List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Enzyme Replacement Therapy for Genetic Disorders Associated with Enzyme Deficiency. Current Medicinal Chemistry, 2022, 29, 489-525.	2.4	12
2	From hemoglobin allostery to hemoglobin-based oxygen carriers. Molecular Aspects of Medicine, 2022, 84, 101050.	6.4	15
3	Targeted Biologics: The New Frontier for Precision Therapy. Current Medicinal Chemistry, 2022, 29, 383-384.	2.4	2
4	Human Serine Racemase Weakly Binds the Third PDZ Domain of PSD-95. International Journal of Molecular Sciences, 2022, 23, 4959.	4.1	1
5	Human serine racemase is inhibited by glyceraldehyde 3-phosphate, but not by glyceraldehyde 3-phosphate dehydrogenase. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2021, 1869, 140544.	2.3	3
6	The allosteric interplay between Sâ€nitrosylation and glycine binding controls the activity of human serine racemase. FEBS Journal, 2021, 288, 3034-3054.	4.7	8
7	Investigational Studies on a Hit Compound Cyclopropane–Carboxylic Acid Derivative Targeting <i>O</i> -Acetylserine Sulfhydrylase as a Colistin Adjuvant. ACS Infectious Diseases, 2021, 7, 281-292.	3.8	13
8	Discovery of Substituted (2-Aminooxazol-4-yl)Isoxazole-3-carboxylic Acids as Inhibitors of Bacterial Serine Acetyltransferase in the Quest for Novel Potential Antibacterial Adjuvants. Pharmaceuticals, 2021, 14, 174.	3.8	5
9	A Competitive O-Acetylserine Sulfhydrylase Inhibitor Modulates the Formation of Cysteine Synthase Complex. Catalysts, 2021, 11, 700.	3.5	4
10	Stability of Maleimide-PEG and Mono-Sulfone-PEG Conjugation to a Novel Engineered Cysteine in the Human Hemoglobin Alpha Subunit. Frontiers in Chemistry, 2021, 9, 707797.	3.6	4
11	A Key Silencing Histone Mark on Chromatin Is Lost When Colorectal Adenocarcinoma Cells Are Depleted of Methionine by Methionine γ-Lyase. Frontiers in Molecular Biosciences, 2021, 8, 735303.	3.5	7
12	Modulation of Oxygen Affinity in Hemoglobin-based Oxygen Carriers. Regenerative Medicine, Artificial Cells and Nanomedicine, 2021, , 375-403.	0.1	1
13	Structural and Functional Characterization of the Globin-Coupled Sensors ofAzotobacter vinelandiiandBordetella pertussis. Antioxidants and Redox Signaling, 2020, 32, 378-395.	5.4	4
14	SP-B and SP-C analogues within CHF5633 synthetic surfactant probed by fluorescence labeling. Journal of Molecular Liquids, 2020, 298, 111983.	4.9	2
15	Rational Design of a User-Friendly Aptamer/Peptide-Based Device for the Detection of Staphylococcus aureus. Sensors, 2020, 20, 4977.	3.8	7
16	Engineering hemoglobin to enable homogenous PEGylation without modifying protein functionality. Biomaterials Science, 2020, 8, 3896-3906.	5.4	16
17	Inhibition of Nonessential Bacterial Targets: Discovery of a Novel Serine <i>O</i> -Acetyltransferase Inhibitor. ACS Medicinal Chemistry Letters, 2020, 11, 790-797.	2.8	17
18	Model-based evaluation of the microhemodynamic effects of PEGylated HBOC molecules in the rat brain cortex: a laser speckle imaging study. Biomedical Optics Express, 2020, 11, 4150.	2.9	2

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19	Is the protein profile of pig Longissimus dorsi affected by gender and diet?. Journal of Proteomics, 2019, 206, 103437.	2.4	8
20	Combination of SAXS and Protein Painting Discloses the Three-Dimensional Organization of the Bacterial Cysteine Synthase Complex, a Potential Target for Enhancers of Antibiotic Action. International Journal of Molecular Sciences, 2019, 20, 5219.	4.1	9
21	Engineering tyrosine residues into hemoglobin enhances heme reduction, decreases oxidative stress and increases vascular retention of a hemoglobin based blood substitute. Free Radical Biology and Medicine, 2019, 134, 106-118.	2.9	19
22	ADIFAB fluorescence data used for the quantification of free fatty acids in media at different pH. Data in Brief, 2019, 22, 158-163.	1.0	1
