

# Stefan Seemayer

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5999291/publications.pdf>

Version: 2024-02-01

5  
papers

1,308  
citations

1684188

5  
h-index

2053705

5  
g-index

6  
all docs

6  
docs citations

6  
times ranked

2088  
citing authors

#	ARTICLE	IF	CITATIONS
1	A large-scale evaluation of computational protein function prediction. <i>Nature Methods</i> , 2013, 10, 221-227.	19.0	789
2	CCMpred – fast and precise prediction of protein residue-residue contacts from correlated mutations. <i>Bioinformatics</i> , 2014, 30, 3128-3130.	4.1	387
3	A structural model of the active ribosome-bound membrane protein insertase YidC. <i>ELife</i> , 2014, 3, e03035.	6.0	69
4	Homology-based inference sets the bar high for protein function prediction. <i>BMC Bioinformatics</i> , 2013, 14, S7.	2.6	38
5	Synthetic protein alignments by CCMgen quantify noise in residue-residue contact prediction. <i>PLoS Computational Biology</i> , 2018, 14, e1006526.	3.2	25