

Mounir Tarek

List of Publications by Year in descending order

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116
papers

6,034
citations

66250

44
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87275

74
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124
all docs

124
docs citations

124
times ranked

6247
citing authors

#	ARTICLE	IF	CITATIONS
1	The role of catechin in electroporation of pancreatic cancer cells – Effects on pore formation and multidrug resistance proteins. <i>Bioelectrochemistry</i> , 2022, 147, 108199.	2.4	4
2	Mechanisms of curcumin-based photodynamic therapy and its effects in combination with electroporation: An in vitro and molecular dynamics study. <i>Bioelectrochemistry</i> , 2021, 140, 107806.	2.4	14
3	Editorial: Molecular Mechanisms of Voltage-Gating in Ion Channels. <i>Frontiers in Pharmacology</i> , 2021, 12, 768153.	1.6	2
4	Molecular dynamics simulations of the effects of lipid oxidation on the permeability of cell membranes. <i>Bioelectrochemistry</i> , 2021, 141, 107869.	2.4	23
5	Oxidative Effects during Irreversible Electroporation of Melanoma Cells – In Vitro Study. <i>Molecules</i> , 2021, 26, 154.	1.7	28
6	Atorvastatin Modulates the Efficacy of Electroporation and Calcium Electrochemotherapy. <i>International Journal of Molecular Sciences</i> , 2021, 22, .	1.8	1
7	Atorvastatin Modulates the Efficacy of Electroporation and Calcium Electrochemotherapy. <i>International Journal of Molecular Sciences</i> , 2021, 22, 11245.	1.8	9
8	In Vitro Study of Calcium Microsecond Electroporation of Prostate Adenocarcinoma Cells. <i>Molecules</i> , 2020, 25, 5406.	1.7	11
9	Cepharanthine induces ROS stress in glioma and neuronal cells via modulation of VDAC permeability. <i>Saudi Pharmaceutical Journal</i> , 2020, 28, 1364-1373.	1.2	8
10	Lipid composition of the cancer cell membrane. <i>Journal of Bioenergetics and Biomembranes</i> , 2020, 52, 321-342.	1.0	190
11	Two-stage –Hand-and-Elbow–Gating Mechanism of a KV Channel. <i>Biophysical Journal</i> , 2020, 118, 113a.	0.2	0
12	New insights on the role of ROS in the mechanisms of sonoporation-mediated gene delivery. <i>Ultrasonics Sonochemistry</i> , 2020, 64, 104998.	3.8	16
13	Two-stage electro –mechanical coupling of a KV channel in voltage-dependent activation. <i>Nature Communications</i> , 2020, 11, 676.	5.8	46
14	The contribution of lipid peroxidation to membrane permeability in electropermeabilization: A molecular dynamics study. <i>Bioelectrochemistry</i> , 2019, 125, 46-57.	2.4	71
15	Membrane Electroporation and Electropermeabilization: Mechanisms and Models. <i>Annual Review of Biophysics</i> , 2019, 48, 63-91.	4.5	417
16	Effects of hydration on the protonation state of a lysine analog crossing a phospholipid bilayer – insights from molecular dynamics and free-energy calculations. <i>Physical Chemistry Chemical Physics</i> , 2018, 20, 9101-9107.	1.3	9
17	Studying Kv Channels Function using Computational Methods. <i>Methods in Molecular Biology</i> , 2018, 1684, 321-341.	0.4	4
18	Electrical Conductance of Lipid Pores. , 2017, , 219-233.		0

