

Stephen J Benkovic

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355
papers

18,565
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72
h-index

116
g-index

368
ext. papers

20,252
ext. citations

10.1
avg, IF

6.81
L-index

#	Paper	IF	Citations
355	A perspective on enzyme catalysis. <i>Science</i> , 2003 , 301, 1196-202	33.3	977
354	Construction and evaluation of the kinetic scheme associated with dihydrofolate reductase from <i>Escherichia coli</i> . <i>Biochemistry</i> , 1987 , 26, 4085-92	3.2	474
353	Reversible compartmentalization of de novo purine biosynthetic complexes in living cells. <i>Science</i> , 2008 , 320, 103-6	33.3	351
352	A dynamic knockout reveals that conformational fluctuations influence the chemical step of enzyme catalysis. <i>Science</i> , 2011 , 332, 234-8	33.3	350
351	Replisome-mediated DNA replication. <i>Annual Review of Biochemistry</i> , 2001 , 70, 181-208	29.1	273
350	Surface sites for engineering allosteric control in proteins. <i>Science</i> , 2008 , 322, 438-42	33.3	269
349	Metallo-beta-lactamase: structure and mechanism. <i>Current Opinion in Chemical Biology</i> , 1999 , 3, 614-22	9.7	262
348	Chemical basis for enzyme catalysis. <i>Biochemistry</i> , 2000 , 39, 6267-74	3.2	252
347	Controlling cell-cell interactions using surface acoustic waves. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015 , 112, 43-8	11.5	247
346	Kinetic characterization of the polymerase and exonuclease activities of the gene 43 protein of bacteriophage T4. <i>Biochemistry</i> , 1992 , 31, 10984-94	3.2	233
345	Free-energy landscape of enzyme catalysis. <i>Biochemistry</i> , 2008 , 47, 3317-21	3.2	218
344	Design and evolution of new catalytic activity with an existing protein scaffold. <i>Science</i> , 2006 , 311, 535-8	33.3	213
343	Interaction of dihydrofolate reductase with methotrexate: ensemble and single-molecule kinetics. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002 , 99, 13481-6	11.5	212
342	A New View into the Regulation of Purine Metabolism: The Purinosome. <i>Trends in Biochemical Sciences</i> , 2017 , 42, 141-154	10.3	202
341	Flexibility, diversity, and cooperativity: pillars of enzyme catalysis. <i>Biochemistry</i> , 2011 , 50, 10422-30	3.2	190
340	Boron-containing inhibitors of synthetases. <i>Chemical Society Reviews</i> , 2011 , 40, 4279-85	58.5	190
339	Tunneling and coupled motion in the <i>Escherichia coli</i> dihydrofolate reductase catalysis. <i>Journal of the American Chemical Society</i> , 2004 , 126, 4778-9	16.4	182

