

Jacek Czub

List of Publications by Year in descending order

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Version: 2024-02-01

61
papers

1,302
citations

411340

20
h-index

445137

33
g-index

65
all docs

65
docs citations

65
times ranked

2012
citing authors

#	ARTICLE	IF	CITATIONS
1	Determinants of Directionality and Efficiency of the ATP Synthase F _o Motor at Atomic Resolution. <i>Journal of Physical Chemistry Letters</i> , 2022, 13, 387-392.	2.1	8
2	Guanosine Dianions Hydrated by One to Four Water Molecules. <i>Journal of Physical Chemistry Letters</i> , 2022, , 3230-3236.	2.1	4
3	Self-assembly, stability and conductance of amphotericin B channels: bridging the gap between structure and function. <i>Nanoscale</i> , 2021, 13, 3686-3697.	2.8	9
4	Defining a novel domain that provides an essential contribution to site-specific interaction of Rep protein with DNA. <i>Nucleic Acids Research</i> , 2021, 49, 3394-3408.	6.5	8
5	Light-Modulated Sunscreen Mechanism in the Retina of the Human Eye. <i>Journal of Physical Chemistry B</i> , 2021, 125, 6090-6102.	1.2	9
6	Two Bacterial Small Heat Shock Proteins, IbpA and IbpB, Form a Functional Heterodimer. <i>Journal of Molecular Biology</i> , 2021, 433, 167054.	2.0	12
7	Role of cholesterol in substrate recognition by γ -secretase. <i>Scientific Reports</i> , 2021, 11, 15213.	1.6	4
8	Mechanism of recognition of parallel G-quadruplexes by DEAH/RHAU helicase DHX36 explored by molecular dynamics simulations. <i>Computational and Structural Biotechnology Journal</i> , 2021, 19, 2526-2536.	1.9	12
9	Molecular mechanism of proton-coupled ligand translocation by the bacterial efflux pump EmrE. <i>PLoS Computational Biology</i> , 2021, 17, e1009454.	1.5	8
10	Sequence-dependent structural properties of B-DNA: what have we learned in 40 years?. <i>Biophysical Reviews</i> , 2021, 13, 995-1005.	1.5	13
11	The Product of Matrix Metalloproteinase Cleavage of Doxorubicin Conjugate for Anticancer Drug Delivery: Calorimetric, Spectroscopic, and Molecular Dynamics Studies on Peptide-Doxorubicin Binding to DNA. <i>International Journal of Molecular Sciences</i> , 2020, 21, 6923.	1.8	7
12	Effect of chemical structure on complexation efficiency of aromatic drugs with cyclodextrins: The example of dibenzazepine derivatives. <i>Carbohydrate Polymers</i> , 2020, 250, 116957.	5.1	5
13	Two-step mechanism of J-domain action in driving Hsp70 function. <i>PLoS Computational Biology</i> , 2020, 16, e1007913.	1.5	18
14	Molywood: streamlining the design and rendering of molecular movies. <i>Bioinformatics</i> , 2020, 36, 4660-4661.	1.8	16
15	Telomere uncapping by common oxidative guanine lesions: Insights from atomistic models. <i>Free Radical Biology and Medicine</i> , 2020, 148, 162-169.	1.3	2
16	Why do G-quadruplexes dimerize through the 5' ends? Driving forces for G4 DNA dimerization examined in atomic detail. <i>PLoS Computational Biology</i> , 2019, 15, e1007383.	1.5	26
17	Structure and evolution of the 4-helix bundle domain of Zuotin, a J-domain protein co-chaperone of Hsp70. <i>PLoS ONE</i> , 2019, 14, e0217098.	1.1	8
18	Molecular modelling of membrane activity of amphotericin B, a polyene macrolide antifungal antibiotic.. <i>Acta Biochimica Polonica</i> , 2019, 52, 655-658.	0.3	69