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19	Notice of Removal: New insights in the role of reactive oxygen species in mechanisms of sonoporation: In-vitro validation and molecular dynamic simulations. , 2017, , .		1
20	Atomistic Simulations of Electroporation of Model Cell Membranes. Advances in Anatomy, Embryology and Cell Biology, 2017, 227, 1-15.	1.0	3
21	A molecular insight into the electro-transfer of small molecules through electropores driven by electric fields. Biochimica Et Biophysica Acta - Biomembranes, 2016, 1858, 2278-2289.	1.4	66
22	In-Silico Electrophysiology: On the Activation of Voltage-Gated Ion Channels using Molecular Dynamics Simulations. Biophysical Journal, 2016, 110, 107a.	0.2	1
23	Properties of lipid electropores II: Comparison of continuum-level modeling of pore conductance to molecular dynamics simulations. Bioelectrochemistry, 2016, 112, 112-124.	2.4	25
24	Properties of lipid electropores I: Molecular dynamics simulations of stabilized pores by constant charge imbalance. Bioelectrochemistry, 2016, 109, 108-116.	2.4	42
25	Electrical Conductance of Lipid Pores. , 2016, , 1-15.		0
26	Molecular Determinants of Voltage Sensor Domain Activation. Biophysical Journal, 2015, 108, 427a.	0.2	0
27	PIP2 Modifies the Free Energy of the Kv1.2 Voltage-Sensor Activation. Biophysical Journal, 2015, 108, 120a.	0.2	0
28	PIP2-dependent coupling is prominent in Kv7.1 due to weakened interactions between S4-S5 and S6. Scientific Reports, 2015, 5, 7474.	1.6	53
29	Free-energy landscape of ion-channel voltage-sensorâ€ˆdomain activation. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 124-129.	3.3	63
30	Membrane Protein Structure, Function, and Dynamics: a Perspective from Experiments and Theory. Journal of Membrane Biology, 2015, 248, 611-640.	1.0	157
31	The Mechanism of KCNE1 Modulation of KCNQ1 Channels. Biophysical Journal, 2015, 108, 349a.	0.2	0
32	Electroporation Threshold of POPC Lipid Bilayers with Incorporated Polyoxyethylene Glycol (C ₁₂ E ₈). Journal of Physical Chemistry B, 2015, 119, 192-200.	1.2	17
33	Functional interaction between S1 and S4 segments in voltage-gated sodium channels revealed by human channelopathies. Channels, 2014, 8, 414-420.	1.5	7
34	Prerequisites to proton transport in the bacterial ClC-ec1 Cl ⁻ /H ⁺ -exchanger: Fig. 1.. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 1668-1669.	3.3	2
35	State-Dependent Lipid Interactions Couple the Conformations of the Voltage-Sensing and Pore-Gate Domains. Biophysical Journal, 2014, 106, 742a.	0.2	0
36	Investigation of Ion Permeation through the Cx26 Hemichannel. Biophysical Journal, 2014, 106, 556a.	0.2	0

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37	Voltage-gated ion channel modulation by lipids: Insights from molecular dynamics simulations. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2014, 1838, 1322-1331.	1.4	32
38	Electroporation of archaeal lipid membranes using MD simulations. <i>Bioelectrochemistry</i> , 2014, 100, 18-26.	2.4	56
39	A molecular dynamic study of cholesterol rich lipid membranes: comparison of electroporation protocols. <i>Bioelectrochemistry</i> , 2014, 100, 11-17.	2.4	75
40	Structural Properties of Archaeal Lipid Bilayers: Small-Angle X-ray Scattering and Molecular Dynamics Simulation Study. <i>Langmuir</i> , 2014, 30, 8308-8315.	1.6	24
41	On the Coupling between the Collective Dynamics of Proteins and Their Hydration Water. <i>Journal of Physical Chemistry Letters</i> , 2014, 5, 1181-1186.	2.1	59
42	Evidence of Conducting Hydrophobic Nanopores Across Membranes in Response to an Electric Field. <i>Journal of Physical Chemistry C</i> , 2014, 118, 6752-6757.	1.5	38
43	Multi-Dimensional Free Energy Landscape of Voltage Sensor Domain Activation. <i>Biophysical Journal</i> , 2014, 106, 233a.	0.2	0
44	Molecular Insights into Electroporation and Electrotransfer through Model Cell Membranes. <i>Biophysical Journal</i> , 2014, 106, 291a.	0.2	0
45	Molecular dynamics simulation of short-wavelength collective dynamics of phospholipid membranes. <i>Physical Review E</i> , 2014, 89, 050301.	0.8	24
46	Lipid Peroxidation in Membranes: The Peroxyl Radical Does Not "Float". <i>Journal of Physical Chemistry Letters</i> , 2014, 5, 1653-1658.	2.1	67
47	High-Yield Nontoxic Gene Transfer through Conjugation of the CM ₁₈ -Tat ₁₁ Chimeric Peptide with Nanosecond Electric Pulses. <i>Molecular Pharmaceutics</i> , 2014, 11, 2466-2474.	2.3	23
48	Domain-domain interactions determine the gating, permeation, pharmacology, and subunit modulation of the IKs ion channel. <i>ELife</i> , 2014, 3, e03606.	2.8	81
49	On the Electroporation Thresholds of Lipid Bilayers: Molecular Dynamics Simulation Investigations. <i>Journal of Membrane Biology</i> , 2013, 246, 843-850.	1.0	54
50	How do functionalized carbon nanotubes land on, bind to and pierce through model and plasma membranes. <i>Nanoscale</i> , 2013, 5, 10242.	2.8	61
51	Voltage Gated Cation Channels Activation: Towards an Ab-Initio Kinetic Model. <i>Biophysical Journal</i> , 2013, 104, 279a.	0.2	0
52	Dual Effect of PIP2 on Shaker K ⁺ Channels. <i>Biophysical Journal</i> , 2013, 104, 464a.	0.2	1
53	Vibrational excitations of proteins and their hydration water in the far-infrared range. <i>Chemical Physics</i> , 2013, 424, 80-83.	0.9	8
54	Omega Currents in Voltage-Gated Ion Channels: What Can We Learn from Uncovering the Voltage-Sensing Mechanism Using MD Simulations?. <i>Accounts of Chemical Research</i> , 2013, 46, 2755-2762.	7.6	20

