

Ulrich Schwaneberg

List of Publications by Year in descending order

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351
papers

11,114
citations

30068

54
h-index

66906

78
g-index

379
all docs

379
docs citations

379
times ranked

7725
citing authors

#	ARTICLE	IF	CITATIONS
1	Laboratory evolution of cytochrome P450 BM-3 monooxygenase for organic cosolvents. <i>Biotechnology and Bioengineering</i> , 2004, 85, 351-358.	3.3	184
2	Advances in ultrahigh-throughput screening for directed enzyme evolution. <i>Chemical Society Reviews</i> , 2020, 49, 233-262.	38.1	182
3	Directed Evolution Empowered Redesign of Natural Proteins for the Sustainable Production of Chemicals and Pharmaceuticals. <i>Angewandte Chemie - International Edition</i> , 2019, 58, 36-40.	13.8	169
4	Directed Evolution of the Fatty-Acid Hydroxylase P450 BM-3 into an Indole-Hydroxylating Catalyst. <i>Chemistry - A European Journal</i> , 2000, 6, 1531-1536.	3.3	167
5	Advances in generating functional diversity for directed protein evolution. <i>Current Opinion in Chemical Biology</i> , 2009, 13, 19-25.	6.1	156
6	Directed Evolution of a Cytochrome P450 Monooxygenase for Alkane Oxidation. <i>Advanced Synthesis and Catalysis</i> , 2001, 343, 601-606.	4.3	148
7	A Continuous Spectrophotometric Assay for P450 BM-3, a Fatty Acid Hydroxylating Enzyme, and Its Mutant F87A. <i>Analytical Biochemistry</i> , 1999, 269, 359-366.	2.4	143
8	A P450 BM-3 mutant hydroxylates alkanes, cycloalkanes, arenes and heteroarenes. <i>Journal of Biotechnology</i> , 2001, 88, 167-171.	3.8	136
9	Protein engineering in bioelectrocatalysis. <i>Current Opinion in Biotechnology</i> , 2003, 14, 590-596.	6.6	132
10	A Statistical Analysis of Random Mutagenesis Methods Used for Directed Protein Evolution. <i>Journal of Molecular Biology</i> , 2006, 355, 858-871.	4.2	132
11	Sequence saturation mutagenesis (SeSaM): a novel method for directed evolution. <i>Nucleic Acids Research</i> , 2004, 32, 26e-26.	14.5	130
12	Directed evolution 2.0: improving and deciphering enzyme properties. <i>Chemical Communications</i> , 2015, 51, 9760-9772.	4.1	122
13	Reengineering CelA2 cellulase for hydrolysis in aqueous solutions of deep eutectic solvents and concentrated seawater. <i>Green Chemistry</i> , 2012, 14, 2719.	9.0	120
14	Applying metagenomics for the identification of bacterial cellulases that are stable in ionic liquids. <i>Green Chemistry</i> , 2009, 11, 957.	9.0	113
15	Asymmetric reduction of ketones with recombinant E. coli whole cells in neat substrates. <i>Chemical Communications</i> , 2011, 47, 12230.	4.1	111
16	Making glucose oxidase fit for biofuel cell applications by directed protein evolution. <i>Biosensors and Bioelectronics</i> , 2006, 21, 2046-2051.	10.1	109
17	Phosphorothioate-based ligase-independent gene cloning (PLICing): An enzyme-free and sequence-independent cloning method. <i>Analytical Biochemistry</i> , 2010, 406, 141-146.	2.4	109
18	Arginine deiminase, a potential anti-tumor drug. <i>Cancer Letters</i> , 2008, 261, 1-11.	7.2	105

