

Dario A Estrin

List of Publications by Year in descending order

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200
papers

6,876
citations

57758

44
h-index

98798

67
g-index

203
all docs

203
docs citations

203
times ranked

6871
citing authors

#	ARTICLE	IF	CITATIONS
1	When Galectins Recognize Glycans: From Biochemistry to Physiology and Back Again. <i>Biochemistry</i> , 2011, 50, 7842-7857.	2.5	238
2	Small Clusters of Water Molecules Using Density Functional Theory. <i>The Journal of Physical Chemistry</i> , 1996, 100, 8701-8711.	2.9	182
3	A Density Functional Study of Tautomerism of Uracil and Cytosine. <i>The Journal of Physical Chemistry</i> , 1994, 98, 5653-5660.	2.9	140
4	The Catalytic Mechanism of Peptidylglycine β -Hydroxylating Monooxygenase Investigated by Computer Simulation. <i>Journal of the American Chemical Society</i> , 2006, 128, 12817-12828.	13.7	137
5	Spectroscopy in Complex Environments from QM \rightarrow MM Simulations. <i>Chemical Reviews</i> , 2018, 118, 4071-4113.	47.7	136
6	A DFT-Based QM-MM Approach Designed for the Treatment of Large Molecular Systems: Application to Chorismate Mutase. <i>Journal of Physical Chemistry B</i> , 2003, 107, 13728-13736.	2.6	116
7	Aromatic \rightarrow Aromatic Interactions in Proteins: Beyond the Dimer. <i>Journal of Chemical Information and Modeling</i> , 2011, 51, 1623-1633.	5.4	115
8	Evidence for a ferryl intermediate in a heme-based dioxygenase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 17371-17376.	7.1	113
9	Multiple-Steering QM \rightarrow MM Calculation of the Free Energy Profile in Chorismate Mutase. <i>Journal of the American Chemical Society</i> , 2005, 127, 6940-6941.	13.7	112
10	Theoretical Study of the Truncated Hemoglobin HbN: Exploring the Molecular Basis of the NO Detoxification Mechanism. <i>Journal of the American Chemical Society</i> , 2005, 127, 4433-4444.	13.7	111
11	pH-Dependent Conformational Changes in Proteins and Their Effect on Experimental pKas: The Case of Nitrophorin 4. <i>PLoS Computational Biology</i> , 2012, 8, e1002761.	3.2	110
12	Discrimination of Nitroxyl and Nitric Oxide by Water-Soluble Mn(III) Porphyrins. <i>Journal of the American Chemical Society</i> , 2005, 127, 4680-4684.	13.7	109
13	Ligand-induced dynamical regulation of NO conversion in Mycobacterium tuberculosis truncated hemoglobin-N. <i>Proteins: Structure, Function and Bioinformatics</i> , 2006, 64, 457-464.	2.6	95
14	Heme Protein Oxygen Affinity Regulation Exerted by Proximal Effects. <i>Journal of the American Chemical Society</i> , 2006, 128, 12455-12461.	13.7	91
15	Dioxygen affinity in heme proteins investigated by computer simulation. <i>Journal of Inorganic Biochemistry</i> , 2006, 100, 761-770.	3.5	89
16	Fast Nitroxyl Trapping by Ferric Porphyrins. <i>Journal of the American Chemical Society</i> , 2003, 125, 15272-15273.	13.7	82
17	Naturally occurring fluorescence in frogs. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 3672-3677.	7.1	81
18	Scavenging of NO by Melatonin. <i>Journal of the American Chemical Society</i> , 2000, 122, 10468-10469.	13.7	80

