

# Donald Hilvert

## List of Publications by Year in descending order

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229  
papers

16,226  
citations

14124

69  
h-index

24511

114  
g-index

254  
all docs

254  
docs citations

254  
times ranked

11783  
citing authors

#	ARTICLE	IF	CITATIONS
1	Whi3 mnemon association with endoplasmic reticulum membranes confines the memory of deceptive courtship to the yeast mother cell. <i>Current Biology</i> , 2022, 32, 963-974.e7.	1.8	7
2	Structure and Function of the $\hat{1}^2$ -Asp-Arg Polymerase Cyanophycin Synthetase 2. <i>ACS Chemical Biology</i> , 2022, 17, 670-679.	1.6	11
3	Protein Cages: From Fundamentals to Advanced Applications. <i>Chemical Reviews</i> , 2022, 122, 9145-9197.	23.0	54
4	The road to fully programmable protein catalysis. <i>Nature</i> , 2022, 606, 49-58.	13.7	126
5	A cryptic third active site in cyanophycin synthetase creates primers for polymerization. <i>Nature Communications</i> , 2022, 13, .	5.8	12
6	Trapping Transient Protein Species by Genetic Code Expansion. <i>ChemBioChem</i> , 2021, 22, 92-99.	1.3	7
7	Efficient Lewis acid catalysis of an abiological reaction in a de novo protein scaffold. <i>Nature Chemistry</i> , 2021, 13, 231-235.	6.6	46
8	Biosynthetic Functionalization of Nonribosomal Peptides. <i>Journal of the American Chemical Society</i> , 2021, 143, 2736-2740.	6.6	13
9	Cell-Specific Delivery Using an Engineered Protein Nanocage. <i>ACS Chemical Biology</i> , 2021, 16, 838-843.	1.6	16
10	The OP Protein Cage: A Versatile Molecular Delivery Platform. <i>Chimia</i> , 2021, 75, 323.	0.3	6
11	Evolution of the Chemical Step in Enzyme Catalysis. <i>ACS Catalysis</i> , 2021, 11, 6726-6732.	5.5	14
12	Analysis of electrostatic coupling throughout the laboratory evolution of a designed retroaldolase. <i>Protein Science</i> , 2021, 30, 1617-1627.	3.1	5
13	Noncanonical Heme Ligands Steer Carbene Transfer Reactivity in an Artificial Metalloenzyme**. <i>Angewandte Chemie</i> , 2021, 133, 15190-15195.	1.6	3
14	Noncanonical Heme Ligands Steer Carbene Transfer Reactivity in an Artificial Metalloenzyme**. <i>Angewandte Chemie - International Edition</i> , 2021, 60, 15063-15068.	7.2	18
15	Evolution of a virus-like architecture and packaging mechanism in a repurposed bacterial protein. <i>Science</i> , 2021, 372, 1220-1224.	6.0	53
16	Structures and function of the amino acid polymerase cyanophycin synthetase. <i>Nature Chemical Biology</i> , 2021, 17, 1101-1110.	3.9	24
17	Evolution of dynamical networks enhances catalysis in a designer enzyme. <i>Nature Chemistry</i> , 2021, 13, 1017-1022.	6.6	60
18	De novo peptide grafting to a self-assembling nanocapsule yields a hepatocyte growth factor receptor agonist. <i>IScience</i> , 2021, 24, 103302.	1.9	9

