## Christian Lw

## List of Publications by Citations

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

43
papers

1,089
citations

15
papers

1,617
ext. papers

1,617
ext. citations

1,617
avg, IF

1,617
L-index

#	Paper	IF	Citations
43	Understanding transport by the major facilitator superfamily (MFS): structures pave the way. <i>Nature Reviews Molecular Cell Biology</i> , <b>2016</b> , 17, 123-32	48.7	214
42	A saposin-lipoprotein nanoparticle system for membrane proteins. <i>Nature Methods</i> , <b>2016</b> , 13, 345-51	21.6	152
41	Structural basis for substrate transport in the GLUT-homology family of monosaccharide transporters. <i>Nature Structural and Molecular Biology</i> , <b>2013</b> , 20, 766-8	17.6	103
40	Selection, biophysical and structural analysis of synthetic nanobodies that effectively neutralize SARS-CoV-2. <i>Nature Communications</i> , <b>2020</b> , 11, 5588	17.4	73
39	Structural insights into substrate recognition in proton-dependent oligopeptide transporters. <i>EMBO Reports</i> , <b>2013</b> , 14, 804-10	6.5	69
38	Selectivity mechanism of a bacterial homolog of the human drug-peptide transporters PepT1 and PepT2. <i>Nature Structural and Molecular Biology</i> , <b>2014</b> , 21, 728-31	17.6	59
37	Crystal structure determination and functional characterization of the metallochaperone SlyD from Thermus thermophilus. <i>Journal of Molecular Biology</i> , <b>2010</b> , 398, 375-90	6.5	54
36	Saposin Lipid Nanoparticles: A Highly Versatile and Modular Tool for Membrane Protein Research. <i>Structure</i> , <b>2018</b> , 26, 345-355.e5	5.2	40
35	High-throughput stability screening for detergent-solubilized membrane proteins. <i>Scientific Reports</i> , <b>2019</b> , 9, 10379	4.9	35
34	Multispecific Substrate Recognition in a Proton-Dependent Oligopeptide Transporter. <i>Structure</i> , <b>2018</b> , 26, 467-476.e4	5.2	26
33	High-throughput analytical gel filtration screening of integral membrane proteins for structural studies. <i>Biochimica Et Biophysica Acta - General Subjects</i> , <b>2013</b> , 1830, 3497-508	4	25
32	Structure of Prototypic Peptide Transporter DtpA from E. coli in Complex with Valganciclovir Provides Insights into Drug Binding of Human PepT1. <i>Journal of the American Chemical Society</i> , <b>2019</b> , 141, 2404-2412	16.4	22
31	NMR relaxation unravels interdomain crosstalk of the two domain prolyl isomerase and chaperone SlyD. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , <b>2011</b> , 1814, 873-81	4	20
30	Optimisation of over-expression in E. coli and biophysical characterisation of human membrane protein synaptogyrin 1. <i>PLoS ONE</i> , <b>2012</b> , 7, e38244	3.7	17
29	Molecular insights into substrate recognition and catalytic mechanism of the chaperone and FKBP peptidyl-prolyl isomerase SlyD. <i>BMC Biology</i> , <b>2016</b> , 14, 82	7.3	17
28	Dissecting the Gene Expression, Localization, Membrane Topology, and Function of the Plasmodium falciparum STEVOR Protein Family. <i>MBio</i> , <b>2019</b> , 10,	7.8	15
27	Nanobody mediated crystallization of an archeal mechanosensitive channel. <i>PLoS ONE</i> , <b>2013</b> , 8, e77984	3.7	15

