

Eva Nordberg Karlsson

List of Publications by Year in descending order

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132
papers

4,432
citations

101543

36
h-index

133252

59
g-index

149
all docs

149
docs citations

149
times ranked

5011
citing authors

#	ARTICLE	IF	CITATIONS
1	Lignocellulose degradation for the bioeconomy: The potential of enzyme synergies between xylanases, ferulic acid esterase and laccase for the production of arabinoxylo-oligosaccharides. <i>Bioresource Technology</i> , 2022, 343, 126114.	9.6	24
2	Crystal structure and initial characterization of a novel archaeal-like Holliday junction-resolving enzyme from <i>Thermus thermophilus</i> phage Tth15-6. <i>Acta Crystallographica Section D: Structural Biology</i> , 2022, 78, 212-227.	2.3	5
3	Investigation of Structural Features of Two Related Lipases and the Impact on Fatty Acid Specificity in Vegetable Fats. <i>International Journal of Molecular Sciences</i> , 2022, 23, 7072.	4.1	3
4	Microwave-assisted xylanase reaction: impact in the production of prebiotic xylooligosaccharides. <i>RSC Advances</i> , 2021, 11, 11882-11888.	3.6	6
5	Xylooligosaccharides Increase <i>Bifidobacteria</i> and <i>Lachnospiraceae</i> in Mice on a High-Fat Diet, with a Concomitant Increase in Short-Chain Fatty Acids, Especially Butyric Acid. <i>Journal of Agricultural and Food Chemistry</i> , 2021, 69, 3617-3625.	5.2	48
6	Ultrasound Assisted Alkaline Pre-treatment Efficiently Solubilises Hemicellulose from Oat Hulls. <i>Waste and Biomass Valorization</i> , 2021, 12, 5371-5381.	3.4	12
7	Modeled 3D-Structures of Proteobacterial Transglycosylases from Glycoside Hydrolase Family 17 Give Insight in Ligand Interactions Explaining Differences in Transglycosylation Products. <i>Applied Sciences (Switzerland)</i> , 2021, 11, 4048.	2.5	3
8	Chemical and biochemical bleaching of oat hulls: The effect of hydrogen peroxide, laccase, xylanase and sonication on optical properties and chemical composition. <i>Biotechnology Reports (Amsterdam)</i> , 2021, 10, 1004.	4.8	10
9	Rational Enzyme Design without Structural Knowledge: A Sequence-Based Approach for Efficient Generation of Transglycosylases. <i>Chemistry - A European Journal</i> , 2021, 27, 10323-10334.	3.3	29
10	Cultivation of the gut bacterium <i>Prevotella copri</i> DSM 18205 using glucose and xylose as carbon sources. <i>MicrobiologyOpen</i> , 2021, 10, e1213.	3.0	13
11	Novel xylan degrading enzymes from polysaccharide utilizing loci of <i>Prevotella copri</i> DSM18205. <i>Glycobiology</i> , 2021, 31, 1330-1349.	2.5	9
12	Going to extremes – a metagenomic journey into the dark matter of life. <i>FEMS Microbiology Letters</i> , 2021, 368, .	1.8	16
13	Exploring Codon Adjustment Strategies towards Escherichia coli-Based Production of Viral Proteins Encoded by HTH1, a Novel Prophage of the Marine Bacterium <i>Hypnocyclus thymotrophus</i> . <i>Viruses</i> , 2021, 13, 1215.	3.3	3
14	Extraction of sugarcane bagasse arabinoxylan, integrated with enzymatic production of xylo-oligosaccharides and separation of cellulose. <i>Biotechnology for Biofuels</i> , 2021, 14, 153.	6.2	28
15	Endo-xylanases from <i>Cohnella</i> sp. AR92 aimed at xylan and arabinoxylan conversion into value-added products. <i>Applied Microbiology and Biotechnology</i> , 2021, 105, 6759-6778.	3.6	5
16	Glucuronosylated and linear xylooligosaccharides from Quinoa stalks xylan as potential prebiotic source for growth of <i>Bifidobacterium adolescentis</i> and <i>Weissella cibaria</i> . <i>LWT - Food Science and Technology</i> , 2021, 152, 112348.	5.2	11
17	Altering the water holding capacity of potato pulp via structural modifications of the pectic polysaccharides. <i>Carbohydrate Polymer Technologies and Applications</i> , 2021, 2, 100153.	2.6	2
18	Engineering CGTase to improve synthesis of alkyl glycosides. <i>Glycobiology</i> , 2021, 31, 603-612.	2.5	7

