Atsushi Nishikawa

List of Publications by Year in descending order

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147566 3,129 119 31 citations h-index papers

51 g-index 122 122 122 2053 docs citations times ranked citing authors all docs

182168

#	Article	IF	CITATIONS
1	Suppression of lung metastasis of B16 mouse melanoma by N-acetylglucosaminyltransferase III gene transfection Proceedings of the National Academy of Sciences of the United States of America, 1995, 92, 8754-8758.	3.3	256
2	Purification and Characterization of UDP-N-Acetylglucosamine: -6-D-Mannoside ß-6N-Acetylglucosaminyltransferase(N-Acetylglucosaminyltransferase V) from a Human Lung Cancer Cell Line1. Journal of Biochemistry, 1993, 113, 614-619.	0.9	147
3	Structure of serum transferrin in carbohydrate-deficient glycoprotein syndrome. Biochemical and Biophysical Research Communications, 1992, 189, 832-836.	1.0	130
4	cDNA Cloning, Expression, and Chromosomal Localization of Human N-Acetylglucosaminyltransferase III (GnT-III)1. Journal of Biochemistry, 1993, 113, 692-698.	0.9	111
5	Determination of N-acetylglucosaminyltransferases III, IV and V in normal and hepatoma tissues of rats. Biochimica Et Biophysica Acta - General Subjects, 1990, 1035, 313-318.	1.1	104
6	CDNA Cloning and Chromosomal Mapping of Human N-Acetylglucosaminyltransferase-V. Biochemical and Biophysical Research Communications, 1994, 198, 318-327.	1.0	104
7	A glycomic approach to the identification and characterization of glycoprotein function in cells transfected with glycosyltransferase genes. Proteomics, 2001, 1, 239-247.	1.3	79
8	[31] Glycosyltransferase assays using pyridylaminated acceptors: N-acetylglucosaminyltransferase III, IV, and V. Methods in Enzymology, 1989, 179, 397-408.	0.4	73
9	Molecular characterization of binding subcomponents of Clostridium botulinum type C progenitor toxin for intestinal epithelial cells and erythrocytes. Microbiology (United Kingdom), 2004, 150, 1529-1538.	0.7	71
10	A Fluorescent Assay Method for GDP-L-Fuc: N-Acetyl-ÂD- Glucosaminide Â1-6Fucosyltransferase Activity, Involving High Performance Liquid Chromatography. Journal of Biochemistry, 1996, 120, 385-392.	0.9	68
11	The receptor and transporter for internalization of Clostridium botulinum type C progenitor toxin into HT-29 cells. Biochemical and Biophysical Research Communications, 2004, 319, 327-333.	1.0	63
12	A method for the determination of N-acetylglucosaminyltransferase III activity in rat tissues involving HPLC. Analytical Biochemistry, 1988, 170, 349-354.	1.1	62
13	Transforming Growth Factor \hat{I}^2 Up-regulates Expression of the N-Acetylglucosaminyltransferase V Gene in Mouse Melanoma Cells. Journal of Biological Chemistry, 1995, 270, 6216-6220.	1.6	59
14	High expression of an N-acetylglucosaminyltransferase III in 3′-methyl DAB-induced hepatoma and ascites hepatoma. Biochemical and Biophysical Research Communications, 1988, 152, 107-112.	1.0	57
15	Remodeling of cell surface glycoproteins by N-acetylglucosaminyltransferase III gene transfection: modulation of metastatic potentials and down regulation of hepatitis B virus replication. Glycobiology, 1996, 6, 691-694.	1.3	56
16	Enzymatic basis of sugar structures of \hat{l} ±-fetoprotein in hepatoma and hepatoblastoma cell lines: Correlation with activities of \hat{l} ±1 \hat{a} €"6 fucosyltransferase and N-acetylglucosaminyltransferases III and V. International Journal of Cancer, 1992, 51, 315-317.	2.3	54
17	N-Acetylglucosaminyltransferase III in human serum, and liver and hepatoma tissues: Increased activity in liver cirrhosis and hepatoma patients. Clinica Chimica Acta, 1989, 185, 325-332.	0.5	52
18	Modulation of N-acetylglucosaminyltransferase III, IV and V activities and alteration of the surface oligosaccharide structure of a myeloma cell line by interleukin 6. Biochemical and Biophysical Research Communications, 1990, 172, 1260-1266.	1.0	50