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55	Effects of Phospholipid Composition on the Transfer of a Small Cationic Peptide Across a Model Biological Membrane. <i>Journal of Chemical Theory and Computation</i> , 2013, 9, 5675-5684.	2.3	25
56	Computational tools to investigate genetic cardiac channelopathies. <i>Frontiers in Physiology</i> , 2013, 4, 390.	1.3	6
57	Dual effect of phosphatidylinositol (4,5)-bisphosphate PIP2 on Shaker K+ channels.. <i>Journal of Biological Chemistry</i> , 2013, 288, 10951.	1.6	2
58	Mechanisms of Ion Channels Voltage-Dependency: All about Molecular Sensors, Gates, Levers, Locks, and Grease. <i>Frontiers in Pharmacology</i> , 2012, 3, 174.	1.6	2
59	An emerging consensus on voltage-dependent gating from computational modeling and molecular dynamics simulations. <i>Journal of General Physiology</i> , 2012, 140, 587-594.	0.9	179
60	Cell membrane electroporation- Part 1: The phenomenon. <i>IEEE Electrical Insulation Magazine</i> , 2012, 28, 14-23.	1.1	375
61	Dual Effect of Phosphatidyl (4,5)-Bisphosphate PIP2 on Shaker K+ Channels. <i>Journal of Biological Chemistry</i> , 2012, 287, 36158-36167.	1.6	37
62	Transport of siRNA through Lipid Membranes Driven by Nanosecond Electric Pulses: An Experimental and Computational Study. <i>Journal of the American Chemical Society</i> , 2012, 134, 13938-13941.	6.6	85
63	Molecular-Level Characterization of Lipid Membrane Electroporation using Linearly Rising Current. <i>Journal of Membrane Biology</i> , 2012, 245, 651-659.	1.0	36
64	Molecular Dynamics Simulations of Voltage-Gated Cation Channels: Insights on Voltage-Sensor Domain Function and Modulation. <i>Frontiers in Pharmacology</i> , 2012, 3, 97.	1.6	26
65	Molecular Dynamics Simulations of Lipid Membrane Electroporation. <i>Journal of Membrane Biology</i> , 2012, 245, 531-543.	1.0	158
66	Insertion of Short Amino-Functionalized Single-Walled Carbon Nanotubes into Phospholipid Bilayer Occurs by Passive Diffusion. <i>PLoS ONE</i> , 2012, 7, e40703.	1.1	67
67	Intermediate states of the Kv1.2 voltage sensor from atomistic molecular dynamics simulations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 6109-6114.	3.3	171
68	Uptake and Translocation Mechanisms of Cationic Amino Derivatives Functionalized on Pristine C60 by Lipid Membranes: A Molecular Dynamics Simulation Study. <i>ACS Nano</i> , 2011, 5, 8571-8578.	7.3	36
69	Structural, hydration, and phase transition properties of phosphatidylcholine from salmon heads. <i>European Journal of Lipid Science and Technology</i> , 2011, 113, 744-755.	1.0	6
70	Ethanol enhances collective dynamics of lipid membranes. <i>Physical Review E</i> , 2011, 83, 050907.	0.8	21
71	Influence of multiple double bonds on the structural and dynamical properties of a multi-component PC membrane extracted from Salmon heads. <i>Chemistry and Physics of Lipids</i> , 2010, 163, S30.	1.5	0
72	On the Antibacterial Action of Cyclic Peptides: Insights from Coarse-Grained MD Simulations. <i>Journal of Physical Chemistry B</i> , 2010, 114, 2676-2684.	1.2	42