23	Editorial: Enzymes Regulating the Homeostasis of Agonists and Antagonists of the N-Methyl D-Aspartate Receptors. Frontiers in Molecular Biosciences, 2019, 6, 37.	3.5	0
24	Covalent Inhibitors of Plasmodium falciparum Glyceraldehyde 3-Phosphate Dehydrogenase with Antimalarial Activity in Vitro. ACS Medicinal Chemistry Letters, 2019, 10, 590-595.	2.8	13
25	More than a Confinement: "Soft―and "Hard―Enzyme Entrapment Modulates Biological Catalyst Function. Catalysts, 2019, 9, 1024.	3.5	12
26	Refining the structureâ^'activity relationships of 2-phenylcyclopropane carboxylic acids as inhibitors of O-acetylserine sulfhydrylase isoforms. Journal of Enzyme Inhibition and Medicinal Chemistry, 2019, 34, 31-43.	5.2	12
27	Proteomics of Meat Products. , 2018, , 297-309.		1
28	Human serine racemase is nitrosylated at multiple sites. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2018, 1866, 813-821.	2.3	11
29	Inhibition of <i>O</i> -acetylserine sulfhydrylase by fluoroalanine derivatives. Journal of Enzyme Inhibition and Medicinal Chemistry, 2018, 33, 1343-1351.	5.2	12
30	Engineering methionine Î ³ -lyase from Citrobacter freundii for anticancer activity. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2018, 1866, 1260-1270.	2.3	11
31	Phospholipid components of the synthetic pulmonary surfactant CHF5633 probed by fluorescence spectroscopy. International Journal of Pharmaceutics, 2018, 553, 290-297.	5.2	2
32	Insight into GFPmut2 pH Dependence by Single Crystal Microspectrophotometry and X-ray Crystallography. Journal of Physical Chemistry B, 2018, 122, 11326-11337.	2.6	3
33	Discovery of novel fragments inhibiting O-acetylserine sulphhydrylase by combining scaffold hopping and ligand–based drug design. Journal of Enzyme Inhibition and Medicinal Chemistry, 2018, 33, 1444-1452.	5.2	17
34	Protein carbonylation detection methods: A comparison. Data in Brief, 2018, 19, 2215-2220.	1.0	20
35	Higher expression of miR-133b is associated with better efficacy of erlotinib as the second or third line in non-small cell lung cancer patients. PLoS ONE, 2018, 13, e0196350.	2.5	15
36	Soluble and Nanoporous Silica Gel-Entrapped <i>C. freundii</i> Methionine <i>γ</i> -Lyase. Journal of Nanoscience and Nanotechnology, 2018, 18, 2210-2219.	0.9	8

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37	Glutamine 89 is a key residue in the allosteric modulation of human serine racemase activity by ATP. Scientific Reports, 2018, 8, 9016.	3.3	12
38	Comparison of the oxidative reactivity of recombinant fetal and adult human hemoglobin: implications for the design of hemoglobin-based oxygen carriers. Bioscience Reports, 2018, 38, .	2.4	22
39	The Energy Landscape of Human Serine Racemase. Frontiers in Molecular Biosciences, 2018, 5, 112.	3.5	28
40	High- and low-affinity PEGylated hemoglobin-based oxygen carriers: Differential oxidative stress in a Guinea pig transfusion model. Free Radical Biology and Medicine, 2018, 124, 299-310.	2.9	13
41	Magnesium and calcium ions differentially affect human serine racemase activity and modulate its quaternary equilibrium toward a tetrameric form. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2017, 1865, 381-387.	2.3	17
42	The Roles of Water in the Protein Matrix: A Largely Untapped Resource for Drug Discovery. Journal of Medicinal Chemistry, 2017, 60, 6781-6827.	6.4	111
43	Modulation of <i>Escherichia coli</i> serine acetyltransferase catalytic activity in the cysteine synthase complex. FEBS Letters, 2017, 591, 1212-1224.	2.8	15
44	Activation of an anti-bacterial toxin by the biosynthetic enzyme CysK: mechanism of binding, interaction specificity and competition with cysteine synthase. Scientific Reports, 2017, 7, 8817.	3.3	7
45	Proteomics of Parma Dry-Cured Ham: Analysis of Salting Exudates. Journal of Agricultural and Food Chemistry, 2017, 65, 6307-6316.	5.2	27