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19	Free Energy Calculations with Non-Equilibrium Methods: Applications of the Jarzynski Relationship. <i>Theoretical Chemistry Accounts</i> , 2006, 116, 338-346.	1.4	79
20	Modeling heme proteins using atomistic simulations. <i>Physical Chemistry Chemical Physics</i> , 2006, 8, 5611-5628.	2.8	77
21	Solvent effects in density functional calculations of uracil and cytosine tautomerism. <i>International Journal of Quantum Chemistry</i> , 1995, 56, 615-625.	2.0	74
22	Structure and spin-state energetics of an iron porphyrin model: An assessment of theoretical methods. <i>International Journal of Quantum Chemistry</i> , 2002, 87, 158-166.	2.0	74
23	Reactions of Melatonin and Related Indoles with Free Radicals: A Computational Study. <i>Journal of Medicinal Chemistry</i> , 1998, 41, 3684-3689.	6.4	73
24	Complete Reaction Mechanism of Indoleamine 2,3-Dioxygenase as Revealed by QM/MM Simulations. <i>Journal of Physical Chemistry B</i> , 2012, 116, 1401-1413.	2.6	68
25	Catalysis of Peroxide Reduction by Fast Reacting Protein Thiols. <i>Chemical Reviews</i> , 2019, 119, 10829-10855.	47.7	68
26	Acidity and nucleophilic reactivity of glutathione persulfide. <i>Journal of Biological Chemistry</i> , 2020, 295, 15466-15481.	3.4	68
27	Molecular Basis for the Electric Field Modulation of Cytochrome <i>c</i> Structure and Function. <i>Journal of the American Chemical Society</i> , 2009, 131, 16248-16256.	13.7	66
28	Inhibitory effect of quercetin on matrix metalloproteinase 9 activity Molecular mechanism and structure-activity relationship of the flavonoid-enzyme interaction. <i>European Journal of Pharmacology</i> , 2010, 644, 138-145.	3.5	65
29	Nitric Oxide Interaction with Cytochrome <i>c</i> and Its Relevance to Guanylate Cyclase. Why Does the Iron Histidine Bond Break?. <i>Journal of the American Chemical Society</i> , 2005, 127, 7721-7728.	13.7	64
30	A Microscopic Study of the Deoxyhemoglobin-Catalyzed Generation of Nitric Oxide from Nitrite Anion. <i>Biochemistry</i> , 2008, 47, 9793-9802.	2.5	62
31	Molecular Basis of the Mechanism of Thiol Oxidation by Hydrogen Peroxide in Aqueous Solution: Challenging the S _N ² Paradigm. <i>Chemical Research in Toxicology</i> , 2012, 25, 741-746.	3.3	61
32	Theoretical and Experimental Study of Medium Effects on the Structure and Spectroscopy of the [Fe(CN)5NO]2- Ion. <i>Inorganic Chemistry</i> , 1996, 35, 3897-3903.	4.0	59
33	Role of Pre-A Motif in Nitric Oxide Scavenging by Truncated Hemoglobin, HbN, of Mycobacterium tuberculosis. <i>Journal of Biological Chemistry</i> , 2009, 284, 14457-14468.	3.4	59
34	Exploring the molecular basis of heme coordination in human neuroglobin. <i>Proteins: Structure, Function and Bioinformatics</i> , 2008, 71, 695-705.	2.6	54
35	The first step of the dioxygenation reaction carried out by tryptophan dioxygenase and indoleamine 2,3-dioxygenase as revealed by quantum mechanical/molecular mechanical studies. <i>Journal of Biological Inorganic Chemistry</i> , 2010, 15, 811-823.	2.6	53
36	Transnitrosation of Nitrosothiols: Characterization of an Elusive Intermediate. <i>Journal of the American Chemical Society</i> , 2005, 127, 486-487.	13.7	51