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19	Self-Assembly of Proteinaceous Shells around Positively Charged Gold Nanomaterials Enhances Colloidal Stability in High-Ionic-Strength Buffers. <i>ChemBioChem</i> , 2020, 21, 74-79.	1.3	11
20	How directed evolution reshapes the energy landscape in an enzyme to boost catalysis. <i>Science</i> , 2020, 370, 1442-1446.	6.0	101
21	Tight and specific lanthanide binding in a de novo TIM barrel with a large internal cavity designed by symmetric domain fusion. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 30362-30369.	3.3	31
22	Enhancing promiscuous chemistries of a Schiff-base forming enzyme by divergent evolution. <i>Methods in Enzymology</i> , 2020, 644, 95-120.	0.4	2
23	An evolution-based model for designing chorismate mutase enzymes. <i>Science</i> , 2020, 369, 440-445.	6.0	195
24	Two-tier supramolecular encapsulation of small molecules in a protein cage. <i>Nature Communications</i> , 2020, 11, 5410.	5.8	42
25	Engineered Artificial Carboligases Facilitate Regioselective Preparation of Enantioenriched Aldol Adducts. <i>Journal of the American Chemical Society</i> , 2020, 142, 10250-10254.	6.6	15
26	Contribution of Oxyanion Stabilization to Kemp Eliminase Efficiency. <i>ACS Catalysis</i> , 2020, 10, 4460-4464.	5.5	16
27	Syntheses of Cyanophycin Segments for Investigations of Cell-Penetration. <i>Synthesis</i> , 2019, 51, 31-39.	1.2	10
28	Emergence of a Negative Activation Heat Capacity during Evolution of a Designed Enzyme. <i>Journal of the American Chemical Society</i> , 2019, 141, 11745-11748.	6.6	42
29	Expanding the mass range for UVPD-based native top-down mass spectrometry. <i>Chemical Science</i> , 2019, 10, 7163-7171.	3.7	29
30	A computational method for design of connected catalytic networks in proteins. <i>Protein Science</i> , 2019, 28, 2036-2041.	3.1	28
31	Reprogramming Nonribosomal Peptide Synthesis by Surgical Mutation. <i>Synlett</i> , 2019, 30, 2123-2130.	1.0	7
32	Ultrahigh-throughput screening enables efficient single-round oxidase remodelling. <i>Nature Catalysis</i> , 2019, 2, 740-747.	16.1	74
33	Virus-Inspired Function in Engineered Protein Cages. <i>Journal of the American Chemical Society</i> , 2019, 141, 9432-9443.	6.6	46
34	Cytoplasmic glycoengineering enables biosynthesis of nanoscale glycoprotein assemblies. <i>Nature Communications</i> , 2019, 10, 5403.	5.8	36
35	Directed Evolution of Protein Catalysts. <i>Annual Review of Biochemistry</i> , 2018, 87, 131-157.	5.0	330
36	Stereodivergent Evolution of Artificial Enzymes for the Michael Reaction. <i>Angewandte Chemie</i> , 2018, 130, 5386-5389.	1.6	6

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37	Stereodivergent Evolution of Artificial Enzymes for the Michael Reaction. <i>Angewandte Chemie - International Edition</i> , 2018, 57, 5288-5291.	7.2	34
38	Speeding up enzyme discovery and engineering with ultrahigh-throughput methods. <i>Current Opinion in Structural Biology</i> , 2018, 48, 149-156.	2.6	103
39	Modular Protein Cages for Size-Selective RNA Packaging in Vivo. <i>Journal of the American Chemical Society</i> , 2018, 140, 566-569.	6.6	37
40	Substrate Sorting by a Supercharged Nanoreactor. <i>Journal of the American Chemical Society</i> , 2018, 140, 860-863.	6.6	48
41	Diversification of Protein Cage Structure Using Circularly Permuted Subunits. <i>Journal of the American Chemical Society</i> , 2018, 140, 558-561.	6.6	32
42	A Noncanonical Proximal Heme Ligand Affords an Efficient Peroxidase in a Globin Fold. <i>Journal of the American Chemical Society</i> , 2018, 140, 1535-1543.	6.6	79
43	Tailoring lumazine synthase assemblies for bionanotechnology. <i>Chemical Society Reviews</i> , 2018, 47, 3543-3557.	18.7	92
44	Nonribosomal biosynthesis of backbone-modified peptides. <i>Nature Chemistry</i> , 2018, 10, 282-287.	6.6	92
45	Evolution of a highly active and enantiospecific metalloenzyme from short peptides. <i>Science</i> , 2018, 362, 1285-1288.	6.0	116
46	Engineered Metalloenzymes with Noncanonical Coordination Environments. <i>Chemistry - A European Journal</i> , 2018, 24, 11821-11830.	1.7	33
47	Cell Penetration, Herbicidal Activity, and <i>in vivo</i> Toxicity of Oligoarginine Derivatives and of Novel Guanidinium-Rich Compounds Derived from the Biopolymer Cyanophycin. <i>Helvetica Chimica Acta</i> , 2018, 101, e1800112.	1.0	17
48	Capture and characterization of a reactive haem-carbenoid complex in an artificial metalloenzyme. <i>Nature Catalysis</i> , 2018, 1, 578-584.	16.1	93
49	Laboratory evolution of virus-like nucleocapsids from nonviral protein cages. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 5432-5437.	3.3	61
50	Frontispiece: Engineered Metalloenzymes with Noncanonical Coordination Environments. <i>Chemistry - A European Journal</i> , 2018, 24, .	1.7	0
51	Rational Engineering of a Designed Protein Cage for siRNA Delivery. <i>Journal of the American Chemical Society</i> , 2018, 140, 10439-10442.	6.6	86
52	Enzyme Encapsulation in an Engineered Lumazine Synthase Protein Cage. <i>Methods in Molecular Biology</i> , 2018, 1798, 39-55.	0.4	13
53	TOP-DOWN AND BOTTOM-UP APPROACHES FOR ELUCIDATING THE ORIGINS OF ENZYME EFFICIENCY. , 2018, , .		0
54	Chemoselective Henry Condensations Catalyzed by Artificial Carbolygases. <i>Chemistry - A European Journal</i> , 2017, 23, 6001-6003.	1.7	21