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19	Ectopic expression of N-acetylglucosaminyltransferase III in transgenic hepatocytes disrupts apolipoprotein B secretion and induces aberrant cellular morphology with lipid storage. Proceedings of the National Academy of Sciences of the United States of America, 1998, 95, 2526-2530.	3.3	48
20	Hydrophobic Man-1-P derivatives correct abnormal glycosylation in Type I congenital disorder of glycosylation fibroblasts. Glycobiology, 2005, 15, 1084-1093.	1.3	48
21	Sugar-binding Sites of the HA1 Subcomponent of Clostridium botulinum Type C Progenitor Toxin. Journal of Molecular Biology, 2008, 376, 854-867.	2.0	48
22	Organization of the Human N-Acetylglucosaminyltransferase V Gene. FEBS Journal, 1995, 233, 18-26.	0.2	44
23	Structural Insights into Substrate Specificity and Function of Glucodextranase. Journal of Biological Chemistry, 2004, 279, 10575-10583.	1.6	43
24	Structural Insights into the Substrate Specificity and Function of Escherichia coli K12 YgjK, a Glucosidase Belonging to the Glycoside Hydrolase Family 63. Journal of Molecular Biology, 2008, 381, 116-128.	2.0	42
25	The Phosphorylation of Bovine DNase I Asn-linked Oligosaccharides Is Dependent on Specific Lysine and Arginine Residues. Journal of Biological Chemistry, 1997, 272, 19408-19412.	1.6	39
26	Gene expression of N-acetylglucosaminyltransferases III and V: A possible implication for liver regeneration. Hepatology, 1995, 22, 1847-1855.	3.6	38
27	Structural Characterization of Chemically Derivatized Oligosaccharides by Nanoflow Electrospray Ionization Mass Spectrometry. Analytical Chemistry, 1999, 71, 4100-4106.	3.2	37
28	Crystal Structure of the HA3 Subcomponent of Clostridium botulinum Type C Progenitor Toxin. Journal of Molecular Biology, 2009, 385, 1193-1206.	2.0	37
29	High expression of udp-n-acetylglucosamine: \hat{l}^2 -d mannoside \hat{l}^2 -1, 4-n-acetylglucosaminyltransferase III (GnT-III) in chronic myelogenous leukemia in blast crisis. International Journal of Cancer, 1995, 60, 443-449.	2.3	35
30	Crystal structure of a lactosucrose-producing enzyme, Arthrobacter sp. K-1 \hat{I}^2 -fructofuranosidase. Enzyme and Microbial Technology, 2012, 51, 359-365.	1.6	33
31	Hypogalactosylation of immunoglobulin G sugar chains and elevated serum interleukin 6 in Castleman's disease. Clinica Chimica Acta, 1991, 197, 221-228.	0.5	32
32	Bisecting GlcNAc Structures Act as Negative Sorting Signals for Cell Surface Glycoproteins in Forskolin-treated Rat Hepatoma Cells. Journal of Biological Chemistry, 1997, 272, 2866-2872.	1.6	32
33	Cell internalization and traffic pathway of Clostridium botulinum type C neurotoxin in HT-29 cells. Biochimica Et Biophysica Acta - Molecular Cell Research, 2006, 1763, 120-128.	1.9	32
34	Carbohydrate recognition mechanism of HA70 from <i>Clostridium botulinum</i> deduced from Xâ€ray structures in complexes with sialylated oligosaccharides. FEBS Letters, 2012, 586, 2404-2410.	1.3	29
35	Crystal structure of a glycoside hydrolase family 6 enzyme, CcCel6C, a cellulase constitutively produced by <i>Coprinopsisâ€∫cinerea</i> . FEBS Journal, 2010, 277, 1532-1542.	2.2	28
36	Molecular diversity of the two sugar-binding sites of the \hat{l}^2 -trefoil lectin HA33/C (HA1) from Clostridium botulinum type C neurotoxin. Archives of Biochemistry and Biophysics, 2011, 512, 69-77.	1.4	27

