

Bo Li

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6519619/publications.pdf>

Version: 2024-02-01

18
papers

10,505
citations

623188

14
h-index

752256

20
g-index

22
all docs

22
docs citations

22
times ranked

11848
citing authors

#	ARTICLE	IF	CITATIONS
1	Selective delivery of low-affinity IL-2 to PD-1+ T cells rejuvenates antitumor immunity with reduced toxicity. <i>Journal of Clinical Investigation</i> , 2022, 132, .	3.9	38
2	Concurrent delivery of immune checkpoint blockade modulates T cell dynamics to enhance neoantigen vaccine-generated antitumor immunity. <i>Nature Cancer</i> , 2022, 3, 437-452.	5.7	19
3	Comprehensive Characterizations of Immune Receptor Repertoire in Tumors and Cancer Immunotherapy Studies. <i>Cancer Immunology Research</i> , 2022, 10, 788-799.	1.6	10
4	GIANA allows computationally-efficient TCR clustering and multi-disease repertoire classification by isometric transformation. <i>Nature Communications</i> , 2021, 12, 4699.	5.8	30
5	AutoCAT: automated cancer-associated TCRs discovery from TCR-seq data. <i>Bioinformatics</i> , 2021, , .	1.8	1
6	Investigation of Antigen-Specific T-Cell Receptor Clusters in Human Cancers. <i>Clinical Cancer Research</i> , 2020, 26, 1359-1371.	3.2	90
7	A peripheral immune signature of responsiveness to PD-1 blockade in patients with classical Hodgkin lymphoma. <i>Nature Medicine</i> , 2020, 26, 1468-1479.	15.2	87
8	De novo prediction of cancer-associated T cell receptors for noninvasive cancer detection. <i>Science Translational Medicine</i> , 2020, 12, .	5.8	59
9	TIMER2.0 for analysis of tumor-infiltrating immune cells. <i>Nucleic Acids Research</i> , 2020, 48, W509-W514.	6.5	2,546
10	A PoleP286R mouse model of endometrial cancer recapitulates high mutational burden and immunotherapy response. <i>JCI Insight</i> , 2020, 5, .	2.3	25
11	Landscape of B cell immunity and related immune evasion in human cancers. <i>Nature Genetics</i> , 2019, 51, 560-567.	9.4	115
12	Immune receptor repertoires in pediatric and adult acute myeloid leukemia. <i>Genome Medicine</i> , 2019, 11, 73.	3.6	38
13	Evaluation of immune repertoire inference methods from RNA-seq data. <i>Nature Biotechnology</i> , 2018, 36, 1034-1034.	9.4	7
14	Defining T Cell States Associated with Response to Checkpoint Immunotherapy in Melanoma. <i>Cell</i> , 2018, 175, 998-1013.e20.	13.5	1,260
15	Ultrasensitive detection of TCR hypervariable-region sequences in solid-tissue RNA-seq data. <i>Nature Genetics</i> , 2017, 49, 482-483.	9.4	66
16	TIMER: A Web Server for Comprehensive Analysis of Tumor-Infiltrating Immune Cells. <i>Cancer Research</i> , 2017, 77, e108-e110.	0.4	4,049
17	Comprehensive analyses of tumor immunity: implications for cancer immunotherapy. <i>Genome Biology</i> , 2016, 17, 174.	3.8	1,768
18	Landscape of tumor-infiltrating T cell repertoire of human cancers. <i>Nature Genetics</i> , 2016, 48, 725-732.	9.4	288