## Ioannis Vakonakis

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4658081/publications.pdf

Version: 2024-02-01

37 papers

1,761 citations

331538 21 h-index 35 g-index

42 all docs 42 docs citations

42 times ranked 2619 citing authors

#	Article	IF	CITATIONS
1	The human cognition-enhancing CORD7 mutation increases active zone number and synaptic release. Brain, 2022, 145, 3787-3802.	3.7	8
2	Structures of SAS-6 coiled coil hold implications for the polarity of the centriolar cartwheel. Structure, 2022, 30, 671-684.e5.	1.6	4
3	The centriolar cartwheel structure: symmetric, stacked, and polarized. Current Opinion in Structural Biology, 2021, 66, 1-7.	2.6	10
4	A COVID moonshot: assessment of ligand binding to the SARS-CoV-2 main protease by saturation transfer difference NMR spectroscopy. Journal of Biomolecular NMR, 2021, 75, 167-178.	1.6	9
5	Structures of the <i>Plasmodium falciparum</i> heat-shock protein 70-x ATPase domain in complex with chemical fragments identify conserved and unique binding sites. Acta Crystallographica Section F, Structural Biology Communications, 2021, 77, 262-268.	0.4	3
6	The 3-phosphoinositide–dependent protein kinase 1 is an essential upstream activator of protein kinase A in malaria parasites. PLoS Biology, 2021, 19, e3001483.	2.6	9
7	Allosteric Inhibition of the SARSâ€CoVâ€2 Main Protease: Insights from Mass Spectrometry Based Assays**. Angewandte Chemie, 2020, 132, 23750-23754.	1.6	10
8	Allosteric Inhibition of the SARSâ€CoVâ€2 Main Protease: Insights from Mass Spectrometry Based Assays**. Angewandte Chemie - International Edition, 2020, 59, 23544-23548.	7.2	92
9	Identification of compounds that bind the centriolar protein SAS-6 and inhibit its oligomerization. Journal of Biological Chemistry, 2020, 295, 17922-17934.	1.6	2
10	Structure of the substrate-binding domain of <i>Plasmodium falciparum</i> heat-shock protein 70-x. Acta Crystallographica Section F, Structural Biology Communications, 2020, 76, 495-500.	0.4	8
11	The <i>Plasmodium falciparum</i> Hsp70â€x chaperone assists the heat stress response of the malaria parasite. FASEB Journal, 2019, 33, 14611-14624.	0.2	39
12	A dynamically interacting flexible loop assists oligomerisation of the Caenorhabditis elegans centriolar protein SAS-6. Scientific Reports, 2019, 9, 3526.	1.6	3
13	The complex of Plasmodium falciparum falcipain-2 protease with an (E)-chalcone-based inhibitor highlights a novel, small, molecule-binding site. Malaria Journal, 2019, 18, 388.	0.8	19
14	Interaction between the <i>Caenorhabditis elegans </i> centriolar protein SAS-5 and microtubules facilitates organelle assembly. Molecular Biology of the Cell, 2018, 29, 722-735.	0.9	8
15	Coupling Form and Function: How the Oligomerisation Symmetry of the SAS-6 Protein Contributes to the Architecture of Centriole Organelles. Symmetry, 2017, 9, 74.	1.1	О
16	Structural analysis of P. falciparum KAHRP and PfEMP1 complexes with host erythrocyte spectrin suggests a model for cytoadherent knob protrusions. PLoS Pathogens, 2017, 13, e1006552.	2.1	26
17	A spiral scaffold underlies cytoadherent knobs in Plasmodium falciparum–infected erythrocytes. Blood, 2016, 127, 343-351.	0.6	50
18	Plasmodium Helical Interspersed Subtelomeric (PHIST) Proteins, at the Center of Host Cell Remodeling. Microbiology and Molecular Biology Reviews, 2016, 80, 905-927.	2.9	49

#	Article	IF	Citations
19	<i>Plasmodium falciparum Plasmodium <math>\langle   i \rangle</math> helical interspersed subtelomeric proteins contribute to cytoadherence and anchor <math>\langle i \rangle</math>P. falciparum <math>\langle   i \rangle</math> erythrocyte membrane protein 1 to the host cell cytoskeleton. Cellular Microbiology, 2016, 18, 1415-1428.</i>	1.1	37
20	The centriolar protein CPAP G-box: an amyloid fibril in a single domain. Biochemical Society Transactions, 2015, 43, 838-843.	1.6	7
21	The Caenorhabditis elegans protein SAS-5 forms large oligomeric assemblies critical for centriole formation. ELife, 2015, 4, e07410.	2.8	37
22	A <i>Plasmodium falciparum</i> PHIST protein binds the virulence factor PfEMP1 and comigrates to knobs on the host cell surface. FASEB Journal, 2014, 28, 4420-4433.	0.2	78
23	Structural Analysis of the G-Box Domain of the Microcephaly Protein CPAP Suggests a Role in Centriole Architecture. Structure, 2013, 21, 2069-2077.	1.6	66
24	<i>Caenorhabditis elegans</i> centriolar protein SAS-6 forms a spiral that is consistent with imparting a ninefold symmetry. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 11373-11378.	3.3	54
25	Structural Analysis of Collagen Type I Interactions with Human Fibronectin Reveals a Cooperative Binding Mode. Journal of Biological Chemistry, 2013, 288, 17441-17450.	1.6	67
26	Structural Analysis of the Plasmodium falciparum Erythrocyte Membrane Protein 1 (PfEMP1) Intracellular Domain Reveals a Conserved Interaction Epitope. Journal of Biological Chemistry, 2012, 287, 7182-7189.	1.6	53
27	The GPS Motif Is a Molecular Switch for Bimodal Activities of Adhesion Class G Protein-Coupled Receptors. Cell Reports, 2012, 2, 321-331.	2.9	123
28	Structural Basis of the 9-Fold Symmetry of Centrioles. Cell, 2011, 144, 364-375.	13.5	317
29	Multi-factorial modulation of IGD motogenic potential in MSF (Migration Stimulating Factor). Experimental Cell Research, 2010, 316, 2465-2476.	1.2	10
30	Implications for Collagen Binding from the Crystallographic Structure of Fibronectin 6FnI1–2FnII7FnI. Journal of Biological Chemistry, 2010, 285, 33764-33770.	1.6	30
31	The Streptococcal Binding Site in the Gelatin-binding Domain of Fibronectin Is Consistent with a Non-linear Arrangement of Modules. Journal of Biological Chemistry, 2010, 285, 36977-36983.	1.6	15
32	Motogenic Sites in Human Fibronectin Are Masked by Long Range Interactions. Journal of Biological Chemistry, 2009, 284, 15668-15675.	1.6	46
32		3.3	77
	Chemistry, 2009, 284, 15668-15675.  Identification and structural analysis of type I collagen sites in complex with fibronectin fragments.		
33	Chemistry, 2009, 284, 15668-15675.  Identification and structural analysis of type I collagen sites in complex with fibronectin fragments. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 4195-4200.  Latrophilin Signaling Links Anterior-Posterior Tissue Polarity and Oriented Cell Divisions in the	3.3	77

#	‡	Article	lF	CITATIONS
3	37	Extracellular matrix: from atomic resolution to ultrastructure. Current Opinion in Cell Biology, 2007, 19, 578-583.	2.6	67