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55	Structure and assembly of scalable porous protein cages. <i>Nature Communications</i> , 2017, 8, 14663.	5.8	102
56	Irreversible Cysteine-Selective Protein Labeling Employing Modular Electrophilic Tetrafluoroethylation Reagents. <i>Chemistry - A European Journal</i> , 2017, 23, 6490-6494.	1.7	37
57	The C-terminal peptide of Aquifex aeolicus riboflavin synthase directs encapsulation of native and foreign guests by a cage-forming lumazine synthase. <i>Journal of Biological Chemistry</i> , 2017, 292, 10321-10327.	1.6	20
58	Synthesis and characterization of catalytically active thiazolium gold( <i>i</i> )-carbenes. <i>Chemical Communications</i> , 2017, 53, 7585-7587.	2.2	6
59	Enantiocomplementary Synthesis of $\hat{1}^3$ -Nitroketones Using Designed and Evolved Carboligases. <i>Journal of the American Chemical Society</i> , 2017, 139, 103-106.	6.6	37
60	Efficient laboratory evolution of computationally designed enzymes with low starting activities using fluorescence-activated droplet sorting. <i>Protein Engineering, Design and Selection</i> , 2017, 30, 531-531.	1.0	4
61	High-Mass MALDI-MS Analysis for the Investigation of Protein Encapsulation within an Engineered Capsid Forming Protein. <i>Helvetica Chimica Acta</i> , 2017, 100, e1700166.	1.0	3
62	Surface-Engineered Cationic Nanocrystals Stable in Biological Buffers and High Ionic Strength Solutions. <i>Chemistry of Materials</i> , 2017, 29, 9416-9428.	3.2	31
63	Enzyme Encapsulation by a Ferritin Cage. <i>Angewandte Chemie</i> , 2017, 129, 15129-15132.	1.6	72
64	Enzyme Encapsulation by a Ferritin Cage. <i>Angewandte Chemie - International Edition</i> , 2017, 56, 14933-14936.	7.2	89
65	Optimization of Enzyme Mechanism along the Evolutionary Trajectory of a Computationally Designed (Retro-)Aldolase. <i>Journal of the American Chemical Society</i> , 2017, 139, 12541-12549.	6.6	45
66	Emergence of a catalytic tetrad during evolution of a highly active artificial aldolase. <i>Nature Chemistry</i> , 2017, 9, 50-56.	6.6	248
67	Evaluation of the Ser-His Dipeptide, a Putative Catalyst of Amide and Ester Hydrolysis. <i>Organic Letters</i> , 2016, 18, 3518-3521.	2.4	21
68	Chiral recognition in amyloid fiber growth. <i>Journal of Peptide Science</i> , 2016, 22, 290-304.	0.8	25
69	Quantitative Beladung eines Proteinkäfigs mit aktiven Enzymen. <i>Angewandte Chemie</i> , 2016, 128, 1555-1558.	1.6	14
70	DNA Nanoparticles for Improved Protein Synthesis In Vitro. <i>Angewandte Chemie - International Edition</i> , 2016, 55, 3120-3123.	7.2	19
71	Fast Knoevenagel Condensations Catalyzed by an Artificial Schiff-Base-Forming Enzyme. <i>Journal of the American Chemical Society</i> , 2016, 138, 6972-6974.	6.6	83
72	DNA Nanoparticles for Improved Protein Synthesis In Vitro. <i>Angewandte Chemie</i> , 2016, 128, 3172-3175.	1.6	8