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37	Dynamical Characterization of the Heme NO Oxygen Binding (HNOX) Domain. Insight into Soluble Guanylate Cyclase Allosteric Transition. <i>Biochemistry</i> , 2008, 47, 9416-9427.	2.5	49
38	Role of Heme Distortion on Oxygen Affinity in Heme Proteins: The Protoglobin Case. <i>Journal of Physical Chemistry B</i> , 2010, 114, 8536-8543.	2.6	49
39	A Surprisingly Stable S-Nitrosothiol Complex. <i>Journal of the American Chemical Society</i> , 2006, 128, 2512-2513.	13.7	48
40	Exploring the molecular basis of human manganese superoxide dismutase inactivation mediated by tyrosine 34 nitration. <i>Archives of Biochemistry and Biophysics</i> , 2011, 507, 304-309.	3.0	48
41	Physiological concentrations of melatonin inhibit the nitridergic pathway in the Syrian hamster retina. <i>Journal of Pineal Research</i> , 2002, 33, 31-36.	7.4	47
42	Structural determinants of ligand migration in <i>Mycobacterium tuberculosis</i> truncated hemoglobin O. <i>Proteins: Structure, Function and Bioinformatics</i> , 2008, 73, 372-379.	2.6	47
43	3-Nitrotyrosine and related derivatives in proteins: precursors, radical intermediates and impact in function. <i>Essays in Biochemistry</i> , 2020, 64, 111-133.	4.7	47
44	Solvent-induced symmetry breaking of nitrate ion in aqueous clusters: A quantum-classical simulation study. <i>Journal of Chemical Physics</i> , 2002, 117, 2718-2725.	3.0	46
45	Dynamical Regulation of Ligand Migration by a Gate-Opening Molecular Switch in Truncated Hemoglobin-N from <i>Mycobacterium tuberculosis</i> . <i>Journal of the American Chemical Society</i> , 2007, 129, 6782-6788.	13.7	46
46	Electronic energy shifts of a sodium atom in argon clusters by simulated annealing. <i>Journal of Chemical Physics</i> , 1990, 93, 7187-7200.	3.0	45
47	Copper-Transfer Mechanism from the Human Chaperone Atox1 to a Metal-Binding Domain of Wilson Disease Protein. <i>Journal of Physical Chemistry B</i> , 2010, 114, 3698-3706.	2.6	44
48	Structural basis of redox-dependent modulation of galectin-1 dynamics and function. <i>Glycobiology</i> , 2014, 24, 428-441.	2.5	44
49	AM1 Study of the Ground and Excited State Potential Energy Surfaces of Symmetric Carbocyanines. <i>Journal of Physical Chemistry A</i> , 1997, 101, 6998-7006.	2.5	43
50	High pressure reveals structural determinants for globin hexacoordination: Neuroglobin and myoglobin cases. <i>Proteins: Structure, Function and Bioinformatics</i> , 2009, 75, 885-894.	2.6	43
51	The extraordinary catalytic ability of peroxiredoxins: a combined experimental and QM/MM study on the fast thiol oxidation step. <i>Chemical Communications</i> , 2014, 50, 10070-10073.	4.1	43
52	Molecular Basis for the Substrate Stereoselectivity in Tryptophan Dioxygenase. <i>Biochemistry</i> , 2011, 50, 10910-10918.	2.5	42
53	Insights into the mechanism of the reaction between hydrogen sulfide and peroxynitrite. <i>Free Radical Biology and Medicine</i> , 2015, 80, 93-100.	2.9	41
54	Nitrosation of melatonin by nitric oxide: a computational study. <i>Journal of Pineal Research</i> , 2001, 31, 97-101.	7.4	40

