

Jan Antosiewicz

List of Publications by Year in descending order

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93
papers

4,191
citations

172457

29
h-index

114465

63
g-index

108
all docs

108
docs citations

108
times ranked

2770
citing authors

#	ARTICLE	IF	CITATIONS
1	Editorial: Intracellular Molecular Processes Affected by pH. <i>Frontiers in Molecular Biosciences</i> , 2022, 9, 891533.	3.5	0
2	Circular Dichroism Spectra of $\hat{\pm}$ -Chymotrypsin $\hat{\pm}$ SDS Solutions Depend on the Procedure of Their Preparation. <i>ACS Omega</i> , 2022, 7, 23782-23789.	3.5	4
3	Searching for Hydrodynamic Orienting Effects in the Association of Tri-N-acetylglucosamine with Hen Egg-White Lysozyme. <i>Journal of Physical Chemistry B</i> , 2021, 125, 10701-10709.	2.6	2
4	Constant-pH Brownian Dynamics Simulations of a Protein near a Charged Surface. <i>ACS Omega</i> , 2020, 5, 30282-30298.	3.5	7
5	Diffusional Encounter Rate Constants for Xanthone and 2-Naphthoic Acid by Flash Photolysis Experiments and Brownian Dynamics Simulations: Substantial Effects of Polarizability of the Triplet State. <i>Journal of Physical Chemistry B</i> , 2019, 123, 9328-9342.	2.6	2
6	Effects of Hydrodynamic Interactions on the Near-Surface Diffusion of Spheroidal Molecules. <i>ACS Omega</i> , 2019, 4, 17016-17030.	3.5	7
7	Does Ionic Screening Lower Activation Barriers for Conformational Transitions in Proteins?. <i>Journal of Physical Chemistry B</i> , 2018, 122, 11817-11826.	2.6	4
8	Hydrodynamic Steering in Protein Association Revisited: Surprisingly Minuscule Effects of Considerable Torques. <i>Journal of Physical Chemistry B</i> , 2017, 121, 8475-8491.	2.6	3
9	UV-Vis spectroscopy of tyrosine side-groups in studies of protein structure. Part 1: basic principles and properties of tyrosine chromophore. <i>Biophysical Reviews</i> , 2016, 8, 151-161.	3.2	52
10	Effects of Spatially Dependent Mobilities on the Kinetics of the Diffusion-Controlled Association Derived from the First-Passage-Time Approach. <i>Journal of Physical Chemistry B</i> , 2016, 120, 7114-7127.	2.6	4
11	UV-Vis spectroscopy of tyrosine side-groups in studies of protein structure. Part 2: selected applications. <i>Biophysical Reviews</i> , 2016, 8, 163-177.	3.2	140
12	Toward an Accurate Modeling of Hydrodynamic Effects on the Translational and Rotational Dynamics of Biomolecules in Many-Body Systems. <i>Journal of Physical Chemistry B</i> , 2015, 119, 8425-8439.	2.6	14
13	8-Azapurines as isosteric purine fluorescent probes for nucleic acid and enzymatic research. <i>Molecular BioSystems</i> , 2014, 10, 2756-2774.	2.9	40
14	Evaluation of Proteins $\hat{\pm}$ Rotational Diffusion Coefficients from Simulations of Their Free Brownian Motion in Volume-Occupied Environments. <i>Journal of Chemical Theory and Computation</i> , 2014, 10, 481-491.	5.3	16
15	Transient Effects of Excluded Volume Interactions on the Translational Diffusion of Hydrodynamically Anisotropic Molecules. <i>Journal of Chemical Theory and Computation</i> , 2014, 10, 2583-2590.	5.3	6
16	Hydrodynamic Effects on the Relative Rotational Velocity of Associating Proteins. <i>Journal of Physical Chemistry B</i> , 2013, 117, 6165-6174.	2.6	10
17	Anisotropic Diffusion Effects on the Barnase $\hat{\pm}$ Barstar Encounter Kinetics. <i>Journal of Chemical Theory and Computation</i> , 2013, 9, 1667-1677.	5.3	5
18	Resolving Differences in Substrate Specificities between Human and Parasite Phosphoribosyltransferases via Analysis of Functional Groups of Substrates and Receptors. <i>Current Pharmaceutical Design</i> , 2013, 19, 4226-4240.	1.9	3