338	On the mechanism of the metallo-beta-lactamase from <i>Bacteroides fragilis</i> . <i>Biochemistry</i> , 1999 , 38, 10013-23	3.2	181
337	Evidence for a functional role of the dynamics of glycine-121 of <i>Escherichia coli</i> dihydrofolate reductase obtained from kinetic analysis of a site-directed mutant. <i>Biochemistry</i> , 1997 , 36, 15792-800	3.2	178
336	A combinatorial approach to hybrid enzymes independent of DNA homology. <i>Nature Biotechnology</i> , 1999 , 17, 1205-9	44.5	176
335	Coordinated effects of distal mutations on environmentally coupled tunneling in dihydrofolate reductase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006 , 103, 15753-8	11.5	160
334	Coupling interactions of distal residues enhance dihydrofolate reductase catalysis: mutational effects on hydride transfer rates. <i>Biochemistry</i> , 2002 , 41, 12618-28	3.2	156
333	Dynamics of the dihydrofolate reductase-folate complex: catalytic sites and regions known to undergo conformational change exhibit diverse dynamical features. <i>Biochemistry</i> , 1995 , 34, 11037-48	3.2	151
332	Coupled motions in enzyme catalysis. <i>Current Opinion in Chemical Biology</i> , 2010 , 14, 644-51	9.7	146
331	Dynamics of a flexible loop in dihydrofolate reductase from <i>Escherichia coli</i> and its implication for catalysis. <i>Biochemistry</i> , 1994 , 33, 439-42	3.2	142
330	A perspective on biological catalysis. <i>Nature Structural and Molecular Biology</i> , 1996 , 3, 821-33	17.6	138
329	Solvation, reorganization energy, and biological catalysis. <i>Journal of Biological Chemistry</i> , 1998 , 273, 26254-60	54.60	134
328	Impact of distal mutations on the network of coupled motions correlated to hydride transfer in dihydrofolate reductase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005 , 102, 6807-12	11.5	131
327	Split-intein mediated circular ligation used in the synthesis of cyclic peptide libraries in <i>E. coli</i> . <i>Nature Protocols</i> , 2007 , 2, 1126-33	18.8	120
326	Real-time observation of bacteriophage T4 gp41 helicase reveals an unwinding mechanism. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007 , 104, 19790-5	11.5	119
325	Inhibition of HIV budding by a genetically selected cyclic peptide targeting the Gag-TSG101 interaction. <i>ACS Chemical Biology</i> , 2008 , 3, 757-64	4.9	115
324	Transition-state stabilization as a measure of the efficiency of antibody catalysis. <i>Nature</i> , 1995 , 375, 388-91	50.4	113
323	The dynamic processivity of the T4 DNA polymerase during replication. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004 , 101, 8289-94	11.5	110
322	Purification and characterization of human immunodeficiency virus type 1 reverse transcriptase. <i>Methods in Enzymology</i> , 1995 , 262, 130-44	1.7	109
321	Spatial colocalization and functional link of purinosomes with mitochondria. <i>Science</i> , 2016 , 351, 733-7	33.3	106

320	Perspectives on electrostatics and conformational motions in enzyme catalysis. <i>Accounts of Chemical Research</i> , 2015 , 48, 482-9	24.3	106
319	Regulation of polymerase exchange between Poleta and Poldelta by monoubiquitination of PCNA and the movement of DNA polymerase holoenzyme. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008 , 105, 5361-6	11.5	104
318	Mechanism of oxygen activation by pteridine-dependent monooxygenases. <i>Accounts of Chemical Research</i> , 1988 , 21, 101-107	24.3	103
317	Evolution of cyclic peptide protease inhibitors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 11052-6	11.5	100
316	Electron spin-echo studies of the copper binding site in phenylalanine hydroxylase from <i>Chromobacterium violaceum</i> . <i>Journal of the American Chemical Society</i> , 1988 , 110, 1069-1074	16.4	100
315	A systematic method for identifying small-molecule modulators of protein-protein interactions. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004 , 101, 15591-6	11.5	99
314	NMR characterization of the metallo-beta-lactamase from <i>Bacteroides fragilis</i> and its interaction with a tight-binding inhibitor: role of an active-site loop. <i>Biochemistry</i> , 1999 , 38, 14507-14	3.2	99
313	Catalytic antibodies. <i>Annual Review of Biochemistry</i> , 1992 , 61, 29-54	29.1	97
312	Functional role of a mobile loop of <i>Escherichia coli</i> dihydrofolate reductase in transition-state stabilization. <i>Biochemistry</i> , 1992 , 31, 7826-33	3.2	97
311	Direct Observation of an Enzyme-Bound Intermediate in the Catalytic Cycle of the Metallo-β-Lactamase from <i>Bacteroides fragilis</i> . <i>Journal of the American Chemical Society</i> , 1998 , 120, 10788-10789 ^{16.4}	16.4	95
310	Effects of the donor-acceptor distance and dynamics on hydride tunneling in the dihydrofolate reductase catalyzed reaction. <i>Journal of the American Chemical Society</i> , 2012 , 134, 1738-45	16.4	90
309	Protein-DNA cross-linking demonstrates stepwise ATP-dependent assembly of T4 DNA polymerase and its accessory proteins on the primer-template. <i>Cell</i> , 1991 , 65, 249-58	56.2	90
308	Metabolomics and mass spectrometry imaging reveal channeled de novo purine synthesis in cells. <i>Science</i> , 2020 , 368, 283-290	33.3	90
307	Reaction mechanisms displayed by catalytic antibodies. <i>Accounts of Chemical Research</i> , 1993 , 26, 396-404	24.3	89
306	Bait and switch strategy for obtaining catalytic antibodies with acyl-transfer capabilities. <i>Journal of the American Chemical Society</i> , 1990 , 112, 1274-1275	16.4	89
305	Ring Structure and Aromatic Substituent Effects on the pK _a of the Benzoxaborole Pharmacophore. <i>ACS Medicinal Chemistry Letters</i> , 2012 , 3, 48-52	4.3	88
304	Substrate-driven chemotactic assembly in an enzyme cascade. <i>Nature Chemistry</i> , 2018 , 10, 311-317	17.6	87
303	Identification of borinic esters as inhibitors of bacterial cell growth and bacterial methyltransferases, CcrM and MenH. <i>Journal of Medicinal Chemistry</i> , 2005 , 48, 7468-76	8.3	86