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55	Theoretical insight into the hydroxylamine oxidoreductase mechanism. <i>Journal of Inorganic Biochemistry</i> , 2008, 102, 1523-1530.	3.5	40
56	NMR and molecular dynamics studies of the interaction of melatonin with calmodulin. <i>Protein Science</i> , 2008, 13, 2925-2938.	7.6	40
57	Tetrachlorocarbonyliridates: Water-Soluble Carbon Monoxide Releasing Molecules Rate-Modulated by the Sixth Ligand. <i>Inorganic Chemistry</i> , 2011, 50, 2334-2345.	4.0	40
58	Modulation of the NO trans effect in heme proteins: implications for the activation of soluble guanylate cyclase. <i>Journal of Biological Inorganic Chemistry</i> , 2003, 8, 595-600.	2.6	39
59	QM ²⁺ MM Study of Nitrite Reduction by Nitrite Reductase of <i>Pseudomonas aeruginosa</i> . <i>Journal of Physical Chemistry B</i> , 2004, 108, 18073-18080.	2.6	39
60	Comparing and combining implicit ligand sampling with multiple steered molecular dynamics to study ligand migration processes in heme proteins. <i>Journal of Computational Chemistry</i> , 2011, 32, 2219-2231.	3.3	39
61	Design, synthesis and biological evaluation of quinoxaline compounds as anti-HIV agents targeting reverse transcriptase enzyme. <i>European Journal of Medicinal Chemistry</i> , 2020, 188, 111987.	5.5	39
62	Bond or Cage Effect: How Nitrophorins Transport and Release Nitric Oxide. <i>Journal of the American Chemical Society</i> , 2008, 130, 1611-1618.	13.7	38
63	Theoretical Investigation on the Electronic Structure of Pentacyano(L)ferrate(II) Complexes with NO ⁺ , NO, and NO-Ligands. Redox Interconversion, Protonation, and Cyanide-Releasing Reactions. <i>Inorganic Chemistry</i> , 2001, 40, 4127-4133.	4.0	37
64	pH-Dependent Mechanism of Nitric Oxide Release in Nitrophorins 2 and 4. <i>Journal of Physical Chemistry B</i> , 2009, 113, 1192-1201.	2.6	37
65	Small ligand-globin interactions: Reviewing lessons derived from computer simulation. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2013, 1834, 1722-1738.	2.3	37
66	Unraveling the molecular basis for ligand binding in truncated hemoglobins: The trHbO <i>Bacillus subtilis</i> case. <i>Proteins: Structure, Function and Bioinformatics</i> , 2010, 78, 962-970.	2.6	36
67	Reactivity of Inorganic Sulfide Species toward a Heme Protein Model. <i>Inorganic Chemistry</i> , 2015, 54, 527-533.	4.0	36
68	Evolutionary and Functional Relationships in the Truncated Hemoglobin Family. <i>PLoS Computational Biology</i> , 2016, 12, e1004701.	3.2	36
69	Electric field effects on the reactivity of heme model systems. <i>Chemical Physics Letters</i> , 2007, 434, 121-126.	2.6	35
70	HNO trapping and assisted decomposition of nitroxyl donors by ferric hemes. <i>Polyhedron</i> , 2007, 26, 4673-4679.	2.2	35
71	Mechanism of Product Release in NO Detoxification from <i>Mycobacterium tuberculosis</i> Truncated Hemoglobin N. <i>Journal of the American Chemical Society</i> , 2008, 130, 1688-1693.	13.7	35
72	Substrate stereo-specificity in tryptophan dioxygenase and indoleamine 2,3-dioxygenase. <i>Proteins: Structure, Function and Bioinformatics</i> , 2010, 78, 2961-2972.	2.6	35

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73	Oxygen affinity controlled by dynamical distal conformations: The soybean leghemoglobin and the <i>Paramecium caudatum</i> hemoglobin cases. <i>Proteins: Structure, Function and Bioinformatics</i> , 2007, 68, 480-487.	2.6	33
74	Carbohydrate-Binding Proteins: Dissecting Ligand Structures through Solvent Environment Occupancy. <i>Journal of Physical Chemistry B</i> , 2009, 113, 8717-8724.	2.6	33
75	Characterization of the Galectin-1 Carbohydrate Recognition Domain in Terms of Solvent Occupancy. <i>Journal of Physical Chemistry B</i> , 2007, 111, 7360-7366.	2.6	31
76	Ligand Migration in <i>Methanosarcina acetivorans</i> Protoglobin: Effects of Ligand Binding and Dimeric Assembly. <i>Journal of Physical Chemistry B</i> , 2011, 115, 13771-13780.	2.6	31
77	Molecular basis of intramolecular electron transfer in proteins during radical-mediated oxidations: Computer simulation studies in model tyrosine-cysteine peptides in solution. <i>Archives of Biochemistry and Biophysics</i> , 2012, 525, 82-91.	3.0	31
78	Protein dynamics and ligand migration interplay as studied by computer simulation. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2011, 1814, 1054-1064.	2.3	30
79	Mechanism of cysteine oxidation by peroxyxynitrite: An integrated experimental and theoretical study. <i>Archives of Biochemistry and Biophysics</i> , 2013, 539, 81-86.	3.0	30
80	Fluoride as a Probe for H-Bonding Interactions in the Active Site of Heme Proteins: The Case of <i>Thermobifida fusca</i> Hemoglobin. <i>Journal of the American Chemical Society</i> , 2011, 133, 20970-20980.	13.7	29
81	Thiol redox biochemistry: insights from computer simulations. <i>Biophysical Reviews</i> , 2014, 6, 27-46.	3.2	29
82	Electronic spectra of indolyl radicals: a time-dependent DFT study. <i>Chemical Physics Letters</i> , 2002, 365, 15-21.	2.6	28
83	Mechanisms of NO Release by N1-Nitrosomelatonin: Nucleophilic Attack versus Reducing Pathways. <i>Journal of Organic Chemistry</i> , 2005, 70, 5790-5798.	3.2	28
84	Hydrophobic Effect Drives Oxygen Uptake in Myoglobin via Histidine E7. <i>Journal of Biological Chemistry</i> , 2013, 288, 6754-6762.	3.4	28
85	Access and Binding of H ₂ S to Hemeproteins: The Case of Hbl of <i>Lucina pectinata</i> . <i>Journal of Physical Chemistry B</i> , 2016, 120, 9642-9653.	2.6	28
86	Biophysical Characterisation of Neuroglobin of the Icefish, a Natural Knockout for Hemoglobin and Myoglobin. Comparison with Human Neuroglobin. <i>PLoS ONE</i> , 2012, 7, e44508.	2.5	28
87	Concertedness and solvent effects in multiple proton transfer reactions: The formic acid dimer in solution. <i>Journal of Chemical Physics</i> , 2000, 112, 9498-9508.	3.0	27
88	Hydrogen Bonding and O ₂ Affinity of Hemoglobins. <i>Journal of the American Chemical Society</i> , 2001, 123, 8436-8437.	13.7	27
89	The hemoglobins of the sub-Antarctic fish <i>Cottoperca gobio</i> , a phylogenetically basal species: oxygen-binding equilibria, kinetics and molecular dynamics. <i>FEBS Journal</i> , 2009, 276, 2266-2277.	4.7	27
90	Nitric Oxide Binding to Ferric Cytochrome P450: A Computational Study. <i>Inorganic Chemistry</i> , 2000, 39, 2352-2359.	4.0	26