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73	Self-Assembly of Proteinaceous Multishell Structures Mediated by a Supercharged Protein. <i>Journal of Physical Chemistry B</i> , 2016, 120, 6089-6095.	1.2	22
74	Enzyme-mediated polymerization inside engineered protein cages. <i>Chemical Communications</i> , 2016, 52, 10423-10426.	2.2	30
75	Bottom-up Construction of a Primordial Carboxysome Mimic. <i>Journal of the American Chemical Society</i> , 2016, 138, 10072-10075.	6.6	73
76	A Chemically Programmed Proximal Ligand Enhances the Catalytic Properties of a Heme Enzyme. <i>Journal of the American Chemical Society</i> , 2016, 138, 11344-11352.	6.6	64
77	Efficient laboratory evolution of computationally designed enzymes with low starting activities using fluorescence-activated droplet sorting. <i>Protein Engineering, Design and Selection</i> , 2016, 29, 355-366.	1.0	58
78	Upregulation of an Artificial Zymogen by Proteolysis. <i>Angewandte Chemie - International Edition</i> , 2016, 55, 11587-11590.	7.2	29
79	Upregulation of an Artificial Zymogen by Proteolysis. <i>Angewandte Chemie</i> , 2016, 128, 11759-11762.	1.6	7
80	Quantitative Packaging of Active Enzymes into a Protein Cage. <i>Angewandte Chemie - International Edition</i> , 2016, 55, 1531-1534.	7.2	110
81	Construction of Matryoshka-Type Structures from Supercharged Protein Nanocages. <i>Angewandte Chemie - International Edition</i> , 2015, 54, 937-940.	7.2	60
82	A Subdomain Swap Strategy for Reengineering Nonribosomal Peptides. <i>Chemistry and Biology</i> , 2015, 22, 640-648.	6.2	90
83	Diffusion-Limited Cargo Loading of an Engineered Protein Container. <i>Journal of the American Chemical Society</i> , 2015, 137, 16121-16132.	6.6	62
84	Substitution of Proline <sup>32</sup> by $\beta$ -Methylproline Preorganizes $\beta$ -2-Microglobulin for Oligomerization but Not for Aggregation into Amyloids. <i>Journal of the American Chemical Society</i> , 2015, 137, 2524-2535.	6.6	17
85	Comparative Laboratory Evolution of Ordered and Disordered Enzymes. <i>Journal of Biological Chemistry</i> , 2015, 290, 9310-9320.	1.6	12
86	A Promiscuous De $\alpha$ -Novo Retro $\alpha$ -Aldolase Catalyzes Asymmetric Michael Additions via Schiff Base Intermediates. <i>Angewandte Chemie - International Edition</i> , 2015, 54, 5609-5612.	7.2	53
87	Kinetic Consequences of Introducing a Proximal Selenocysteine Ligand into Cytochrome P450cam. <i>Biochemistry</i> , 2015, 54, 6692-6703.	1.2	14
88	Harnessing selenocysteine reactivity for oxidative protein folding. <i>Chemical Science</i> , 2015, 6, 322-325.	3.7	50
89	Impact of scaffold rigidity on the design and evolution of an artificial Diels-Alderase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 8013-8018.	3.3	111
90	Electrostatic transition state stabilization rather than reactant destabilization provides the chemical basis for efficient chorismate mutase catalysis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 17516-17521.	3.3	31

