

# Motomitsu Kitaoka

## List of Publications by Citations

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#	Paper	IF	Citations
209	Physiology of consumption of human milk oligosaccharides by infant gut-associated bifidobacteria. <i>Journal of Biological Chemistry</i> , <b>2011</b> , 286, 34583-92	5.4	278
208	Bifidobacterium bifidum lacto-N-biosidase, a critical enzyme for the degradation of human milk oligosaccharides with a type 1 structure. <i>Applied and Environmental Microbiology</i> , <b>2008</b> , 74, 3996-4004	4.8	176
207	Novel putative galactose operon involving lacto-N-biose phosphorylase in Bifidobacterium longum. <i>Applied and Environmental Microbiology</i> , <b>2005</b> , 71, 3158-62	4.8	169
206	Carbohydrate-Processing Phosphorolytic Enzymes.. <i>Trends in Glycoscience and Glycotechnology</i> , <b>2002</b> , 14, 35-50	0.1	153
205	Identification of N-acetylhexosamine 1-kinase in the complete lacto-N-biose I/galacto-N-biose metabolic pathway in Bifidobacterium longum. <i>Applied and Environmental Microbiology</i> , <b>2007</b> , 73, 6444-9	4.8	149
204	Practical preparation of lacto-N-biose I, a candidate for the bifidus factor in human milk. <i>Bioscience, Biotechnology and Biochemistry</i> , <b>2007</b> , 71, 2101-4	2.1	126
203	Recent development of phosphorylases possessing large potential for oligosaccharide synthesis. <i>Current Opinion in Chemical Biology</i> , <b>2013</b> , 17, 301-9	9.7	102
202	Bifidobacterium longum subsp. infantis uses two different $\beta$ -galactosidases for selectively degrading type-1 and type-2 human milk oligosaccharides. <i>Glycobiology</i> , <b>2012</b> , 22, 361-8	5.8	96
201	Distribution of in vitro fermentation ability of lacto-N-biose I, a major building block of human milk oligosaccharides, in bifidobacterial strains. <i>Applied and Environmental Microbiology</i> , <b>2010</b> , 76, 54-9	4.8	95
200	Structural and thermodynamic analyses of solute-binding Protein from Bifidobacterium longum specific for core 1 disaccharide and lacto-N-biose I. <i>Journal of Biological Chemistry</i> , <b>2008</b> , 283, 13165-73	5.4	92
199	Synthesis of highly ordered cellulose II in vitro using cellodextrin phosphorylase. <i>Carbohydrate Research</i> , <b>2009</b> , 344, 2468-73	2.9	91
198	1,2- $\alpha$ -L-Fucosynthase: a glycosynthase derived from an inverting $\alpha$ -glycosidase with an unusual reaction mechanism. <i>FEBS Letters</i> , <b>2008</b> , 582, 3739-43	3.8	88
197	Cooperation of $\beta$ -galactosidase and N-acetylhexosaminidase from bifidobacteria in assimilation of human milk oligosaccharides with type 2 structure. <i>Glycobiology</i> , <b>2010</b> , 20, 1402-9	5.8	87
196	Role of a PA14 domain in determining substrate specificity of a glycoside hydrolase family 3 $\beta$ -glucosidase from Kluyveromyces marxianus. <i>Biochemical Journal</i> , <b>2010</b> , 431, 39-49	3.8	83
195	Chitobiose phosphorylase from Vibrio proteolyticus, a member of glycosyl transferase family 36, has a clan GH-L-like ( $\alpha/\alpha$ )(6) barrel fold. <i>Structure</i> , <b>2004</b> , 12, 937-47	5.2	83
194	Bifidobacterium bifidum Lacto- N -Biosidase, a Critical Enzyme for the Degradation of Human Milk Oligosaccharides with a Type 1 Structure. <i>Applied and Environmental Microbiology</i> , <b>2009</b> , 75, 6414-6414	4.8	78
