Stefano Vanni

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8208352/publications.pdf

Version: 2024-02-01

49 papers	3,501 citations	212478 28 h-index	242451 47 g-index
58 all docs	58 docs citations	58 times ranked	5253 citing authors

#	Article	IF	Citations
1	The Choice of Nanoparticle Surfaceâ€Coupled Fluorescent Dyes Impacts Cellular Interaction. ChemNanoMat, 2022, 8, .	1.5	3
2	Estimating the accuracy of the MARTINI model towards the investigation of peripheral proteinâ€“membrane interactions. Faraday Discussions, 2021, 232, 131-148.	1.6	25
3	Investigating the structural properties of hydrophobic solvent-rich lipid bilayers. Soft Matter, 2021, 17, 5329-5335.	1.2	8
4	Pre-existing bilayer stresses modulate triglyceride accumulation in the ER versus lipid droplets. ELife, 2021, 10, .	2.8	55
5	Seipin accumulates and traps diacylglycerols and triglycerides in its ring-like structure. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	52
6	Protonation Equilibrium in the Active Site of the Photoactive Yellow Protein. Molecules, 2021, 26, 2025.	1.7	0
7	Recharging your fats: CHARMM36 parameters for neutral lipids triacylglycerol and diacylglycerol. Biophysical Reports, 2021, 1, 100034.	0.7	10
8	Computational Approaches to Investigate and Design Lipid-binding Domains for Membrane Biosensing. Chimia, 2021, 75, 1031.	0.3	1
9	How Do Ethanolamine Plasmalogens Contribute to Order and Structure of Neurological Membranes?. Journal of Physical Chemistry B, 2020, 124, 828-839.	1.2	23
10	Conserved Functions of Ether Lipids and Sphingolipids in the Early Secretory Pathway. Current Biology, 2020, 30, 3775-3787.e7.	1.8	59
11	Accurate Estimation of Membrane Capacitance from Atomistic Molecular Dynamics Simulations of Zwitterionic Lipid Bilayers. Journal of Physical Chemistry B, 2020, 124, 8278-8286.	1.2	11
12	Ceramide chain lengthâ€“dependent protein sorting into selective endoplasmic reticulum exit sites. Science Advances, 2020, 6, .	4.7	38
13	Structure and Dynamics of the Acyl Chains in the Membrane Trafficking and Enzymatic Processing of Lipids. Accounts of Chemical Research, 2019, 52, 3087-3096.	7.6	35
14	Local accumulation of diacylglycerol alters membrane properties nonlinearly due to its transbilayer activity. Communications Chemistry, 2019, 2, .	2.0	37
15	Proteomics-Based Monitoring of Pathway Activity Reveals that Blocking Diacylglycerol Biosynthesis Rescues from Alpha-Synuclein Toxicity. Cell Systems, 2019, 9, 309-320.e8.	2.9	12
16	An Atomistic Look into Bio-inspired Nanoparticles and their Molecular Interactions with Cells. Chimia, 2019, 73, 78.	0.3	4
17	To Bud or Not to Bud: A Perspective on Molecular Simulations of Lipid Droplet Budding. Frontiers in Molecular Biosciences, 2019, 6, 124.	1.6	27
18	Templated Assembly of Pore-forming Peptides in Lipid Membranes. Chimia, 2019, 73, 59.	0.3	3