302	The unique chemistry of benzoxaboroles: current and emerging applications in biotechnology and therapeutic treatments. <i>Bioorganic and Medicinal Chemistry</i> , 2014 , 22, 4462-73	3.4	84
301	DNA polymerase as a molecular motor and pump. <i>ACS Nano</i> , 2014 , 8, 2410-8	16.7	84
300	Hybrid enzymes: manipulating enzyme design. <i>Trends in Biotechnology</i> , 1998 , 16, 258-64	15.1	82
299	Stretching exercises--flexibility in dihydrofolate reductase catalysis. <i>Chemistry and Biology</i> , 1998 , 5, R105-13		81
298	Purification, characterization, and kinetic studies of a soluble <i>Bacteroides fragilis</i> metallo-beta-lactamase that provides multiple antibiotic resistance. <i>Journal of Biological Chemistry</i> , 1998 , 273, 22402-8	5.4	81
297	Biochemistry. Enzyme motions inside and out. <i>Science</i> , 2006 , 312, 208-9	33.3	80
296	The control mechanism for lagging strand polymerase recycling during bacteriophage T4 DNA replication. <i>Molecular Cell</i> , 2006 , 21, 153-64	17.6	80
295	Genetically selected cyclic-peptide inhibitors of AICAR transformylase homodimerization. <i>Angewandte Chemie - International Edition</i> , 2005 , 44, 2760-2763	16.4	78
294	Coupling DNA unwinding activity with primer synthesis in the bacteriophage T4 primosome. <i>Nature Chemical Biology</i> , 2009 , 5, 904-12	11.7	77
293	Structural requirements for the biosynthesis of backbone cyclic peptide libraries. <i>Chemistry and Biology</i> , 2001 , 8, 801-15		77
292	Evaluation of the importance of hydrophobic interactions in drug binding to dihydrofolate reductase. <i>Journal of Medicinal Chemistry</i> , 1988 , 31, 129-37	8.3	76
291	Studies on Sulfate Esters. I. Nucleophilic Reactions of Amines with p-Nitrophenyl Sulfate. <i>Journal of the American Chemical Society</i> , 1966 , 88, 5504-5511	16.4	76
290	Towards structure-based drug design: crystal structure of a multisubstrate adduct complex of glycinamide ribonucleotide transformylase at 1.96 Å resolution. <i>Journal of Molecular Biology</i> , 1995 , 249, 153-75	6.5	75
289	Tunable, pulsatile chemical gradient generation via acoustically driven oscillating bubbles. <i>Lab on a Chip</i> , 2013 , 13, 328-31	7.2	74
288	Replication clamps and clamp loaders. <i>Cold Spring Harbor Perspectives in Biology</i> , 2013 , 5, a010165	10.2	74
287	A Comparison of the Bimolecular and Intramolecular Nucleophilic Catalysis of the Hydrolysis of Substituted Phenyl Acylates by the Dimethylamino Group. <i>Journal of the American Chemical Society</i> , 1963 , 85, 1-8	16.4	73
286	GPCRs regulate the assembly of a multienzyme complex for purine biosynthesis. <i>Nature Chemical Biology</i> , 2011 , 7, 909-15	11.7	72
285	Accessory proteins function as matchmakers in the assembly of the T4 DNA polymerase holoenzyme. <i>Current Biology</i> , 1995 , 5, 149-57	6.3	72

