## Tomas Malinauskas

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Single-particle cryo-EM at atomic resolution. Nature, 2020, 587, 152-156.	13.7	572
2	Neutralizing nanobodies bind SARS-CoV-2 spike RBD and block interaction with ACE2. Nature Structural and Molecular Biology, 2020, 27, 846-854.	3.6	434
3	GABAA receptor signalling mechanisms revealed by structural pharmacology. Nature, 2019, 565, 454-459.	13.7	386
4	Neutralization of SARS-CoV-2 by Destruction of the Prefusion Spike. Cell Host and Microbe, 2020, 28, 445-454.e6.	5.1	298
5	Structural basis for the neutralization of SARS-CoV-2 by an antibody from a convalescent patient. Nature Structural and Molecular Biology, 2020, 27, 950-958.	3.6	268
6	Cryo-EM structure of the human $\hat{1}\pm 1\hat{1}^23\hat{1}^32$ GABAA receptor in a lipid bilayer. Nature, 2019, 565, 516-520.	13.7	264
7	A COVID-19 vaccine candidate using SpyCatcher multimerization of the SARS-CoV-2 spike protein receptor-binding domain induces potent neutralising antibody responses. Nature Communications, 2021, 12, 542.	5.8	200
8	Combined sequence-based and genetic mapping analysis of complex traits in outbred rats. Nature Genetics, 2013, 45, 767-775.	9.4	176
9	Whole-genome sequencing reveals host factors underlying critical COVID-19. Nature, 2022, 607, 97-103.	13.7	174
10	Neuropilins lock secreted semaphorins onto plexins in a ternary signaling complex. Nature Structural and Molecular Biology, 2012, 19, 1293-1299.	3.6	160
11	Structure and function of the Smoothened extracellular domain in vertebrate Hedgehog signaling. ELife, 2013, 2, e01340.	2.8	140
12	Modular mechanism of Wnt signaling inhibition by Wnt inhibitory factor 1. Nature Structural and Molecular Biology, 2011, 18, 886-893.	3.6	135
13	Structural and Functional Studies of LRP6 Ectodomain Reveal a Platform for Wnt Signaling. Developmental Cell, 2011, 21, 848-861.	3.1	109
14	Application of whole genome and RNA sequencing to investigate the genomic landscape of common variable immunodeficiency disorders. Clinical Immunology, 2015, 160, 301-314.	1.4	100
15	Extracellular modulators of Wnt signalling. Current Opinion in Structural Biology, 2014, 29, 77-84.	2.6	96
16	Glypicans shield the Wnt lipid moiety to enable signalling at a distance. Nature, 2020, 585, 85-90.	13.7	90
17	Structural Basis for Plexin Activation and Regulation. Neuron, 2016, 91, 548-560.	3.8	89
18	Genetic Control over mtDNA and Its Relationship to Major Depressive Disorder. Current Biology, 2015, 25, 3170-3177.	1.8	84

Tomas Malinauskas

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19	Structural insights into proteoglycan-shaped Hedgehog signaling. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 16420-16425.	3.3	79
20	Site-Specific Steric Control of SARS-CoV-2 Spike Glycosylation. Biochemistry, 2021, 60, 2153-2169.	1.2	54
21	Efficacy of a Plasmodium vivax Malaria Vaccine Using ChAd63 and Modified Vaccinia Ankara Expressing Thrombospondin-Related Anonymous Protein as Assessed with Transgenic Plasmodium berghei Parasites. Infection and Immunity, 2014, 82, 1277-1286.	1.0	53
22	The morphogen Sonic hedgehog inhibits its receptor Patched by a pincer grasp mechanism. Nature Chemical Biology, 2019, 15, 975-982.	3.9	52
23	Structural Insights into the Inhibition of Wnt Signaling by Cancer Antigen 5T4/Wnt-Activated Inhibitory Factor 1. Structure, 2014, 22, 612-620.	1.6	42
24	R-spondins engage heparan sulfate proteoglycans to potentiate WNT signaling. ELife, 2020, 9, .	2.8	37
25	Differential assembly diversifies GABAA receptor structures and signalling. Nature, 2022, 604, 190-194.	13.7	36
26	A calcium-sensing receptor mutation causing hypocalcemia disrupts a transmembrane salt bridge to activate β-arrestin–biased signaling. Science Signaling, 2018, 11, .	1.6	32
27	Anti-prothrombin autoantibodies enriched after infection with SARS-CoV-2 and influenced by strength of antibody response against SARS-CoV-2 proteins. PLoS Pathogens, 2021, 17, e1010118.	2.1	30
28	Genomes and phenomes of a population of outbred rats and its progenitors. Scientific Data, 2014, 1, 140011.	2.4	25
29	Calcium-sensing receptor residues with loss- and gain-of-function mutations are located in regions of conformational change and cause signalling bias. Human Molecular Genetics, 2018, 27, 3720-3733.	1.4	23
30	Simultaneous binding of Guidance Cues NET1 and RGM blocks extracellular NEO1 signaling. Cell, 2021, 184, 2103-2120.e31.	13.5	20
31	Tailoring a Combination Preerythrocytic Malaria Vaccine. Infection and Immunity, 2016, 84, 622-634.	1.0	18
32	Repulsive guidance molecules lock growth differentiation factor 5 in an inhibitory complex. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 15620-15631.	3.3	18
33	Docking of Fatty Acids into the WIF Domain of the Human Wnt Inhibitory Factorâ€1. Lipids, 2008, 43, 227-230.	0.7	17
34	Hedgehog-Interacting Protein is a multimodal antagonist of Hedgehog signalling. Nature Communications, 2021, 12, 7171.	5.8	16
35	Diversity of oligomerization in Drosophila semaphorins suggests a mechanism of functional fine-tuning. Nature Communications, 2019, 10, 3691.	5.8	10
36	Structure dynamics of HIV-1 Env trimers on native virions engaged with living T cells. Communications Biology, 2021, 4, 1228.	2.0	4

#	Article	IF	CITATIONS
37	High-throughput Molecular Docking Now in Reach for a Wider Biochemical Community. , 2012, , .		2
38	Production of Heteromeric Transmembrane Receptors with Defined Subunit Stoichiometry. Structure, 2016, 24, 653-655.	1.6	1