Stephen G Withers

List of Publications by Year in descending order

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434 papers 24,726 citations

4960 84 h-index 130 g-index

460 all docs

460 docs citations

460 times ranked

13401 citing authors

#	Article	IF	CITATIONS
1	Catalysis by hen egg-white lysozyme proceeds via a covalent intermediate. Nature, 2001, 412, 835-838.	27.8	588
2	Glycosynthases:  Mutant Glycosidases for Oligosaccharide Synthesis. Journal of the American Chemical Society, 1998, 120, 5583-5584.	13.7	513
3	Glycosidase mechanisms. Current Opinion in Chemical Biology, 2000, 4, 573-580.	6.1	447
4	X-ray structures along the reaction pathway of cyclodextrin glycosyltransferase elucidate catalysis in the alpha-amylase family. Nature Structural Biology, 1999, 6, 432-436.	9.7	348
5	The structure of human pancreatic (i> $\hat{i}\pm\langle i\rangle$ -amylase at 1.8 \tilde{A} resolution and comparisons with related enzymes. Protein Science, 1995, 4, 1730-1742.	7.6	333
6	Crystal structure of the retaining galactosyltransferase LgtC from Neisseria meningitidis in complex with donor and acceptor sugar analogs. Nature Structural Biology, 2001, 8, 166-175.	9.7	313
7	Mechanism of Agrobacterium .betaglucosidase: kinetic studies. Biochemistry, 1992, 31, 9961-9969.	2.5	304
8	Mutagenesis of Glycosidases. Annual Review of Biochemistry, 1999, 68, 487-522.	11.1	280
9	The pKaof the General Acid/Base Carboxyl Group of a Glycosidase Cycles during Catalysis:Â A13C-NMR Study ofBacillus circulansXylanaseâ€. Biochemistry, 1996, 35, 9958-9966.	2.5	269
10	Engineering of glycosidases and glycosyltransferases. Current Opinion in Chemical Biology, 2006, 10, 509-519.	6.1	267
11	Snapshots along an Enzymatic Reaction Coordinate: Analysis of a Retaining β-Glycoside Hydrolaseâ€,‡. Biochemistry, 1998, 37, 11707-11713.	2.5	255
12	NAG-thiazoline, An N-Acetyl- \hat{l}^2 -hexosaminidase Inhibitor That Implicates Acetamido Participation. Journal of the American Chemical Society, 1996, 118, 6804-6805.	13.7	248
13	Crystallographic Evidence for Substrate-assisted Catalysis in a Bacterial β-Hexosaminidase. Journal of Biological Chemistry, 2001, 276, 10330-10337.	3.4	239
14	Unequivocal demonstration of the involvement of a glutamate residue as a nucleophile in the mechanism of a retaining glycosidase. Journal of the American Chemical Society, 1990, 112, 5887-5889.	13.7	236
15	Subsite Mapping of the Human Pancreatic α-Amylase Active Site through Structural, Kinetic, and Mutagenesis Techniquesâ€,‡. Biochemistry, 2000, 39, 4778-4791.	2.5	231
16	Hydrogen bonding and catalysis: a novel explanation for how a single amino acid substitution can change the ph optimum of a glycosidase 1 1Edited by M. F. Summers. Journal of Molecular Biology, 2000, 299, 255-279.	4.2	214
17	High-throughput screening methodology for the directed evolution of glycosyltransferases. Nature Methods, 2006, 3, 609-614.	19.0	211
18	Hydrogen bonding and specificity. Fluorodeoxy sugars as probes of hydrogen bonding in the glycogen phosphorylase-glucose complex. Biochemistry, 1986, 25, 6021-6027.	2.5	207