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73	Effect of Sensor Domain Mutations on the Properties of Voltage-Gated Ion Channels: Molecular Dynamics Studies of the Potassium Channel Kv1.2. <i>Biophysical Journal</i> , 2010, 99, L72-L74.	0.2	48
74	Affinity of C ₆₀ Neat Fullerenes with Membrane Proteins: A Computational Study on Potassium Channels. <i>ACS Nano</i> , 2010, 4, 4158-4164.	7.3	63
75	Water and Membranes. , 2010, , 139-164.		0
76	Initial Response of the Potassium Channel Voltage Sensor to a Transmembrane Potential. <i>Journal of the American Chemical Society</i> , 2009, 131, 2107-2109.	6.6	55
77	Self assembly of peptides near or within membranes using coarse grained MD simulations. <i>Chemical Physics</i> , 2009, 358, 161-170.	0.9	28
78	Hydration dynamics of purple membranes. <i>Faraday Discussions</i> , 2009, 141, 99-116.	1.6	31
79	The role of protein's solvent hydrogen bond dynamics in the structural relaxation of a protein in glycerol versus water. <i>European Biophysics Journal</i> , 2008, 37, 701-709.	1.2	23
80	Modeling Membranes under a Transmembrane Potential. <i>Journal of Physical Chemistry B</i> , 2008, 112, 5547-5550.	1.2	94
81	Gating Motions in Voltage-Gated Potassium Channels Revealed by Coarse-Grained Molecular Dynamics Simulations. <i>Journal of Physical Chemistry B</i> , 2008, 112, 3277-3282.	1.2	64
82	Fingerprints of Amorphous Icelike Behavior in the Vibrational Density of States of Protein Hydration Water. <i>Physical Review Letters</i> , 2008, 101, 148104.	2.9	45
83	Energetics of Ion Transport in a Peptide Nanotube. <i>Journal of Physical Chemistry B</i> , 2007, 111, 10633-10635.	1.2	49
84	Large-scale molecular dynamics of a G protein-coupled receptor, the human 5-HT ₄ serotonin receptor, in a lipid bilayer. <i>Computational and Theoretical Chemistry</i> , 2007, 817, 19-26.	1.5	37
85	Probing a Model of a GPCR/Ligand Complex in an Explicit Membrane Environment: The Human Cholecystokinin-1 Receptor. <i>Biophysical Journal</i> , 2006, 90, 1232-1240.	0.2	57
86	Environment of the Gating Charges in the Kv1.2 Shaker Potassium Channel. <i>Biophysical Journal</i> , 2006, 90, L64-L66.	0.2	90
87	K ⁺ Conduction in the Selectivity Filter of Potassium Channels Is Monitored by the Charge Distribution along Their Sequence. <i>Biophysical Journal</i> , 2006, 91, L81-L83.	0.2	30
88	Molecular Restraints in the Permeation Pathway of Ion Channels. <i>Biophysical Journal</i> , 2006, 91, L26-L28.	0.2	31
89	Interaction of a peptide nanotube with a water's membrane interface. <i>Physical Biology</i> , 2006, 3, S20-S25.	0.8	22
90	Modeling Lipid Membranes. , 2005, , 929-958.		5

