Regulation and Torque Generation in Bacterial Flagellar Motor. Physical Review Letters, 2012, 108, 178105.	7.8	28
79	Probing Meiotic Recombination and Aneuploidy of Single Sperm Cells by Whole-Genome Sequencing. Science, 2012, 338, 1627-1630.	12.6	273
80	Steps and Bumps: Precision Extraction of Discrete States of Molecular Machines. Biophysical Journal, 2011, 101, 477-485.	0.5	29
81	Non-genetic individuality inEscherichia colimotor switching. Physical Biology, 2011, 8, 024001.	1.8	7
82	Conformational Spread as a Mechanism for Cooperativity in the Bacterial Flagellar Switch. Science, 2010, 327, 685-689.	12.6	176
83	Model Studies of the Dynamics of Bacterial Flagellar Motors. Biophysical Journal, 2009, 96, 3154-3167.	0.5	22
84	A Programmable Optical Angle Clamp for Rotary Molecular Motors. Biophysical Journal, 2007, 93, 264-275.	0.5	30
85	Stoichiometry and turnover in single, functioning membrane protein complexes. Nature, 2006, 443, 355-358.	27.8	559
86	Torque-speed relationship of the bacterial flagellar motor. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 1260-1265.	7.1	103
87	Slow Protein Conformational Change, Allostery and Network Dynamics. , 0, , .		1