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19	A nanocompartment system (Synthosome) designed for biotechnological applications. <i>Journal of Biotechnology</i> , 2006, 123, 50-59.	3.8	104
20	Challenges and advances in the field of self-assembled membranes. <i>Chemical Society Reviews</i> , 2013, 42, 6578.	38.1	96
21	Rational evolution of a medium chain-specific cytochrome P-450 BM-3 variant. <i>BBA - Proteins and Proteomics</i> , 2001, 1545, 114-121.	2.1	94
22	P450 in biotechnology: zinc driven β -hydroxylation of p-nitrophenoxydodecanoic acid using P450 BM-3 F87A as a catalyst. <i>Journal of Biotechnology</i> , 2000, 84, 249-257.	3.8	92
23	The Diversity Challenge in Directed Protein Evolution. <i>Combinatorial Chemistry and High Throughput Screening</i> , 2006, 9, 271-288.	1.1	92
24	Biofunctional Microgel-Based Fertilizers for Controlled Foliar Delivery of Nutrients to Plants. <i>Angewandte Chemie - International Edition</i> , 2017, 56, 7380-7386.	13.8	89
25	Structural Insight into Enantioselective Inversion of an Alcohol Dehydrogenase Reveals a β -Polar Gate in Stereorecognition of Diaryl Ketones. <i>Journal of the American Chemical Society</i> , 2018, 140, 12645-12654.	13.7	87
26	OmniChange: The Sequence Independent Method for Simultaneous Site-Saturation of Five Codons. <i>PLoS ONE</i> , 2011, 6, e26222.	2.5	83
27	Advances in protease engineering for laundry detergents. <i>New Biotechnology</i> , 2015, 32, 629-634.	4.4	82
28	A rhodium complex-linked β -barrel protein as a hybrid biocatalyst for phenylacetylene polymerization. <i>Chemical Communications</i> , 2012, 48, 9756.	4.1	78
29	Regioselective β -Hydroxylation of Monosubstituted Benzenes by P450 BM3. <i>Angewandte Chemie - International Edition</i> , 2013, 52, 8459-8462.	13.8	77
30	Cloning and characterization of a thermostable and halo-tolerant endoglucanase from <i>Thermoanaerobacter tengcongensis</i> MB4. <i>Applied Microbiology and Biotechnology</i> , 2011, 89, 315-326.	3.6	76
31	Directed Evolution of Oxygenases: Screening Systems, Success Stories and Challenges. <i>Combinatorial Chemistry and High Throughput Screening</i> , 2007, 10, 197-217.	1.1	72
32	Increasing activity and thermal resistance of <i>Bacillus gibsonii</i> alkaline protease (BgAP) by directed evolution. <i>Biotechnology and Bioengineering</i> , 2013, 110, 711-720.	3.3	72
33	Targeting microplastic particles in the void of diluted suspensions. <i>Environment International</i> , 2019, 123, 428-435.	10.0	72
34	Stereoselective epoxidation of the last double bond of polyunsaturated fatty acids by human cytochromes P450. <i>Journal of Lipid Research</i> , 2010, 51, 1125-1133.	4.2	71
35	Cellulolytic RoboLector β towards an automated high-throughput screening platform for recombinant cellulase expression. <i>Journal of Biological Engineering</i> , 2017, 11, 1.	4.7	71
36	A Whole Cell <i>E. coli</i> Display Platform for Artificial Metalloenzymes: Poly(phenylacetylene) Production with a Rhodium β -Nitrobindin Metalloprotein. <i>ACS Catalysis</i> , 2018, 8, 2611-2614.	11.2	71