46	Gene cloning, characterization, and cytotoxic activity of methionine γâ€ŀyase from <i>Clostridium novyi</i> . IUBMB Life, 2017, 69, 668-676.	3.4	12
47	Insights on O-acetylserine sulfhydrylase structure, function and biopharmaceutical applications , 2017, , 211-222.		0
48	Insight of Saffron Proteome by Gel-Electrophoresis. Molecules, 2016, 21, 167.	3.8	12
49	Structural insight into the interaction of <i>Oâ€</i> acetylserine sulfhydrylase with competitive, peptidic inhibitors by saturation transfer differenceâ€< scp>NMR. FEBS Letters, 2016, 590, 943-953.	2.8	10
50	Selectivity of 3-bromo-isoxazoline inhibitors between human and Plasmodium falciparum glyceraldehyde-3-phosphate dehydrogenases. Bioorganic and Medicinal Chemistry, 2016, 24, 2654-2659.	3.0	18
51	Cyclopropane-1,2-dicarboxylic acids as new tools for the biophysical investigation of <i>O</i> -acetylserine sulfhydrylases by fluorimetric methods and saturation transfer difference (STD) NMR. Journal of Enzyme Inhibition and Medicinal Chemistry, 2016, 31, 78-87.	5.2	21
52	Human serine racemase is allosterically modulated by NADH and reduced nicotinamide derivatives. Biochemical Journal, 2016, 473, 3505-3516.	3.7	11
53	Engineering tyrosine electron transfer pathways decreases oxidative toxicity in hemoglobin: implications for blood substitute design. Biochemical Journal, 2016, 473, 3371-3383.	3.7	23
54	Rational Design, Synthesis, and Preliminary Structure–Activity Relationships of α-Substituted-2-Phenylcyclopropane Carboxylic Acids as Inhibitors of <i>Salmonella typhimurium</i> <i>O</i> Acetylserine Sulfhydrylase. Journal of Medicinal Chemistry, 2016, 59, 2567-2578.	6.4	28

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55	Cyclopropane derivatives as potential human serine racemase inhibitors: unveiling novel insights into a difficult target. Journal of Enzyme Inhibition and Medicinal Chemistry, 2016, 31, 645-652.	5.2	12
56	EGFR-related miRNAs as potential biomarkers of response to Erlotinib in metastatic NSCLC patients. Annals of Oncology, 2015, 26, vi81.	1.2	0
57	Experiments on Hemoglobin in Single Crystals and Silica Gels Distinguish among Allosteric Models. Biophysical Journal, 2015, 109, 1264-1272.	0.5	33
58	Understanding Water and Its Many Roles in Biological Structure: Ways to Exploit a Resource for Drug Discovery. Methods in Pharmacology and Toxicology, 2015, , 85-110.	0.2	0
59	Regulation of human serine racemase activity and dynamics by halides, ATP and malonate. Amino Acids, 2015, 47, 163-173.	2.7	21
60	Moonlighting O-acetylserine sulfhydrylase: New functions for an old protein. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2015, 1854, 1184-1193.	2.3	35
61	Special Issue on "Cofactor-dependent proteins: Evolution, chemical diversity and bio-applications― Biochimica Et Biophysica Acta - Proteins and Proteomics, 2015, 1854, 1071-1072.	2.3	Ο
62	From protein structure to function via single crystal optical spectroscopy. Frontiers in Molecular Biosciences, 2015, 2, 12.	3.5	14
63	MediaChrom: Discovering a Class of Pyrimidoindolone-Based Polarity-Sensitive Dyes. Journal of Organic Chemistry, 2015, 80, 10939-10954.	3.2	24
64	Expanding the chemical space of human serine racemase inhibitors. Bioorganic and Medicinal Chemistry Letters, 2015, 25, 4297-4303.	2.2	22
65	Immobilization of Proteins in Silica Gel: Biochemical and Biophysical Properties. Current Organic Chemistry, 2015, 19, 1653-1668.	1.6	20
66	Inhibitors of the Sulfur Assimilation Pathway in Bacterial Pathogens as Enhancers of Antibiotic Therapy. Current Medicinal Chemistry, 2014, 22, 187-213.	2.4	42
67	Ormosil gels doped with engineered catechol 1,2 dioxygenases for chlorocatechol bioremediation. Biotechnology and Applied Biochemistry, 2014, 61, 297-303.	3.1	1