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19	Contributions of Far-Field Hydrodynamic Interactions to the Kinetics of Electrostatically Driven Molecular Association. <i>Journal of Physical Chemistry B</i> , 2012, 116, 5437-5447.	2.6	12
20	Poisson-Boltzmann continuum-solvation models: applications to pH-dependent properties of biomolecules. <i>Molecular BioSystems</i> , 2011, 7, 2923.	2.9	23
21	p <i>K</i> _a 's of Ionizable Groups and Energetics of Protein Conformational Transitions. <i>Journal of Physical Chemistry B</i> , 2010, 114, 1393-1406.	2.6	2
22	Poisson-Boltzmann model analysis of binding mRNA cap analogues to the translation initiation factor eIF4E. <i>Biophysical Chemistry</i> , 2009, 140, 16-23.	2.8	5
23	pH-Dependent Association of Proteins. The Test Case of Monoclonal Antibody HyHEL-5 and Its Antigen Hen Egg White Lysozyme. <i>Journal of Physical Chemistry B</i> , 2009, 113, 15662-15669.	2.6	4
24	Effects of Hydrodynamic Coupling on Electro-Optical Transients. <i>Journal of Physical Chemistry B</i> , 2009, 113, 13988-13992.	2.6	7
25	Protonation free energy levels in complex molecular systems. <i>Biopolymers</i> , 2008, 89, 262-269.	2.4	7
26	Multiple Protonation Equilibria in Electrostatics of Protein-Protein Binding. <i>Journal of Physical Chemistry B</i> , 2008, 112, 15074-15085.	2.6	9
27	Association of Aminoglycosidic Antibiotics with the Ribosomal A-Site Studied with Brownian Dynamics. <i>Journal of Chemical Theory and Computation</i> , 2008, 4, 549-559.	5.3	19
28	On the analysis of fluorimetric titration curves of purine nucleoside phosphorylase. <i>Nucleic Acids Symposium Series</i> , 2008, 52, 671-672.	0.3	0
29	Simulation of pH-Dependent Properties of Proteins Using Mesoscopic Models. <i>Reviews in Computational Chemistry</i> , 2007, , 249-311.	1.5	9
30	Biophysical Approach to Studies of Cap-eIF4E Interaction by Synthetic Cap Analogs. <i>Methods in Enzymology</i> , 2007, 430, 209-245.	1.0	33
31	Kinetics of Binding of Multisubstrate Analogue Inhibitor (2-Amino-9-[2-(Phosphonomethoxy)Ethyl]-6-Sulfanylpurine) with Trimeric Purine Nucleoside Phosphorylase. <i>Nucleosides, Nucleotides and Nucleic Acids</i> , 2007, 26, 969-974.	1.1	0
32	Brownian Dynamics Simulations of Binding mRNA Cap Analogues to eIF4E Protein. <i>Journal of Physical Chemistry B</i> , 2007, 111, 13107-13115.	2.6	10
33	Kinetics of binding the mRNA cap analogues to the translation initiation factor eIF4E under second-order reaction conditions. <i>Biophysical Chemistry</i> , 2007, 129, 289-297.	2.8	6
34	Towards the mechanism of trimeric purine nucleoside phosphorylases: Stopped-flow studies of binding of multisubstrate analogue inhibitor 2-amino-9-[2-(phosphonomethoxy)ethyl]-6-sulfanylpurine. <i>Biophysical Chemistry</i> , 2007, 125, 260-268.	2.8	11
35	Stopped-flow studies of guanine binding by calf spleen purine nucleoside phosphorylase. <i>Biophysical Chemistry</i> , 2005, 115, 67-76.	2.8	6
36	The impact of protonation equilibria on protein structure. <i>Journal of Physics Condensed Matter</i> , 2005, 17, S1607-S1616.	1.8	5