284	Functional significance of evolving protein sequence in dihydrofolate reductase from bacteria to humans. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 10159-64	11.5	71
283	The Compensation in Δ [UNK] and δ [UNK] Accompanying the Conversion of Lower Order Nucleophilic Displacement Reactions to Higher Order Catalytic Processes. The Temperature Dependence of the Hydrazinolysis and Imidazole-Catalyzed Hydrolysis of Substituted Phenyl Acetates. <i>Journal of the American Chemical Society</i> , 1964 , 86, 418-426	16.4	71
282	Peptide bond formation via catalytic antibodies: Synthesis of a novel phosphonate diester hapten. <i>Tetrahedron Letters</i> , 1994 , 35, 6853-6856	2	70
281	Probing the electrostatics of active site microenvironments along the catalytic cycle for Escherichia coli dihydrofolate reductase. <i>Journal of the American Chemical Society</i> , 2014 , 136, 10349-60	16.4	69
280	Catalytic antibody model and mutagenesis implicate arginine in transition-state stabilization. <i>Journal of Molecular Biology</i> , 1994 , 235, 1098-116	6.5	69
279	A distal mutation perturbs dynamic amino acid networks in dihydrofolate reductase. <i>Biochemistry</i> , 2013 , 52, 4605-19	3.2	68
278	Mapping Protein-Protein Interactions in the Bacteriophage T4 DNA Polymerase Holoenzyme Using a Novel Trifunctional Photo-cross-linking and Affinity Reagent. <i>Journal of the American Chemical Society</i> , 2000 , 122, 6126-6127	16.4	68
277	Direct observation of stalled fork restart via fork regression in the T4 replication system. <i>Science</i> , 2012 , 338, 1217-20	33.3	67
276	Crystal structure of a bifunctional transformylase and cyclohydrolase enzyme in purine biosynthesis. <i>Nature Structural Biology</i> , 2001 , 8, 402-6		67
275	Quantitative analysis of purine nucleotides indicates that purinosomes increase de novo purine biosynthesis. <i>Journal of Biological Chemistry</i> , 2015 , 290, 6705-13	5.4	66
274	Elucidation of the mechanism of the reaction between phenylboronic acid and a model diol, Alizarin Red S. <i>Journal of Organic Chemistry</i> , 2012 , 77, 2098-106	4.2	66
273	Sliding clamp of the bacteriophage T4 polymerase has open and closed subunit interfaces in solution. <i>Biochemistry</i> , 1999 , 38, 7696-709	3.2	65
272	Regulation of Rad6/Rad18 Activity During DNA Damage Tolerance. <i>Annual Review of Biophysics</i> , 2015 , 44, 207-28	21.1	64
271	Microtubule-assisted mechanism for functional metabolic macromolecular complex formation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010 , 107, 12872-6	11.5	64
270	Subcloning, characterization, and affinity labeling of Escherichia coli glycinamide ribonucleotide transformylase. <i>Biochemistry</i> , 1990 , 29, 1436-43	3.2	64
269	On the cofactor specificity of glycinamide ribonucleotide and 5-aminoimidazole-4-carboxamide ribonucleotide transformylase from chicken liver. <i>Biochemistry</i> , 1981 , 20, 1241-5	3.2	64
268	Single-molecule and transient kinetics investigation of the interaction of dihydrofolate reductase with NADPH and dihydrofolate. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004 , 101, 2764-9	11.5	63
267	Deletion of a highly motional residue affects formation of the Michaelis complex for Escherichia coli dihydrofolate reductase. <i>Biochemistry</i> , 1998 , 37, 6327-35	3.2	63