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19	Identification of Phlorotannins in the Brown Algae, <i>Saccharina latissima</i> and <i>Ascophyllum nodosum</i> by Ultra-High-Performance Liquid Chromatography Coupled to High-Resolution Tandem Mass Spectrometry. <i>Molecules</i> , 2021, 26, 43.	3.8	15
20	Engineering the carotenoid biosynthetic pathway in <i>Rhodothermus marinus</i> for lycopene production. <i>Metabolic Engineering Communications</i> , 2020, 11, e00140.	3.6	5
21	Warming weather changes the chemical composition of oat hulls. <i>Plant Biology</i> , 2020, 22, 1086-1091.	3.8	32
22	Enzyme synergy for the production of arabinoxylo-oligosaccharides from highly substituted arabinoxytan and evaluation of their prebiotic potential. <i>LWT - Food Science and Technology</i> , 2020, 131, 109762.	5.2	11
23	Opportunities for seaweed biorefinery. , 2020, , 3-31.		10
24	Extraction and Modification of Macroalgal Polysaccharides for Current and Next-Generation Applications. <i>Molecules</i> , 2020, 25, 930.	3.8	125
25	The Catalytic Acidâ€‘Base in GH109 Resides in a Conserved GGHG Loop and Allows for Comparable β -Retaining and β -Inverting Activity in an <i>N</i> -Acetylgalactosaminidase from <i>Akkermansia muciniphila</i> . <i>ACS Catalysis</i> , 2020, 10, 3809-3819.	11.2	15
26	Characterization and diversity of the complete set of GH family 3 enzymes from <i>Rhodothermus marinus</i> DSM 4253. <i>Scientific Reports</i> , 2020, 10, 1329.	3.3	9
27	Taxogenomic assessment and genomic characterisation of <i>Weissella cibaria</i> strain 92 able to metabolise oligosaccharides derived from dietary fibres. <i>Scientific Reports</i> , 2020, 10, 5853.	3.3	15
28	Composition analysis and minimal treatments to solubilize polysaccharides from the brown seaweed <i>Laminaria digitata</i> for microbial growth of thermophiles. <i>Journal of Applied Phycology</i> , 2020, 32, 1933-1947.	2.8	13
29	Evaluation of Sequential Processing for the Extraction of Starch, Lipids, and Proteins From Wheat Bran. <i>Frontiers in Bioengineering and Biotechnology</i> , 2019, 7, 413.	4.1	30
30	Crystal structures of the <i>Bacillus subtilis</i> prophage lytic cassette proteins XepA and YomS. <i>Acta Crystallographica Section D: Structural Biology</i> , 2019, 75, 1028-1039.	2.3	9
31	β -Mannanase-catalyzed synthesis of alkyl mannoooligosides. <i>Applied Microbiology and Biotechnology</i> , 2018, 102, 5149-5163.	3.6	19
32	Xylo- and arabinoxylooligosaccharides from wheat bran by endoxylanases, utilisation by probiotic bacteria, and structural studies of the enzymes. <i>Applied Microbiology and Biotechnology</i> , 2018, 102, 3105-3120.	3.6	36
33	Valorization of Brewerâ€™s spent grain to prebiotic oligosaccharide: Production, xylanase catalyzed hydrolysis, in-vitro evaluation with probiotic strains and in a batch human fecal fermentation model. <i>Journal of Biotechnology</i> , 2018, 268, 61-70.	3.8	48
34	Arabinoxytanase from glycoside hydrolase family 5 is a selective enzyme for production of specific arabinoxylooligosaccharides. <i>Food Chemistry</i> , 2018, 242, 579-584.	8.2	28
35	Structural insights of Rm Xyn10A â€‘ A prebiotic-producing GH10 xylanase with a non-conserved aglycone binding region. <i>Biochimica Et Biophysica Acta - Proteins and Proteomics</i> , 2018, 1866, 292-306.	2.3	14
36	Marine Poly- and Oligosaccharides as Prebiotics. <i>Journal of Agricultural and Food Chemistry</i> , 2018, 66, 11544-11549.	5.2	42