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37	Strong Effect of Hydrodynamic Coupling on the Electric Dichroism of Bent Rods. <i>Journal of Physical Chemistry B</i> , 2005, 109, 1034-1038.	2.6	14
38	Effects of Solute-Solvent Proton Exchange on Polypeptide Chain Dynamics: A Constant-pH Molecular Dynamics Study. <i>Journal of Physical Chemistry B</i> , 2005, 109, 13777-13784.	2.6	19
39	Constant-pH molecular dynamics study of protonation-structure relationship in a heptapeptide derived from ovomucoid third domain. <i>Physical Review E</i> , 2004, 69, 021915.	2.1	58
40	Constant-pH molecular dynamics simulations: a test case of succinic acid. <i>Chemical Physics</i> , 2004, 302, 161-170.	1.9	76
41	Effects of pH on kinetics of binding of mRNA-cap analogs by translation initiation factor eIF4E. <i>European Biophysics Journal</i> , 2003, 31, 608-616.	2.2	10
42	Charge-Charge Interactions are Key Determinants of the pK Values of Ionizable Groups in Ribonuclease Sa (pI=3.5) and a Basic Variant (pI=10.2). <i>Journal of Molecular Biology</i> , 2003, 325, 1077-1092.	4.2	96
43	pK Values of Histidine Residues in Ribonuclease Sa: Effect of Salt and Net Charge. <i>Journal of Molecular Biology</i> , 2003, 325, 1093-1105.	4.2	37
44	Prediction of Secondary Ionization of the Phosphate Group in Phosphotyrosine Peptides. <i>Biophysical Journal</i> , 2003, 84, 750-756.	0.5	24
45	Langevin dynamics of proteins at constant pH. <i>Physical Review E</i> , 2002, 66, 051911.	2.1	34
46	A procedure for analysis of stopped-flow transients for protein-ligand association. <i>Journal of Proteomics</i> , 2002, 51, 179-193.	2.4	7
47	Empirical relationships between protein structure and carboxyl pKa values in proteins. <i>Proteins: Structure, Function and Bioinformatics</i> , 2002, 48, 388-403.	2.6	191
48	Stopped-flow and Brownian dynamics studies of electrostatic effects in the kinetics of binding of 7-methyl-GpppG to the protein eIF4E. <i>European Biophysics Journal</i> , 2000, 29, 487-498.	2.2	42
49	Poisson-Boltzmann model studies of molecular electrostatic properties of the cAMP-dependent protein kinase. <i>European Biophysics Journal</i> , 1999, 28, 457-467.	2.2	10
50	Thermodynamic linkage between the binding of protons and inhibitors to HIV-1 protease. <i>Protein Science</i> , 1999, 8, 180-195.	7.6	55
51	Prediction of pKas of Titratable Residues in Proteins Using a Poisson-Boltzmann Model of the Solute-Solvent System. <i>Lecture Notes in Computational Science and Engineering</i> , 1999, , 176-196.	0.3	2
52	Theoretical and Experimental Analysis of Ionization Equilibria in Ovomuroid Third Domain. <i>Biochemistry</i> , 1998, 37, 8643-8652.	2.5	63
53	Brownian Dynamics of the Polarization of Rodlike Polyelectrolytes: Anisotropy and the Effect of Hydrodynamic Interactions. <i>Journal of Physical Chemistry B</i> , 1997, 101, 4478-4484.	2.6	8
54	On the Mechanism of Acetylcholinesterase Action: The Electrostatically Induced Acceleration of the Catalytic Acylation Step. <i>Journal of the American Chemical Society</i> , 1997, 119, 8159-8165.	13.7	53

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55	pKaMeasurements from Nuclear Magnetic Resonance for the B1 and B2 Immunoglobulin G-Binding Domains of Protein G: A Comparison with Calculated Values for Nuclear Magnetic Resonance and X-ray Structures. Biochemistry, 1997, 36, 3580-3589.	2.5	62
56	Prediction of titration properties of structures of a protein derived from molecular dynamics trajectories. Protein Science, 1997, 6, 373-382.	7.6	40
57	Simulation of electrostatic and hydrodynamic properties of Serratia endonuclease. , 1997, 41, 443-450.		10
58	HIV-1 Protease and its Inhibitors. , 1997, , 237-254.		0
59	The Determinants of pKas in Proteins. Biochemistry, 1996, 35, 7819-7833.	2.5	439
60	Computing ionization states of proteins with a detailed charge model. Journal of Computational Chemistry, 1996, 17, 1633-1644.	3.3	139
61	Orientational steering in enzyme-substrate association: Ionic strength dependence of hydrodynamic torque effects. European Biophysics Journal, 1996, 24, 137-41.	2.2	37
62	Acetylcholinesterase: Role of the enzyme's charge distribution in steering charged ligands toward the active site. Biopolymers, 1996, 39, 85-94.	2.4	31
63	Acetylcholinesterase: role of the enzyme's charge distribution in steering charged ligands toward the active site. Biopolymers, 1996, 39, 85-94.	2.4	16
64	Acetylcholinesterase: Role of the enzyme's charge distribution in steering charged ligands toward the active site. Biopolymers, 1996, 39, 85-94.	2.4	28
65	Electrostatics and diffusion of molecules in solution: simulations with the University of Houston Brownian Dynamics program. Computer Physics Communications, 1995, 91, 57-95.	7.5	622
66	Simulation of charge-mutant acetylcholinesterases. Biochemistry, 1995, 34, 4211-4219.	2.5	63
67	Electrostatic and hydrodynamic orientational steering effects in enzyme-substrate association. Biophysical Journal, 1995, 69, 57-65.	0.5	60
68	Computation of the dipole moments of proteins. Biophysical Journal, 1995, 69, 1344-1354.	0.5	71
69	Acetylcholinesterase: diffusional encounter rate constants for dumbbell models of ligand. Biophysical Journal, 1995, 68, 62-68.	0.5	54
70	Electrostatics of hemoglobins from measurements of the electric dichroism and computer simulations. Biophysical Journal, 1995, 68, 655-664.	0.5	28
71	Binding of Cations and Protons in the Active Site of Acetylcholinesterase. Jerusalem Symposia on Quantum Chemistry and Biochemistry, 1995, , 25-37.	0.2	5
72	Prediction of Ph-dependent Properties of Proteins. Journal of Molecular Biology, 1994, 238, 415-436.	4.2	807