193	Sharing of human milk oligosaccharides degradants within bifidobacterial communities in faecal cultures supplemented with Bifidobacterium bifidum. <i>Scientific Reports</i> , <b>2018</b> , 8, 13958	4.9	78

192	A family 8 glycoside hydrolase from <i>Bacillus halodurans</i> C-125 (BH2105) is a reducing end xylose-releasing exo-oligoxyranase. <i>Journal of Biological Chemistry</i> , <b>2004</b> , 279, 55097-103	5.4	77
191	The first glycosynthase derived from an inverting glycoside hydrolase. <i>Journal of Biological Chemistry</i> , <b>2006</b> , 281, 1426-31	5.4	76
190	EN-acetylgalactosaminidase from infant-associated bifidobacteria belonging to novel glycoside hydrolase family 129 is implicated in alternative mucin degradation pathway. <i>Journal of Biological Chemistry</i> , <b>2012</b> , 287, 693-700	5.4	73
189	Bifidobacterial enzymes involved in the metabolism of human milk oligosaccharides. <i>Advances in Nutrition</i> , <b>2012</b> , 3, 422S-9S	10	70
188	Evolutionary adaptation in fucosyllactose uptake systems supports bifidobacteria-infant symbiosis. <i>Science Advances</i> , <b>2019</b> , 5, eaaw7696	14.3	68
187	1,3-1,4- $\beta$ -fucosynthase that specifically introduces Lewis a/x antigens into type-1/2 chains. <i>Journal of Biological Chemistry</i> , <b>2012</b> , 287, 16709-19	5.4	65
186	A chemoenzymatic route to N-acetylglucosamine-1-phosphate analogues: substrate specificity investigations of N-acetylhexosamine 1-kinase. <i>Chemical Communications</i> , <b>2009</b> , 2944-6	5.8	65
185	Structural dissection of the reaction mechanism of cellobiose phosphorylase. <i>Biochemical Journal</i> , <b>2006</b> , 398, 37-43	3.8	64
184	Molecular Insight into Evolution of Symbiosis between Breast-Fed Infants and a Member of the Human Gut Microbiome <i>Bifidobacterium longum</i> . <i>Cell Chemical Biology</i> , <b>2017</b> , 24, 515-524.e5	8.2	62
183	One-step random mutagenesis by error-prone rolling circle amplification. <i>Nucleic Acids Research</i> , <b>2004</b> , 32, e145	20.1	62
182	Lacto-N-biosidase encoded by a novel gene of <i>Bifidobacterium longum</i> subspecies <i>longum</i> shows unique substrate specificity and requires a designated chaperone for its active expression. <i>Journal of Biological Chemistry</i> , <b>2013</b> , 288, 25194-25206	5.4	61
181	Discovery of $\beta$ 1,4-D-mannosyl-N-acetyl-D-glucosamine phosphorylase involved in the metabolism of N-glycans. <i>Journal of Biological Chemistry</i> , <b>2013</b> , 288, 27366-27374	5.4	57
180	One-pot enzymatic production of beta-D-galactopyranosyl-(1 $\rightarrow$ 3)-2-acetamido-2-deoxy-D-galactose (galacto-N-biose) from sucrose and 2-acetamido-2-deoxy-D-galactose (N-acetylgalactosamine). <i>Carbohydrate Research</i> , <b>2009</b> , 344, 2573-6	2.9	57
179	Transglycosylation of naringin by <i>Bacillus stearothermophilus</i> Maltogenic amylase to give glycosylated naringin. <i>Journal of Agricultural and Food Chemistry</i> , <b>1999</b> , 47, 3669-74	5.7	54
178	Adsorption of Bisphenol A by Cross-Linked $\beta$ Cyclodextrin Polymer. <i>Journal of Inclusion Phenomena and Macrocyclic Chemistry</i> , <b>2002</b> , 44, 429-431		53