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37	Data on saponins, xylan and cellulose yield obtained from quinoa stalks after pressurized hot water extraction. Data in Brief, 2018, 20, 289-292.	1.0	3
38	Endo-xylanases as tools for production of substituted xylooligosaccharides with prebiotic properties. Applied Microbiology and Biotechnology, 2018, 102, 9081-9088.	3.6	116
39	Integrated process for sequential extraction of saponins, xylan and cellulose from quinoa stalks (<i>Chenopodium quinoa</i> Willd.). Industrial Crops and Products, 2018, 121, 54-65.	5.2	47
40	Crystal structure of β -glucosidase 1A from <i>Thermotoga neapolitana</i> and comparison of active site mutants for hydrolysis of flavonoid glucosides. Proteins: Structure, Function and Bioinformatics, 2017, 85, 872-884.	2.6	7
41	Eliminating hydrolytic activity without affecting the transglycosylation of a GH1 β -glucosidase. Applied Microbiology and Biotechnology, 2017, 101, 1121-1131.	3.6	39
42	Three-dimensional structures and functional studies of two GH43 arabinofuranosidases from <i>Weissella</i> sp. strain 142 and <i>Lactobacillus brevis</i> . FEBS Journal, 2017, 284, 2019-2036.	4.7	16
43	Extraction of soluble arabinoxylan from enzymatically pretreated wheat bran and production of short xylo-oligosaccharides and arabinoxylo-oligosaccharides from arabinoxylan by glycoside hydrolase family 10 and 11 endoxylanases. Journal of Biotechnology, 2017, 260, 53-61.	3.8	35
44	Cover Image, Volume 85, Issue 5. Proteins: Structure, Function and Bioinformatics, 2017, 85, C4-C4.	2.6	0
45	Cover Image, Volume 85, Issue 6. Proteins: Structure, Function and Bioinformatics, 2017, 85, C4.	2.6	0
46	Extraction of Glucuronoarabinoxylan from Quinoa Stalks (<i>Chenopodium quinoa</i> Willd.) and Evaluation of Xylooligosaccharides Produced by GH10 and GH11 Xylanases. Journal of Agricultural and Food Chemistry, 2017, 65, 8663-8673.	5.2	30
47	Rational design of a thermostable glycoside hydrolase from family 3 introduces β -glycosynthase activity. Glycobiology, 2017, 27, 165-175.	2.5	9
48	Evaluation of the production of exopolysaccharides by two strains of the thermophilic bacterium <i>Rhodothermus marinus</i> . Carbohydrate Polymers, 2017, 156, 1-8.	10.2	109
49	Structural Considerations on the Use of Endo-Xylanases for the Production of prebiotic Xylooligosaccharides from Biomass. Current Protein and Peptide Science, 2017, 19, 48-67.	1.4	73
50	Characterization of a family 43 β -xylosidase from the xylooligosaccharide utilizing putative probiotic <i>Weissella</i> sp. strain 92. Glycobiology, 2016, 26, 193-202.	2.5	23
51	Development of the Nordic Bioeconomy. TemaNord, 2016, , .	1.3	4
52	A CGTase with high coupling activity using β -cyclodextrin isolated from a novel strain clustering under the genus <i>Carboxydocella</i> . Glycobiology, 2015, 25, 514-523.	2.5	8
53	Complexation of alkyl glycosides with β -cyclodextrin can have drastically different effects on their conversion by glycoside hydrolases. Journal of Biotechnology, 2015, 200, 52-58.	3.8	3
54	A GH57 β -glucanotransferase of hyperthermophilic origin with potential for alkyl glycoside production. Applied Microbiology and Biotechnology, 2015, 99, 7101-7113.	3.6	8

