Qingbo S Wang

List of Publications by Year
in descending order

14 papers	8,701 citations	758635 12 h-index	940134 16 g-index
22 all docs	22 docs citations	22 times ranked	20342 citing authors

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