177	Structural basis for the specificity of the reducing end xylose-releasing exo-oligoxyranase from <i>Bacillus halodurans</i> C-125. <i>Journal of Biological Chemistry</i> , <b>2005</b> , 280, 17180-6	5.4	53
176	Characterization of a hyperthermostable glycogen phosphorylase from <i>Aquifex aeolicus</i> expressed in <i>Escherichia coli</i> . <i>Journal of Molecular Catalysis B: Enzymatic</i> , <b>2003</b> , 22, 173-180		50
175	Prebiotic effect of lacto-N-biose I on bifidobacterial growth. <i>Bioscience, Biotechnology and Biochemistry</i> , <b>2009</b> , 73, 1175-9	2.1	49

174	Varied Pathways of Infant Gut-Associated to Assimilate Human Milk Oligosaccharides: Prevalence of the Gene Set and Its Correlation with Bifidobacteria-Rich Microbiota Formation. <i>Nutrients</i> , <b>2019</b> , 12,	6.7	49
173	Alternative strategy for converting an inverting glycoside hydrolase into a glycosynthase. <i>Glycobiology</i> , <b>2008</b> , 18, 325-30	5.8	48
172	Characterization of a thermostable family 10 endo-xylanase (XynB) from <i>Thermotoga maritima</i> that cleaves p-nitrophenyl-β-D-xyloside. <i>Journal of Bioscience and Bioengineering</i> , <b>2001</b> , 92, 423-428	3.3	48
171	Hydrolysis of beta-1,3/1,6-glucan by glycoside hydrolase family 16 endo-1,3(4)-beta-glucanase from the basidiomycete <i>Phanerochaete chrysosporium</i> . <i>Applied Microbiology and Biotechnology</i> , <b>2006</b> , 71, 898-906	5.7	47
170	Kinetics of substrate transglycosylation by glycoside hydrolase family 3 glucan (1→3)-beta-glucosidase from the white-rot fungus <i>Phanerochaete chrysosporium</i> . <i>Carbohydrate Research</i> , <b>2004</b> , 339, 2851-7	2.9	47
169	Fusion of family 2b carbohydrate-binding module increases the catalytic activity of a xylanase from <i>Thermotoga maritima</i> to soluble xylan. <i>FEBS Letters</i> , <b>2003</b> , 549, 147-51	3.8	47
168	Diversity and similarity of microbial communities in petroleum crude oils produced in Asia. <i>Bioscience, Biotechnology and Biochemistry</i> , <b>2008</b> , 72, 2831-9	2.1	44
167	A cycloamylose-forming hyperthermostable 4-β-glucanotransferase of <i>Aquifex aeolicus</i> expressed in <i>Escherichia coli</i> . <i>Journal of Molecular Catalysis B: Enzymatic</i> , <b>2003</b> , 22, 45-53		44
166	Synthetic reaction of <i>Cellvibrio gilvus</i> cellobiose phosphorylase. <i>Journal of Biochemistry</i> , <b>1992</b> , 112, 40-43	3.1	44
165	Reaction mechanism of chitobiose phosphorylase from <i>Vibrio proteolyticus</i> : identification of family 36 glycosyltransferase in <i>Vibrio</i> . <i>Biochemical Journal</i> , <b>2004</b> , 377, 225-32	3.8	42
164	Phosphorolytic Reaction of <i>Cellvibrio gilvus</i> Cellobiose Phosphorylase. <i>Bioscience, Biotechnology and Biochemistry</i> , <b>1992</b> , 56, 652-5	2.1	42
163	Kinetic studies of a recombinant cellobiose phosphorylase (CBP) of the <i>Clostridium thermocellum</i> YM4 strain expressed in <i>Escherichia coli</i> . <i>Journal of Biochemistry</i> , <b>2002</b> , 132, 197-203	3.1	41
162	Crystal structure of glycoside hydrolase family 55 {beta}-1,3-glucanase from the basidiomycete <i>Phanerochaete chrysosporium</i> . <i>Journal of Biological Chemistry</i> , <b>2009</b> , 284, 10100-9	5.4	40