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55	Characterization of cyclodextrin glycosyltransferases (CGTases) and their application for synthesis of alkyl glycosides with oligomeric head group. <i>Process Biochemistry</i> , 2015, 50, 722-728.	3.7	19
56	Production of prebiotic xylooligosaccharides from alkaline extracted wheat straw using the K80R-variant of a thermostable alkali-tolerant xylanase. <i>Food and Bioproducts Processing</i> , 2015, 93, 1-10.	3.6	59
57	Characterization of the substitution pattern of cellulose derivatives using carbohydrate-binding modules. <i>BMC Biotechnology</i> , 2014, 14, 113.	3.3	17
58	Production of arabinoxylan-oligosaccharide mixtures of varying composition from rye bran by a combination of process conditions and type of xylanase. <i>Bioresource Technology</i> , 2014, 174, 118-125.	9.6	47
59	Extraction of water-soluble xylan from wheat bran and utilization of enzymatically produced xylooligosaccharides by <i>Lactobacillus</i> , <i>Bifidobacterium</i> and <i>Weissella</i> spp.. <i>LWT - Food Science and Technology</i> , 2014, 56, 321-327.	5.2	65
60	Preparation of two glycoside hydrolases for use in micro-aqueous media. <i>Journal of Molecular Catalysis B: Enzymatic</i> , 2014, 108, 1-6.	1.8	9
61	Cereal Byproducts Have Prebiotic Potential in Mice Fed a High-Fat Diet. <i>Journal of Agricultural and Food Chemistry</i> , 2014, 62, 8169-8178.	5.2	43
62	Substituent Effects on in Vitro Antioxidizing Properties, Stability, and Solubility in Flavonoids. <i>Journal of Agricultural and Food Chemistry</i> , 2014, 62, 3321-3333.	5.2	176
63	Glycosynthases from <i>Thermotoga neapolitana</i> β -glucosidase 1A: A comparison of β -glucosyl fluoride and in situ-generated β -glucosyl formate donors. <i>Journal of Molecular Catalysis B: Enzymatic</i> , 2014, 107, 132-139.	1.8	15
64	Carbohydrate binding module recognition of xyloglucan defined by polar contacts with branching xyloses and β -1,4 interactions. <i>Proteins: Structure, Function and Bioinformatics</i> , 2014, 82, 3466-3475.	2.6	13
65	Xylooligosaccharides from Hardwood and Cereal Xylans Produced by a Thermostable Xylanase as Carbon Sources for <i>Lactobacillus brevis</i> and <i>Bifidobacterium adolescentis</i> . <i>Journal of Agricultural and Food Chemistry</i> , 2013, 61, 7333-7340.	5.2	99
66	An on-line method for pressurized hot water extraction and enzymatic hydrolysis of quercetin glucosides from onions. <i>Analytica Chimica Acta</i> , 2013, 785, 50-59.	5.4	31
67	Microbial Glycoside Hydrolases for Biomass Utilization in Biofuels Applications. , 2013, , 171-188.		11
68	Phylogenetic analysis and substrate specificity of GH2 β -mannosidases from <i>Aspergillus</i> species. <i>FEBS Letters</i> , 2013, 587, 3444-3449.	2.8	15
69	Bioresource utilisation by sustainable technologies in new value-added biorefinery concepts – two case studies from food and forest industry. <i>Journal of Cleaner Production</i> , 2013, 57, 46-58.	9.3	66
70	Evidence for xylooligosaccharide utilization in <i>Weissella</i> strains isolated from Indian fermented foods and vegetables. <i>FEMS Microbiology Letters</i> , 2013, 346, 20-28.	1.8	48
71	Improved Transferase/Hydrolase Ratio through Rational Design of a Family 1 β -Glucosidase from <i>Thermotoga neapolitana</i> . <i>Applied and Environmental Microbiology</i> , 2013, 79, 3400-3405.	3.1	40
72	Glycoside Hydrolases for Extraction and Modification of Polyphenolic Antioxidants. , 2013, , 9-21.		4