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37	The efficiency of N-linked glycosylation of bovine DNase I depends on the Asn-Xaa-Ser/Thr sequence and the tissue of origin. Biochemical Journal, 2001, 355, 245-248.	1.7	26
38	Structural and biochemical characterization of novel bacterial \hat{l} ±-galactosidases belonging to glycoside hydrolase family 31. Biochemical Journal, 2015, 469, 145-158.	1.7	26
39	Crystal structure of a β-fructofuranosidase with high transfructosylation activity from <i>Aspergillus kawachii</i> . Bioscience, Biotechnology and Biochemistry, 2017, 81, 1786-1795.	0.6	26
40	Crystal structures of open and closed forms of cyclo/maltodextrinâ€binding protein. FEBS Journal, 2009, 276, 3008-3019.	2,2	25
41	Crystal structure of the catalytic domain of a GH16 \hat{l}^2 -agarase from a deep-sea bacterium, Microbulbifer thermotolerans JAMB-A94. Bioscience, Biotechnology and Biochemistry, 2015, 79, 625-632.	0.6	25
42	Nitric Oxide Synthase from Rat Colorectum: Purification, Peptide Sequencing, Partial PCR Cloning, and Immunohistochemistry1. Journal of Biochemistry, 1994, 115, 602-607.	0.9	24
43	Jaw1/LRMP has a role in maintaining nuclear shape via interaction with SUN proteins. Journal of Biochemistry, 2018, 164, 303-311.	0.9	24
44	Cell Spreading in Colo 201 by Staurosporin Is 3 1 Integrin-mediated with Tyrosine Phosphorylation of Src and Tensin. Journal of Biological Chemistry, 1995, 270, 2298-2304.	1.6	23
45	Binding properties of Clostridium botulinum type C progenitor toxin to mucins. Biochimica Et Biophysica Acta - General Subjects, 2007, 1770, 551-555.	1.1	23
46	Crystal Structure of Aspergillus niger Isopullulanase, a Member of Glycoside Hydrolase Family 49. Journal of Molecular Biology, 2008, 376, 210-220.	2.0	23
47	Human N-Acetylglucosaminyltransferase III Gene is Transcribed from Multiple Promoters. FEBS Journal, 1996, 238, 853-861.	0.2	21
48	The crystal structure of Thermoactinomyces vulgaris R-47 alpha-amylase II (TVA II) complexed with transglycosylated product. FEBS Journal, 2004, 271, 2530-2538.	0.2	21
49	Crystal Structure and Mutational Analysis of Isomalto-dextranase, a Member of Glycoside Hydrolase Family 27. Journal of Biological Chemistry, 2015, 290, 26339-26349.	1.6	21
50	Substrate specificity and distribution of UDP-GalNAc:sialylparagloboside N-acetylgalactosaminyltransferase in the human stomach. Biochemical Journal, 1992, 288, 161-165.	1.7	20
51	Structural basis for cyclodextrin recognition by Thermoactinomyces vulgaris cyclo/maltodextrin-binding protein. FEBS Journal, 2007, 274, 2109-2120.	2.2	20
52	The relative contribution of mannose salvage pathways to glycosylation in PMIâ€deficient mouse embryonic fibroblast cells. FEBS Journal, 2008, 275, 788-798.	2,2	20
53	A glycoside hydrolase family 31 dextranase with high transglucosylation activity from Flavobacterium johnsoniae. Bioscience, Biotechnology and Biochemistry, 2016, 80, 1562-1567.	0.6	20
54	Clinical application of Clostridium botulinum type A neurotoxin purified by a simple procedure for patients with urinary incontinence caused by refractory destrusor overactivity. FEMS Immunology and Medical Microbiology, 2007, 51, 201-211.	2.7	19