161	Crystallographic and mutational analyses of substrate recognition of endo-α-N-acetylgalactosaminidase from <i>Bifidobacterium longum</i> . <i>Journal of Biochemistry</i> , <b>2009</b> , 146, 389-98	3.1	40
160	RAISE: a simple and novel method of generating random insertion and deletion mutations. <i>Nucleic Acids Research</i> , <b>2006</b> , 34, e30	20.1	39
159	Purification and properties of laminaribiose phosphorylase (EC 2.4.1.31) from <i>Euglena gracilis</i> Z. <i>Archives of Biochemistry and Biophysics</i> , <b>1993</b> , 304, 508-14	4.1	39
158	Identification of the putative proton donor residue of lacto-N-biose phosphorylase (EC 2.4.1.211). <i>Bioscience, Biotechnology and Biochemistry</i> , <b>2007</b> , 71, 1587-91	2.1	38
157	Fusion of family VI cellulose binding domains to <i>Bacillus halodurans</i> xylanase increases its catalytic activity and substrate-binding capacity to insoluble xylan. <i>Journal of Molecular Catalysis B: Enzymatic</i> , <b>2003</b> , 21, 221-230		38

156	Discovery of nigerose phosphorylase from <i>Clostridium phytofermentans</i> . <i>Applied Microbiology and Biotechnology</i> , <b>2012</b> , 93, 1513-22	5.7	37
155	The crystal structure of galacto-N-biose/lacto-N-biose I phosphorylase: a large deformation of a TIM barrel scaffold. <i>Journal of Biological Chemistry</i> , <b>2009</b> , 284, 7273-83	5.4	37
154	General function of N-terminal propeptide on assisting protein folding and inhibiting catalytic activity based on observations with a chimeric thermolysin-like protease. <i>Biochemical and Biophysical Research Communications</i> , <b>2003</b> , 301, 1093-8	3.4	37
153	Enzymatic synthesis of a library of beta-(1-->4) hetero- D-glucose and D-xylose-based oligosaccharides employing cellodextrin phosphorylase. <i>Carbohydrate Research</i> , <b>2003</b> , 338, 1981-90	2.9	36
152	Characterization of a cellobiose phosphorylase from a hyperthermophilic eubacterium, <i>Thermotoga maritima</i> MSB8. <i>Bioscience, Biotechnology and Biochemistry</i> , <b>2002</b> , 66, 2578-86	2.1	36
151	Characterization of three beta-galactoside phosphorylases from <i>Clostridium phytofermentans</i> : discovery of d-galactosyl-beta1->4-l-rhamnose phosphorylase. <i>Journal of Biological Chemistry</i> , <b>2009</b> , 284, 19220-7	5.4	35
150	Characterization of glycosynthase mutants derived from glycoside hydrolase family 10 xylanases. <i>Bioscience, Biotechnology and Biochemistry</i> , <b>2006</b> , 70, 1210-7	2.1	35
149	Substrate specificity of N-acetylhexosamine kinase towards N-acetylgalactosamine derivatives. <i>Bioorganic and Medicinal Chemistry Letters</i> , <b>2009</b> , 19, 5433-5	2.9	34
148	Characterization of a bacterial laminaribiose phosphorylase. <i>Bioscience, Biotechnology and Biochemistry</i> , <b>2012</b> , 76, 343-8	2.1	33
147	Practical preparation of D-galactosyl-beta1-->4-L-rhamnose employing the combined action of phosphorylases. <i>Bioscience, Biotechnology and Biochemistry</i> , <b>2010</b> , 74, 1652-5	2.1	33
146	Purification, crystallization and preliminary X-ray analysis of the galacto-N-biose-/lacto-N-biose I-binding protein (GL-BP) of the ABC transporter from <i>Bifidobacterium longum</i> JCM1217. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , <b>2007</b> , 63, 751-3		32