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91	Solvation and Structure of LiAlH ₄ in Ethereal Solvents. <i>Inorganic Chemistry</i> , 2005, 44, 5286-5292.	4.0	26
92	Nitric Oxide Reactivity with Globins as Investigated Through Computer Simulation. <i>Methods in Enzymology</i> , 2008, 437, 477-498.	1.0	26
93	Structural Model for p75NTR ^{TrkA} Intracellular Domain Interaction: A Combined FRET and Bioinformatics Study. <i>Journal of Molecular Biology</i> , 2011, 414, 681-698.	4.2	26
94	Role of PheE15 Gate in Ligand Entry and Nitric Oxide Detoxification Function of <i>Mycobacterium tuberculosis</i> Truncated Hemoglobin N. <i>PLoS ONE</i> , 2012, 7, e49291.	2.5	26
95	Chemical Reactivity and Spectroscopy Explored From QM/MM Molecular Dynamics Simulations Using the LIO Code. <i>Frontiers in Chemistry</i> , 2018, 6, 70.	3.6	26
96	Electronic spectra of NaAr ₄ and NaAr ₆ : Isomerization and melting. <i>Journal of Chemical Physics</i> , 1992, 96, 7977-7991.	3.0	25
97	Structure, solvation, and bonding in pentacyano(L)ferrate(II) ions (L=aliphatic amine): a density functional study. <i>Journal of Molecular Modeling</i> , 2001, 7, 201-206.	1.8	25
98	Heme Pocket Structural Properties of a Bacterial Truncated Hemoglobin from <i>Thermobifida fusca</i> . <i>Biochemistry</i> , 2010, 49, 10394-10402.	2.5	25
99	Ligand Uptake Modulation by Internal Water Molecules and Hydrophobic Cavities in Hemoglobins. <i>Journal of Physical Chemistry B</i> , 2014, 118, 1234-1245.	2.6	25
100	Structure and Bonding in Pentacyano(L)ferrate(II) and Pentacyano(L)ruthenate(II) Complexes (L = Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 6832-6837.	4.0	24
101	Coordination of peroxide to the CuM center of peptidylglycine β -hydroxylating monooxygenase (PHM): structural and computational study. <i>Journal of Biological Inorganic Chemistry</i> , 2013, 18, 223-232.	2.6	24
102	Structural flexibility of the heme cavity in the cold-adapted truncated hemoglobin from the Antarctic marine bacterium <i>Pseudoalteromonas haloplanktis</i> TAC 125. <i>FEBS Journal</i> , 2015, 282, 2948-2965.	4.7	24
103	Environment effects on chemical reactivity of heme proteins. <i>International Journal of Quantum Chemistry</i> , 2002, 90, 1505-1514.	2.0	23
104	DFT study on the reactivity of iron porphyrins tuned by ring substitution. <i>Journal of Inorganic Biochemistry</i> , 2008, 102, 70-76.	3.5	23
105	A Unique Family of Stable and Water-Soluble S-Nitrosothiol Complexes. <i>Inorganic Chemistry</i> , 2008, 47, 4723-4733.	4.0	23
106	An Integrated Computational Analysis of the Structure, Dynamics, and Ligand Binding Interactions of the Human Galectin Network. <i>Journal of Chemical Information and Modeling</i> , 2011, 51, 1918-1930.	5.4	23
107	Role of the Distal Hydrogen-Bonding Network in Regulating Oxygen Affinity in the Truncated Hemoglobin III from <i>Campylobacter jejuni</i> . <i>Biochemistry</i> , 2011, 50, 3946-3956.	2.5	23
108	Structural basis for ligand recognition in a mushroom lectin: solvent structure as specificity predictor. <i>Carbohydrate Research</i> , 2011, 346, 939-948.	2.3	23

