

Binbin Wang

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8562847/publications.pdf>

Version: 2024-02-01

14
papers

5,683
citations

840119

11
h-index

1125271

13
g-index

20
all docs

20
docs citations

20
times ranked

6419
citing authors

#	ARTICLE	IF	CITATIONS
1	TIMER: A Web Server for Comprehensive Analysis of Tumor-Infiltrating Immune Cells. <i>Cancer Research</i> , 2017, 77, e108-e110.	0.4	4,049
2	TISCH: a comprehensive web resource enabling interactive single-cell transcriptome visualization of tumor microenvironment. <i>Nucleic Acids Research</i> , 2021, 49, D1420-D1430.	6.5	481
3	Landscape of tumor-infiltrating T cell repertoire of human cancers. <i>Nature Genetics</i> , 2016, 48, 725-732.	9.4	288
4	Integrative analysis of pooled CRISPR genetic screens using MAGeCKFlute. <i>Nature Protocols</i> , 2019, 14, 756-780.	5.5	260
5	Cistrome Cancer: A Web Resource for Integrative Gene Regulation Modeling in Cancer. <i>Cancer Research</i> , 2017, 77, e19-e22.	0.4	130
6	Landscape of B cell immunity and related immune evasion in human cancers. <i>Nature Genetics</i> , 2019, 51, 560-567.	9.4	115
7	Synthetic Lethal and Resistance Interactions with BET Bromodomain Inhibitors in Triple-Negative Breast Cancer. <i>Molecular Cell</i> , 2020, 78, 1096-1113.e8.	4.5	114
8	In vivo CRISPR screens identify the E3 ligase Cop1 as a modulator of macrophage infiltration and cancer immunotherapy target. <i>Cell</i> , 2021, 184, 5357-5374.e22.	13.5	79
9	Ultrasensitive detection of TCR hypervariable-region sequences in solid-tissue RNA-seq data. <i>Nature Genetics</i> , 2017, 49, 482-483.	9.4	66
10	Immune receptor repertoires in pediatric and adult acute myeloid leukemia. <i>Genome Medicine</i> , 2019, 11, 73.	3.6	38
11	Systematic characterization of mutations altering protein degradation in human cancers. <i>Molecular Cell</i> , 2021, 81, 1292-1308.e11.	4.5	36
12	CRISPR Screens Identify Essential Cell Growth Mediators in BRAF Inhibitor-resistant Melanoma. <i>Genomics, Proteomics and Bioinformatics</i> , 2020, 18, 26-40.	3.0	14
13	Abstract LB-264: Landscape of tumor-infiltrating T-cell repertoire of human cancers. , 2016, , .		2
14	<i>In vivo</i> CRISPR Screens Identify E3 Ligase Cop1 as a Modulator of Macrophage Infiltration and Cancer Immunotherapy Target. <i>SSRN Electronic Journal</i> , 0, , .	0.4	0