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19	Glycosyl fluorides in enzymatic reactions. Carbohydrate Research, 2000, 327, 27-46.	2.3	207
20	A Structural View of the Action ofEscherichia coli(lacZ) β-Galactosidaseâ€,‡. Biochemistry, 2001, 40, 14781-14794.	2.5	207
21	Structural Insights into the Catalytic Mechanism of Trypanosoma cruzi trans-Sialidase. Structure, 2004, 12, 775-784.	3.3	197
22	Structural analysis of the sialyltransferase Cstll from Campylobacter jejuni in complex with a substrate analog. Nature Structural and Molecular Biology, 2004, 11, 163-170.	8.2	196
23	Trypanosoma cruziTrans-sialidase Operates through a Covalent Sialylâ^'Enzyme Intermediate:Â Tyrosine Is the Catalytic Nucleophile. Journal of the American Chemical Society, 2003, 125, 7532-7533.	13.7	188
24	Pharmacological Enhancement of \hat{l}^2 -Hexosaminidase Activity in Fibroblasts from Adult Tay-Sachs and Sandhoff Patients. Journal of Biological Chemistry, 2004, 279, 13478-13487.	3.4	186
25	Mechanism of Agrobacterium .betaglucosidase: kinetic analysis of the role of noncovalent enzyme/substrate interactions. Biochemistry, 1995, 34, 16194-16202.	2.5	182
26	Sugar Ring Distortion in the Glycosyl-Enzyme Intermediate of a Family G/11 Xylanaseâ€,‡. Biochemistry, 1999, 38, 5346-5354.	2.5	182
27	The Role of Sugar Substituents in Glycoside Hydrolysis. Journal of the American Chemical Society, 2000, 122, 1270-1277.	13.7	175
28	Mechanism-Based Covalent Neuraminidase Inhibitors with Broad-Spectrum Influenza Antiviral Activity. Science, 2013, 340, 71-75.	12.6	175
29	Crystallographic Structure of Human β-Hexosaminidase A: Interpretation of Tay-Sachs Mutations and Loss of GM2 Ganglioside Hydrolysis. Journal of Molecular Biology, 2006, 359, 913-929.	4.2	169
30	Covalent inhibitors of glycosidases and their applications in biochemistry and biology. Glycobiology, 2008, 18, 570-586.	2.5	167
31	Crystal structure of the catalytic domain of the .beta1,4-glycanase Cex from Cellulomonas fimi. Biochemistry, 1994, 33, 12546-12552.	2.5	166
32	The Mechanism of Cellulose Hydrolysis by a Two-Step, Retaining Cellobiohydrolase Elucidated by Structural and Transition Path Sampling Studies. Journal of the American Chemical Society, 2014, 136, 321-329.	13.7	164
33	The Search for Novel Human Pancreatic αâ€Amylase Inhibitors: Highâ€Throughput Screening of Terrestrial and Marine Natural Product Extracts. ChemBioChem, 2008, 9, 433-438.	2.6	163
34	2-Deoxy-2-fluoroglucosides: a novel class of mechanism-based glucosidase inhibitors. Journal of the American Chemical Society, 1987, 109, 7530-7531.	13.7	161
35	Mechanistic and Structural Analysis of a Family 31 α-Glycosidase and Its Glycosyl-enzyme Intermediate. Journal of Biological Chemistry, 2005, 280, 2105-2115.	3.4	156
36	Crystallographic observation of a covalent catalytic intermediate in a \hat{l}^2 -glycosidase. Nature Structural Biology, 1996, 3, 149-154.	9.7	153