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109	Ligand Migration in the Apolar Tunnel of <i>Cerebratulus lacteus</i> Mini-Hemoglobin. <i>Journal of Biological Chemistry</i> , 2011, 286, 5347-5358.	3.4	23
110	Molecular Mechanism of Myoglobin Autoxidation: Insights from Computer Simulations. <i>Journal of Physical Chemistry B</i> , 2015, 119, 1802-1813.	2.6	23
111	PrxQ B from <i>Mycobacterium tuberculosis</i> is a monomeric, thioredoxin-dependent and highly efficient fatty acid hydroperoxide reductase. <i>Free Radical Biology and Medicine</i> , 2016, 101, 249-260.	2.9	23
112	Tertiary and quaternary structural basis of oxygen affinity in human hemoglobin as revealed by multiscale simulations. <i>Scientific Reports</i> , 2017, 7, 10926.	3.3	23
113	Stabilization of Aliphatic and Aromatic Diazonium Ions by Coordination: An Experimental and Theoretical Study. <i>Organometallics</i> , 2000, 19, 3810-3817.	2.3	22
114	Exploring the Molecular Basis of Action of the Passive Antiglucocorticoid 21-Hydroxy-6,19-epoxyprogesterone. <i>Journal of Medicinal Chemistry</i> , 2008, 51, 1352-1360.	6.4	22
115	Interaction between Proteins and Ir Based CO Releasing Molecules: Mechanism of Adduct Formation and CO Release. <i>Inorganic Chemistry</i> , 2014, 53, 10456-10462.	4.0	22
116	Following Ligand Migration Pathways from Picoseconds to Milliseconds in Type II Truncated Hemoglobin from <i>Thermobifida fusca</i> . <i>PLoS ONE</i> , 2012, 7, e39884.	2.5	22
117	Two distinct heme distal site states define <i>Cerebratulus lacteus</i> mini-hemoglobin oxygen affinity. <i>Proteins: Structure, Function and Bioinformatics</i> , 2005, 62, 641-648.	2.6	21
118	The peculiar heme pocket of the 2/2 hemoglobin of cold-adapted <i>Pseudoalteromonas haloplanktis</i> TAC125. <i>Journal of Biological Inorganic Chemistry</i> , 2011, 16, 299-311.	2.6	21
119	H-bonding networks of the distal residues and water molecules in the active site of <i>Thermobifida fusca</i> hemoglobin. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2013, 1834, 1901-1909.	2.3	21
120	Coupling of tyrosine deprotonation and axial ligand exchange in nitrocytochrome c. <i>Chemical Communications</i> , 2014, 50, 2592-2594.	4.1	21
121	Structural Study of a Flexible Active Site Loop in Human Indoleamine 2,3-Dioxygenase and Its Functional Implications. <i>Biochemistry</i> , 2016, 55, 2785-2793.	2.5	21
122	Kinetics, subcellular localization, and contribution to parasite virulence of a <i>Trypanosoma cruzi</i> hybrid type A heme peroxidase (<i>Tc</i> APx-CcP). <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E1326-E1335.	7.1	21
123	Engineered chimeras reveal the structural basis of hexacoordination in globins: A case study of neuroglobin and myoglobin. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2015, 1850, 169-177.	2.4	20
124	Thermal Stability of Globins: Implications of Flexibility and Heme Coordination Studied by Molecular Dynamics Simulations. <i>Journal of Chemical Information and Modeling</i> , 2019, 59, 441-452.	5.4	20
125	Hybrid Quantum Classical Molecular Dynamics Simulation of the Proton-Transfer Reaction of HO ₂ with HBr in Aqueous Clusters. <i>Journal of Physical Chemistry A</i> , 1999, 103, 5105-5112.	2.5	19
126	Molecular structure effects on the kinetics of hydroxyl radical addition to azo dyes. <i>Journal of Physical Organic Chemistry</i> , 2002, 15, 287-292.	1.9	19