68	Experimental basis for a new allosteric model for multisubunit proteins. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 12758-12763.	7.1	46
69	Discovery of Covalent Inhibitors of Glyceraldehyde-3-phosphate Dehydrogenase, A Target for the Treatment of Malaria. Journal of Medicinal Chemistry, 2014, 57, 7465-7471.	6.4	47
70	Molecular insights into dimerization inhibition of c-Maf transcription factor. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2014, 1844, 2108-2115.	2.3	13
71	Identification of a Small Molecule that Increases Hemoglobin Oxygen Affinity and Reduces SS Erythrocyte Sickling. ACS Chemical Biology, 2014, 9, 2318-2325.	3.4	44
72	Targeting Cystalysin, a Virulence Factor of <i>Treponema denticolaâ€</i> Supported Periodontitis. ChemMedChem, 2014, 9, 1501-1511.	3.2	26

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73	<scp>ATP</scp> binding to human serine racemase is cooperative and modulated by glycine. FEBS Journal, 2013, 280, 5853-5863.	4.7	33
74	Unintended consequences? Water molecules at biological and crystallographic protein–protein interfaces. Computational Biology and Chemistry, 2013, 47, 126-141.	2.3	14
75	Fine tuning of the active site modulates specificity in the interaction of O-acetylserine sulfhydrylase isozymes with serine acetyltransferase. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2013, 1834, 169-181.	2.3	35
76	Role of tertiary structures on the Root effect in fish hemoglobins. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2013, 1834, 1885-1893.	2.3	9
77	Oxygen binding and sensing proteins. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2013, 1834, 1683.	2.3	0
78	Muscle and meat: New horizons and applications for proteomics on a farm to fork perspective. Journal of Proteomics, 2013, 88, 58-82.	2.4	53
79	Tertiary and Quaternary Allostery in Tetrameric Hemoglobin from <i>Scapharca inaequivalvis</i> . Biochemistry, 2013, 52, 2108-2117.	2.5	7
80	Chemogenomics of pyridoxal 5â€2-phosphate dependent enzymes. Journal of Enzyme Inhibition and Medicinal Chemistry, 2013, 28, 183-194.	5.2	12
81	Asymmetry of the Active Site Loop Conformation between Subunits of Glutamate-1-semialdehyde Aminomutase in Solution. BioMed Research International, 2013, 2013, 1-10.	1.9	15
82	Development of a novel, hemolysis-resistant reagent for assessment of α-amylase in biological fluids. Clinical Chemistry and Laboratory Medicine, 2013, 51, 1409-15.	2.3	2
83	Serine racemase: a key player in neuron activity and in neuropathologies. Frontiers in Bioscience - Landmark, 2013, 18, 1112.	3.0	34
84	CO Rebinding Kinetics and Molecular Dynamics Simulations Highlight Dynamic Regulation of Internal Cavities in Human Cytoglobin. PLoS ONE, 2013, 8, e49770.	2.5	28
85	Isozyme-Specific Ligands for O-acetylserine sulfhydrylase, a Novel Antibiotic Target. PLoS ONE, 2013, 8, e77558.	2.5	43
86	The role of salt in dry cured ham processing characterized by LC-MS/MS-based proteomics. , 2013, , 274-277.		0
87	Biochemistry of Hemoglobin. , 2013, , 55-73.		1
88	International Consortium for Development of Hemoglobin-Based Oxygen Carriers, Oxygen Therapeutics and Multifunctional Resuscitation Fluids–A White Paper. , 2013, , 737-746.		2
89	"Muscle to meat―molecular events and technological transformations: The proteomics insight. Journal of Proteomics, 2012, 75, 4275-4289.	2.4	115
90	Design and synthesis of trans-2-substituted-cyclopropane-1-carboxylic acids as the first non-natural small molecule inhibitors of O-acetylserine sulfhydrylase. MedChemComm, 2012, 3, 1111.	3.4	36

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91	The proteomic insight of the Italian dry cured ham manufacturing. , 2012, , 138-140.		0
92	From meat to food: the proteomics assessment. , 2012, , 31-34.		0