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91	Active Site Plasticity of a Computationally Designed Retroaldolase Enzyme. <i>ChemCatChem</i> , 2014, 6, 1043-1050.	1.8	23
92	Building Proficient Enzymes with Foldamer Prostheses. <i>Angewandte Chemie - International Edition</i> , 2014, 53, 6978-6981.	7.2	54
93	Exploration of Alternate Catalytic Mechanisms and Optimization Strategies for Retroaldolase Design. <i>Journal of Molecular Biology</i> , 2014, 426, 256-271.	2.0	33
94	Reprogramming Nonribosomal Peptide Synthetases for "Clickable" Amino Acids. <i>Angewandte Chemie</i> , 2014, 126, 10269-10272.	1.6	20
95	Natural and synthetic selenoproteins. <i>Current Opinion in Chemical Biology</i> , 2014, 22, 27-34.	2.8	80
96	Reprogramming Nonribosomal Peptide Synthetases for "Clickable" Amino Acids. <i>Angewandte Chemie - International Edition</i> , 2014, 53, 10105-10108.	7.2	102
97	Affinity maturation of a computationally designed binding protein affords a functional but disordered polypeptide. <i>Journal of Structural Biology</i> , 2014, 185, 168-177.	1.3	10
98	Precision is essential for efficient catalysis in an evolved Kemp eliminase. <i>Nature</i> , 2013, 503, 418-421.	13.7	281
99	De novo enzymes by computational design. <i>Current Opinion in Chemical Biology</i> , 2013, 17, 221-228.	2.8	221
100	Evolution of a designed retro-aldolase leads to complete active site remodeling. <i>Nature Chemical Biology</i> , 2013, 9, 494-498.	3.9	220
101	A Genetically Encodable Ligand for Transfer Hydrogenation. <i>European Journal of Organic Chemistry</i> , 2013, 2013, 3427-3431.	1.2	17
102	Design of Protein Catalysts. <i>Annual Review of Biochemistry</i> , 2013, 82, 447-470.	5.0	174
103	Protein Conformational Disorder and Enzyme Catalysis. <i>Topics in Current Chemistry</i> , 2013, 337, 41-67.	4.0	47
104	Directed Evolution of a Model Primordial Enzyme Provides Insights into the Development of the Genetic Code. <i>PLoS Genetics</i> , 2013, 9, e1003187.	1.5	27
105	<i>cis</i> → <i>trans</i> Peptide Bond Isomerization in <i>trans</i> -Methylproline Derivatives. <i>Helvetica Chimica Acta</i> , 2012, 95, 2411-2420.	1.0	9
106	Iterative approach to computational enzyme design. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 3790-3795.	3.3	291
107	A novel genetic selection system for PLP-dependent threonine aldolases. <i>Tetrahedron</i> , 2012, 68, 7549-7557.	1.0	17
108	Harnessing Protein Symmetry for Enzyme Design. <i>ACS Catalysis</i> , 2012, 2, 982-985.	5.5	14