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37	Directed evolution of a thermophilic endoglucanase (Cel5A) into highly active Cel5A variants with an expanded temperature profile. <i>Journal of Biotechnology</i> , 2011, 154, 46-53.	3.8	70
38	A Hybrid Ring-Opening Metathesis Polymerization Catalyst Based on an Engineered Variant of the β -Barrel Protein FhuA. <i>Chemistry - A European Journal</i> , 2013, 19, 13865-13871.	3.3	70
39	Laboratory evolution of P450 BM3 for mediated electron transfer yielding an activity-improved and reductase-independent variant. <i>Protein Engineering, Design and Selection</i> , 2007, 21, 29-35.	2.1	68
40	A Highly Active Biohybrid Catalyst for Olefin Metathesis in Water: Impact of a Hydrophobic Cavity in a β -Barrel Protein. <i>ACS Catalysis</i> , 2015, 5, 7519-7522.	11.2	68
41	Engineering Robust Cellulases for Tailored Lignocellulosic Degradation Cocktails. <i>International Journal of Molecular Sciences</i> , 2020, 21, 1589.	4.1	68
42	P450 monooxygenase in biotechnology. <i>Journal of Chromatography A</i> , 1999, 848, 149-159.	3.7	66
43	Direct Oxidation of Cycloalkanes to Cycloalkanones with Oxygen in Water. <i>Angewandte Chemie - International Edition</i> , 2013, 52, 2359-2363.	13.8	65
44	Directed evolution of a highly active <i>Yersinia mollaretii</i> phytase. <i>Applied Microbiology and Biotechnology</i> , 2012, 95, 405-418.	3.6	64
45	Directed laccase evolution for improved ionic liquid resistance. <i>Green Chemistry</i> , 2013, 15, 1348.	9.0	64
46	Towards the Evolution of Artificial Metalloenzymes – A Protein Engineer's Perspective. <i>Angewandte Chemie - International Edition</i> , 2019, 58, 4454-4464.	13.8	64
47	Functionalized Nanocompartments (Synthosomes) with a Reduction-Triggered Release System. <i>Angewandte Chemie - International Edition</i> , 2008, 47, 7029-7031.	13.8	63
48	Directed evolution of glucose oxidase from <i>Aspergillus niger</i> for ferrocenemethanol-mediated electron transfer. <i>Biotechnology Journal</i> , 2007, 2, 241-248.	3.5	61
49	A Screening System for the Directed Evolution of Epoxigenases: Importance of Position 184 in P450 ^{BM3} for Stereoselective Styrene Epoxidation. <i>Angewandte Chemie - International Edition</i> , 2006, 45, 5380-5383.	13.8	59
50	Steering directed protein evolution: strategies to manage combinatorial complexity of mutant libraries. <i>Environmental Microbiology</i> , 2007, 9, 2645-2659.	3.8	59
51	Protein Engineering – An Option for Enzymatic Biofuel Cell Design. <i>Electroanalysis</i> , 2010, 22, 765-775.	2.9	59
52	To get what we aim for – Progress in diversity generation methods. <i>FEBS Journal</i> , 2013, 280, 2961-2978.	4.7	59
53	PTEN-inhibition by zinc ions augments interleukin-2-mediated Akt phosphorylation. <i>Metallomics</i> , 2014, 6, 1277.	2.4	59
54	Machine learning-assisted enzyme engineering. <i>Methods in Enzymology</i> , 2020, 643, 281-315.	1.0	59

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55	A roadmap to directed enzyme evolution and screening systems for biotechnological applications. <i>Biological Research</i> , 2013, 46, 395-405.	3.4	57
56	Directed Evolution of a Bacterial Laccase (CueO) for Enzymatic Biofuel Cells. <i>Angewandte Chemie - International Edition</i> , 2019, 58, 4562-4565.	13.8	57
57	Computer-Assisted Recombination (CompassR) Teaches us How to Recombine Beneficial Substitutions from Directed Evolution Campaigns. <i>Chemistry - A European Journal</i> , 2020, 26, 643-649.	3.3	57
58	Anchor peptides: A green and versatile method for polypropylene functionalization. <i>Polymer</i> , 2017, 116, 124-132.	3.8	55
59	An efficient transformation method for <i>Bacillus subtilis</i> DB104. <i>Applied Microbiology and Biotechnology</i> , 2012, 94, 487-493.	3.6	53
60	Enzyme Hydration Determines Resistance in Organic Cosolvents. <i>ACS Catalysis</i> , 2020, 10, 14847-14856.	11.2	53
61	COMPUTER-AIDED PROTEIN DIRECTED EVOLUTION: A REVIEW OF WEB SERVERS, DATABASES AND OTHER COMPUTATIONAL TOOLS FOR PROTEIN ENGINEERING. <i>Computational and Structural Biotechnology Journal</i> , 2012, 2, e201209008.	4.1	52
62	Multi-step biocatalytic depolymerization of lignin. <i>Applied Microbiology and Biotechnology</i> , 2017, 101, 6277-6287.	3.6	51
63	Directed Evolution of the Fatty-Acid Hydroxylase P450 BM-3 into an Indole-Hydroxylating Catalyst. <i>Chemistry - A European Journal</i> , 2000, 6, 1531-1536.	3.3	49
64	Sortase-Mediated Surface Functionalization of Stimuli-Responsive Microgels. <i>Biomacromolecules</i> , 2017, 18, 2789-2798.	5.4	49
65	A Flow Cytometry-Based Screening System for Directed Evolution of Proteases. <i>Journal of Biomolecular Screening</i> , 2011, 16, 285-294.	2.6	47
66	Ionic liquid and deep eutectic solvent-activated CelA2 variants generated by directed evolution. <i>Applied Microbiology and Biotechnology</i> , 2014, 98, 5775-5785.	3.6	47
67	In vitro flow cytometry-based screening platform for cellulase engineering. <i>Scientific Reports</i> , 2016, 6, 26128.	3.3	47
68	KnowVolution of the Polymer-Binding Peptide LCI for Improved Polypropylene Binding. <i>Polymers</i> , 2018, 10, 423.	4.5	47
69	Sensitive Assay for Laboratory Evolution of Hydroxylases toward Aromatic and Heterocyclic Compounds. <i>Journal of Biomolecular Screening</i> , 2005, 10, 246-252.	2.6	46
70	Ionic liquid effects on the activity of monooxygenase P450 BM-3. <i>Green Chemistry</i> , 2008, 10, 117-123.	9.0	46
71	Reengineered glucose oxidase for amperometric glucose determination in diabetes analytics. <i>Biosensors and Bioelectronics</i> , 2013, 50, 84-90.	10.1	46
72	Tunable Enzymatic Activity and Enhanced Stability of Cellulase Immobilized in Biohybrid Nanogels. <i>Biomacromolecules</i> , 2016, 17, 3619-3631.	5.4	46