145	Diversity of phosphorylases in glycoside hydrolase families. <i>Applied Microbiology and Biotechnology</i> , <b>2015</b> , 99, 8377-90	5.7	31
144	Distinct substrate specificities of three glycoside hydrolase family 42 $\beta$ -galactosidases from <i>Bifidobacterium longum</i> subsp. <i>infantis</i> ATCC 15697. <i>Glycobiology</i> , <b>2014</b> , 24, 208-16	5.8	31
143	1,2- $\beta$ -Oligoglucan phosphorylase from <i>Listeria innocua</i> . <i>PLoS ONE</i> , <b>2014</b> , 9, e92353	3.7	31
142	Crystal structure of an Exo-1,5- $\alpha$ -L-arabinofuranosidase from <i>Streptomyces avermitilis</i> provides insights into the mechanism of substrate discrimination between exo- and endo-type enzymes in glycoside hydrolase family 43. <i>Journal of Biological Chemistry</i> , <b>2010</b> , 285, 34134-43	5.4	31
141	Error-prone rolling circle amplification: the simplest random mutagenesis protocol. <i>Nature Protocols</i> , <b>2006</b> , 1, 2493-7	18.8	31
140	Cellodextrin Phosphorylase from <i>Clostridium thermocellum</i> YM4 Strain Expressed in <i>Escherichia coli</i> . <i>Journal of Applied Glycoscience (1999)</i> , <b>2002</b> , 49, 1-8	1	30
139	Synthesis of cellobiose from starch by the successive actions of two phosphorylases. <i>New Biotechnology</i> , <b>2009</b> , 26, 137-42	6.4	28

138	2-O- $\beta$ -glucosylglycerol phosphorylase from <i>Bacillus selenitireducens</i> MLS10 possessing hydrolytic activity on $\beta$ -glucose 1-phosphate. <i>PLoS ONE</i> , <b>2014</b> , 9, e86548	3.7	28
137	Introduction of H-antigens into oligosaccharides and sugar chains of glycoproteins using highly efficient 1,2- $\beta$ -fucosyltransferase. <i>Glycobiology</i> , <b>2016</b> , 26, 1235-1247	5.8	27
136	Large-scale Preparation of 1,2- $\beta$ -Glucan Using 1,2- $\beta$ -Oligoglucan Phosphorylase. <i>Journal of Applied Glycoscience (1999)</i> , <b>2015</b> , 62, 47-52	1	27
135	Identification of galacto-N-biose phosphorylase from <i>Clostridium perfringens</i> ATCC13124. <i>Applied Microbiology and Biotechnology</i> , <b>2008</b> , 78, 465-71	5.7	27
134	Discovery of two $\beta$ 1,2-mannoside phosphorylases showing different chain-length specificities from <i>Thermoanaerobacter</i> sp. X-514. <i>PLoS ONE</i> , <b>2014</b> , 9, e114882	3.7	27
133	Functional and Structural Analysis of a $\beta$ -Glucosidase Involved in $\beta$ 1,2-Glucan Metabolism in <i>Listeria innocua</i> . <i>PLoS ONE</i> , <b>2016</b> , 11, e0148870	3.7	27
132	A $\beta$ 1-6/ $\beta$ 1-3 galactosidase from <i>Bifidobacterium animalis</i> subsp. <i>lactis</i> BI-04 gives insight into sub-specificities of $\beta$ -galactoside catabolism within <i>Bifidobacterium</i> . <i>Molecular Microbiology</i> , <b>2014</b> , 94, 1024	4.1	26
131	In vitro comparative evaluation of the impact of lacto-N-biose I, a major building block of human milk oligosaccharides, on the fecal microbiota of infants. <i>Anaerobe</i> , <b>2013</b> , 19, 50-7	2.8	26
130	Conversion of Sucrose into Cellobiose Using Sucrose Phosphorylase, Xylose Isomerase and Cellobiose Phosphorylase.. <i>Journal of the Japanese Society of Starch Science</i> , <b>1992</b> , 39, 281-283		26
129	Discovery of cellobionic acid phosphorylase in cellulolytic bacteria and fungi. <i>FEBS Letters</i> , <b>2013</b> , 587, 3556-61	3.8	25
