

# Matthias Haffke

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1498148/publications.pdf>

Version: 2024-02-01

13  
papers

311  
citations

840776

11  
h-index

1199594

12  
g-index

13  
all docs

13  
docs citations

13  
times ranked

534  
citing authors

#	ARTICLE	IF	CITATIONS
1	Development of a biochemical and biophysical suite for integral membrane protein targets: A review. <i>Protein Expression and Purification</i> , 2020, 167, 105545.	1.3	7
2	Super Potent Bispecific Llama VHH Antibodies Neutralize HIV via a Combination of gp41 and gp120 Epitopes. <i>Antibodies</i> , 2019, 8, 38.	2.5	25
3	Chaperonin CCT checkpoint function in basal transcription factor TFIID assembly. <i>Nature Structural and Molecular Biology</i> , 2018, 25, 1119-1127.	8.2	43
4	Zooming in on Transcription Preinitiation. <i>Journal of Molecular Biology</i> , 2016, 428, 2581-2591.	4.2	36
5	ACEMBL Tool-Kits for High-Throughput Multigene Delivery and Expression in Prokaryotic and Eukaryotic Hosts. <i>Advances in Experimental Medicine and Biology</i> , 2016, 896, 27-42.	1.6	17
6	Cytoplasmic TAF2â€“TAF8â€“TAF10 complex provides evidence for nuclear holoâ€“TFIID assembly from preformed submodules. <i>Nature Communications</i> , 2015, 6, 6011.	12.8	77
7	Characterization and Production of Protein Complexes by Co-expression in <i>Escherichia coli</i> . <i>Methods in Molecular Biology</i> , 2015, 1261, 63-89.	0.9	14
8	More pieces to the puzzle: recent structural insights into class II transcription initiation. <i>Current Opinion in Structural Biology</i> , 2014, 24, 91-97.	5.7	22
9	Crystal structure of human TAF5/TAF6/TAF9 complex at 2.7 Å resolution. <i>Acta Crystallographica Section A: Foundations and Advances</i> , 2014, 70, C1165-C1165.	0.1	0
10	The MultiBac Protein Complex Production Platform at the EMBL. <i>Journal of Visualized Experiments</i> , 2013, , e50159.	0.3	22
11	Tandem Recombineering by SLIC Cloning and Cre-LoxP Fusion to Generate Multigene Expression Constructs for Protein Complex Research. <i>Methods in Molecular Biology</i> , 2013, 1073, 131-140.	0.9	23
12	Structure of the effector-binding domain of the LysR-type transcription factor RovM from <i>Yersinia pseudotuberculosis</i> . <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2011, 67, 81-90.	2.5	13
13	Structures of the nucleotide-binding domain of the human ABCB6 transporter and its complexes with nucleotides. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2010, 66, 979-987.	2.5	12