

Binbin Wang

List of Publications by Year in descending order

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

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14
papers

5,683
citations

840776
11
h-index

1125743
13
g-index

20
all docs

20
docs citations

20
times ranked

6419
citing authors

#	ARTICLE	IF	CITATIONS
1	TISCH: a comprehensive web resource enabling interactive single-cell transcriptome visualization of tumor microenvironment. Nucleic Acids Research, 2021, 49, D1420-D1430.	14.5	481
2	Systematic characterization of mutations altering protein degradation in human cancers. Molecular Cell, 2021, 81, 1292-1308.e11.	9.7	36
3	InÂvivo CRISPR screens identify the E3 ligase Cop1 as a modulator of macrophage infiltration and cancer immunotherapy target. Cell, 2021, 184, 5357-5374.e22.	28.9	79
4	CRISPR Screens Identify Essential Cell Growth Mediators in BRAF Inhibitor-resistant Melanoma. Genomics, Proteomics and Bioinformatics, 2020, 18, 26-40.	6.9	14
5	Synthetic Lethal and Resistance Interactions with BET Bromodomain Inhibitors in Triple-Negative Breast Cancer. Molecular Cell, 2020, 78, 1096-1113.e8.	9.7	114
6	Integrative analysis of pooled CRISPR genetic screens using MAGeCKFlute. Nature Protocols, 2019, 14, 756-780.	12.0	260
7	Landscape of B cell immunity and related immune evasion in human cancers. Nature Genetics, 2019, 51, 560-567.	21.4	115
8	Immune receptor repertoires in pediatric and adult acute myeloid leukemia. Genome Medicine, 2019, 11, 73.	8.2	38
9	Ultrasensitive detection of TCR hypervariable-region sequences in solid-tissue RNAâ€seq data. Nature Genetics, 2017, 49, 482-483.	21.4	66
10	Cistrome Cancer: A Web Resource for Integrative Gene Regulation Modeling in Cancer. Cancer Research, 2017, 77, e19-e22.	0.9	130
11	TIMER: A Web Server for Comprehensive Analysis of Tumor-Infiltrating Immune Cells. Cancer Research, 2017, 77, e108-e110.	0.9	4,049
12	Landscape of tumor-infiltrating T cell repertoire of human cancers. Nature Genetics, 2016, 48, 725-732.	21.4	288
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 <i>Cop1</i> as a Modulator of Macrophage Infiltration and Cancer Immunotherapy Target. SSRN Electronic Journal, 0, , .	0.4	0