

Qiaozhen Ye

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4283296/publications.pdf>

Version: 2024-02-01

10
papers

1,120
citations

1040056

9
h-index

1474206

9
g-index

18
all docs

18
docs citations

18
times ranked

1513
citing authors

#	ARTICLE	IF	CITATIONS
1	Control of bacterial immune signaling by a WYL domain transcription factor. <i>Nucleic Acids Research</i> , 2022, 50, 5239-5250.	14.5	23
2	Structure and activity of a bacterial defense-associated 3'→5' exonuclease. <i>Protein Science</i> , 2022, 31, .	7.6	2
3	The SARS-CoV-2 nucleocapsid phosphoprotein forms mutually exclusive condensates with RNA and the membrane-associated M protein. <i>Nature Communications</i> , 2021, 12, 502.	12.8	307
4	Structural Basis for SARS-CoV-2 Nucleocapsid Protein Recognition by Single-Domain Antibodies. <i>Frontiers in Immunology</i> , 2021, 12, 719037.	4.8	35
5	HORMA Domain Proteins and a Trip13-like ATPase Regulate Bacterial cGAS-like Enzymes to Mediate Bacteriophage Immunity. <i>Molecular Cell</i> , 2020, 77, 709-722.e7.	9.7	116
6	Structure and Mechanism of a Cyclic Trinucleotide-Activated Bacterial Endonuclease Mediating Bacteriophage Immunity. <i>Molecular Cell</i> , 2020, 77, 723-733.e6.	9.7	148
7	Architecture and self-assembly of the SARS-CoV-2 nucleocapsid protein. <i>Protein Science</i> , 2020, 29, 1890-1901.	7.6	218
8	A conserved filamentous assembly underlies the structure of the meiotic chromosome axis. <i>ELife</i> , 2019, 8, .	6.0	100
9	TRIP13 and APC15 drive mitotic exit by turnover of interphase- and unattached kinetochore-produced MCC. <i>Nature Communications</i> , 2018, 9, 4354.	12.8	39
10	The AAA + ATPase TRIP13 remodels HORMA domains through N-terminal engagement and unfolding. <i>EMBO Journal</i> , 2017, 36, 2419-2434.	7.8	69