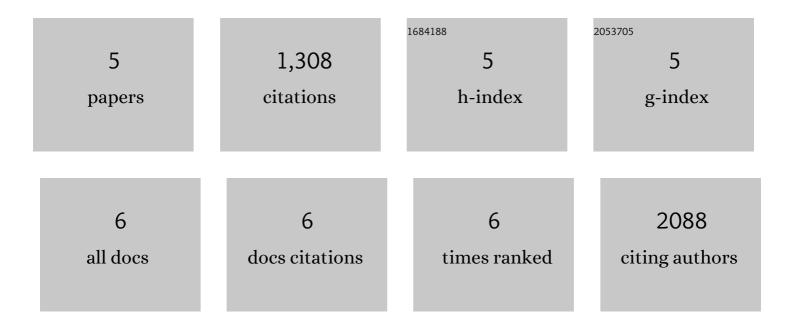
## Stefan Seemayer

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5999291/publications.pdf Version: 2024-02-01



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