128	Characterization of a laminaribiose phosphorylase from <i>Acholeplasma laidlawii</i> PG-8A and production of 1,3- $\beta$ -glucosyl disaccharides. <i>Carbohydrate Research</i> , <b>2012</b> , 361, 49-54	2.9	25
127	A cellobiose phosphorylase from <i>Cellvibrio gilvus</i> recognizes only the beta-D-form of 5a-carba-glucopyranose. <i>Carbohydrate Research</i> , <b>1993</b> , 247, 355-9	2.9	25
126	Characterization of a thermostable family 10 endo-xylanase (XynB) from <i>Thermotoga maritima</i> that cleaves p-nitrophenyl-beta-D-xyloside. <i>Journal of Bioscience and Bioengineering</i> , <b>2001</b> , 92, 423-8	3.3	25
125	Facile enzymatic synthesis of sugar 1-phosphates as substrates for phosphorylases using anomeric kinases. <i>Carbohydrate Research</i> , <b>2015</b> , 401, 1-4	2.9	24
124	A glycosynthase derived from an inverting GH19 chitinase from the moss <i>Bryum coronatum</i> . <i>Biochemical Journal</i> , <b>2012</b> , 444, 437-43	3.8	24
123	3-O- $\beta$ -glucopyranosyl-L-rhamnose phosphorylase from <i>Clostridium phytofermentans</i> . <i>Carbohydrate Research</i> , <b>2012</b> , 350, 94-7	2.9	24
122	Kinetic Studies on p-Nitrophenyl-cellobioside Hydrolyzing Xylanase from <i>Cellvibrio gilvus</i> . <i>Bioscience, Biotechnology and Biochemistry</i> , <b>1993</b> , 57, 1987-1989	2.1	24
121	Identification of lacto-N-Biose I phosphorylase from <i>Vibrio vulnificus</i> CMCP6. <i>Applied and Environmental Microbiology</i> , <b>2008</b> , 74, 6333-7	4.8	23



120	Use of a Microtiter Plate Screening Method for Obtaining <i>Leuconostoc mesenteroides</i> Mutants Constitutive for Glucansucrase. <i>Enzyme and Microbial Technology</i> , <b>1998</b> , 22, 527-531	3.8	22
119	A reducing-end-acting chitinase from <i>Vibrio proteolyticus</i> belonging to glycoside hydrolase family 19. <i>Applied Microbiology and Biotechnology</i> , <b>2008</b> , 78, 627-34	5.7	22
118	Computational analyses of the conformational itinerary along the reaction pathway of GH94 cellobiose phosphorylase. <i>Carbohydrate Research</i> , <b>2008</b> , 343, 1023-33	2.9	21
117	A thermostable non-xylanolytic alpha-glucuronidase of <i>Thermotoga maritima</i> MSB8. <i>Bioscience, Biotechnology and Biochemistry</i> , <b>2003</b> , 67, 2359-64	2.1	21
116	Kinetic studies on the hydrolysis of N-acetylated and N-deacetylated derivatives of 4-methylumbelliferyl chitobioside by the family 18 chitinases ChiA and ChiB from <i>Serratia marcescens</i> . <i>Journal of Biochemistry</i> , <b>2003</b> , 133, 253-8	3.1	21
115	Characterization of a thermophilic 4-O- $\beta$ -mannosyl-D-glucose phosphorylase from <i>Rhodothermus marinus</i> . <i>Bioscience, Biotechnology and Biochemistry</i> , <b>2014</b> , 78, 263-70	2.1	20
114	Structural explanation for the acquisition of glycosynthase activity. <i>Journal of Biochemistry</i> , <b>2010</b> , 147, 237-44	3.1	20
113	Thermal decomposition of beta-D-galactopyranosyl-(1 $\rightarrow$ 3)-2-acetamido-2-deoxy-D-hexopyranoses under neutral conditions. <i>Carbohydrate Research</i> , <b>2010</b> , 345, 1901-8	2.9	20
112	The role of the N-terminal propeptide of the pro-aminopeptidase processing protease: refolding, processing, and enzyme inhibition. <i>Biochemical and Biophysical Research Communications</i> , <b>2002</b> , 296, 78-84	3.4	20