93	Histidine E7 Dynamics Modulates Ligand Exchange between Distal Pocket and Solvent in AHb1 from <i>Arabidopsis thaliana</i> . Journal of Physical Chemistry B, 2011, 115, 4138-4146.	2.6	20
94	Proteomic analysis of pork meat in the production of cooked ham. Molecular BioSystems, 2011, 7, 2252.	2.9	32
95	Modulation of expression and polymerization of hemoglobin Polytaur, a potential blood substitute. Archives of Biochemistry and Biophysics, 2011, 505, 42-47.	3.0	14
96	Exploring O-acetylserine sulfhydrylase-B isoenzyme from Salmonella typhimurium by fluorescence spectroscopy. Archives of Biochemistry and Biophysics, 2011, 505, 178-185.	3.0	8
97	The multifaceted pyridoxal 5â€2-phosphate-dependent O-acetylserine sulfhydrylase. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2011, 1814, 1497-1510.	2.3	39
98	Drug Discovery Targeting Amino Acid Racemases. Chemical Reviews, 2011, 111, 6919-6946.	47.7	97
99	Human kynurenine aminotransferase Il â€ʿ reactivity with substrates and inhibitors. FEBS Journal, 2011, 278, 1882-1900.	4.7	25
100	Low affinity PEGylated hemoglobin from Trematomus bernacchii, a model for hemoglobin-based blood substitutes. BMC Biochemistry, 2011, 12, 66.	4.4	9
101	Oxygen binding to <i>Arabidopsis thaliana</i> AHb2 nonsymbiotic hemoglobin: evidence for a role in oxygen transport. IUBMB Life, 2011, 63, 355-362.	3.4	19
102	Oxygen and nitric oxide rebinding kinetics in nonsymbiotic hemoglobin AHb1 from <i>Arabidopsis thaliana</i> . IUBMB Life, 2011, 63, 1094-1100.	3.4	16
103	Electrophoretic analysis of PEGylated hemoglobin-based blood substitutes. Analytical Biochemistry, 2011, 408, 118-123.	2.4	9
104	Exploring methionine Î ³ -lyase structure-function relationship via microspectrophotometry and X-ray crystallography. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2011, 1814, 834-842.	2.3	18
105	Ligand migration and hexacoordination in type 1 non-symbiotic rice hemoglobin. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2011, 1814, 1042-1053.	2.3	15
106	Exploring and exploiting allostery: Models, evolution, and drug targeting. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2011, 1814, 922-933.	2.3	60
107	Protein crystal microspectrophotometry. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2011, 1814, 734-741.	2.3	11
108	X-ray crystallography marries spectroscopy to unveil structure and function of biological macromolecules. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2011, 1814, 731-733.	2.3	12

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109	Bound Water at Protein-Protein Interfaces: Partners, Roles and Hydrophobic Bubbles as a Conserved Motif. PLoS ONE, 2011, 6, e24712.	2.5	57
110	Polymerized and polyethylene glycol-conjugated hemoglobins: A globin-based calibration curve for dynamic light scattering analysis. Analytical Biochemistry, 2010, 401, 266-270.	2.4	5
111	A Two-step Process Controls the Formation of the Bienzyme Cysteine Synthase Complex. Journal of Biological Chemistry, 2010, 285, 12813-12822.	3.4	35
112	Pyridoxal 5′-Phosphate-Dependent Enzymes: Catalysis, Conformation, and Genomics. , 2010, , 273-350.		12
113	Identification of the Structural Determinants for the Stability of Substrate and Aminoacrylate External Schiff Bases in <i>O</i> -Acetylserine Sulfhydrylase-A. Biochemistry, 2010, 49, 6093-6103.	2.5	25
114	Design of <i>O</i> -Acetylserine Sulfhydrylase Inhibitors by Mimicking Nature. Journal of Medicinal Chemistry, 2010, 53, 345-356.	6.4	75
115	Haemoglobin-based oxygen carriers: research and reality towards an alternative to blood transfusions. Blood Transfusion, 2010, 8 Suppl 3, s59-68.	0.4	24
116	Ligand migration through the internal hydrophobic cavities in human neuroglobin. Proceedings of the United States of America, 2009, 106, 18984-18989.	7.1	47
117	Tryptophan synthase: a mine for enzymologists. Cellular and Molecular Life Sciences, 2009, 66, 2391-2403.	5.4	83