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73	Expression of the Zn ²⁺ -containing hydroxynitrile lyase from flax (<i>Linum usitatissimum</i>) in <i>Pichia pastoris</i> utilization of the recombinant enzyme for enzymatic analysis and site-directed mutagenesis. <i>Plant Science</i> , 1998, 139, 19-27.	3.6	45
74	Cost-Effective Whole-Cell Assay for Laboratory Evolution of Hydroxylases in <i>Escherichia coli</i> . <i>Journal of Biomolecular Screening</i> , 2001, 6, 111-117.	2.6	45
75	A Fluorescent Hydrogel-Based Flow Cytometry High-Throughput Screening Platform for Hydrolytic Enzymes. <i>Chemistry and Biology</i> , 2014, 21, 1733-1742.	6.0	45
76	Towards Understanding Directed Evolution: More than Half of All Amino Acid Positions Contribute to Ionic Liquid Resistance of <i>Bacillus subtilis</i> Lipase A. <i>ChemBioChem</i> , 2015, 16, 937-945.	2.6	45
77	Electron transfer pathways in a light, oxygen, voltage (LOV) protein devoid of the photoactive cysteine. <i>Scientific Reports</i> , 2017, 7, 13346.	3.3	45
78	Disulfide Bond Engineering of an Endoglucanase from <i>Penicillium verruculosum</i> to Improve Its Thermostability. <i>International Journal of Molecular Sciences</i> , 2019, 20, 1602.	4.1	45
79	How to Engineer Organic Solvent Resistant Enzymes: Insights from Combined Molecular Dynamics and Directed Evolution Study. <i>ChemCatChem</i> , 2020, 12, 4073-4083.	3.7	45
80	Less Unfavorable Salt Bridges on the Enzyme Surface Result in More Organic Cosolvent Resistance. <i>Angewandte Chemie - International Edition</i> , 2021, 60, 11448-11456.	13.8	45
81	Understanding a Mechanism of Organic Cosolvent Inactivation in Heme Monooxygenase P450 BM-3. <i>Journal of the American Chemical Society</i> , 2007, 129, 5786-5787.	13.7	44
82	Engineering of the <i>E. coli</i> Outer Membrane Protein FhuA to overcome the Hydrophobic Mismatch in Thick Polymeric Membranes. <i>Journal of Nanobiotechnology</i> , 2011, 9, 8.	9.1	44
83	Flow Cytometer-Based High-Throughput Screening System for Accelerated Directed Evolution of P450 Monooxygenases. <i>ACS Catalysis</i> , 2012, 2, 2724-2728.	11.2	44
84	Exploring the Protein Stability Landscape: <i>Bacillus subtilis</i> Lipase A as a Model for Detergent Tolerance. <i>ChemBioChem</i> , 2015, 16, 930-936.	2.6	44
85	Hybrid Ruthenium ROMP Catalysts Based on an Engineered Variant of β -Barrel Protein FhuA β -CVF ^{teV} : Effect of Spacer Length. <i>Chemistry - an Asian Journal</i> , 2015, 10, 177-182.	3.3	44
86	Multi-site saturation by OmniChange yields a pH- and thermally improved phytase. <i>Journal of Biotechnology</i> , 2014, 170, 68-72.	3.8	43
87	Directed evolution of polypropylene and polystyrene binding peptides. <i>Biotechnology and Bioengineering</i> , 2018, 115, 321-330.	3.3	42
88	Modification of the fatty acid specificity of cytochrome P450 BM-3 from <i>Bacillus megaterium</i> by directed evolution: a validated assay. <i>Journal of Molecular Catalysis B: Enzymatic</i> , 2001, 15, 123-133.	1.8	41
89	Laboratory Evolution of P450 BM-3 for Mediated Electron Transfer. <i>ChemBioChem</i> , 2006, 7, 638-644.	2.6	41
90	Photophysics of the LOV-Based Fluorescent Protein Variant iLOV-Q489K Determined by Simulation and Experiment. <i>Journal of Physical Chemistry B</i> , 2016, 120, 3344-3352.	2.6	41