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37	The Acid/Base Catalyst in the Exoglucanase/Xylanase from Cellulomonas fimi Is Glutamic Acid 127: Evidence from Detailed Kinetic Studies of Mutants. Biochemistry, 1994, 33, 6371-6376.	2.5	152
38	Ultrahighâ€Throughput FACSâ€Based Screening for Directed Enzyme Evolution. ChemBioChem, 2009, 10, 2704-2715.	2.6	151
39	Inactivation of a .betaglucosidase through the accumulation of a stable 2-deoxy-2-fluoroalphaD-glucopyranosyl-enzyme intermediate: a detailed investigation. Biochemistry, 1992, 31, 9970-9978.	2.5	150
40	Approaches to labeling and identification of active site residues in glycosidases. Protein Science, 1995, 4, 361-372.	7.6	148
41	Dissection of nucleophilic and acid–base catalysis in glycosidases. Current Opinion in Chemical Biology, 2001, 5, 643-649.	6.1	146
42	Thioglycoligases: Mutant Glycosidases for Thioglycoside Synthesis. Angewandte Chemie - International Edition, 2003, 42, 352-354.	13.8	143
43	Crystal Structure of Thermotoga maritima α-l-Fucosidase. Journal of Biological Chemistry, 2004, 279, 13119-13128.	3.4	141
44	Advances in Enzymatic Glycoside Synthesis. ACS Chemical Biology, 2016, 11, 1784-1794.	3.4	140
45	Carbohydrate-active enzymes (CAZymes) in the gut microbiome. Nature Reviews Microbiology, 2022, 20, 542-556.	28.6	139
46	Identification of a covalent .alphaD-glucopyranosyl enzyme intermediate formed on a .betaglucosidase. Journal of the American Chemical Society, 1988, 110, 8551-8553.	13.7	135
47	Emerging methods for the production of homogeneous human glycoproteins. Nature Chemical Biology, 2009, 5, 206-215.	8.0	133
48	Observing cellulose biosynthesis and membrane translocation in crystallo. Nature, 2016, 531, 329-334.	27.8	133
49	Identification of Glutamic Acid 78 as the Active Site Nucleophile in Bacillus subtilis Xylanase Using Electrospray Tandem Mass Spectrometry. Biochemistry, 1994, 33, 7027-7032.	2.5	130
50	Dissecting the Electrostatic Interactions and pH-Dependent Activity of a Family 11 Glycosidaseâ€,‡. Biochemistry, 2001, 40, 10115-10139.	2.5	128
51	Substrate Distortion by a -Mannanase: Snapshots of the Michaelis and Covalent-Intermediate Complexes Suggest a B2,5 Conformation for the Transition State. Angewandte Chemie - International Edition, 2002, 41, 2824-2827.	13.8	127
52	Mechanisms of Cellulases and Xylanases: A Detailed Kinetic Study of the Exobeta1,4-glycanase from Cellulomonas Fimi. Biochemistry, 1994, 33, 6363-6370.	2.5	126
53	Pre-Steady State Kinetic Analysis of an Enzymatic Reaction Monitored by Time-Resolved Electrospray Ionization Mass Spectrometryâ€. Biochemistry, 1998, 37, 7664-7669.	2.5	126
54	Aspartate 313 in the Streptomyces plicatusHexosaminidase Plays a Critical Role in Substrate-assisted Catalysis by Orienting the 2-Acetamido Group and Stabilizing the Transition State. Journal of Biological Chemistry, 2002, 277, 40055-40065.	3.4	126

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55	Unequivocal Identification of Asp-214 as the Catalytic Nucleophile of Saccharomyces cerevisiae α-Glucosidase Using 5-Fluoro Glycosyl Fluorides. Journal of Biological Chemistry, 1996, 271, 6889-6894.	3.4	124
56	An Unusual Mechanism of Glycoside Hydrolysis Involving Redox and Elimination Steps by a Family 4 β-Glycosidase fromThermotoga maritima. Journal of the American Chemical Society, 2004, 126, 8354-8355.	13.7	119
57	Mechanistic Analyses of Catalysis in Human Pancreatic α-Amylase:  Detailed Kinetic and Structural Studies of Mutants of Three Conserved Carboxylic Acids. Biochemistry, 2002, 41, 4492-4502.	2.5	116
58	Directed Evolution of a Glycosynthase from Agrobacterium sp. Increases Its Catalytic Activity Dramatically and Expands Its Substrate Repertoire. Journal of Biological Chemistry, 2004, 279, 42787-42793.	3.4	116
59	Cloning, Expression, Characterization, and Nucleophile Identification of Family 3, Aspergillus nigerî ² -Glucosidase. Journal of Biological Chemistry, 2000, 275, 4973-4980.	3.4	115
60	5-Fluoro Glycosides: Â A New Class of Mechanism-Based Inhibitors of Both \hat{l}_{\pm} - and \hat{l}^2 -Glucosidases. Journal of the American Chemical Society, 1996, 118, 241-242.	13.7	113
61	The E358S mutant of Agrobacterium sp. \hat{l}^2 -glucosidase is a greatly improved glycosynthase. FEBS Letters, 2000, 466, 40-44.	2.8	113
62	The amylase inhibitor montbretin A reveals a new glycosidase inhibition motif. Nature Chemical Biology, 2015, 11, 691-696.	8.0	113
63	Alternative Catalytic Anions Differentially Modulate Human α-Amylase Activity and Specificity [,] . Biochemistry, 2008, 47, 3332-3344.	2.5	111
64	Mechanism of Action and Identification of Asp242 as the Catalytic Nucleophile of Vibrio furnisii N-Acetyl-Î ² -d-glucosaminidase Using 2-Acetamido-2-deoxy-5-fluoro-α-l-idopyranosyl Fluoride. Biochemistry, 2000, 39, 117-126.	2.5	106
65	Structural insight into mammalian sialyltransferases. Nature Structural and Molecular Biology, 2009, 16, 1186-1188.	8.2	105
66	The Donor Subsite of Trehalose-6-phosphate Synthase. Journal of Biological Chemistry, 2004, 279, 1950-1955.	3.4	104
67	Teaching old enzymes new tricks: engineering and evolution of glycosidases and glycosyl transferases for improved glycoside synthesisThis paper is one of a selection of papers published in this Special Issue, entitled CSBMCB — Systems and Chemical Biology, and has undergone the Journal's usual peer review process Biochemistry and Cell Biology, 2008, 86, 169-177.	2.0	104
68	The Structure of Clostridium perfringens Nanl Sialidase and Its Catalytic Intermediates. Journal of Biological Chemistry, 2008, 283, 9080-9088.	3.4	102
69	\hat{l}^2 -Mannosynthase: Synthesis of \hat{l}^2 -Mannosides with a Mutant \hat{l}^2 -Mannosidase. Angewandte Chemie - International Edition, 2001, 40, 417-420.	13.8	101
70	High-Throughput Screening for Human Lysosomal \hat{l}^2 -N-Acetyl Hexosaminidase Inhibitors Acting as Pharmacological Chaperones. Chemistry and Biology, 2007, 14, 153-164.	6.0	99
71	Detailed Comparative Analysis of the Catalytic Mechanisms of \hat{l}^2 -N-Acetylglucosaminidases from Families 3 and 20 of Glycoside Hydrolases. Biochemistry, 2005, 44, 12809-12818.	2.5	98
72	Insights into transition state stabilization of the \hat{l}^2 -1,4-glycosidase Cex by covalent intermediate accumulation in active site mutants. Nature Structural Biology, 1998, 5, 812-818.	9.7	97