266	Synthesis and application of derivatizable oligonucleotides. <i>Nucleic Acids Research</i> , 1987 , 15, 6455-67	20.1	63
265	FamClash: a method for ranking the activity of engineered enzymes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2004 , 101, 4142-7	11.5	61
264	Crystal structure of glycinamide ribonucleotide transformylase from <i>Escherichia coli</i> at 3.0 Å resolution. A target enzyme for chemotherapy. <i>Journal of Molecular Biology</i> , 1992 , 227, 283-92	6.5	61
263	A multifunctional protein possessing glycinamide ribonucleotide synthetase, glycinamide ribonucleotide transformylase, and aminoimidazole ribonucleotide synthetase activities in de novo purine biosynthesis. <i>Biochemistry</i> , 1985 , 24, 7059-62	3.2	61
262	Phenylalanine Hydroxylase Stimulator Protein Is a 4E ₂ Carbinolamine Dehydratase. <i>Journal of Biological Chemistry</i> , 1983 , 258, 10960-10962	5.4	61
261	Hsp70/Hsp90 chaperone machinery is involved in the assembly of the purinosome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 2528-33	11.5	60
260	Incremental truncation as a strategy in the engineering of novel biocatalysts. <i>Bioorganic and Medicinal Chemistry</i> , 1999 , 7, 2139-44	3.4	60
259	How a holoenzyme for DNA replication is formed. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 99-104	11.5	59
258	Preorganization and protein dynamics in enzyme catalysis. <i>Chemical Record</i> , 2002 , 2, 24-36	6.6	59
257	A clamp-like biohybrid catalyst for DNA oxidation. <i>Nature Chemistry</i> , 2013 , 5, 945-51	17.6	58
256	Strength of an interloop hydrogen bond determines the kinetic pathway in catalysis by <i>Escherichia coli</i> dihydrofolate reductase. <i>Biochemistry</i> , 1998 , 37, 6336-42	3.2	58
255	Structure-reactivity correlation for the hydrolysis of phosphoramidate monoanions. <i>Journal of the American Chemical Society</i> , 1971 , 93, 4009-4016	16.4	58
254	Identification of a novel boron-containing antibacterial agent (AN0128) with anti-inflammatory activity, for the potential treatment of cutaneous diseases. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2006 , 16, 5963-7	2.9	57
253	Role of adenosine 5′-triphosphate hydrolysis in the assembly of the bacteriophage T4 DNA replication holoenzyme complex. <i>Biochemistry</i> , 1996 , 35, 9253-65	3.2	57
252	Structural basis for amide hydrolysis catalyzed by the 43C9 antibody. <i>Journal of Molecular Biology</i> , 1999 , 291, 329-45	6.5	56
251	Probing cell-cell communication with microfluidic devices. <i>Lab on A Chip</i> , 2013 , 13, 3152-62	7.2	55
250	Purinosome formation as a function of the cell cycle. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015 , 112, 1368-73	11.5	55
249	The structure of a ring-opened proliferating cell nuclear antigen-replication factor C complex revealed by fluorescence energy transfer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006 , 103, 2546-51	11.5	54

