

Thomas R Weikl

List of Publications by Year in descending order

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77
papers

5,098
citations

109321

35
h-index

91884

69
g-index

82
all docs

82
docs citations

82
times ranked

5011
citing authors

#	ARTICLE	IF	CITATIONS
1	A litmus test for classifying recognition mechanisms of transiently binding proteins. <i>Nature Communications</i> , 2022, 13, .	12.8	13
2	Membrane morphologies induced by mixtures of arc-shaped particles with opposite curvature. <i>Soft Matter</i> , 2021, 17, 268-275.	2.7	6
3	Cooperative Stabilization of Close-Contact Zones Leads to Sensitivity and Selectivity in T-Cell Recognition. <i>Cells</i> , 2021, 10, 1023.	4.1	3
4	Structural variability and concerted motions of the T cell receptor " CD3 complex. <i>ELife</i> , 2021, 10, .	6.0	7
5	Interplay of Trans- and Cis-Interactions of Glycolipids in Membrane Adhesion. <i>Frontiers in Molecular Biosciences</i> , 2021, 8, 754654.	3.5	4
6	Accessory mutations balance the marginal stability of the HIV-1 protease in drug resistance. <i>Proteins: Structure, Function and Bioinformatics</i> , 2020, 88, 476-484.	2.6	6
7	Weak carbohydrate-carbohydrate interactions in membrane adhesion are fuzzy and generic. <i>Nanoscale</i> , 2020, 12, 17342-17353.	5.6	10
8	Theoretical modeling of interactions at the bio-nano interface. <i>Nanoscale</i> , 2020, 12, 10426-10429.	5.6	7
9	On the relationship between docking scores and protein conformational changes in HIV-1 protease. <i>Journal of Molecular Graphics and Modelling</i> , 2019, 91, 186-193.	2.4	4
10	Membrane Morphologies Induced by Arc-Shaped Scaffolds Are Determined by Arc Angle and Coverage. <i>Biophysical Journal</i> , 2019, 116, 1239-1247.	0.5	18
11	Binding and segregation of proteins in membrane adhesion: theory, modeling, and simulations. <i>Advances in Biomembranes and Lipid Self-Assembly</i> , 2019, 30, 159-194.	0.6	6
12	Particle-based membrane model for mesoscopic simulation of cellular dynamics. <i>Journal of Chemical Physics</i> , 2018, 148, 044901.	3.0	33
13	Identifying Conformational-Selection and Induced-Fit Aspects in the Binding-Induced Folding of PMI from Markov State Modeling of Atomistic Simulations. <i>Journal of Physical Chemistry B</i> , 2018, 122, 5649-5656.	2.6	24
14	Curvature-Mediated Assembly of Janus Nanoparticles on Membrane Vesicles. <i>Nano Letters</i> , 2018, 18, 1259-1263.	9.1	41
15	Membrane-Mediated Cooperativity of Proteins. <i>Annual Review of Physical Chemistry</i> , 2018, 69, 521-539.	10.8	41
16	Membrane fluctuations and acidosis regulate cooperative binding of "marker of self-CD47 with macrophage checkpoint receptor SIRP". <i>Journal of Cell Science</i> , 2018, 132, .	2.0	45
17	The 2018 biomembrane curvature and remodeling roadmap. <i>Journal Physics D: Applied Physics</i> , 2018, 51, 343001.	2.8	212
18	Protein-peptide association kinetics beyond the seconds timescale from atomistic simulations. <i>Nature Communications</i> , 2017, 8, 1095.	12.8	137