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127	Molecular Basis for the pH Dependent Structural Transition of Nitrophorin 4. <i>Journal of Physical Chemistry B</i> , 2009, 113, 2135-2142.	2.6	19
128	Mechanism of Sulfide Binding by Ferric Heme proteins. <i>Inorganic Chemistry</i> , 2018, 57, 7591-7600.	4.0	19
129	Structural and Energetic Study of Cisplatin and Derivatives: Comparison of the Performance of Density Functional Theory Implementations. <i>Journal of Chemical Theory and Computation</i> , 2008, 4, 740-750.	5.3	18
130	Linking the Structure and Thermal Stability of β -Galactoside-Binding Protein Galectin-1 to Ligand Binding and Dimerization Equilibria. <i>Biochemistry</i> , 2010, 49, 7652-7658.	2.5	18
131	Molecular Basis of Hydroperoxide Specificity in Peroxiredoxins: The Case of AhpE from <i>Mycobacterium tuberculosis</i> . <i>Biochemistry</i> , 2015, 54, 7237-7247.	2.5	18
132	Gating in plant plasma membrane aquaporins: the involvement of leucine in the formation of a pore constriction in the closed state. <i>FEBS Journal</i> , 2019, 286, 3473-3487.	4.7	18
133	Mapping the protein-binding sites for iridium(III)-based CO-releasing molecules. <i>Dalton Transactions</i> , 2016, 45, 12206-12214.	3.3	18
134	Solvent Effects on Peroxynitrite Structure and Properties from QM/MM Simulations. <i>Journal of Physical Chemistry A</i> , 2005, 109, 9598-9604.	2.5	17
135	Lessons learned about steered molecular dynamics simulations and free energy calculations. <i>Chemical Biology and Drug Design</i> , 2019, 93, 1129-1138.	3.2	17
136	Proximal effects in the modulation of nitric oxide synthase reactivity: a QM-MM study. <i>Journal of Biological Inorganic Chemistry</i> , 2005, 10, 595-604.	2.6	16
137	Hemisuccinate of 21 α -Hydroxy-19 α -Epoxyprogesterone: A Tissue-Specific Modulator of the Glucocorticoid Receptor. <i>ChemMedChem</i> , 2008, 3, 1869-1877.	3.2	16
138	Impact of human galectin-1 binding to saccharide ligands on dimer dissociation kinetics and structure. <i>Glycobiology</i> , 2016, 26, 1317-1327.	2.5	16
139	Short-range and long-range solvent effects on charge-transfer-to-solvent transitions of $\text{I}^{\bullet-}$ and $\text{K}+\text{I}^{\bullet-}$ contact ion pair dissolved in supercritical ammonia. <i>Journal of Chemical Physics</i> , 2007, 126, 174504.	3.0	15
140	Mechanistic Insight into the Enzymatic Reduction of Truncated Hemoglobin N of <i>Mycobacterium tuberculosis</i> . <i>Journal of Biological Chemistry</i> , 2014, 289, 21573-21583.	3.4	15
141	Interplay of the H-Bond Donor-Acceptor Role of the Distal Residues in Hydroxyl Ligand Stabilization of <i>Thermobifida fusca</i> Truncated Hemoglobin. <i>Biochemistry</i> , 2014, 53, 8021-8030.	2.5	15
142	A quantitative model for oxygen uptake and release in a family of heme proteins. <i>Bioinformatics</i> , 2016, 32, 1805-1813.	4.1	15
143	Role of Core Electrons in Quantum Dynamics Using TDDFT. <i>Journal of Chemical Theory and Computation</i> , 2017, 13, 77-85.	5.3	15
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