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91	Membrane Electroporation: A Molecular Dynamics Simulation. <i>Biophysical Journal</i> , 2005, 88, 4045-4053.	0.2	438
92	Methyl Group Dynamics as a Probe of the Protein Dynamical Transition. <i>Journal of the American Chemical Society</i> , 2004, 126, 15928-15929.	6.6	43
93	Coupled Motions between Pore and Voltage-Sensor Domains: A Model for Shaker B, a Voltage-Gated Potassium Channel. <i>Biophysical Journal</i> , 2004, 87, 2365-2379.	0.2	45
94	Characterization of sub-nanosecond dynamics of the molten globule state of α -lactalbumin using quasielastic neutron scattering and molecular dynamics simulations. <i>Chemical Physics</i> , 2003, 292, 435-443.	0.9	17
95	Influence of Anesthetic and Nonimmobilizer Molecules on the Physical Properties of a Polyunsaturated Lipid Bilayer. <i>Journal of Physical Chemistry B</i> , 2003, 107, 14500-14508.	1.2	41
96	Molecular Dynamics Investigation of an Oriented Cyclic Peptide Nanotube in DMPC Bilayers. <i>Biophysical Journal</i> , 2003, 85, 2287-2298.	0.2	87
97	Membrane Structural Perturbations Caused by Anesthetics and Nonimmobilizers: A Molecular Dynamics Investigation. <i>Biophysical Journal</i> , 2001, 81, 3339-3345.	0.2	46
98	Intermolecular Interactions and the Structure of Fatty Acid Soap Crystals. <i>Journal of Physical Chemistry B</i> , 2001, 105, 552-561.	1.2	39
99	Effects of solvent damping on side chain and backbone contributions to the protein boson peak. <i>Journal of Chemical Physics</i> , 2001, 115, 1607-1612.	1.2	54
100	Molecular Dynamics Study of the Poly(oxyethylene) Surfactant C12E2 and Water. <i>Langmuir</i> , 2000, 16, 942-946.	1.6	64
101	The Dynamics of Protein Hydration Water: A Quantitative Comparison of Molecular Dynamics Simulations and Neutron-scattering Experiments. <i>Biophysical Journal</i> , 2000, 79, 3244-3257.	0.2	295
102	Distribution of Halothane in a Dipalmitoylphosphatidylcholine Bilayer from Molecular Dynamics Calculations. <i>Biophysical Journal</i> , 2000, 78, 800-811.	0.2	104
103	Amplitudes and Frequencies of Protein Dynamics: Analysis of Discrepancies between Neutron Scattering and Molecular Dynamics Simulations. <i>Journal of the American Chemical Society</i> , 2000, 122, 10450-10451.	6.6	52
104	Molecular Dynamics Study of a Lipid-DNA Complex. <i>Journal of Physical Chemistry B</i> , 1999, 103, 10075-10080.	1.2	97
105	Molecular Dynamics Simulations of Supported Phospholipid/Alkanethiol Bilayers on a Gold(111) Surface. <i>Biophysical Journal</i> , 1999, 77, 964-972.	0.2	49
106	Environmental Dependence of the Dynamics of Protein Hydration Water. <i>Journal of the American Chemical Society</i> , 1999, 121, 9740-9741.	6.6	63
107	Surfactant Aggregation at a Hydrophobic Surface. <i>Journal of Physical Chemistry B</i> , 1998, 102, 6318-6322.	1.2	81
108	Computer simulation studies of amphiphilic interfaces. <i>Current Opinion in Colloid and Interface Science</i> , 1998, 3, 242-246.	3.4	25

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109	Molecular dynamics studies of aqueous surfactants systems. Journal of Molecular Liquids, 1998, 78, 1-6.	2.3	40
110	Effects of Anesthetics on the Structure of a Phospholipid Bilayer: Molecular Dynamics Investigation of Halothane in the Hydrated Liquid Crystal Phase of Dipalmitoylphosphatidylcholine. Biophysical Journal, 1998, 75, 2123-2134.	0.2	86
111	Molecular dynamics studies of the hexagonal mesophase of sodium dodecylsulphate in aqueous solution. Molecular Physics, 1998, 95, 377-384.	0.8	20
112	Molecular Dynamics Study of Two-Component Systems: The Shape and Surface Structure of Water/Ethanol Droplets. Journal of Physical Chemistry A, 1997, 101, 8639-8642.	1.1	29
113	Molecular dynamics investigation of the surface/bulk equilibrium in an ethanol-water solution. Journal of the Chemical Society, Faraday Transactions, 1996, 92, 559-563.	1.7	64
114	Molecular dynamics investigation of an ethanol-water solution. Physica A: Statistical Mechanics and Its Applications, 1996, 231, 117-122.	1.2	38
115	Modelization of experimental isotherms of n-alkanes in NaX zeolite. Zeolites, 1995, 15, 67-72.	0.9	30
116	Molecular Dynamics Simulation of Tetradecyltrimethylammonium Bromide Monolayers at the Air/Water Interface. The Journal of Physical Chemistry, 1995, 99, 1393-1402.	2.9	120