248	Using an AraC-based three-hybrid system to detect biocatalysts in vivo. <i>Nature Biotechnology</i> , 2000 , 18, 544-7	44.5	54
247	Role of a solvent-exposed tryptophan in the recognition and binding of antibiotic substrates for a metallo-beta-lactamase. <i>Protein Science</i> , 2003 , 12, 1368-75	6.3	53
246	Bacteriophage T4 Dda helicase translocates in a unidirectional fashion on single-stranded DNA. <i>Journal of Biological Chemistry</i> , 1995 , 270, 22236-42	5.4	53
245	Cloning and characterization of a new purine biosynthetic enzyme: a non-folate glycinamide ribonucleotide transformylase from <i>E. coli</i> . <i>Biochemistry</i> , 1994 , 33, 2531-7	3.2	53
244	Mechanism of strand displacement synthesis by DNA replicative polymerases. <i>Nucleic Acids Research</i> , 2012 , 40, 6174-86	20.1	52
243	Collaborative coupling between polymerase and helicase for leading-strand synthesis. <i>Nucleic Acids Research</i> , 2012 , 40, 6187-98	20.1	52
242	Discovery of antibacterial cyclic peptides that inhibit the ClpXP protease. <i>Protein Science</i> , 2007 , 16, 1535-42	6.3	52
241	Tracking sliding clamp opening and closing during bacteriophage T4 DNA polymerase holoenzyme assembly. <i>Biochemistry</i> , 2000 , 39, 3076-90	3.2	52
240	Detection of dihydrofolate reductase conformational change by FRET using two fluorescent amino acids. <i>Journal of the American Chemical Society</i> , 2013 , 135, 12924-7	16.4	51
239	Interloop contacts modulate ligand cycling during catalysis by <i>Escherichia coli</i> dihydrofolate reductase. <i>Biochemistry</i> , 2001 , 40, 867-75	3.2	51
238	Stoichiometry and DNA unwinding by the bacteriophage T4 41:59 helicase. <i>Journal of Biological Chemistry</i> , 1996 , 271, 14074-81	5.4	51
237	The interconversion of the 5,6,7,8-tetrahydro-, 6,7,8-dihydro-, and radical forms of 6,6,7,7-tetramethyldihydropterin. A model for the biopterin center of aromatic amino acid mixed function oxidases. <i>Journal of the American Chemical Society</i> , 1984 , 106, 7916-7924	16.4	50
236	Dissecting the order of bacteriophage T4 DNA polymerase holoenzyme assembly. <i>Biochemistry</i> , 1998 , 37, 7749-56	3.2	49
235	Mechanistic aspects of DNA polymerases: <i>Escherichia coli</i> DNA polymerase I (Klenow fragment) as a paradigm. <i>Chemical Reviews</i> , 1990 , 90, 1291-1307	68.1	49
234	Dynamic regulation of a metabolic multi-enzyme complex by protein kinase CK2. <i>Journal of Biological Chemistry</i> , 2010 , 285, 11093-9	5.4	48
233	Evolution of highly active enzymes by homology-independent recombination. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005 , 102, 10082-7	11.5	48
232	Dual role of the 44/62 protein as a matchmaker protein and DNA polymerase chaperone during assembly of the bacteriophage T4 holoenzyme complex. <i>Biochemistry</i> , 1996 , 35, 1084-92	3.2	48
231	Phenylalanine hydroxylase: structural determination of the tetrahydropterin intermediates by carbon-13 NMR spectroscopy. <i>Journal of the American Chemical Society</i> , 1982 , 104, 6869-6871	16.4	48

230	Enhanced crossover SCRATCHY: construction and high-throughput screening of a combinatorial library containing multiple non-homologous crossovers. <i>Nucleic Acids Research</i> , 2003 , 31, e126	20.1	47
229	A zinc ribbon protein in DNA replication: primer synthesis and macromolecular interactions by the bacteriophage T4 primase. <i>Biochemistry</i> , 2001 , 40, 15074-85	3.2	47
228	Evolution of protein function by domain swapping. <i>Advances in Protein Chemistry</i> , 2000 , 55, 29-77		46
227	Truncating alpha-helix E of p66 human immunodeficiency virus reverse transcriptase modulates RNase H function and impairs DNA strand transfer. <i>Journal of Biological Chemistry</i> , 1995 , 270, 7068-76	5.4	46
226	Nonadditivity of mutational effects at the folate binding site of Escherichia coli dihydrofolate reductase. <i>Biochemistry</i> , 1994 , 33, 11576-85	3.2	46
225	A multisubstrate adduct inhibitor of a purine biosynthetic enzyme with a picomolar dissociation constant. <i>Journal of Medicinal Chemistry</i> , 1989 , 32, 937-40	8.3	46
224	Targeting tumour proliferation with a small-molecule inhibitor of AICAR transformylase homodimerization. <i>ChemBioChem</i> , 2012 , 13, 1628-34	3.8	45
223	Mechanism of action of fructose 1,6-bisphosphatase. <i>Advances in Enzymology and Related Areas of Molecular Biology</i> , 1982 , 53, 45-82		45
222	IPro: an iterative computational protein library redesign and optimization procedure. <i>Biophysical Journal</i> , 2006 , 90, 4167-80	2.9	45
221	Substituent effects of an antibody-catalyzed hydrolysis of phenyl esters: further evidence for an acyl-antibody intermediate. <i>Journal of the American Chemical Society</i> , 1992 , 114, 3528-3534	16.4	45
220	Principles of antibody catalysis. <i>BioEssays</i> , 1988 , 9, 107-12	4.1	45
219	Cyanotryptophans as Novel Fluorescent Probes for Studying Protein Conformational Changes and DNA-Protein Interaction. <i>Biochemistry</i> , 2015 , 54, 7457-69	3.2	44
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217	Biochemical characterization of bacteriophage T4 Mre11-Rad50 complex. <i>Journal of Biological Chemistry</i> , 2011 , 286, 2382-92	5.4	44
216	Mapping protein-protein proximity in the purinosome. <i>Journal of Biological Chemistry</i> , 2012 , 287, 36201-3	3.4	44
215	Cyclic peptides, a chemical genetics tool for biologists. <i>Cell Cycle</i> , 2005 , 4, 552-5	4.7	44
214	Electrostatic Characterization of Enzyme Complexes: Evaluation of the Mechanism of Catalysis of Dihydrofolate Reductase. <i>Journal of the American Chemical Society</i> , 1997 , 119, 2386-2395	16.4	43
213	Unexpected formation of an epoxide-derived multisubstrate adduct inhibitor on the active site of GAR transformylase. <i>Biochemistry</i> , 2001 , 40, 13538-47	3.2	43

