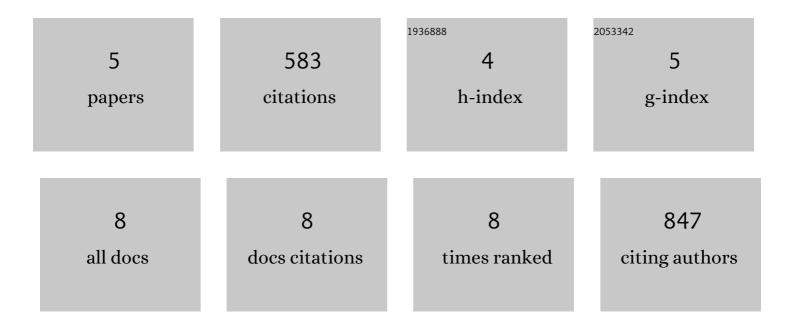
## Andreas Scheck

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3412334/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Macromolecular modeling and design in Rosetta: recent methods and frameworks. Nature Methods, 2020, 17, 665-680.	9.0	513
2	Rosetta FunFolDes – A general framework for the computational design of functional proteins. PLoS Computational Biology, 2018, 14, e1006623.	1.5	32
3	rstoolbox - a Python library for large-scale analysis of computational protein design data and structural bioinformatics. BMC Bioinformatics, 2019, 20, 240.	1.2	15
4	Computational design of anti-CRISPR proteins with improved inhibition potency. Nature Chemical Biology, 2020, 16, 725-730.	3.9	14
5	RosettaSurf—A surface-centric computational design approach. PLoS Computational Biology, 2022, 18, e1009178.	1.5	5