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91	Directed Evolution of P _{450 BM3} into a <i>pXyl</i> Xylene Hydroxylase. <i>ChemCatChem</i> , 2012, 4, 771-773.	3.7	40
92	Lessons from diversity of directed evolution experiments by an analysis of 3,000 mutations. <i>Biotechnology and Bioengineering</i> , 2014, 111, 2380-2389.	3.3	40
93	KnowVolution of a Fungal Laccase toward Alkaline pH. <i>ChemBioChem</i> , 2019, 20, 1458-1466.	2.6	40
94	Transversion-enriched sequence saturation mutagenesis (SeSaTv): A random mutagenesis method with consecutive nucleotide exchanges that complements the bias of error-prone PCR. <i>Biotechnology Journal</i> , 2008, 3, 74-82.	3.5	39
95	Surface charge engineering of a <i>Bacillus gibsonii</i> subtilisin protease. <i>Applied Microbiology and Biotechnology</i> , 2013, 97, 6793-6802.	3.6	39
96	QM/MM Calculations Revealing the Resting and Catalytic States in Zinc-Dependent Medium-Chain Dehydrogenases/Reductases. <i>ACS Catalysis</i> , 2015, 5, 3207-3215.	11.2	39
97	Cavity Size Engineering of a β -Barrel Protein Generates Efficient Biohybrid Catalysts for Olefin Metathesis. <i>ACS Catalysis</i> , 2018, 8, 3358-3364.	11.2	39
98	<i>In Vitro</i> Double Oxidation of <i>n</i> -Heptane with Direct Cofactor Regeneration. <i>Advanced Synthesis and Catalysis</i> , 2013, 355, 1787-1798.	4.3	38
99	A loop engineering strategy improves laccase lcc2 activity in ionic liquid and aqueous solution. <i>Green Chemistry</i> , 2018, 20, 2801-2812.	9.0	38
100	Membrane-Mimetic Dendrimerosomes Engulf Living Bacteria via Endocytosis. <i>Nano Letters</i> , 2019, 19, 5732-5738.	9.1	38
101	How To Engineer Ionic Liquids Resistant Enzymes: Insights from Combined Molecular Dynamics and Directed Evolution Study. <i>ACS Sustainable Chemistry and Engineering</i> , 2019, 7, 11293-11302.	6.7	38
102	Engineering and emerging applications of artificial metalloenzymes with whole cells. <i>Nature Catalysis</i> , 2021, 4, 814-827.	34.4	38
103	Casting epPCR (cepPCR): A simple random mutagenesis method to generate high quality mutant libraries. <i>Biotechnology and Bioengineering</i> , 2017, 114, 1921-1927.	3.3	36
104	Directed Evolution of an Antitumor Drug (Arginine Deiminase PpADI) for Increased Activity at Physiological pH. <i>ChemBioChem</i> , 2010, 11, 691-697.	2.6	35
105	Extending the substrate scope of a Baeyer-Villiger monooxygenase by multiple-site mutagenesis. <i>Applied Microbiology and Biotechnology</i> , 2014, 98, 4009-4020.	3.6	35
106	Functionalized nanocompartments (Synthosomes): Limitations and prospective applications in industrial biotechnology. <i>Biotechnology Journal</i> , 2006, 1, 795-805.	3.5	34
107	Mediated electron transfer with P450cin. <i>Electrochemistry Communications</i> , 2010, 12, 1547-1550.	4.7	34
108	Nanocompartments with a pH release system based on an engineered OmpF channel protein. <i>Soft Matter</i> , 2011, 7, 532-539.	2.7	34