118	Hemoglobin, an "evergreen―red protein. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2009, 1794, 1317-1324.	2.3	15
119	Correlation of protein functional properties in the crystal and in solution: The case study of T-state hemoglobin. Protein Science, 2009, 11, 1845-1849.	7.6	10
120	PEGylation Promotes Hemoglobin Tetramer Dissociation. Bioconjugate Chemistry, 2009, 20, 1356-1366.	3.6	45
121	Identification of Xenoestrogens in Food Additives by an Integrated in Silico and in Vitro Approach. Chemical Research in Toxicology, 2009, 22, 52-63.	3.3	74
122	Cofactor Chemogenomics. Methods in Molecular Biology, 2009, 575, 93-122.	0.9	6
123	Structural Plasticity and Functional Implications of Internal Cavities in Distal Mutants of Type 1 Non-Symbiotic Hemoglobin AHb1 fromArabidopsis thaliana. Journal of Physical Chemistry B, 2009, 113, 16028-16038.	2.6	20
124	Chemogenomic Strategies to Expand the Bioactive Chemical Space. Current Medicinal Chemistry, 2009, 16, 4374-4381.	2.4	23
125	Energyâ€based prediction of amino acidâ€nucleotide base recognition. Journal of Computational Chemistry, 2008, 29, 1955-1969.	3.3	44
126	Towards a novel haemoglobin-based oxygen carrier: Euro-PEG-Hb, physico-chemical properties, vasoactivity and renal filtration. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2008, 1784, 1402-1409.	2.3	42

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127	Hemoglobin-based oxygen carriers as blood substitutes. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2008, 1784, 1363-1364.	2.3	12
128	High and low oxygen affinity conformations of T state hemoglobin. Protein Science, 2008, 10, 2401-2407.	7.6	74
129	Ligand-Induced Tertiary Relaxations During the T-to-R Quaternary Transition in Hemoglobin. Journal of Physical Chemistry B, 2008, 112, 12790-12794.	2.6	10
130	Characterization of Ligand Migration Mechanisms inside Hemoglobins from the Analysis of Geminate Rebinding Kinetics. Methods in Enzymology, 2008, 437, 329-345.	1.0	4
131	Robust Classification of "Relevant―Water Molecules in Putative Protein Binding Sites. Journal of Medicinal Chemistry, 2008, 51, 1063-1067.	6.4	93
132	Trapping of the Thioacylglyceraldehyde-3-phosphate Dehydrogenase Intermediate from Bacillus stearothermophilus. Journal of Biological Chemistry, 2008, 283, 21693-21702.	3.4	35
133	Oxygen Binding to Heme Proteins in Solution, Encapsulated in Silica Gels, and in the Crystalline State. Methods in Enzymology, 2008, 437, 311-328.	1.0	29
134	Trapping Hemoglobin in Rigid Matrices: Fine Tuning of Oxygen Binding Properties by Modulation of Encapsulation Protocols. Artificial Cells, Blood Substitutes, and Biotechnology, 2007, 35, 69-79.	0.9	10
135	Pyridoxal 5-Phosphate Enzymes as Targets for Therapeutic Agents. Current Medicinal Chemistry, 2007, 14, 1291-1324.	2.4	177
136	Hemocyanin from E. californicum encapsulated in silica gels: Oxygen binding and conformational states. Gene, 2007, 398, 202-207.	2.2	7
137	Different roles of protein dynamics and ligand migration in non-symbiotic hemoglobins AHb1 and AHb2 from Arabidopsis thaliana. Gene, 2007, 398, 224-233.	2.2	32
138	The Reactivity with CO of AHb1 and AHb2 from Arabidopsis thaliana is Controlled by the Distal HisE7 and Internal Hydrophobic Cavities. Journal of the American Chemical Society, 2007, 129, 2880-2889.	13.7	54
139	Structure, Mechanism, and Conformational Dynamics of O-Acetylserine Sulfhydrylase from Salmonella typhimurium:  Comparison of A and B Isozymes. Biochemistry, 2007, 46, 8315-8330.	2.5	58
140	Ligand Migration in Nonsymbiotic Hemoglobin AHb1 from Arabidopsis thaliana. Journal of Physical Chemistry B, 2007, 111, 12582-12590.	2.6	27
141	Control of Ionizable Residues in the Catalytic Mechanism of Tryptophan Synthase from <i>Salmonella typhimurium</i> . Biochemistry, 2007, 46, 13223-13234.	2.5	12