111	Conversion of Sucrose into Laminaribiose Using Sucrose Phosphorylase, Xylose Isomerase and Laminaribiose Phosphorylase.. <i>Journal of the Japanese Society of Starch Science</i> , <b>1993</b> , 40, 311-314		20
110	Facile preparation of highly crystalline lamellae of (1 $\rightarrow$ 3)- $\beta$ -D-glucan using an extract of <i>Euglena gracilis</i> . <i>International Journal of Biological Macromolecules</i> , <b>2014</b> , 64, 415-9	7.9	19
109	One Pot Enzymatic Production of Nigerose from Common Sugar Resources Employing Nigerose Phosphorylase. <i>Journal of Applied Glycoscience (1999)</i> , <b>2014</b> , 61, 75-80	1	19
108	Characterization of raffinose synthase from rice ( <i>Oryza sativa</i> L. var. Nipponbare). <i>Biotechnology Letters</i> , <b>2007</b> , 29, 635-40	3	19
107	Enhancement of transglycosylation activity by construction of chimeras between mesophilic and thermophilic beta-glucosidase. <i>Archives of Biochemistry and Biophysics</i> , <b>2002</b> , 407, 125-34	4.1	19
106	Evidence that the putative alpha-glucosidase of <i>Thermotoga maritima</i> MSB8 is a pNP alpha-D-glucuronopyranoside hydrolyzing alpha-glucuronidase. <i>FEBS Letters</i> , <b>2002</b> , 517, 159-62	3.8	19
105	Large-scale preparation of highly purified dextransucrase from a high-producing constitutive mutant of <i>Leuconostoc mesenteroides</i> B-512FMC. <i>Enzyme and Microbial Technology</i> , <b>1998</b> , 23, 386-391	3.8	18
104	Production of glucosyl-xylose using <i>Cellvibrio gilvus</i> cells and its properties. <i>Applied Microbiology and Biotechnology</i> , <b>1990</b> , 34, 178-182	5.7	18
103	Crystal Structure and Substrate Recognition of Cellobionic Acid Phosphorylase, Which Plays a Key Role in Oxidative Cellulose Degradation by Microbes. <i>Journal of Biological Chemistry</i> , <b>2015</b> , 290, 18281-92	5.4	17

102	Kinetic evidence related to substrate-assisted catalysis of family 18 chitinases. <i>FEBS Letters</i> , <b>2004</b> , 567, 307-10	3.8	17
101	Characterization and crystal structure determination of $\beta$ 1,2-mannobiose phosphorylase from <i>Listeria innocua</i> . <i>FEBS Letters</i> , <b>2015</b> , 589, 3816-21	3.8	16
100	Functional reassignment of <i>Cellvibrio vulgaris</i> EpiA to cellobiose 2-epimerase and an evaluation of the biochemical functions of the 4-O- $\beta$ -mannosyl-D-glucose phosphorylase-like protein, UnkA. <i>Bioscience, Biotechnology and Biochemistry</i> , <b>2015</b> , 79, 969-77	2.1	16
99	Characterization of beta-1,3-galactosyl-N-acetylhexosamine phosphorylase from <i>Propionibacterium acnes</i> . <i>Applied Microbiology and Biotechnology</i> , <b>2009</b> , 83, 109-15	5.7	16
98	Identification of <i>Bacillus selenitireducens</i> MLS10 maltose phosphorylase possessing synthetic ability for branched $\beta$ -D-glucosyl trisaccharides. <i>Carbohydrate Research</i> , <b>2012</b> , 360, 25-30	2.9	15
97	A New Method of Carbohydrate Synthesis in Both Solution and Solid Phases Using a Special Hydroxy Protecting Group. <i>European Journal of Organic Chemistry</i> , <b>2005</b> , 2005, 5313-5329	3.2	15
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