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19	Membrane Tubulation by Elongated and Patchy Nanoparticles. <i>Advanced Materials Interfaces</i> , 2017, 4, 1600325.	3.7	9
20	Binding equilibrium and kinetics of membrane-anchored receptors and ligands in cell adhesion: Insights from computational model systems and theory. <i>Cell Adhesion and Migration</i> , 2016, 10, 576-589.	2.7	29
21	The role of membrane curvature for the wrapping of nanoparticles. <i>Soft Matter</i> , 2016, 12, 581-587.	2.7	71
22	How to Distinguish Conformational Selection and Induced Fit Based on Chemical Relaxation Rates. <i>PLoS Computational Biology</i> , 2016, 12, e1005067.	3.2	74
23	Binding kinetics of membrane-anchored receptors and ligands: Molecular dynamics simulations and theory. <i>Journal of Chemical Physics</i> , 2015, 143, 243137.	3.0	27
24	Binding constants of membrane-anchored receptors and ligands: A general theory corroborated by Monte Carlo simulations. <i>Journal of Chemical Physics</i> , 2015, 143, 243136.	3.0	54
25	Modeling nanoparticle wrapping or translocation in bilayer membranes. <i>Nanoscale</i> , 2015, 7, 14505-14514.	5.6	49
26	Wrapping of nanoparticles by membranes. <i>Advances in Colloid and Interface Science</i> , 2014, 208, 214-224.	14.7	186
27	Conformational selection in protein binding and function. <i>Protein Science</i> , 2014, 23, 1508-1518.	7.6	99
28	Cooperative wrapping of nanoparticles by membrane tubes. <i>Soft Matter</i> , 2014, 10, 3570.	2.7	72
29	How conformational changes can affect catalysis, inhibition and drug resistance of enzymes with induced-fit binding mechanism such as the HIV-1 protease. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2013, 1834, 867-873.	2.3	16
30	Binding constants of membrane-anchored receptors and ligands depend strongly on the nanoscale roughness of membranes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 15283-15288.	7.1	117
31	Domain formation in cholesterol- ϕ phospholipid membranes exposed to adhesive surfaces or environments. <i>Soft Matter</i> , 2013, 9, 8438.	2.7	22
32	Adhesion-Induced Phase Behavior of Two-Component Membranes and Vesicles. <i>International Journal of Molecular Sciences</i> , 2013, 14, 2203-2229.	4.1	9
33	Direct Observation of Membrane Insertion by Enveloped Virus Matrix Proteins by Phosphate Displacement. <i>PLoS ONE</i> , 2013, 8, e57916.	2.5	5
34	Tubulation and Aggregation of Spherical Nanoparticles Adsorbed on Vesicles. <i>Physical Review Letters</i> , 2012, 109, 188102.	7.8	144
35	Adhesion of surfaces mediated by adsorbed particles: Monte Carlo simulations and a general relationship between adsorption isotherms and effective adhesion energies. <i>Soft Matter</i> , 2012, 8, 11737.	2.7	2
36	Conformational selection and induced changes along the catalytic cycle of <i>Escherichia coli</i> dihydrofolate reductase. <i>Proteins: Structure, Function and Bioinformatics</i> , 2012, 80, 2369-2383.	2.6	20

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37	Vesicles with multiple membrane domains. <i>Soft Matter</i> , 2011, 7, 6092.	2.7	65
38	Line Tension and Stability of Domains in Cell-Adhesion Zones Mediated by Long and Short Receptor-Ligand Complexes. <i>PLoS ONE</i> , 2011, 6, e23284.	2.5	28
39	Segregation of receptor–ligand complexes in cell adhesion zones: phase diagrams and the role of thermal membrane roughness. <i>New Journal of Physics</i> , 2010, 12, 095003.	2.9	42
40	Transition States in Protein Folding. <i>Communications in Computational Physics</i> , 2010, 7, 283-300.	1.7	0
41	Adhesion of surfaces via particle adsorption: exact results for a lattice of fluid columns. <i>Journal of Statistical Mechanics: Theory and Experiment</i> , 2009, 2009, P11006.	2.3	1
42	In vivo folding efficiencies for mutants of the P22 tailspike β -helix protein correlate with predicted stability changes. <i>Biophysical Chemistry</i> , 2009, 141, 186-192.	2.8	7
43	Selected–fit versus induced–fit protein binding: Kinetic differences and mutational analysis. <i>Proteins: Structure, Function and Bioinformatics</i> , 2009, 75, 104-110.	2.6	144
44	Adhesion of membranes via receptor–ligand complexes: Domain formation, binding cooperativity, and active processes. <i>Soft Matter</i> , 2009, 5, 3213.	2.7	92
45	Binding cooperativity of membrane adhesion receptors. <i>Soft Matter</i> , 2009, 5, 3354.	2.7	71
46	Constructing the equilibrium ensemble of folding pathways from short off-equilibrium simulations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 19011-19016.	7.1	730
47	Transition States in Protein Folding Kinetics: Modeling ΔG^\ddagger -Values of Small β -Sheet Proteins. <i>Biophysical Journal</i> , 2008, 94, 929-937.	0.5	17
48	Loop-closure principles in protein folding. <i>Archives of Biochemistry and Biophysics</i> , 2008, 469, 67-75.	3.0	24
49	The Protein Folding Problem. <i>Annual Review of Biophysics</i> , 2008, 37, 289-316.	10.0	916
50	Effective surface interactions mediated by adhesive particles. <i>Europhysics Letters</i> , 2008, 84, 26004.	2.0	13
51	Stable Patterns of Membrane Domains at Corrugated Substrates. <i>Physical Review Letters</i> , 2008, 100, 098103.	7.8	38
52	Lateral diffusion of receptor-ligand bonds in membrane adhesion zones: Effect of thermal membrane roughness. <i>Europhysics Letters</i> , 2007, 78, 38003.	2.0	38
53	Transition-States in Protein Folding Kinetics: The Structural Interpretation of ΔG^\ddagger values. <i>Journal of Molecular Biology</i> , 2007, 365, 1578-1586.	4.2	32
54	The protein folding problem: when will it be solved?. <i>Current Opinion in Structural Biology</i> , 2007, 17, 342-346.	5.7	208