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55	Identification of Amino Acids That Modulate Mannose Phosphorylation of Mouse DNase I, a Secretory Glycoprotein. Journal of Biological Chemistry, 1999, 274, 19309-19315.	1.6	18
56	Enhancing thermostability and the structural characterization of Microbacterium saccharophilum K-1 \hat{l}^2 -fructofuranosidase. Applied Microbiology and Biotechnology, 2014, 98, 6667-6677.	1.7	18
57	Melanosome degradation in epidermal keratinocytes related to lysosomal protease cathepsin V. Biochemical and Biophysical Research Communications, 2018, 500, 339-343.	1.0	18
58	The efficiency of N-linked glycosylation of bovine DNase I depends on the Asn-Xaa-Ser/Thr sequence and the tissue of origin. Biochemical Journal, 2001, 355, 245.	1.7	18
59	Comparison of the structural changes in two cellobiohydrolases, CcCel6A and CcCel6C, from <i>Coprinopsisâ€fcinerea</i> â€f–â€fa tweezerâ€like motion in the structure of CcCel6C. FEBS Journal, 2012, 1871-1882.	27.9,	17
60	Structural insights into polysaccharide recognition by <i>FlavobacteriumÂjohnsoniae</i> dextranase, a member of glycoside hydrolase family 31. FEBS Journal, 2020, 287, 1195-1207.	2,2	17
61	Insights into the reaction mechanism of glycosyl hydrolase family 49. Site-directed mutagenesis and substrate preference of isopullulanase. FEBS Journal, 2004, 271, 4420-4427.	0.2	16
62	Heterologous expression and characterization of processing \hat{l}_{\pm} -glucosidase I from Aspergillus brasiliensis ATCC 9642. Glycoconjugate Journal, 2011, 28, 563-571.	1.4	16
63	Purification, Characterization, and Subsite Affinities of Thermoactinomyces vulgaris R-47 Maltooligosaccharide-metabolizing Enzyme Homologous to Glucoamylases. Bioscience, Biotechnology and Biochemistry, 2004, 68, 413-420.	0.6	15
64	Structure of the Catalytic Domain of α-l-Arabinofuranosidase from Coprinopsis cinerea, CcAbf62A, Provides Insights into Structure–Function Relationships in Glycoside Hydrolase Family 62. Applied Biochemistry and Biotechnology, 2017, 181, 511-525.	1.4	14
65	The phosphorylation of sorting nexin 5 at serine 226 regulates retrograde transport and macropinocytosis. PLoS ONE, 2018, 13, e0207205.	1.1	14
66	Small-angle X-ray scattering reveals structural dynamics of the botulinum neurotoxin associating protein, nontoxic nonhemagglutinin. Biochemical and Biophysical Research Communications, 2012, 425, 256-260.	1.0	12
67	Mechanisms mediating metabolic abnormalities in the livers of Ehrlich ascites tumor-bearing mice. Archives of Biochemistry and Biophysics, 2003, 412, 216-222.	1.4	11
68	Construction of an Efficient Expression System for Aspergillus Isopullulanase in Pichia pastoris, and a Simple Purification Method. Bioscience, Biotechnology and Biochemistry, 2003, 67, 1149-1153.	0.6	11
69	Exogenous mannose does not raise steady state mannose-6-phosphate pools of normal or N-glycosylation-deficient human fibroblasts. Molecular Genetics and Metabolism, 2009, 96, 268-272.	0.5	11
70	Genomics of Clostridium botulinum group III strains. Research in Microbiology, 2015, 166, 318-325.	1.0	11
71	Crystal structure of a glycoside hydrolase family $68 \hat{l}^2$ -fructosyltransferase from Beijerinckia indica subsp. indica in complex with fructose. Bioscience, Biotechnology and Biochemistry, 2020, 84, 2508-2520.	0.6	11
72	Effects of dibutyryl cAMP and bromodeoxyuridine on expression of N-acetylglucosaminyltransferases III and V in GOTO neuroblastoma cells. Glycoconjugate Journal, 1995, 12, 787-794.	1.4	10