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109	Structural Analyses of Covalent Enzyme-Substrate Analog Complexes Reveal Strengths and Limitations of De Novo Enzyme Design. <i>Journal of Molecular Biology</i> , 2012, 415, 615-625.	2.0	54
110	Computational Design of Catalytic Dyads and Oxyanion Holes for Ester Hydrolysis. <i>Journal of the American Chemical Society</i> , 2012, 134, 16197-16206.	6.6	138
111	Robust design and optimization of retroaldol enzymes. <i>Protein Science</i> , 2012, 21, 717-726.	3.1	137
112	Efficient in Vitro Encapsulation of Protein Cargo by an Engineered Protein Container. <i>Journal of the American Chemical Society</i> , 2012, 134, 909-911.	6.6	109
113	Strategic Use of Non-Native Diselenide Bridges to Steer Oxidative Protein Folding. <i>Angewandte Chemie - International Edition</i> , 2012, 51, 5585-5588.	7.2	89
114	An N-Terminal Protein Degradation Tag Enables Robust Selection of Highly Active Enzymes. <i>Biochemistry</i> , 2011, 50, 8594-8602.	1.2	20
115	An artificial metalloenzyme for olefin metathesis. <i>Chemical Communications</i> , 2011, 47, 12068.	2.2	143
116	Tailor-Made Peptide Synthetases. <i>Chemistry and Biology</i> , 2011, 18, 1206-1207.	6.2	4
117	Selenogluthathione-Mediated Rescue of Kinetically Trapped Intermediates in Oxidative Protein Folding. <i>Israel Journal of Chemistry</i> , 2011, 51, 953-959.	1.0	25
118	Directed Evolution of a Protein Container. <i>Science</i> , 2011, 331, 589-592.	6.0	280
119	Photolyase-Like Repair of Psoralen-Crosslinked Nucleic Acids. <i>Angewandte Chemie - International Edition</i> , 2011, 50, 9483-9486.	7.2	11
120	Modulating PNA/DNA Hybridization by Light. <i>Angewandte Chemie - International Edition</i> , 2010, 49, 9998-10001.	7.2	55
121	Diselenides as universal oxidative folding catalysts of diverse proteins. <i>Journal of Biotechnology</i> , 2010, 150, 481-489.	1.9	43
122	Biocatalysts by evolution. <i>Current Opinion in Biotechnology</i> , 2010, 21, 753-759.	3.3	120
123	Design, selection, and characterization of a split chorismate mutase. <i>Protein Science</i> , 2010, 19, 1000-1010.	3.1	19
124	Computational Design of an Enzyme Catalyst for a Stereoselective Bimolecular Diels-Alder Reaction. <i>Science</i> , 2010, 329, 309-313.	6.0	776
125	Small-Molecule Diselenides Catalyze Oxidative Protein Folding <i>in Vivo</i> . <i>ACS Chemical Biology</i> , 2010, 5, 177-182.	1.6	28
126	Consensus Protein Design without Phylogenetic Bias. <i>Journal of Molecular Biology</i> , 2010, 399, 541-546.	2.0	73



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127	An aspartate and a water molecule mediate efficient acid-base catalysis in a tailored antibody pocket. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 18539-18544.	3.3	23
128	Probing the role of the proximal heme ligand in cytochrome P450cam by recombinant incorporation of selenocysteine. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 5481-5486.	3.3	68
129	A Rationally Designed Aldolase Foldamer. <i>Angewandte Chemie - International Edition</i> , 2009, 48, 922-925.	7.2	150
130	Selenogluta-redoxin as a Glutathione Peroxidase Mimic. <i>ChemBioChem</i> , 2008, 9, 1623-1631.	1.3	48
131	Synthesis of $\beta$ -hydroxy- $\alpha$ -amino acids with a reengineered alanine racemase. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2008, 18, 5987-5990.	1.0	26
132	Protein Design by Directed Evolution. <i>Annual Review of Biophysics</i> , 2008, 37, 153-173.	4.5	344
133	Switching Antibody Specificity through Minimal Mutation. <i>Journal of Molecular Biology</i> , 2008, 377, 993-1001.	2.0	5
134	Kinetics and Thermodynamics of Ligand Binding to a Molten Globular Enzyme and Its Native Counterpart. <i>Journal of Molecular Biology</i> , 2008, 382, 971-977.	2.0	67
135	The Putative Diels-Alderase Macrophomate Synthase is an Efficient Aldolase. <i>Journal of the American Chemical Society</i> , 2008, 130, 7798-7799.	6.6	70
136	Catalysis of Oxidative Protein Folding by Small-Molecule Diselenides. <i>Biochemistry</i> , 2008, 47, 6985-6987.	1.2	68
137	Relative Tolerance of an Enzymatic Molten Globule and Its Thermostable Counterpart to Point Mutation. <i>Biochemistry</i> , 2008, 47, 13489-13496.	1.2	30
138	De Novo Computational Design of Retro-Aldol Enzymes. <i>Science</i> , 2008, 319, 1387-1391.	6.0	1,031
139	Closely related antibody receptors exploit fundamentally different strategies for steroid recognition. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 11725-11730.	3.3	8
140	Conformational Isomerism Can Limit Antibody Catalysis. <i>Journal of Biological Chemistry</i> , 2008, 283, 16554-16560.	1.6	12
141	On the relationship between folding and chemical landscapes in enzyme catalysis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 13877-13882.	3.3	82
142	Metabolic engineering of a genetic selection system with tunable stringency. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 13907-13912.	3.3	16
143	Reinvestigation of a Selenopeptide with Purportedly High Glutathione Peroxidase Activity. <i>Journal of Biological Chemistry</i> , 2007, 282, 30518-30522.	1.6	16
144	Stereoselective Reactions with Catalytic Antibodies. <i>Topics in Stereochemistry</i> , 2007, , 83-135.	2.0	7

