

Isabel Moraes

List of Publications by Year in descending order

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Version: 2024-02-01

37
papers

1,166
citations

516215

16
h-index

433756

31
g-index

39
all docs

39
docs citations

39
times ranked

2013
citing authors

#	ARTICLE	IF	CITATIONS
1	Two states of a light-sensitive membrane protein captured at room temperature using thin-film sample mounts. <i>Acta Crystallographica Section D: Structural Biology</i> , 2022, 78, 52-58.	1.1	2
2	Structures of the archaerhodopsin-3 transporter reveal that disordering of internal water networks underpins receptor sensitization. <i>Biophysical Journal</i> , 2022, 121, 316a.	0.2	0
3	Measuring Protein Aggregation and Stability Using High-Throughput Biophysical Approaches. <i>Frontiers in Molecular Biosciences</i> , 2022, 9, .	1.6	7
4	Structures of the archaerhodopsin-3 transporter reveal that disordering of internal water networks underpins receptor sensitization. <i>Nature Communications</i> , 2021, 12, 629.	5.8	22
5	Structural Biology and Structureâ€“Function Relationships of Membrane Proteins. <i>Biology</i> , 2021, 10, 245.	1.3	2
6	In Situ Measurements of Polypeptide Samples by Dynamic Light Scattering: Membrane Proteins, a Case Study. <i>Methods in Molecular Biology</i> , 2021, 2208, 189-202.	0.4	1
7	Structures of the archaerhodopsin-3 transporter reveal that disordering of internal water networks underpins receptor sensitization. <i>Acta Crystallographica Section A: Foundations and Advances</i> , 2021, 77, C478-C478.	0.0	0
8	High-throughput approach to prepare high-density microcrystals in lipidic cubic phase for serial crystallography and fragment screening. <i>Acta Crystallographica Section A: Foundations and Advances</i> , 2021, 77, C566-C566.	0.0	0
9	Molecular basis for <sc>GTP</sc> recognition by lightâ€“activated guanylate cyclase Rh<sc>GC</sc>. <i>FEBS Journal</i> , 2020, 287, 2797-2807.	2.2	9
10	Insights on the Quest for the Structureâ€“Function Relationship of the Mitochondrial Pyruvate Carrier. <i>Biology</i> , 2020, 9, 407.	1.3	4
11	Probing Membrane Protein Assembly into Nanodiscs by In Situ Dynamic Light Scattering: A2A Receptor as a Case Study. <i>Biology</i> , 2020, 9, 400.	1.3	4
12	Super-Resolution Fluorescence Microscopy Reveals Clustering Behaviour of Chlamydia pneumoniaeâ€™s Major Outer Membrane Protein. <i>Biology</i> , 2020, 9, 344.	1.3	5
13	Membrane protein crystallography in the era of modern structural biology. <i>Biochemical Society Transactions</i> , 2020, 48, 2505-2524.	1.6	9
14	High-throughput stability screening for detergent-solubilized membrane proteins. <i>Scientific Reports</i> , 2019, 9, 10379.	1.6	79
15	Selection of Biophysical Methods for Characterisation of Membrane Proteins. <i>International Journal of Molecular Sciences</i> , 2019, 20, 2605.	1.8	21
16	Structural biology and structureâ€“function relationships of membrane proteins. <i>Biochemical Society Transactions</i> , 2019, 47, 47-61.	1.6	24
17	Toward G protein-coupled receptor structure-based drug design using X-ray lasers. <i>IUCrJ</i> , 2019, 6, 1106-1119.	1.0	53
18	Crystallisation for serial crystallography in lipidic cubic phase (LCP) made simple. <i>Acta Crystallographica Section A: Foundations and Advances</i> , 2019, 75, e658-e658.	0.0	0

#	ARTICLE	IF	CITATIONS
19	Secrets to a successful collaboration. <i>Acta Crystallographica Section A: Foundations and Advances</i> , 2019, 75, e718-e718.	0.0	0
20	Human mitochondrial pyruvate carrier 2 as an autonomous membrane transporter. <i>Scientific Reports</i> , 2018, 8, 3510.	1.6	39
21	The fine art of integral membrane protein crystallisation. <i>Methods</i> , 2018, 147, 150-162.	1.9	45
22	Structure of a lipid A phosphoethanolamine transferase suggests how conformational changes govern substrate binding. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 2218-2223.	3.3	113
23	A Novel Approach to Data Collection for Difficult Structures: Data Management for Large Numbers of Crystals with the BLEND Software. <i>Crystals</i> , 2017, 7, 242.	1.0	6
24	Membrane proteins involved in bacterial phospholipid biosynthesis as drug targets?. <i>Acta Crystallographica Section A: Foundations and Advances</i> , 2017, 73, C397-C397.	0.0	0
25	Crystal Dehydration in Membrane Protein Crystallography. <i>Advances in Experimental Medicine and Biology</i> , 2016, 922, 73-89.	0.8	4
26	Green Fluorescent Protein-based Expression Screening of Membrane Proteins in <i>Escherichia coli</i> . <i>Journal of Visualized Experiments</i> , 2015, , e52357.	0.2	21
27	Lipidic cubic phase serial millisecond crystallography using synchrotron radiation. <i>IUCr</i> , 2015, 2, 168-176.	1.0	196
28	GPCR structure, function, drug discovery and crystallography: report from Academia-Industry International Conference (UK Royal Society) Chicheley Hall, 1-2 September 2014. <i>Naunyn-Schmiedeberg's Archives of Pharmacology</i> , 2015, 388, 883-903.	1.4	34
29	Methods for the Successful Crystallization of Membrane Proteins. <i>Methods in Molecular Biology</i> , 2015, 1261, 211-230.	0.4	8
30	Membrane protein structure determination – The next generation. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2014, 1838, 78-87.	1.4	190
31	X-ray structure of a CDP-alcohol phosphatidyltransferase membrane enzyme and insights into its catalytic mechanism. <i>Nature Communications</i> , 2014, 5, 4169.	5.8	39
32	Amphipol-Trapped ExbA-ExbD Membrane Protein Complex from <i>Escherichia coli</i> : A Biochemical and Structural Case Study. <i>Journal of Membrane Biology</i> , 2014, 247, 1005-1018.	1.0	18
33	Using high-throughput <i>in situ</i> plate screening to evaluate the effect of dehydration on protein crystals. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2013, 69, 920-923.	2.5	17
34	Challenges and opportunities in structure determination of membrane proteins. <i>Acta Crystallographica Section A: Foundations and Advances</i> , 2013, 69, s362-s362.	0.3	0
35	<i>In situ</i> plate screening to evaluate the dehydration effect on protein crystals. <i>Acta Crystallographica Section A: Foundations and Advances</i> , 2013, 69, s39-s39.	0.3	0
36	<i>In situ</i> macromolecular crystallography using microbeams. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2012, 68, 592-600.	2.5	113

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37	The Structure of Mammalian Serine Racemase. Journal of Biological Chemistry, 2010, 285, 12873-12881.	1.6	76