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73	Mechanistic consequences of mutation of the active site nucleophile Glu 358 in Agrobacterium .betaglucosidase. Biochemistry, 1992, 31, 9979-9985.	2.5	96
74	Mechanistic analogies amongst carbohydrate modifying enzymes. Chemical Communications, 2004, , 2243.	4.1	95
75	Order and Disorder: Differential Structural Impacts of Myricetin and Ethyl Caffeate on Human Amylase, an Antidiabetic Target. Journal of Medicinal Chemistry, 2012, 55, 10177-10186.	6.4	95
76	Mechanism-based Inhibition of Yeast \hat{l}_{\pm} -Glucosidase and Human Pancreatic \hat{l}_{\pm} -Amylase by a New Class of Inhibitors. Journal of Biological Chemistry, 1995, 270, 26778-26781.	3.4	94
77	Exploring the Cellulose/Xylan Specificity of the \hat{l}^2 -1,4-Glycanase Cex from Cellulomonas fimi through Crystallography and Mutation,. Biochemistry, 1998, 37, 4751-4758.	2.5	94
78	Breakdown of oligosaccharides by the process of elimination. Current Opinion in Chemical Biology, 2006, 10, 147-155.	6.1	92
79	The synthesis and hydrolysis of a series of deoxyfluoro-d-glucopyranosyl phosphates. Carbohydrate Research, 1986, 154, 127-144.	2.3	91
80	Insights into the Mechanism of Drosophila melanogaster Golgi \hat{l} ±-Mannosidase II through the Structural Analysis of Covalent Reaction Intermediates. Journal of Biological Chemistry, 2003, 278, 48074-48083.	3.4	91
81	Structural Analysis of the α-2,3-Sialyltransferase Cst-I from Campylobacter jejuni in Apo and Substrate-Analogue Bound Forms,. Biochemistry, 2007, 46, 7196-7204.	2.5	90
82	Intermediate Trapping on a Mutant Retaining α-Galactosyltransferase Identifies an Unexpected Aspartate Residue. Journal of Biological Chemistry, 2004, 279, 28339-28344.	3.4	89
83	Mechanistic Consequences of Mutation of Active Site Carboxylates in a Retaining β-1,4-Glycanase fromCellulomonas fimiâ€. Biochemistry, 1996, 35, 13165-13172.	2.5	88
84	Novel Catalytic Mechanism of Glycoside Hydrolysis Based on the Structure of an NAD+/Mn2+-Dependent Phospho-α-Glucosidase from Bacillus subtilis. Structure, 2004, 12, 1619-1629.	3.3	88
85	Directed evolution of new glycosynthases from Agrobacterium \hat{l}^2 -glucosidase: a general screen to detect enzymes for oligosaccharide synthesis. Chemistry and Biology, 2001, 8, 437-443.	6.0	87
86	Direct 1H N.M.R. determination of the stereochemical course of hydrolyses catalysed by glucanase components of the cellulase complex. Biochemical and Biophysical Research Communications, 1986, 139, 487-494.	2.1	84
87	Direct Observation of the Protonation State of an Imino Sugar Glycosidase Inhibitor upon Binding. Journal of the American Chemical Society, 2003, 125, 7496-7497.	13.7	84
88	Configurationally selective transition state analogue inhibitors of glycosidases. A study with nojiritetrazoles, a new class of glycosidase inhibitors. Carbohydrate Research, 1993, 250, 113-128.	2.3	83
89	Site-Directed Mutation of the Putative Catalytic Residues of Endoglucanase CenA from Cellulomonas fimi. Biochemistry, 1995, 34, 2220-2224.	2.5	83
90	Self″mmobilizing Fluorogenic Imaging Agents of Enzyme Activity. Angewandte Chemie - International Edition, 2011, 50, 300-303.	13.8	81