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145	Bifunctional Catalysis of Proton Transfer at an Antibody Active Site. <i>Journal of the American Chemical Society</i> , 2007, 129, 460-461.	6.6	21
146	Selenoglutathione: Efficient Oxidative Protein Folding by a Diselenide. <i>Biochemistry</i> , 2007, 46, 5382-5390.	1.2	136
147	Minimalist Active-Site Redesign: Teaching Old Enzymes New Tricks. <i>Angewandte Chemie - International Edition</i> , 2007, 46, 3212-3236.	7.2	244
148	Enhancing Activity and Controlling Stereoselectivity in a Designed PLP-Dependent Aldolase. <i>Angewandte Chemie - International Edition</i> , 2007, 46, 4468-4470.	7.2	39
149	A simple selection strategy for evolving highly efficient enzymes. <i>Nature Biotechnology</i> , 2007, 25, 1145-1147.	9.4	59
150	Synthesis and application of an azobenzene amino acid as a light-switchable turn element in polypeptides. <i>Nature Protocols</i> , 2007, 2, 161-167.	5.5	31
151	Structure and dynamics of a molten globular enzyme. <i>Nature Structural and Molecular Biology</i> , 2007, 14, 1202-1206.	3.6	102
152	Novel Enzymes Through Design and Evolution. <i>Advances in Enzymology and Related Areas of Molecular Biology</i> , 2007, 75, 241-294.	1.3	14
153	A Monofunctional and Thermostable Prephenate Dehydratase from the Archaeon <i>Methanocaldococcus jannaschii</i> . <i>Biochemistry</i> , 2006, 45, 14101-14110.	1.2	20
154	A Simple Tagging System for Protein Encapsulation. <i>Journal of the American Chemical Society</i> , 2006, 128, 4516-4517.	6.6	145
155	Mimicking Enzymes with Antibodies. , 2006, , 89-107.		1
156	Relative tolerance of mesostable and thermostable protein homologs to extensive mutation. <i>Proteins: Structure, Function and Bioinformatics</i> , 2006, 66, 500-506.	1.5	57
157	Toward bifunctional antibody catalysis. <i>Bioorganic and Medicinal Chemistry</i> , 2006, 14, 6189-6196.	1.4	19
158	Tunnel plasticity and quaternary structural integrity of a pentameric protein ring. <i>Protein Science</i> , 2006, 15, 1106-1114.	3.1	10
159	A Photoswitchable Miniprotein Based on the Sequence of Avian Pancreatic Polypeptide. <i>Angewandte Chemie - International Edition</i> , 2006, 45, 6297-6300.	7.2	36
160	Stereoselectivity and Expanded Substrate Scope of an Engineered PLP-Dependent Aldolase. <i>Angewandte Chemie - International Edition</i> , 2006, 45, 6824-6826.	7.2	32
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