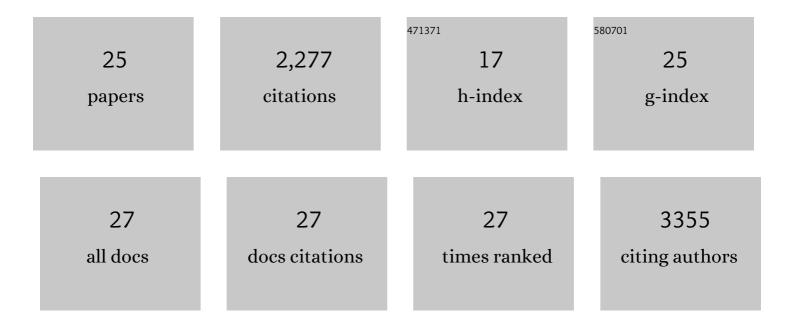
Matthias Zebisch

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8677676/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Cellular function and molecular structure of ecto-nucleotidases. Purinergic Signalling, 2012, 8, 437-502.	1.1	850
2	Notum deacylates Wnt proteins to suppress signalling activity. Nature, 2015, 519, 187-192.	13.7	348
3	Crystal Structure of the Human Ecto-5′-Nucleotidase (CD73): Insights into the Regulation of Purinergic Signaling. Structure, 2012, 20, 2161-2173.	1.6	164
4	Structural and molecular basis of ZNRF3/RNF43 transmembrane ubiquitin ligase inhibition by the Wnt agonist R-spondin. Nature Communications, 2013, 4, 2787.	5.8	161
5	Notum Is Required for Neural and Head Induction via Wnt Deacylation, Oxidation, and Inactivation. Developmental Cell, 2015, 32, 719-730.	3.1	155
6	Structure and functional properties of Norrin mimic Wnt for signalling with Frizzled4, Lrp5/6, and proteoglycan. ELife, 2015, 4, .	2.8	90
7	Crystallographic Evidence for a Domain Motion in Rat Nucleoside Triphosphate Diphosphohydrolase (NTPDase) 1. Journal of Molecular Biology, 2012, 415, 288-306.	2.0	73
8	Structural insight into signal conversion and inactivation by NTPDase2 in purinergic signaling. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 6882-6887.	3.3	71
9	Crystallographic Snapshots along the Reaction Pathway of Nucleoside Triphosphate Diphosphohydrolases. Structure, 2013, 21, 1460-1475.	1.6	44
10	Crystal structure of R-spondin 2 in complex with the ectodomains of its receptors LGR5 and ZNRF3. Journal of Structural Biology, 2015, 191, 149-155.	1.3	43
11	ZNRF3/RNF43 – A direct linkage of extracellular recognition and E3 ligase activity to modulate cell surface signalling. Progress in Biophysics and Molecular Biology, 2015, 118, 112-118.	1.4	40
12	Characterization of Rat NTPDase1, -2, and -3 Ectodomains Refolded from Bacterial Inclusion Bodies. Biochemistry, 2007, 46, 11945-11956.	1.2	34
13	The proâ€form of BMPâ€2 interferes with BMPâ€2 signalling by competing with BMPâ€2 for IA receptor binding. FEBS Journal, 2009, 276, 6386-6398.	2.2	34
14	Structure of the Dual-Mode Wnt Regulator Kremen1 and Insight into Ternary Complex Formation with LRP6 and Dickkopf. Structure, 2016, 24, 1599-1605.	1.6	32
15	Structures of <i>Legionella pneumophila</i> NTPDase1 in complex with polyoxometallates. Acta Crystallographica Section D: Biological Crystallography, 2014, 70, 1147-1154.	2.5	25
16	Crystal structure of NTPDase2 in complex with the sulfoanthraquinone inhibitor PSB-071. Journal of Structural Biology, 2014, 185, 336-341.	1.3	25
17	Structural Insight into Activation Mechanism of Toxoplasma gondii Nucleoside Triphosphate Diphosphohydrolases by Disulfide Reduction*. Journal of Biological Chemistry, 2012, 287, 3051-3066.	1.6	21
18	The Biochemical Properties of the Arabidopsis Ecto-Nucleoside Triphosphate Diphosphohydrolase AtAPY1 Contradict a Direct Role in Purinergic Signaling, PLoS ONE, 2015, 10, e0115832.	1.1	14

MATTHIAS ZEBISCH

#	Article	IF	CITATIONS
19	Contribution of the two domains of <i>E. coli</i> 5′â€nucleotidase to substrate specificity and catalysis. FEBS Letters, 2013, 587, 460-466.	1.3	13
20	New crystal forms of NTPDase1 from the bacterium <i>Legionella pneumophila</i> . Acta Crystallographica Section F: Structural Biology Communications, 2013, 69, 257-262.	0.7	13
21	Crystal structure and substrate binding mode of ectonucleotide phosphodiesterase/pyrophosphatase-3 (NPP3). Scientific Reports, 2018, 8, 10874.	1.6	9
22	Crystallization and preliminary X-ray analysis of the open form of human ecto-5′-nucleotidase (CD73). Acta Crystallographica Section F: Structural Biology Communications, 2012, 68, 1545-1549.	0.7	6
23	The ATP/ADP Substrate Specificity Switch between <i>Toxoplasma gondii</i> NTPDase1 and NTPDase3 is Caused by an Altered Mode of Binding of the Substrate Base. ChemBioChem, 2013, 14, 2292-2300.	1.3	5
24	Development of a robust crystallization platform for immune receptor TREM2 using a crystallization chaperone strategy. Protein Expression and Purification, 2021, 179, 105796.	0.6	4
25	Crystallization of ectonucleotide phosphodiesterase/pyrophosphatase-3 and orientation of the SMB domains in the full-length ectodomain. Acta Crystallographica Section F, Structural Biology Communications, 2018, 74, 696-703.	0.4	2