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109	Polymersome surface decoration by an EGFP fusion protein employing Cecropin A as peptide "anchor". Journal of Biotechnology, 2012, 157, 31-37.	3.8	34
110	Are Directed Evolution Approaches Efficient in Exploring Nature's Potential to Stabilize a Lipase in Organic Cosolvents?. Catalysts, 2017, 7, 142.	3.5	34
111	Directed Evolution of Hyaluronic Acid Synthase from <i>Pasteurella multocida</i> towards High-Molecular-Weight Hyaluronic Acid. ChemBioChem, 2018, 19, 1414-1423.	2.6	34
112	Cloning, expression, and characterization of a thermostable glucoamylase from <i>Thermoanaerobacter tengcongensis</i> MB4. Applied Microbiology and Biotechnology, 2010, 87, 225-233.	3.6	33
113	Design of an activity and stability improved carbonyl reductase from <i>Candida parapsilosis</i> . Journal of Biotechnology, 2013, 165, 52-62.	3.8	33
114	Benzylic hydroxylation of aromatic compounds by P450 BM3. Green Chemistry, 2013, 15, 2408.	9.0	33
115	Artificial Diels-Alderase based on the transmembrane protein FhuA. Beilstein Journal of Organic Chemistry, 2016, 12, 1314-1321.	2.2	33
116	MIXed plastics biodegradation and UPcycling using microbial communities: EU Horizon 2020 project MIX-UP started January 2020. Environmental Sciences Europe, 2021, 33, 99.	5.5	33
117	Rhodium-Complex-Linked Hybrid Biocatalyst: Stereo-Controlled Phenylacetylene Polymerization within an Engineered Protein Cavity. ChemCatChem, 2014, 6, 1229-1235.	3.7	32
118	Matter: A universal immobilization platform for enzymes on polymers, metals, and silicon-based materials. Biotechnology and Bioengineering, 2020, 117, 49-61.	3.3	32
119	Chemoenzymatic route to β -blockers via 3-hydroxy esters. Tetrahedron: Asymmetry, 1996, 7, 2017-2022.	1.8	31
120	A Competitive Flow Cytometry Screening System for Directed Evolution of Therapeutic Enzyme. ACS Synthetic Biology, 2015, 4, 768-775.	3.8	31
121	Water-Soluble Reactive Copolymers Based on Cyclic <i>N</i> -Vinylamides with Succinimide Side Groups for Bioconjugation with Proteins. Macromolecules, 2015, 48, 4256-4268.	4.8	31
122	A bifunctional dermaseptin-thanatin dipeptide functionalizes the crop surface for sustainable pest management. Green Chemistry, 2019, 21, 2316-2325.	9.0	31
123	Rapid and Oriented Immobilization of Laccases on Electrodes via a Methionine-Rich Peptide. ACS Catalysis, 2021, 11, 2445-2453.	11.2	31
124	Enzyme-Polyelectrolyte Complexes Boost the Catalytic Performance of Enzymes. ACS Catalysis, 2018, 8, 10876-10887.	11.2	30
125	A thermostable flavin-based fluorescent protein from <i>Chloroflexus aggregans</i> : a framework for ultra-high resolution structural studies. Photochemical and Photobiological Sciences, 2019, 18, 1793-1805.	2.9	30
126	A nanophosphor-based method for selective DNA recovery in Synthosomes. Biotechnology Journal, 2006, 1, 828-834.	3.5	29