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73	Gene expression and regulation of N-acetylglucosaminyltransferases III and V in cancer tissues. Advances in Enzyme Regulation, 1998, 38, 223-232.	2.9	10
74	Heterologous Expression, Purification, and Characterization of an α-Mannosidase Belonging to Glycoside Hydrolase Family 99 of <i>Shewanella amazonensis</i> Bioscience, Biotechnology and Biochemistry, 2011, 75, 797-799.	0.6	10
75	Facile construction of 1,2-cis glucosidic linkage using sequential oxidation–reduction route for synthesis of an ER processing l±-glucosidase I substrate. Tetrahedron Letters, 2012, 53, 4452-4456.	0.7	9
76	Preparation and characterization of monoclonal antibodies to an N-linked oligosaccharide. Analytical Biochemistry, 1990, 188, 149-154.	1.1	8
77	Suppression of the xenoantigen GAL $\hat{1}$ +(1,3)Gal by N-Acetylglucosaminyltransferase III (GnT-III) in transgenic mice. Transplantation Proceedings, 1997, 29, 895-896.	0.3	8
78	Sugar-induced conformational change found in the HA-33/HA-17 trimer of the botulinum toxin complex. Biochemical and Biophysical Research Communications, 2013, 438, 483-487.	1.0	8
79	Crystal structure and substrate-binding mode of GH63 mannosylglycerate hydrolase from Thermus thermophilus HB8. Journal of Structural Biology, 2015, 190, 21-30.	1.3	8
80	The side chain of a glycosylated asparagine residue is important for the stability of isopullulanase. Journal of Biochemistry, 2015, 157, 225-234.	0.9	8
81	Gene expression of N-acetylglucosaminyltransferases III and V: A possible implication for liver regeneration*1. Hepatology, 1995, 22, 1847-1855.	3.6	7
82	Molecular cloning and expression of pyruvate kinase from globefish (Fugu rubripes) skeletal muscle. Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2003, 135, 397-405.	0.7	7
83	Cross-Linking of CD45 on Suppressive/Regulatory T Cells Leads to the Abrogation of Their Suppressive Activity In Vitro. Journal of Immunology, 2005, 174, 4090-4097.	0.4	7
84	Structure of a bacterial glycoside hydrolase familyÂ63 enzyme in complex with its glycosynthase product, and insights into the substrate specificity. FEBS Journal, 2013, 280, 4560-4571.	2.2	7
85	A convenient method for the preparation of all of the partially methylated derivatives of methyl $\hat{l}\pm$ -d-mannopyranoside and $\hat{l}\pm$ -d-galactopyranoside. Carbohydrate Research, 1983, 112, 313-319.	1.1	6
86	Deficiency of β1â€6 Nâ€Acetylglucosaminyltransferase Involved in the Biosynthesis of Blood Group I Antigen in the Liver of LEC Rats. Japanese Journal of Cancer Research, 1992, 83, 878-884.	1.7	6
87	Crystal structure of the Nâ€terminal domain of a glycoside hydrolase family 131 protein from <i>Coprinopsis cinerea</i> . FEBS Letters, 2013, 587, 2193-2198.	1.3	6
88	Crystal structure of the enzyme-product complex reveals sugar ring distortion during catalysis by family 63 inverting î±-glycosidase. Journal of Structural Biology, 2016, 196, 479-486.	1.3	6
89	Jaw1/LRMP increases Ca2+ influx upon GPCR stimulation with heterogeneous effect on the activity of each ITPR subtype. Scientific Reports, 2022, 12, .	1.6	6
90	Crystallization and preliminary X-ray analysis of the HA3 component of <i>Clostridium botulinum </i> type C progenitor toxin. Acta Crystallographica Section F: Structural Biology Communications, 2007, 63, 1038-1040.	0.7	5