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73	Brownian Dynamics of the Polarization of Rodlike Polyelectrolytes. <i>The Journal of Physical Chemistry</i> , 1994, 98, 10881-10887.	2.9	13
74	Acetylcholinesterase: Effects of Ionic Strength and Dimerization on the Rate Constants. <i>Israel Journal of Chemistry</i> , 1994, 34, 151-158.	2.3	21
75	Structure and dynamics of curved DNA fragments in solution: Evidence for slow modes of configurational transitions. <i>Biophysical Chemistry</i> , 1993, 47, 179-191.	2.8	23
76	Brownian dynamics simulation of electrooptical transients for complex macrodipoles. <i>The Journal of Physical Chemistry</i> , 1993, 97, 2767-2773.	2.9	20
77	Modes of rotational motion of wormlike chains and the effect of charges on electrooptical transients. <i>Macromolecules</i> , 1992, 25, 6500-6504.	4.8	6
78	Electric moments of rodlike molecules due to asymmetry of ligand binding induced by electric fields. <i>The Journal of Physical Chemistry</i> , 1991, 95, 5983-5988.	2.9	6
79	Brownian dynamics simulation of electrooptical transients for solutions of rigid macromolecules. <i>Journal of Chemical Physics</i> , 1991, 95, 1354-1360.	3.0	16
80	Permanent dipole moment of tRNA's and variation of their structure in solution. <i>Biophysical Journal</i> , 1990, 58, 403-411.	0.5	17
81	An unusual electrooptical effect observed for DNA fragments and its apparent relation to a permanent electric moment associated with bent DNA. <i>Biophysical Chemistry</i> , 1989, 33, 19-30.	2.8	42
82	The nature of protein dipole moments: experimental and calculated permanent dipole of .alpha.-chymotrypsin. <i>Biochemistry</i> , 1989, 28, 10072-10078.	2.5	64
83	Volume correction for bead model simulations of rotational friction coefficients of macromolecules. <i>The Journal of Physical Chemistry</i> , 1989, 93, 5301-5305.	2.9	44
84	Helix-coil dynamics of a Z-helix hairpin. <i>Biopolymers</i> , 1988, 27, 1319-1327.	2.4	7
85	Structure of the Tet repressor and Tet repressor-operator complexes in solution from electrooptical measurements and hydrodynamic simulations. <i>Biochemistry</i> , 1988, 27, 4674-4679.	2.5	23
86	Turn of Promotor DNA by cAMP Receptor Protein Characterized by Bead Model Simulation of Rotational Diffusion. <i>Journal of Biomolecular Structure and Dynamics</i> , 1988, 5, 819-837.	3.5	21
87	Quasichemical interpretation of the ultrasonic velocity in ternary aqueous systems. <i>Journal of Solution Chemistry</i> , 1987, 16, 285-294.	1.2	2
88	Ultrasonic Velocity Studies on Carbohydrates in Aqueous Ethanolic Solutions. <i>Zeitschrift Fur Physikalische Chemie</i> , 1986, 148, 185-195.	2.8	7
89	Hydration of alcohols by ultrasonic velocity measurements in ternary systems. <i>Journal of Solution Chemistry</i> , 1984, 13, 493-503.	1.2	11
90	Dependence of ultrasonic velocity on structure in a homologous series of nonelectrolytes in aqueous medium. <i>Journal of Solution Chemistry</i> , 1983, 12, 123-133.	1.2	6

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91	Hydration of alcohols in aqueous methanol solutions from ultrasonic velocity measurements. Journal of Solution Chemistry, 1983, 12, 783-789.	1.2	6
92	Ultrasonic studies on hydration of pyrimidine nucleosides in aqueous ethanolic solutions. The Journal of Physical Chemistry, 1982, 86, 4831-4834.	2.9	12
93	Methyl esterification of m ⁷ G5p reversibly blocks its activity as an analog of eukaryotic mRNA 5'-caps. Journal of Molecular Biology, 1981, 153, 451-458.	4.2	26