

Yupeng Cun

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4426803/publications.pdf>

Version: 2024-02-01

10
papers

2,407
citations

1039406

9
h-index

1372195

10
g-index

10
all docs

10
docs citations

10
times ranked

4744
citing authors

#	ARTICLE	IF	CITATIONS
1	Characterizing genetic intra-tumor heterogeneity across 2,658 human cancer genomes. <i>Cell</i> , 2021, 184, 2239-2254.e39.	13.5	260
2	Integrative genomic profiling of large-cell neuroendocrine carcinomas reveals distinct subtypes of high-grade neuroendocrine lung tumors. <i>Nature Communications</i> , 2018, 9, 1048.	5.8	254
3	Copy-number analysis and inference of subclonal populations in cancer genomes using ScIust. <i>Nature Protocols</i> , 2018, 13, 1488-1501.	5.5	51
4	Two mouse models reveal an actionable PARP1 dependence in aggressive chronic lymphocytic leukemia. <i>Nature Communications</i> , 2017, 8, 153.	5.8	39
5	Transcriptomic profile of tobacco in response to Tomato zonate spot orthotospovirus infection. <i>Virology Journal</i> , 2017, 14, 153.	1.4	9
6	Comprehensive genomic profiles of small cell lung cancer. <i>Nature</i> , 2015, 524, 47-53.	13.7	1,634
7	netClass: an R-package for network based, integrative biomarker signature discovery. <i>Bioinformatics</i> , 2014, 30, 1325-1326.	1.8	22
8	Network and Data Integration for Biomarker Signature Discovery via Network Smoothed T-Statistics. <i>PLoS ONE</i> , 2013, 8, e73074.	1.1	63
9	Prognostic gene signatures for patient stratification in breast cancer - accuracy, stability and interpretability of gene selection approaches using prior knowledge on protein-protein interactions. <i>BMC Bioinformatics</i> , 2012, 13, 69.	1.2	45
10	Biomarker Gene Signature Discovery Integrating Network Knowledge. <i>Biology</i> , 2012, 1, 5-17.	1.3	30