#	ARTICLE	IF	CITATIONS
19	Title is missing!. , 2019, 15, e1007383.		0
20	Title is missing!. , 2019, 15, e1007383.		0
21	Title is missing!. , 2019, 15, e1007383.		0
22	Title is missing!. , 2019, 15, e1007383.		0
23	Title is missing!. , 2019, 15, e1007383.		0
24	Title is missing!. , 2019, 15, e1007383.		0
25	Effect of osmolytes on the thermal stability of proteins: replica exchange simulations of Trp-cage in urea and betaine solutions. <i>Physical Chemistry Chemical Physics</i> , 2018, 20, 11174-11182.	1.3	18
26	Mechanism of Binding of Antifungal Antibiotic Amphotericin B to Lipid Membranes: An Insight from Combined Single-Membrane Imaging, Microspectroscopy, and Molecular Dynamics. <i>Molecular Pharmaceutics</i> , 2018, 15, 4202-4213.	2.3	31
27	Effect of osmolytes of different type on DNA behavior in aqueous solution. Experimental and theoretical studies. <i>Journal of Molecular Liquids</i> , 2018, 271, 186-201.	2.3	8
28	Mechanochemical Energy Transduction during the Main Rotary Step in the Synthesis Cycle of F ₁ -ATPase. <i>Journal of the American Chemical Society</i> , 2017, 139, 4025-4034.	6.6	23
29	How proteins bind to DNA: target discrimination and dynamic sequence search by the telomeric protein TRF1. <i>Nucleic Acids Research</i> , 2017, 45, 7643-7654.	6.5	19
30	Localization and Orientation of Xanthophylls in a Lipid Bilayer. <i>Scientific Reports</i> , 2017, 7, 9619.	1.6	66
31	Role of the disulfide bond in stabilizing and folding of the fimbrial protein DraE from uropathogenic <i>Escherichia coli</i> . <i>Journal of Biological Chemistry</i> , 2017, 292, 16136-16149.	1.6	9
32	Dominant Pathways of Adenosyl Radical-Induced DNA Damage Revealed by QM/MM Metadynamics. <i>Journal of Chemical Theory and Computation</i> , 2017, 13, 6415-6423.	2.3	6
33	Thermodynamics and kinetics of amphotericin B self-association in aqueous solution characterized in molecular detail. <i>Scientific Reports</i> , 2016, 6, 19109.	1.6	25
34	Molecular basis of the osmolyte effect on protein stability: a lesson from the mechanical unfolding of lysozyme. <i>Biochemical Journal</i> , 2016, 473, 3705-3724.	1.7	19
35	Intramolecular transformation of an antifungal antibiotic nystatin A ₁ into its isomer, <i>iso</i> -nystatin A ₁ – structural and molecular modeling studies. <i>Magnetic Resonance in Chemistry</i> , 2016, 54, 953-961.	1.1	5
36	Correlation between the number of Pro-Ala repeats in the EmrA homologue of <i>Acinetobacter baumannii</i> and resistance to netilmicin, tobramycin, imipenem and ceftazidime. <i>Journal of Global Antimicrobial Resistance</i> , 2016, 7, 145-149.	0.9	4