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91	Solid-Phase Oligosaccharide and Glycopeptide Synthesis Using Glycosynthases. Journal of Organic Chemistry, 2002, 67, 4143-4149.	3.2	79
92	Glycosynthase-Mediated Synthesis of Glycosphingolipids. Journal of the American Chemical Society, 2006, 128, 6300-6301.	13.7	79
93	Crystal structure of the family 7 endoglucanase I (Cel7B) from Humicola insolens at 2.2ÂÃ resolution and identification of the catalytic nucleophile by trapping of the covalent glycosyl-enzyme intermediate. Biochemical Journal, 1998, 335, 409-416.	3.7	77
94	A New Generation of Specific <i>Trypanosoma cruzi trans</i> â€Sialidase Inhibitors. Angewandte Chemie - International Edition, 2008, 47, 2700-2703.	13.8	77
95	Designer enzymes for glycosphingolipid synthesis by directed evolution. Nature Chemical Biology, 2009, 5, 508-514.	8.0	76
96	Trapping and Characterization of the Reaction Intermediate in Cyclodextrin Glycosyltransferase by Use of Activated Substrates and a Mutant Enzymeâ€. Biochemistry, 1997, 36, 9927-9934.	2.5	75
97	Structural and Kinetic Analysis of Two Covalent Sialosyl-Enzyme Intermediates on Trypanosoma rangeli Sialidase. Journal of Biological Chemistry, 2006, 281, 4149-4155.	3.4	75
98	Identification of the Active Site Nucleophile in Jack Bean α-Mannosidase Using 5-Fluoro-β-l-Gulosyl Fluoride. Journal of Biological Chemistry, 1998, 273, 2067-2072.	3.4	74
99	Acarbose Rearrangement Mechanism Implied by the Kinetic and Structural Analysis of Human Pancreatic α-Amylase in Complex with Analogues and Their Elongated Counterpartsâ€,‡. Biochemistry, 2005, 44, 3347-3357.	2.5	74
100	Rapid Assembly of a Library of Lipophilic Iminosugars via the Thiol–Ene Reaction Yields Promising Pharmacological Chaperones for the Treatment of Gaucher Disease. Journal of Medicinal Chemistry, 2012, 55, 2737-2745.	6.4	74
101	Positioning the Acid/Base Catalyst in a Glycosidase: Studies withBacillus circulansXylanaseâ€. Biochemistry, 1997, 36, 2257-2265.	2.5	73
102	Active-site Peptide "Fingerprinting―of Glycosidases in Complex Mixtures by Mass Spectrometry. Journal of Biological Chemistry, 2005, 280, 35126-35135.	3.4	73
103	Expanding the Thioglycoligase Strategy to the Synthesis of α-Linked Thioglycosides Allows Structural Investigation of the Parent Enzyme/Substrate Complex. Journal of the American Chemical Society, 2006, 128, 2202-2203.	13.7	72
104	Identification of the Catalytic Nucleophile of the Family 29 \hat{I}_{\pm} -L-Fucosidase from Thermotoga maritima through Trapping of a Covalent Glycosyl-Enzyme Intermediate and Mutagenesis. Journal of Biological Chemistry, 2003, 278, 47394-47399.	3.4	70
105	Subsite structure of the endo-type chitin deacetylase from a Deuteromycete, Colletotrichum lindemuthianum: an investigation using steady-state kinetic analysis and MS. Biochemical Journal, 2003, 374, 369-380.	3.7	70
106	N-Acetylglucosaminidases from CAZy Family GH3 Are Really Glycoside Phosphorylases, Thereby Explaining Their Use of Histidine as an Acid/Base Catalyst in Place of Glutamic Acid. Journal of Biological Chemistry, 2015, 290, 4887-4895.	3.4	70
107	Crystal Structure of \hat{l}^2 -d-Xylosidase from Thermoanaerobacterium saccharolyticum, a Family 39 Glycoside Hydrolase. Journal of Molecular Biology, 2004, 335, 155-165.	4.2	69
108	Detailed Structural Analysis of Glycosidase/Inhibitor Interactions: Complexes of Cex fromCellulomonas fimiwith Xylobiose-Derived Aza-Sugarsâ€,‡. Biochemistry, 2000, 39, 11553-11563.	2.5	68