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91	Heterologous expression, crystallization and preliminary X-ray characterization of CcCel6C, a glycoside hydrolase family 6 enzyme from the basidiomyceteCoprinopsis cinerea. Acta Crystallographica Section F: Structural Biology Communications, 2009, 65, 140-143.	0.7	5
92	Anti-melanogenic activity of salacinol by inhibition of tyrosinase oligosaccharide processing. Journal of Biochemistry, 2020, 167, 503-511.	0.9	5
93	The N-terminal region of Jaw1 has a role to inhibit the formation of organized smooth endoplasmic reticulum as an intrinsically disordered region. Scientific Reports, 2021, 11, 753.	1.6	5
94	Heterologous Production and Characterization of Arthrobacter globiformis T6 Isomalto-dextranase. Journal of Applied Glycoscience (1999), 2004, 51, 27-32.	0.3	5
95	Crystallization and preliminary X-ray analysis of Escherichia coli K12 Ygj K protein, a member of glycosyl hydrolase family 63. Acta Crystallographica Section D: Biological Crystallography, 2004, 60, 1284-1285.	2.5	4
96	Application of Purified Botulinum Type A Neurotoxin to Treat Experimental Trigeminal Neuropathy in Rats and Patients with Urinary Incontinence and Prostatic Hyperplasia. Journal of Toxicology, 2012, 2012, 1-8.	1.4	4
97	Enzymatic and structural characterization of \hat{l}^2 -fructofuranosidase from the honeybee gut bacterium Frischella perrara. Applied Microbiology and Biotechnology, 2022, 106, 2455-2470.	1.7	4
98	Structural basis for proteolytic processing of Aspergillus sojae \hat{l}_{\pm} -glucosidase L with strong transglucosylation activity. Journal of Structural Biology, 2022, 214, 107874.	1.3	4
99	Purification and characterization of pyruvate kinase from lamprey (Entosphenus japonicus) muscle. Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2001, 128, 159-164.	0.7	3
100	Crystallization and preliminary X-ray analysis of Thermoactinomyces vulgaris R-47 maltooligosaccharide-metabolizing enzyme homologous to glucoamylase. Acta Crystallographica Section F: Structural Biology Communications, 2005, 61, 302-304.	0.7	3
101	Effect of Nicking the C-terminal Region of the Clostridium botulinum Serotype D Neurotoxin Heavy Chain on its Toxicity and Molecular Properties. Protein Journal, 2007, 26, 173-181.	0.7	3
102	Mutagenesis-induced conformational change in domain B of a pullulan-hydrolyzing $\hat{l}\pm$ -amylase TVA I. Amylase, 2018, 2, 1-10.	0.7	3
103	Purification, crystallization and preliminary X-ray analysis of an HA17–HA70 (HA2–HA3) complex from <i>Clostridium botulinum</i> type C progenitor toxin. Acta Crystallographica Section F, Structural Biology Communications, 2014, 70, 64-67.	0.4	3
104	Glycan detecting tools developed from the Clostridium botulinum whole hemagglutinin complex. Scientific Reports, 2021, 11, 21973.	1.6	3
105	Novel protocol to observe the intestinal tuft cell using transmission electron microscopy. Biology Open, 2022, 11, .	0.6	3
106	Clinical application of <i>Clostridium botulinum </i> type A neurotoxin purified by a simple procedure for patients with urinary incontinence caused by refractory destrusor overactivity. FEMS Immunology and Medical Microbiology, 2007, 51, 587-587.	2.7	1
107	Structural Similarity between a Starch-hydrolyzing Enzyme and an N-Glycan-Hydrolyzing Enzyme: Exohydrolases Cleaving $\hat{l}\pm 1$,X-Glucosidic Linkages to Produce \hat{l}^2 -Glucose. Trends in Glycoscience and Glycotechnology, 2011, 23, 93-102.	0.0	1
108	A Surface Loop in the N-Terminal Domain of <i>Pedobacter heparinus </i> Heparin Lyase II is Important for Activity. Journal of Applied Glycoscience (1999), 2016, 63, 7-11.	0.3	1

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109	Lysosomal Enzyme GlcNAc-1- Phosphotransferase. , 2002, , 611-616.		1
110	Mutagenesis and Structural Analysis of Thermoactinomyces vulgaris R-47 .ALPHAAmylase II (TVA II). Journal of Applied Glycoscience (1999), 2005, 52, 225-231.	0.3	1
111	Inactivation of .ALPHAAmylases from Thermoactinomyces vulgaris R-47, TVA I and TVA II, by .OMEGAEpoxyalkyl .ALPHAD-Glucopyranoside. Journal of Applied Glycoscience (1999), 2005, 52, 273-276.	0.3	1
112	ENZYMATIC AND MOLECULAR BASIS FOR CANCER ASSOCIATED CHANGES IN $\hat{\mathbf{I}}^3$ -GLUTAMYL TRANSPEPTIDASE. , 1989, , 211-226.		1
113	Glycosyltransferases: Molecular Cloning of .BETA.1-4 N-Acetylglucosaminyltransferase III (GnT-III) Nippon Nogeikagaku Kaishi, 1993, 67, 1734-1740.	0.0	0
114	Study on the Lysosomal Enzyme Selection System Journal of Applied Glycoscience (1999), 2002, 49, 205-210.	0.3	0
115	X-ray Crystallographic Study of Glucodextranase from a Gram-positive Bacterium, Arthrobacter globiformis I42. Journal of Applied Glycoscience (1999), 2005, 52, 145-151.	0.3	0
116	Site-Directed Mutagenesis of Tryptophan 622 of Thermoactinomyces vulgaris R-47 Glucoamylase: pH Optima and Activities of Five Mutants. Journal of Applied Glycoscience (1999), 2005, 52, 277-279.	0.3	0
117	[Review: Symposium on Applied Glycoscience] Novel Findings of the Structure and Substrate Specificity of Glucosidases Belonging to Glycoside Hydrolase Family 63. Bulletin of Applied Glycoscience, 2013, 3, 151-158.	0.0	0
118	Traffic of Botulinum Toxin Complex: The Crystal Structure and the Role of Toxin Complex., 2014,, 1-5.		0
119	Traffic of Botulinum Toxin Complex: Crystal Structure and Role of Toxin Complex. , 2015, , 955-958.		0