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127	Ultrahigh Throughput Screening System for Directed Glucose Oxidase Evolution in Yeast Cells. <i>Combinatorial Chemistry and High Throughput Screening</i> , 2011, 14, 55-60.	1.1	29
128	Directed arginine deiminase evolution for efficient inhibition of arginine-auxotrophic melanomas. <i>Applied Microbiology and Biotechnology</i> , 2015, 99, 1237-1247.	3.6	29
129	Engineering Enhanced Pore Sizes Using FhuA ¹⁻¹⁶⁰ from <i>E. coli</i> Outer Membrane as Template. <i>ACS Sensors</i> , 2017, 2, 1619-1626.	7.8	29
130	KnowVolution of a GH5 Cellulase from <i>Penicillium verruculosum</i> to Improve Thermal Stability for Biomass Degradation. <i>ACS Sustainable Chemistry and Engineering</i> , 2020, 8, 12388-12399.	6.7	29
131	Challenges of the genetic code for exploring sequence space in directed protein evolution. <i>Biocatalysis and Biotransformation</i> , 2007, 25, 229-241.	2.0	28
132	SeSaM Generates a Protein Sequence Space that is Unobtainable by epPCR. <i>ChemBioChem</i> , 2011, 12, 1595-1601.	2.6	28
133	Iterative key-residues interrogation of a phytase with thermostability increasing substitutions identified in directed evolution. <i>Applied Microbiology and Biotechnology</i> , 2016, 100, 227-242.	3.6	28
134	An Enzymatic Route to Tocopherol Synthons: Aromatic Hydroxylation of Pseudocumene and Mesitylene with P450 BM3. <i>Chemistry - A European Journal</i> , 2017, 23, 17981-17991.	3.3	28
135	Biocatalyst Immobilization by Anchor Peptides on an Additively Manufacturable Material. <i>Organic Process Research and Development</i> , 2019, 23, 1852-1859.	2.7	28
136	CompassR Yields Highly Organic Solvent-Tolerant Enzymes through Recombination of Compatible Substitutions. <i>Chemistry - A European Journal</i> , 2021, 27, 2789-2797.	3.3	28
137	A Potential Antitumor Drug (Arginine Deiminase) Reengineered for Efficient Operation under Physiological Conditions. <i>ChemBioChem</i> , 2010, 11, 2294-2301.	2.6	27
138	Reengineered carbonyl reductase for reducing methyl-substituted cyclohexanones. <i>Protein Engineering, Design and Selection</i> , 2013, 26, 291-298.	2.1	27
139	Grafting PNIPAAm from β -barrel shaped transmembrane nanopores. <i>Biomaterials</i> , 2016, 107, 115-123.	11.4	27
140	Sortase-Mediated High-Throughput Screening Platform for Directed Enzyme Evolution. <i>ACS Combinatorial Science</i> , 2018, 20, 203-211.	3.8	27
141	Exploring the full natural diversity of single amino acid exchange reveals that 40-60% of BSLA positions improve organic solvents resistance. <i>Bioresources and Bioprocessing</i> , 2018, 5, .	4.2	27
142	Toward understanding the inactivation mechanism of monooxygenase P450 BM-3 by organic cosolvents: A molecular dynamics simulation study. <i>Biopolymers</i> , 2006, 83, 467-476.	2.4	26
143	Whole-cell double oxidation of n-heptane. <i>Journal of Biotechnology</i> , 2014, 191, 196-204.	3.8	26
144	Rapid and Robust Coating Method to Render Polydimethylsiloxane Surfaces Cell-Adhesive. <i>ACS Applied Materials & Interfaces</i> , 2019, 11, 41091-41099.	8.0	26

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145	Ultrahigh-throughput screening system for directed polymer binding peptide evolution. <i>Biotechnology and Bioengineering</i> , 2019, 116, 1856-1867.	3.3	26
146	Chemoenzymatic cascade for stilbene production from cinnamic acid catalyzed by ferulic acid decarboxylase and an artificial metatase. <i>Catalysis Science and Technology</i> , 2019, 9, 5572-5576.	4.1	26
147	CompassR-guided recombination unlocks design principles to stabilize lipases in ILs with minimal experimental efforts. <i>Green Chemistry</i> , 2021, 23, 3474-3486.	9.0	26
148	Temperature effects on structure and dynamics of the psychrophilic protease subtilisin S41 and its thermostable mutants in solution. <i>Protein Engineering, Design and Selection</i> , 2011, 24, 533-544.	2.1	25
149	An electrochemical microtiter plate for parallel spectroelectrochemical measurements. <i>Electrochimica Acta</i> , 2013, 89, 98-105.	5.2	25
150	Reactive Copolymers Based on <i>N</i> -Vinyl Lactams with Pyridyl Disulfide Side Groups via RAFT Polymerization and Postmodification via Thiol-Disulfide Exchange Reaction. <i>Macromolecules</i> , 2016, 49, 7141-7154.	4.8	25
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