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109	Elucidation of the Mechanism of Polysaccharide Cleavage by Chondroitin AC Lyase fromFlavobacteriumheparinum. Journal of the American Chemical Society, 2002, 124, 9756-9767.	13.7	68
110	Using substrate engineering to harness enzymatic promiscuity and expand biological catalysis., 2006, 2, 724-728.		68
111	Anatomy of Glycosynthesis. Chemistry and Biology, 2003, 10, 619-628.	6.0	67
112	Unusual Enzymatic Glycoside Cleavage Mechanisms. Accounts of Chemical Research, 2014, 47, 226-235.	15.6	67
113	Expansion of the glycosynthase repertoire to produce defined manno-oligosaccharides. Chemical Communications, 2003, , 1327-1329.	4.1	66
114	The purification and characterization of a \hat{l}^2 -glucosidase from Alcaligenes faecalis. Biochemistry and Cell Biology, 1986, 64, 914-922.	2.0	65
115	Enzymatic Synthesis of Carbonâ^'Fluorine Bonds. Journal of the American Chemical Society, 2001, 123, 4350-4351.	13.7	64
116	Engineering of a thioglycoligase: randomized mutagenesis of the acid-base residue leads to the identification of improved catalysts. Protein Engineering, Design and Selection, 2005, 18, 33-40.	2.1	62
117	Structure of human ST8SiallI sialyltransferase provides insight into cell-surface polysialylation. Nature Structural and Molecular Biology, 2015, 22, 627-635.	8.2	62
118	The synthesis and hydrolysis of a series of deoxy- and deoxyfluoro-α-d- "glucopyranosyl―phosphates. Carbohydrate Research, 1989, 187, 43-66.	2.3	61
119	Effects of both Shortening and Lengthening the Active Site Nucleophile ofBacillus circulansXylanase on Catalytic Activityâ€. Biochemistry, 1996, 35, 10110-10118.	2.5	61
120	Nanomolar versus Millimolar Inhibition by Xylobiose-Derived Azasugars:  Significant Differences between Two Structurally Distinct Xylanases. Journal of the American Chemical Society, 2000, 122, 2223-2235.	13.7	61
121	Mechanistic Studies of a Retaining α-Galactosyltransferase fromNeisseria meningitidisâ€. Biochemistry, 2002, 41, 5075-5085.	2.5	60
122	Characterization of a beta-N-acetylhexosaminidase and a beta-N-acetylglucosaminidase/beta-glucosidase from Cellulomonas fimi. FEBS Journal, 2006, 273, 2929-2941.	4.7	60
123	Assignment of Sweet Almond \hat{l}^2 -Glucosidase as a Family 1 Glycosidase and Identification of Its Active Site Nucleophile. Journal of Biological Chemistry, 1997, 272, 24864-24867.	3.4	59
124	Detailed Dissection of a New Mechanism for Glycoside Cleavage: α-1,4-Glucan Lyaseâ€. Biochemistry, 2003, 42, 13081-13090.	2.5	59
125	Reassessment of the catalytic mechanism of glycogen debranching enzyme. Biochemistry, 1991, 30, 1419-1424.	2.5	58
126	Identification of Asp-130 as the Catalytic Nucleophile in the Main α-Galactosidase from Phanerochaete chrysosporium, a Family 27 Glycosyl Hydrolase. Biochemistry, 2000, 39, 9826-9836.	2.5	58