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55	Stochastic resonance for adhesion of membranes with active stickers. <i>European Physical Journal E</i> , 2007, 22, 97-106.	1.6	9
56	Chapter 4 Membrane Adhesion and Domain Formation. <i>Behavior Research Methods</i> , 2006, , 63-127.	4.0	14
57	Substructural cooperativity and parallel versus sequential events during protein unfolding. <i>Proteins: Structure, Function and Bioinformatics</i> , 2006, 63, 1052-1058.	2.6	13
58	A simple measure of native state topology and chain connectivity predicts the folding rates of two-state proteins with and without crosslinks. <i>Proteins: Structure, Function and Bioinformatics</i> , 2006, 64, 193-197.	2.6	18
59	Adhesion of membranes via switchable molecules. <i>Physical Review E</i> , 2006, 73, 061908.	2.1	12
60	Adhesion of Membranes with Active Stickers. <i>Physical Review Letters</i> , 2006, 96, 048101.	7.8	25
61	Membrane adhesion via competing receptor/ligand bonds. <i>Europhysics Letters</i> , 2006, 76, 703-709.	2.0	23
62	Loop-closure events during protein folding: Rationalizing the shape of τ_1 -value distributions. <i>Proteins: Structure, Function and Bioinformatics</i> , 2005, 60, 701-711.	2.6	13
63	Δ values in protein-folding kinetics have energetic and structural components. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 10171-10175.	7.1	40
64	Cooperativity in two-state protein folding kinetics. <i>Protein Science</i> , 2004, 13, 822-829.	7.6	66
65	Pattern Formation during T-Cell Adhesion. <i>Biophysical Journal</i> , 2004, 87, 3665-3678.	0.5	117
66	Indirect interactions of membrane-adsorbed cylinders. <i>European Physical Journal E</i> , 2003, 12, 265-273.	1.6	60
67	Folding Rates and Low-entropy-loss Routes of Two-state Proteins. <i>Journal of Molecular Biology</i> , 2003, 329, 585-598.	4.2	88
68	Folding Kinetics of Two-state Proteins: Effect of Circularization, Permutation, and Crosslinks. <i>Journal of Molecular Biology</i> , 2003, 332, 953-963.	4.2	51
69	Dynamic phase separation of fluid membranes with rigid inclusions. <i>Physical Review E</i> , 2002, 66, 061915.	2.1	17
70	Pattern formation during adhesion of multicomponent membranes. <i>Europhysics Letters</i> , 2002, 59, 916-922.	2.0	59
71	Adhesion of membranes with competing specific and generic interactions. <i>European Physical Journal E</i> , 2002, 8, 59-66.	1.6	55
72	Two direct methods to calculate fluctuation forces between rigid objects embedded in fluid membranes. <i>European Physical Journal E</i> , 2001, 5, 423-439.	1.6	20

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73	Fluctuation-induced aggregation of rigid membrane inclusions. <i>Europhysics Letters</i> , 2001, 54, 547-553.	2.0	45
74	Adhesion-induced phase behavior of multicomponent membranes. <i>Physical Review E</i> , 2001, 64, 011903.	2.1	65
75	Unbinding transitions and phase separation of multicomponent membranes. <i>Physical Review E</i> , 2000, 62, R45-R48.	2.1	40
76	Local Adhesion of Membranes to Striped Surface Domains. <i>Langmuir</i> , 2000, 16, 9338-9346.	3.5	14
77	Interaction of conical membrane inclusions: Effect of lateral tension. <i>Physical Review E</i> , 1998, 57, 6988-6995.	2.1	140