

# Wolfgang Maier

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6924239/publications.pdf>

Version: 2024-02-01

11  
papers

630  
citations

1040056

9  
h-index

1281871

11  
g-index

15  
all docs

15  
docs citations

15  
times ranked

490  
citing authors

#	ARTICLE	IF	CITATIONS
1	Matrix metalloproteinase 10 is linked to the risk of progression to dementia of the Alzheimer's type. <i>Brain</i> , 2022, 145, 2507-2517.	7.6	16
2	Selection Analysis Identifies Clusters of Unusual Mutational Changes in Omicron Lineage BA.1 That Likely Impact Spike Function. <i>Molecular Biology and Evolution</i> , 2022, 39, .	8.9	84
3	The Galaxy platform for accessible, reproducible and collaborative biomedical analyses: 2022 update. <i>Nucleic Acids Research</i> , 2022, 50, W345-W351.	14.5	328
4	A novel membrane complex is required for docking and regulated exocytosis of lysosome-related organelles in <i>Tetrahymena thermophila</i> . <i>PLoS Genetics</i> , 2022, 18, e1010194.	3.5	6
5	Ready-to-use public infrastructure for global SARS-CoV-2 monitoring. <i>Nature Biotechnology</i> , 2021, 39, 1178-1179.	17.5	21
6	No more business as usual: Agile and effective responses to emerging pathogen threats require open data and open analytics. <i>PLoS Pathogens</i> , 2020, 16, e1008643.	4.7	22
7	Mutual antagonism between Hippo signaling and cyclin E drives intracellular pattern formation. <i>Journal of Cell Biology</i> , 2020, 219, .	5.2	7
8	LF4/MOK and a CDK-related kinase regulate the number and length of cilia in <i>Tetrahymena</i> . <i>PLoS Genetics</i> , 2019, 15, e1008099.	3.5	27
9	Two Antagonistic Hippo Signaling Circuits Set the Division Plane at the Medial Position in the Ciliate <i>Tetrahymena</i> . <i>Genetics</i> , 2019, 211, 651-663.	2.9	12
10	The Hippo Pathway Maintains the Equatorial Division Plane in the Ciliate <i>Tetrahymena</i> . <i>Genetics</i> , 2017, 206, 873-888.	2.9	21
11	<i>C. elegans</i> DAF-16/FOXO interacts with TGF- $\beta$ /BMP signaling to induce germline tumor formation via mTORC1 activation. <i>PLoS Genetics</i> , 2017, 13, e1006801.	3.5	23