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127	Glycosynthase-based synthesis of xylo-oligosaccharides using an engineered retaining xylanase from Cellulomonas fimi. Organic and Biomolecular Chemistry, 2006, 4, 2025.	2.8	58
128	Fluorescence Activated Cell Sorting as a General Ultra-High-Throughput Screening Method for Directed Evolution of Glycosyltransferases. Journal of the American Chemical Society, 2010, 132, 10570-10577.	13.7	58
129	Directed evolution of an $\hat{l}\pm 1,3$ -fucosyltransferase using a single-cell ultrahigh-throughput screening method. Science Advances, 2019, 5, eaaw8451.	10.3	58
130	The Crystal Structure of a 2-Fluorocellotriosyl Complex of theStreptomyceslividansEndoglucanase CelB2 at 1.2 à Resolutionâ€,‡. Biochemistry, 1999, 38, 4826-4833.	2.5	56
131	Thioglycosynthases: double mutant glycosidases that serve as scaffolds for thioglycoside synthesis. Chemical Communications, 2004, , 274-275.	4.1	56
132	An enzymatic pathway in the human gut microbiome that converts A to universal O type blood. Nature Microbiology, 2019, 4, 1475-1485.	13.3	56
133	Mechanism, Mutagenesis, and Chemical Rescue of a β-Mannosidase fromCellulomonas fimiâ€. Biochemistry, 2003, 42, 7195-7204.	2.5	55
134	Kinetic and Mechanistic Analysis of Trypanosoma cruzi Trans-Sialidase Reveals a Classical Ping-Pong Mechanism with Acid/Base Catalysis. Biochemistry, 2008, 47, 3507-3512.	2.5	55
135	A New, Simple, High-Affinity Glycosidase Inhibitor:  Analysis of Binding through X-ray Crystallography, Mutagenesis, and Kinetic Analysis. Journal of the American Chemical Society, 2000, 122, 4229-4230.	13.7	54
136	Trapping Covalent Intermediates on Î ² -Glycosidases. Methods in Enzymology, 2002, 354, 84-105.	1.0	54
137	Syntheses of 2-deoxy-2-fluoro mono- and oligo-saccharide glycosides from glycals and evaluation as glycosidase inhibitors. Carbohydrate Research, 1993, 249, 77-90.	2.3	53
138	Identification of Glu-330 as the Catalytic Nucleophile of Candida albicans Exo- \hat{l}^2 -(1,3)-glucanase. Journal of Biological Chemistry, 1997, 272, 3161-3167.	3.4	53
139	Cloning, mutagenesis, and structural analysis of human pancreatic αâ€amylase expressed in pichia pastoris. Protein Science, 1999, 8, 635-643.	7.6	53
140	Toward Efficient Enzymes for the Generation of Universal Blood through Structure-Guided Directed Evolution. Journal of the American Chemical Society, 2015, 137, 5695-5705.	13.7	53
141	A general and efficient strategy for generating the stable enzymes. Scientific Reports, 2016, 6, 33797.	3. 3	53
142	NAD+ and Metal-ion Dependent Hydrolysis by Family 4 Glycosidases: Structural Insight into Specificity for Phospho-Î ² -d-glucosides. Journal of Molecular Biology, 2005, 346, 423-435.	4.2	52
143	New approaches to enzymatic glycoside synthesis through directed evolution. Carbohydrate Research, 2010, 345, 1272-1279.	2.3	52
144	D-Gluconhydroximo-1,5-lactam and RelatedN-Arylcarbamates Theoretical Calculations, Structure, Synthesis, and Inhibitory Effect on ?-Glucosidases. Helvetica Chimica Acta, 1993, 76, 2666-2686.	1.6	51

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145	Structural and mechanistic studies of chloride induced activation of human pancreatic Â-amylase. Protein Science, 2005, 14, 743-755.	7.6	51
146	Development of New and Selective <i>Trypanosoma cruzi</i> transâ€Sialidase Inhibitors from Sulfonamide Chalcones and Their Derivatives. ChemBioChem, 2009, 10, 2475-2479.	2.6	51
147	Enhancement of biological reactions on cell surfaces via macromolecular crowding. Nature Communications, 2014, 5, 4683.	12.8	51
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S2-6 Identifying and improving glycosidases through metagenomics and directed evolution (Recent) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5