142	Evidence of Discrete Substates and Unfolding Pathways in Green Fluorescent Protein. Biophysical Journal, 2007, 92, 1724-1731.	0.5	16
143	Complexity in Modeling and Understanding Protonation States: Computational Titration of HIVâ€I â€Protease–Inhibitor Complexes. Chemistry and Biodiversity, 2007, 4, 2564-2577.	2.1	10
144	The consequences of scoring docked ligand conformations using free energy correlations. European Journal of Medicinal Chemistry, 2007, 42, 921-933.	5.5	58

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145	Energetics of the protein-DNA-water interaction. BMC Structural Biology, 2007, 7, 4.	2.3	57
146	Evolution of allosteric models for hemoglobin. IUBMB Life, 2007, 59, 586-599.	3.4	103
147	Environment effects on the oscillatory unfolding kinetics of GFP. European Biophysics Journal, 2007, 36, 795-803.	2.2	5
148	Monitoring the Tâ€R transition of human hemoglobin encapsulated in silica gels. FASEB Journal, 2007, 21, A637.	0.5	0
149	Time-resolved methods in Biophysics. 2. Monitoring haem proteins at work with nanosecond laser flash photolysis. Photochemical and Photobiological Sciences, 2006, 5, 1109.	2.9	53
150	Mapping the Energetics of Water–Protein and Water–Ligand Interactions with the "Natural―HINT Forcefield: Predictive Tools for Characterizing the Roles of Water in Biomolecules. Journal of Molecular Biology, 2006, 358, 289-309.	4.2	85
151	Water: How to evaluate its contribution in protein-ligand interactions. International Journal of Quantum Chemistry, 2006, 106, 647-651.	2.0	12
152	Exploring the pyridoxal $5\hat{a}\in^2$ -phosphate-dependent enzymes. Chemical Record, 2006, 6, 275-287.	5.8	52
153	Allosteric communication between alpha and beta subunits of tryptophan synthase: Modelling the open-closed transition of the alpha subunit. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2006, 1764, 1102-1109.	2.3	22
154	Tools for building a comprehensive modeling system for virtual screening under real biological conditions: The Computational Titration algorithm. Journal of Molecular Graphics and Modelling, 2006, 24, 434-439.	2.4	18
155	Circular dichroism spectroscopy of tertiary and quaternary conformations of human hemoglobin entrapped in wet silica gels. Protein Science, 2006, 15, 1961-1967.	7.6	27
156	Is cooperative oxygen binding by hemoglobin really understood?. Rendiconti Lincei, 2006, 17, 147-162.	2.2	7
157	Sulfur Mobilization in Cyanobacteria. Journal of Biological Chemistry, 2006, 281, 38769-38780.	3.4	16
158	Interaction of serine acetyltransferase withO-acetylserine sulfhydrylase active site: Evidence from fluorescence spectroscopy. Protein Science, 2005, 14, 2115-2124.	7.6	83
159	Identification of the Geometric Requirements for Allosteric Communication between the α- and β-Subunits of Tryptophan Synthase. Journal of Biological Chemistry, 2005, 280, 13450-13456.	3.4	24
160	Evidence for Two Geminate Rebinding States Following Laser Photolysis of R State Hemoglobin Encapsulated in Wet Silica Gels. Journal of Physical Chemistry B, 2005, 109, 11411-11413.	2.6	29
161	Kinetics of Acid-Induced Spectral Changes in the GFPmut2 Chromophore. Journal of the American Chemical Society, 2005, 127, 626-635.	13.7	57
162	Determination of Microscopic Rate Constants for CO Binding and Migration in Myoglobin Encapsulated in Silica Gels. Journal of Physical Chemistry B, 2005, 109, 19523-19528.	2.6	29

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163	Geminate Rebinding in R-State Hemoglobin:Â Kinetic and Computational Evidence for Multiple Hydrophobic Pockets. Journal of the American Chemical Society, 2005, 127, 17427-17432.	13.7	29
164	Confinement and crowding effects on tryptophan synthase $\hat{I}\pm2\hat{I}^2$ 2complex. FEBS Letters, 2005, 579, 2197-2202.	2.8	23
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