

# Sagar Bhogaraju

## List of Publications by Year in Descending Order

**Source:** <https://exaly.com/author-pdf/2393653/sagar-bhogaraju-publications-by-year.pdf>

**Version:** 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

23  
papers

1,501  
citations

17  
h-index

24  
g-index

24  
ext. papers

1,866  
ext. citations

18.2  
avg, IF

4.47  
L-index

#	Paper	IF	Citations
23	Structural basis for protein glutamylation by the Legionella pseudokinase SidJ. <i>Nature Communications</i> , <b>2021</b> , 12, 6174	17.4	0
22	Purification and crystal structure of human ODA16: Implications for ciliary import of outer dynein arms by the intraflagellar transport machinery. <i>Protein Science</i> , <b>2020</b> , 29, 1502-1510	6.3	6
21	Regulation of Phosphoribosyl-Linked Serine Ubiquitination by Deubiquitinases DupA and DupB. <i>Molecular Cell</i> , <b>2020</b> , 77, 164-179.e6	17.6	43
20	Inhibition of bacterial ubiquitin ligases by SidJ-calmodulin catalysed glutamylation. <i>Nature</i> , <b>2019</b> , 572, 382-386	50.4	58
19	A General Approach Towards Triazole-Linked Adenosine Diphosphate Ribosylated Peptides and Proteins. <i>Angewandte Chemie</i> , <b>2018</b> , 130, 1675-1678	3.6	3
18	Crystal structure of tetrameric human Rabin8 GEF domain. <i>Proteins: Structure, Function and Bioinformatics</i> , <b>2018</b> , 86, 405-413	4.2	2
17	A General Approach Towards Triazole-Linked Adenosine Diphosphate Ribosylated Peptides and Proteins. <i>Angewandte Chemie - International Edition</i> , <b>2018</b> , 57, 1659-1662	16.4	18
16	Insights into catalysis and function of phosphoribosyl-linked serine ubiquitination. <i>Nature</i> , <b>2018</b> , 557, 734-738	50.4	48
15	Structural basis for the recognition and degradation of host TRIM proteins by Salmonella effector SopA. <i>Nature Communications</i> , <b>2017</b> , 8, 14004	17.4	32
14	Intraflagellar transport proteins 172, 80, 57, 54, 38, and 20 form a stable tubulin-binding IFT-B2 complex. <i>EMBO Journal</i> , <b>2016</b> , 35, 773-90	13	116
13	Bacteria-host relationship: ubiquitin ligases as weapons of invasion. <i>Cell Research</i> , <b>2016</b> , 26, 499-510	24.7	72
12	Phosphoribosylation of Ubiquitin Promotes Serine Ubiquitination and Impairs Conventional Ubiquitination. <i>Cell</i> , <b>2016</b> , 167, 1636-1649.e13	56.2	157
11	Cell biology: Ubiquitination without E1 and E2 enzymes. <i>Nature</i> , <b>2016</b> , 533, 43-4	50.4	17
10	PLEKHM1 regulates autophagosome-lysosome fusion through HOPS complex and LC3/GABARAP proteins. <i>Molecular Cell</i> , <b>2015</b> , 57, 39-54	17.6	311
9	Crystal structure of a Chlamydomonas reinhardtii flagellar RabGAP TBC-domain at 1.8 Å resolution. <i>Proteins: Structure, Function and Bioinformatics</i> , <b>2014</b> , 82, 2282-7	4.2	3
8	A peek into the atomic details of thalidomide's clinical effects. <i>Nature Structural and Molecular Biology</i> , <b>2014</b> , 21, 739-40	17.6	2
7	Getting tubulin to the tip of the cilium: one IFT train, many different tubulin cargo-binding sites?. <i>BioEssays</i> , <b>2014</b> , 36, 463-7	4.1	31

6	Molecular basis of tubulin transport within the cilium by IFT74 and IFT81. <i>Science</i> , <b>2013</b> , 341, 1009-12	33.3	200
5	Intraflagellar transport complex structure and cargo interactions. <i>Cilia</i> , <b>2013</b> , 2, 10	5.5	85
4	Architecture and function of IFT complex proteins in ciliogenesis. <i>Differentiation</i> , <b>2012</b> , 83, S12-22	3.5	136
3	Crystal structure of the intraflagellar transport complex 25/27. <i>EMBO Journal</i> , <b>2011</b> , 30, 1907-18	13	84
2	Biochemical mapping of interactions within the intraflagellar transport (IFT) B core complex: IFT52 binds directly to four other IFT-B subunits. <i>Journal of Biological Chemistry</i> , <b>2011</b> , 286, 26344-52	5.4	58
1	Circularly permuted GTPase YqeH binds 30S ribosomal subunit: Implications for its role in ribosome assembly. <i>Biochemical and Biophysical Research Communications</i> , <b>2009</b> , 386, 602-6	3.4	19