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37	Molecular dynamics simulations reveal the balance of forces governing the formation of a guanine tetrad—a common structural unit of G-quadruplex DNA. <i>Nucleic Acids Research</i> , 2016, 44, 3020-3030.	6.5	30
38	Iron-Sulfur Cluster Biogenesis Chaperones: Evidence for Emergence of Mutational Robustness of a Highly Specific Protein-Protein Interaction. <i>Molecular Biology and Evolution</i> , 2016, 33, 643-656.	3.5	19
39	Membrane Sterols Modulate the Binding Mode of Amphotericin B without Affecting Its Affinity for a Lipid Bilayer. <i>Langmuir</i> , 2016, 32, 3452-3461.	1.6	22
40	Intercalation complex of imidazoacridinone C-1311, a potential anticancer drug, with DNA helix d(CGATCG) ₂ : stereostructural studies by 2D NMR spectroscopy. <i>Journal of Biomolecular Structure and Dynamics</i> , 2016, 34, 653-663.	2.0	8
41	Specific Binding of Cholesterol to the Amyloid Precursor Protein: Structure of the Complex and Driving Forces Characterized in Molecular Detail. <i>Journal of Physical Chemistry Letters</i> , 2015, 6, 784-790.	2.1	23
42	Solvation of ionic liquids based on N-methyl-N-alkyl morpholinium cations in dimethylsulfoxide—volumetric and compressibility studies. <i>Journal of Chemical Thermodynamics</i> , 2015, 88, 36-43.	1.0	8
43	Hydration of amino acids: FTIR spectra and molecular dynamics studies. <i>Amino Acids</i> , 2015, 47, 2265-2278.	1.2	20
44	Self-Association of Amphotericin B: Spontaneous Formation of Molecular Structures Responsible for the Toxic Side Effects of the Antibiotic. <i>Journal of Physical Chemistry B</i> , 2014, 118, 13821-13832.	1.2	53
45	A first-principles study of electron attachment to the fully hydrated bromonucleobases. <i>Chemical Physics Letters</i> , 2014, 595-596, 133-137.	1.2	32
46	Rotation Triggers Nucleotide-Independent Conformational Transition of the Empty $\hat{\nu}^2$ Subunit of F ₁ -ATPase. <i>Journal of the American Chemical Society</i> , 2014, 136, 6960-6968.	6.6	25
47	Molecular Recognition in Complexes of TRF Proteins with Telomeric DNA. <i>PLoS ONE</i> , 2014, 9, e89460.	1.1	5
48	The Effect of Sterols on Amphotericin B Self-Aggregation in a Lipid Bilayer as Revealed by Free Energy Simulations. <i>Biophysical Journal</i> , 2013, 104, 1485-1494.	0.2	34
49	Keep It Flexible: Driving Macromolecular Rotary Motions in Atomistic Simulations with GROMACS. <i>Journal of Chemical Theory and Computation</i> , 2011, 7, 1381-1393.	2.3	42
50	Torsional elasticity and energetics of F ₁ -ATPase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 7408-7413.	3.3	46
51	Spatial Distribution of Elasticity in the F1 Motor of ATP Synthase Reveals the Microscopic Nature of the Coupling Between the Central Shaft and the Catalytic Subunit. <i>Biophysical Journal</i> , 2010, 98, 168a.	0.2	1
52	How Do Sterols Determine the Antifungal Activity of Amphotericin B? Free Energy of Binding between the Drug and Its Membrane Targets. <i>Journal of the American Chemical Society</i> , 2010, 132, 18266-18272.	6.6	71
53	Influence of a lipid bilayer on the conformational behavior of amphotericin B derivatives—A molecular dynamics study. <i>Biophysical Chemistry</i> , 2009, 141, 105-116.	1.5	27
54	On the Possibility of the Amphotericin B-Sterol Complex Formation in Cholesterol- and Ergosterol-Containing Lipid Bilayers: A Molecular Dynamics Study. <i>Journal of Physical Chemistry B</i> , 2009, 113, 15875-15885.	1.2	38

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55	Interactions of amphotericin B derivatives with lipid membranes—A molecular dynamics study. <i>Biochimica Et Biophysica Acta - Biomembranes</i> , 2007, 1768, 2616-2626.	1.4	16
56	The Mechanism of Overcoming Multidrug Resistance (MDR) of Fungi by Amphotericin B and Its Derivatives. <i>Journal of Antibiotics</i> , 2007, 60, 436-446.	1.0	19
57	Comparative Molecular Dynamics Study of Lipid Membranes Containing Cholesterol and Ergosterol. <i>Biophysical Journal</i> , 2006, 90, 2368-2382.	0.2	153
58	Modulation of Amphotericin B Membrane Interaction by Cholesterol and Ergosterol—A Molecular Dynamics Study. <i>Journal of Physical Chemistry B</i> , 2006, 110, 16743-16753.	1.2	46
59	Interaction of amphotericin B and its selected derivatives with membranes: molecular modeling studies. <i>Chemical Record</i> , 2006, 6, 320-332.	2.9	45
60	Monosaccharides as internal probes for the determination of the absolute configuration of 2-butanol. <i>Magnetic Resonance in Chemistry</i> , 2006, 44, 132-138.	1.1	12
61	Molecular aspects of the interaction between amphotericin B and a phospholipid bilayer: molecular dynamics studies. <i>Journal of Molecular Modeling</i> , 2004, 10, 223-232.	0.8	34