Tamo Fukamizo

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	A conserved loop structure of GH19 chitinases assists the enzyme function from behind the core-functional region. Glycobiology, 2022, 32, 356-364.	2.5	4
2	Multi-functionality of a tryptophan residue conserved in substrate-binding groove of GH19 chitinases. Scientific Reports, 2021, 11, 2494.	3.3	5
3	A structural model for (ClcNAc)2 translocation via a periplasmic chitooligosaccharide-binding protein from marine Vibrio bacteria. Journal of Biological Chemistry, 2021, 297, 101071.	3.4	3
4	Chitin/Chitosan-Active Enzymes Involved in Plant–Microbe Interactions. Advances in Experimental Medicine and Biology, 2019, 1142, 253-272.	1.6	6
5	An Introduction to the Book. Advances in Experimental Medicine and Biology, 2019, 1142, 1-4.	1.6	1
6	Periplasmic solute-binding proteins: Structure classification and chitooligosaccharide recognition. International Journal of Biological Macromolecules, 2019, 128, 985-993.	7.5	21
7	Structure and function of a novel periplasmic chitooligosaccharide-binding protein from marine Vibrio bacteria. Journal of Biological Chemistry, 2018, 293, 5150-5159.	3.4	12
8	Resonance assignments for the apo-form of the cellulose-active lytic polysaccharide monooxygenase TaLPMO9A. Biomolecular NMR Assignments, 2018, 12, 357-361.	0.8	2
9	[Review] Protein Engineering Studies on Chitinaseï¼€hitosanase to Create a Novel Enzyme Function. Bulletin of Applied Glycoscience, 2018, 8, 33-44.	0.0	0
10	Enzymatic properties of a GH19 chitinase isolated from rice lacking a major loop structure involved in chitin binding. Glycobiology, 2017, 27, 477-485.	2.5	11
11	Crystal structure of a "loopless―GH19 chitinase in complex with chitin tetrasaccharide spanning the catalytic center. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2014, 1844, 793-802.	2.3	31
12	Complete subsite mapping of a "loopful―GH19 chitinase from rye seeds based on its crystal structure. FEBS Letters, 2013, 587, 2691-2697.	2.8	30
13	Chitin-Related Enzymes in Agro-Biosciences. Current Drug Targets, 2012, 13, 442-470.	2.1	43
14	Crystal structure and chitin oligosaccharideâ€binding mode of a †loopful' family GH19 chitinase from rye, <i>Secale cereale</i> , seeds. FEBS Journal, 2012, 279, 3639-3651.	4.7	42
15	Chitin oligosaccharide binding to a family GH19 chitinase from the moss <i>Bryum coronatum</i> . FEBS Journal, 2011, 278, 3991-4001.	4.7	40
16	Cloning and characterization of a small family 19 chitinase from moss (Bryum coronatum). Glycobiology, 2011, 21, 644-654.	2.5	49
17	Novel β-N-acetylglucosaminidases from Vibrio harveyi 650: Cloning, expression, enzymatic properties, and subsite identification. BMC Biochemistry, 2010, 11, 40.	4.4	53
18	A flexible loop controlling the enzymatic activity and specificity in a glycosyl hydrolase family 19 endochitinase from barley seeds (Hordeum vulgare L.). Biochimica Et Biophysica Acta - Proteins and Proteomics, 2009, 1794, 1159-1167.	2.3	14

ΤΑΜΟ ΕυκΑΜΙΖΟ

#	Article	IF	CITATIONS
19	Role of the Loop Structure of the Catalytic Domain in Rice Class I Chitinase. Journal of Biochemistry, 2007, 143, 487-495.	1.7	26
20	Family 19 chitinase from rice (Oryza sativa L.): substrate-binding subsites demonstrated by kinetic and molecular modeling studies. Plant Molecular Biology, 2003, 52, 43-52.	3.9	42
21	Kinetic Analysis of Barley Chitinase. Archives of Biochemistry and Biophysics, 1997, 344, 335-342.	3.0	70
22	Separation and Mutarotation of Anomers of Chitooligosaccharides1. Journal of Biochemistry, 1982, 91, 619-626.	1.7	30