#	Article	IF	Citations
19	Understanding Conformational Dynamics of Complex Lipid Mixtures Relevant to Biology. Journal of Membrane Biology, 2018, 251, 609-631.	1.0	33
20	PackMem: A Versatile Tool to Compute and Visualize Interfacial Packing Defects in Lipid Bilayers. Biophysical Journal, 2018, 115, 436-444.	0.2	57
21	Interdigitation between Triglycerides and Lipids Modulates Surface Properties of Lipid Droplets. Biophysical Journal, 2017, 112, 1417-1430.	0.2	102
22	ER Membrane Phospholipids and Surface Tension Control Cellular Lipid Droplet Formation. Developmental Cell, 2017, 41, 591-604.e7.	3.1	213
23	Toward Chemically Resolved Computer Simulations of Dynamics and Remodeling of Biological Membranes. Journal of Physical Chemistry Letters, 2017, 8, 3586-3594.	2.1	35
24	Intracellular Lipid Droplets: From Structure to Function. Lipid Insights, 2017, 10, 117863531774551.	1.0	19
25	A filter at the entrance of the Golgi that selects vesicles according to size and bulk lipid composition. ELife, 2016, 5, .	2.8	57
26	Methyl-branched lipids promote the membrane adsorption of Î±-synuclein by enhancing shallow lipid-packing defects. Physical Chemistry Chemical Physics, 2015, 17, 15589-15597.	1.3	42
27	Membrane Protein Structure, Function, and Dynamics: a Perspective from Experiments and Theory. Journal of Membrane Biology, 2015, 248, 611-640.	1.0	157
28	Phosphatidylserine transport by ORP/Osh proteins is driven by phosphatidylinositol 4-phosphate. Science, 2015, 349, 432-436.	6.0	301
29	From zero to six double bonds: phospholipid unsaturation and organelle function. Trends in Cell Biology, 2015, 25, 427-436.	3.6	168
30	A phosphatidylinositol-4-phosphate powered exchange mechanism to create a lipid gradient between membranes. Nature Communications, 2015, 6, 6671.	5.8	166
31	Toward accurate coarse-graining approaches for protein and membrane simulations. Chemical Modelling, 2015, , 1-52.	0.2	10
32	BIN1/M-Amphiphysin2 induces clustering of phosphoinositides to recruit its downstream partner dynamin. Nature Communications, 2014, 5, 5647.	5.8	94
33	A sub-nanometre view of how membrane curvature and composition modulate lipid packing and protein recruitment. Nature Communications, 2014, 5, 4916.	5.8	230
34	Origin of the Spectral Shifts among the Early Intermediates of the Rhodopsin Photocycle. Journal of the American Chemical Society, 2014, 136, 3842-3851.	6.6	42
35	Polyunsaturated phospholipids facilitate membrane deformation and fission by endocytic proteins. Science, 2014, 345, 693-697.	6.0	291
36	Conical Lipids in Flat Bilayers Induce Packing Defects Similar to that Induced by Positive Curvature. Biophysical Journal, 2013, 104, 585-593.	0.2	149

#	Article	IF	Citations
37	Amphipathic Lipid Packing Sensor Motifs: Probing Bilayer Defects with Hydrophobic Residues. Biophysical Journal, 2013, 104, 575-584.	0.2	171
38	A Closer Look into G Protein Coupled Receptor Activation: X-Ray Crystallography and Long-Scale Molecular Dynamics Simulations. Current Medicinal Chemistry, 2012, 19, 1135-1145.	1.2	21
39	Ion Binding and Internal Hydration in the Multidrug Resistance Secondary Active Transporter NorM Investigated by Molecular Dynamics Simulations. Biochemistry, 2012, 51, 1281-1287.	1.2	32
40	Identification of clustering artifacts in photoactivated localization microscopy. Nature Methods, 2011, 8, 527-528.	9.0	197
41	Quantitative Photo Activated Localization Microscopy: Unraveling the Effects of Photoblinking. PLoS ONE, 2011, 6, e22678.	1.1	252
42	Mechanical (QM/MM) Simulations of Adiabatic and Nonadiabatic Ultrafast Phenomena. Chimia, 2011, 65, 330-333.	0.3	5
43	Pushing the Frontiers of First-Principles Based Computer Simulations of Chemical and Biological Systems. Chimia, 2011, 65, 667.	0.3	22
44	Predicting Novel Binding Modes of Agonists to Î² Adrenergic Receptors Using All-Atom Molecular Dynamics Simulations. PLoS Computational Biology, 2011, 7, e1001053.	1.5	38
45	A Conserved Protonation-Induced Switch can Trigger â€œIonic-Lockâ€•Formation in Adrenergic Receptors. Journal of Molecular Biology, 2010, 397, 1339-1349.	2.0	36
46	Role of Aggregation in Rhodopsin Signal Transduction. Biochemistry, 2010, 49, 4827-4832.	1.2	49
47	Observation of â€œIonic Lockâ€•Formation in Molecular Dynamics Simulations of Wild-Type Î²1 and Î²2 Adrenergic Receptors. Biochemistry, 2009, 48, 4789-4797.	1.2	65
48	A new formulation of the phase change approach in the theory of conical intersections. Chemical Physics, 2008, 347, 46-56.	0.9	13
49	Lipid Droplet Biogenesis is Driven by Liquid-Liquid Phase Separation. SSRN Electronic Journal, 0, , .	0.4	10