

# Dongli Wang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1232544/publications.pdf>

Version: 2024-02-01

12  
papers

599  
citations

840728

11  
h-index

1199563

12  
g-index

12  
all docs

12  
docs citations

12  
times ranked

1265  
citing authors

#	ARTICLE	IF	CITATIONS
1	Structural insights into the assembly and activation of IL-1 $\beta$ with its receptors. <i>Nature Immunology</i> , 2010, 11, 905-911.	14.5	136
2	Structural basis for R-spondin recognition by LGR4/5/6 receptors. <i>Genes and Development</i> , 2013, 27, 1339-1344.	5.9	110
3	Structural basis for the neutralization of MERS-CoV by a human monoclonal antibody MERS-27. <i>Scientific Reports</i> , 2015, 5, 13133.	3.3	63
4	Structural Definition of a Unique Neutralization Epitope on the Receptor-Binding Domain of MERS-CoV Spike Glycoprotein. <i>Cell Reports</i> , 2018, 24, 441-452.	6.4	57
5	Structural basis of dimerization and dual W-box DNA recognition by rice WRKY domain. <i>Nucleic Acids Research</i> , 2019, 47, 4308-4318.	14.5	56
6	A designer rice NLR immune receptor confers resistance to the rice blast fungus carrying noncorresponding avirulence effectors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	48
7	Identification of residues on human receptor DPP4 critical for MERS-CoV binding and entry. <i>Virology</i> , 2014, 471-473, 49-53.	2.4	44
8	Targeted inhibition of activated protein C by a non-active-site inhibitory antibody to treat hemophilia. <i>Nature Communications</i> , 2020, 11, 2992.	12.8	23
9	Structural basis for the specific recognition of IL-18 by its alpha receptor. <i>FEBS Letters</i> , 2014, 588, 3838-3843.	2.8	21
10	Structure-guided analysis of Arabidopsis JASMONATE-INDUCED OXYGENASE (JOX) 2 reveals key residues for recognition of jasmonic acid substrate by plant JOXs. <i>Molecular Plant</i> , 2021, 14, 820-828.	8.3	20
11	Crystal Structure of Human ISG15 Protein in Complex with Influenza B Virus NS1B. <i>Journal of Biological Chemistry</i> , 2011, 286, 30258-30262.	3.4	13
12	Crystal structures of <i>Magnaporthe oryzae</i> trehalose-6-phosphate synthase (MoTps1) suggest a model for catalytic process of Tps1. <i>Biochemical Journal</i> , 2019, 476, 3227-3240.	3.7	8