

Qingbo S Wang

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6755047/publications.pdf>

Version: 2024-02-01

14
papers

8,701
citations

759055

12
h-index

940416

16
g-index

22
all docs

22
docs citations

22
times ranked

20342
citing authors

#	ARTICLE	IF	CITATIONS
1	Methods for statistical fine-mapping and their applications to auto-immune diseases. <i>Seminars in Immunopathology</i> , 2022, 44, 101-113.	2.8	7
2	Centers for Mendelian Genomics: A decade of facilitating gene discovery. <i>Genetics in Medicine</i> , 2022, 24, 784-797.	1.1	44
3	High-resolution Slide-seqV2 spatial transcriptomics enables discovery of disease-specific cell neighborhoods and pathways. <i>iScience</i> , 2022, 25, 104097.	1.9	32
4	Leveraging supervised learning for functionally informed fine-mapping of cis-eQTLs identifies an additional 20,913 putative causal eQTLs. <i>Nature Communications</i> , 2021, 12, 3394.	5.8	44
5	Mapping the human genetic architecture of COVID-19. <i>Nature</i> , 2021, 600, 472-477.	13.7	640
6	Addendum: The mutational constraint spectrum quantified from variation in 141,456 humans. <i>Nature</i> , 2021, 597, E3-E4.	13.7	45
7	A massive effort links protein-coding gene variants to health. <i>Nature</i> , 2021, 599, 561-563.	13.7	1
8	Principles of Spatial Transcriptomics Analysis: A Practical Walk-Through in Kidney Tissue. <i>Frontiers in Physiology</i> , 2021, 12, 809346.	1.3	14
9	Evaluating drug targets through human loss-of-function genetic variation. <i>Nature</i> , 2020, 581, 459-464.	13.7	115
10	The mutational constraint spectrum quantified from variation in 141,456 humans. <i>Nature</i> , 2020, 581, 434-443.	13.7	6,140
11	Landscape of multi-nucleotide variants in 125,748 human exomes and 15,708 genomes. <i>Nature Communications</i> , 2020, 11, 2539.	5.8	98
12	A structural variation reference for medical and population genetics. <i>Nature</i> , 2020, 581, 444-451.	13.7	614
13	Transcript expression-aware annotation improves rare variant interpretation. <i>Nature</i> , 2020, 581, 452-458.	13.7	142
14	The effect of LRRK2 loss-of-function variants in humans. <i>Nature Medicine</i> , 2020, 26, 869-877.	15.2	79