212	Acoustofluidic chemical waveform generator and switch. <i>Analytical Chemistry</i> , 2014 , 86, 11803-10	7.8	42
211	Identification and mapping of protein-protein interactions between gp32 and gp59 by cross-linking. <i>Journal of Biological Chemistry</i> , 2001 , 276, 25236-42	5.4	42
210	Molecular basis for nonadditive mutational effects in Escherichia coli dihydrofolate reductase. <i>Biochemistry</i> , 1995 , 34, 15671-80	3.2	42
209	Assembly of the bacteriophage T4 primosome: single-molecule and ensemble studies. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005 , 102, 3254-9	11.5	41
208	Protein-protein interactions in the bacteriophage T4 replisome. The leading strand holoenzyme is physically linked to the lagging strand holoenzyme and the primosome. <i>Journal of Biological Chemistry</i> , 2003 , 278, 3145-52	5.4	40
207	Stability of the human polymerase I holoenzyme and its implications in lagging strand DNA synthesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, E1777-86	11.5	39
206	Multimeric structure of the secreted meprin A metalloproteinase and characterization of the functional protomer. <i>Journal of Biological Chemistry</i> , 2001 , 276, 23207-11	5.4	39
205	Molecular structure of Escherichia coli PurT-encoded glycinamide ribonucleotide transformylase. <i>Biochemistry</i> , 2000 , 39, 8791-802	3.2	39
204	Catalytic mechanism of Escherichia coli glycinamide ribonucleotide transformylase probed by site-directed mutagenesis and pH-dependent studies. <i>Biochemistry</i> , 1999 , 38, 10024-31	3.2	39
203	Stereoselective synthesis and biological activity of .beta.- and .alpha.-D-arabinose 1,5-diphosphate: analogs of a potent metabolic regulator. <i>Journal of the American Chemical Society</i> , 1984 , 106, 7851-7853	16.4	39
202	Interaction between the T4 helicase loading protein (gp59) and the DNA polymerase (gp43): unlocking of the gp59-gp43-DNA complex to initiate assembly of a fully functional replisome. <i>Biochemistry</i> , 2005 , 44, 7747-56	3.2	38
201	Effect of accessory proteins on T4 DNA polymerase replication fidelity. <i>Journal of Molecular Biology</i> , 1998 , 278, 135-46	6.5	38
200	Kinetic analysis of nucleotide incorporation and misincorporation by Klenow fragment of Escherichia coli DNA polymerase I. <i>Methods in Enzymology</i> , 1995 , 262, 257-69	1.7	38
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