## (2021-2017)

26	Structure determination of a major facilitator peptide transporter: Inward facing PepTSt from Streptococcus thermophilus crystallized in space group P3121. <i>PLoS ONE</i> , <b>2017</b> , 12, e0173126	3.7	15
25	Structural and biochemical characterization of human PR70 in isolation and in complex with the scaffolding subunit of protein phosphatase 2A. <i>PLoS ONE</i> , <b>2014</b> , 9, e101846	3.7	11
24	Selection, biophysical and structural analysis of synthetic nanobodies that effectively neutralize SARS-C	CoV-2	11
23	Membrane Chemistry Tunes the Structure of a Peptide Transporter. <i>Angewandte Chemie - International Edition</i> , <b>2020</b> , 59, 19121-19128	16.4	11
22	Lipid-like Peptides can Stabilize Integral Membrane Proteins for Biophysical and Structural Studies. <i>ChemBioChem</i> , <b>2017</b> , 18, 1735-1742	3.8	10
21	High-resolution insights into binding of unfolded polypeptides by the PPIase chaperone SlpA. <i>FASEB Journal</i> , <b>2012</b> , 26, 4003-13	0.9	9
20	Tripeptide binding in a proton-dependent oligopeptide transporter. FEBS Letters, 2018, 592, 3239-3247	3.8	9
19	Structural Insights Into PfARO and Characterization of its Interaction With PfAIP. <i>Journal of Molecular Biology</i> , <b>2020</b> , 432, 878-896	6.5	7
18	Probing the Architecture of a Multi-PDZ Domain Protein: Structure of PDZK1 in Solution. <i>Structure</i> , <b>2018</b> , 26, 1522-1533.e5	5.2	7
17	Metal-mediated crystallization of the xylose transporter XylE from Escherichia coli in three different crystal forms. <i>Journal of Structural Biology</i> , <b>2013</b> , 184, 375-8	3.4	6
16	Structural role of essential light chains in the apicomplexan glideosome. <i>Communications Biology</i> , <b>2020</b> , 3, 568	6.7	6
15	Molecular basis of mRNA transport by a kinesin-1-atypical tropomyosin complex. <i>Genes and Development</i> , <b>2021</b> , 35, 976-991	12.6	6
14	In-depth interrogation of protein thermal unfolding data with MoltenProt. <i>Protein Science</i> , <b>2021</b> , 30, 201-217	6.3	6
13	Structural snapshots of human PepT1 and PepT2 reveal mechanistic insights into substrate and drug transport across epithelial membranes. <i>Science Advances</i> , <b>2021</b> , 7, eabk3259	14.3	3
12	Single-Molecule FRET of Membrane Transport Proteins. <i>ChemBioChem</i> , <b>2021</b> , 22, 2657-2671	3.8	3
11	Transient Expression of Recombinant Membrane-eGFP Fusion Proteins in HEK293 Cells. <i>Methods in Molecular Biology</i> , <b>2018</b> , 1850, 17-31	1.4	3
10	Structural role of essential light chains in the apicomplexan glideosome		2
9	Identification of novel inner membrane complex and apical annuli proteins of the malaria parasite Plasmodium falciparum. <i>Cellular Microbiology</i> , <b>2021</b> , 23, e13341	3.9	2

8	Moltenprot: A High-Throughput Analysis Platform to Assess Thermodynamic Stability of Membrane Proteins and Complexes. <i>Biophysical Journal</i> , <b>2019</b> , 116, 191a	2.9	2
7	Completing the family of human EH domains: Solution structure of the internal EH domain of Esynergin <i>Protein Science</i> , <b>2021</b> ,	6.3	1
6	Characterization of the complex of the lysosomal membrane transporter MFSD1 and its accessory subunit GLMP. <i>FASEB Journal</i> , <b>2020</b> , 34, 14695-14709	0.9	1
5	Membrane Chemistry Tunes the Structure of a Peptide Transporter. <i>Angewandte Chemie</i> , <b>2020</b> , 132, 19283-19290	3.6	1
4	Deamidation drives molecular aging of the SARS-CoV-2 spike protein receptor-binding motif. <i>Journal of Biological Chemistry</i> , <b>2021</b> , 297, 101175	5.4	O
3	PMRT1, a -Specific Parasite Plasma Membrane Transporter, Is Essential for Asexual and Sexual Blood Stage Development <i>MBio</i> , <b>2022</b> , e0062322	7.8	O
2	REktitelbild: Membrane Chemistry Tunes the Structure of a Peptide Transporter (Angew. Chem. 43/2020). <i>Angewandte Chemie</i> , <b>2020</b> , 132, 19528-19528	3.6	
1	Impact of distant peptide substrate residues on enzymatic activity of SlyD <i>Cellular and Molecular Life Sciences</i> , <b>2022</b> , 79, 138	10.3	