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73	Caloramator boliviensis sp. nov., a thermophilic, ethanol-producing bacterium isolated from a hot spring. International Journal of Systematic and Evolutionary Microbiology, 2012, 62, 1679-1686.	1.7	31
74	Structural basis for carbohydrate-binding specificity—A comparative assessment of two engineered carbohydrate-binding modules. Glycobiology, 2012, 22, 948-961.	2.5	35
75	A cellulolytic Hypocrea strain isolated from South American brave straw produces a modular xylanase. Carbohydrate Research, 2012, 356, 215-223.	2.3	7
76	Immobilization of thermostable β -glucosidase variants on acrylic supports for biocatalytic processes in hot water. Journal of Molecular Catalysis B: Enzymatic, 2012, 80, 28-38.	1.8	22
77	Development of a Sustainable Method for Modification of polyphenolic glucosides. , 2012, , .		0
78	Aglycone specificity of Thermotoga neapolitana β -glucosidase 1A modified by mutagenesis, leading to increased catalytic efficiency in quercetin-3-glucoside hydrolysis. BMC Biochemistry, 2011, 12, 11.	4.4	29
79	The crystal structure of XG4, an evolved xyloglucan-specific carbohydrate-binding module. Proteins: Structure, Function and Bioinformatics, 2010, 78, 785-789.	2.6	11
80	A novel direct screening method for alkyl glucoside production by glucosidases expressed in E. coli in 96-well plates. Journal of Biotechnology, 2010, 145, 186-192.	3.8	5
81	Characterization of the Properties of for Probiotic or Protective Culture Use. Journal of Food Protection, 2010, 73, 960-966.	1.7	18
82	Mutational Tuning of Galectin-3 Specificity and Biological Function. Journal of Biological Chemistry, 2010, 285, 35079-35091.	3.4	98
83	Structural and Functional Analyses of β -Glucosidase 3B from Thermotoga neapolitana: A Thermostable Three-Domain Representative of Glycoside Hydrolase 3. Journal of Molecular Biology, 2010, 397, 724-739.	4.2	117
84	Exploring the possibility of using a thermostable mutant of β -glucosidase for rapid hydrolysis of quercetin glucosides in hot water. Green Chemistry, 2010, 12, 159-168.	9.0	47
85	Affinity maturation generates greatly improved xyloglucan-specific carbohydrate binding modules. BMC Biotechnology, 2009, 9, 92.	3.3	24
86	Differences and similarities in enzymes from the neopullulanase subfamily isolated from thermophilic species. Biologia (Poland), 2008, 63, 1006-1014.	1.5	11
87	Novel Members of Glycoside Hydrolase Family 13 Derived from Environmental DNA. Applied and Environmental Microbiology, 2008, 74, 1914-1921.	3.1	28
88	Novel xylan-binding properties of an engineered family 4 carbohydrate-binding module. Biochemical Journal, 2007, 406, 209-214.	3.7	26
89	A novel variant of Thermotoga neapolitana β -glucosidase B is an efficient catalyst for the synthesis of alkyl glucosides by transglycosylation. Journal of Biotechnology, 2007, 130, 67-74.	3.8	65
90	Potential and utilization of thermophiles and thermostable enzymes in biorefining. Microbial Cell Factories, 2007, 6, 9.	4.0	459

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91	Capture of bacteriocins directly from non-clarified fermentation broth using macroporous monolithic cryogels with phenyl ligands. <i>Enzyme and Microbial Technology</i> , 2007, 40, 786-793.	3.2	26
92	Expression, purification, crystallization and preliminary X-ray diffraction analysis of <i>Thermotoga neapolitana</i> α -D-glucosidase B. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2007, 63, 802-806.	0.7	7
93	Production and physicochemical characterization of acidocin D20079, a bacteriocin produced by <i>Lactobacillus acidophilus</i> DSM 20079. <i>World Journal of Microbiology and Biotechnology</i> , 2007, 23, 911-921.	3.6	5
94	Mode of action of acidocin D20079, a bacteriocin produced by the potential probiotic strain, <i>Lactobacillus acidophilus</i> DSM 20079. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2007, 34, 373-379.	3.0	42
95	Engineered xyloglucan specificity in a carbohydrate-binding module. <i>Glycobiology</i> , 2006, 16, 1171-1180.	2.5	37
96	Subcritical water extraction and α -D-glucosidase-catalyzed hydrolysis of quercetin glycosides in onion waste. <i>Green Chemistry</i> , 2006, 8, 949-959.	9.0	114
97	Molecular engineering of a thermostable carbohydrate-binding module. <i>Biocatalysis and Biotransformation</i> , 2006, 24, 31-37.	2.0	2
98	Dimerisation and an Increase in Active Site Aromatic Groups as Adaptations to High Temperatures: X-ray Solution Scattering and Substrate-bound Crystal Structures of <i>Rhodothermus marinus</i> Endoglucanase Cel12A. <i>Journal of Molecular Biology</i> , 2006, 356, 57-71.	4.2	21
99	Production of a lipolytic enzyme originating from <i>Bacillus halodurans</i> LBB2 in the methylotrophic yeast <i>Pichia pastoris</i> . <i>Applied Microbiology and Biotechnology</i> , 2006, 71, 463-472.	3.6	15
100	Characterisation of two novel cyclodextrinases using on-line microdialysis sampling with high-performance anion exchange chromatography. <i>Analytical and Bioanalytical Chemistry</i> , 2006, 385, 1421-1429.	3.7	6
101	Evolution of a carbohydrate binding module into a protein-specific binder. <i>New Biotechnology</i> , 2006, 23, 111-117.	2.7	18
102	The methylotrophic yeast as a host for the expression and production of thermostable xylanase from the bacterium. <i>FEMS Yeast Research</i> , 2005, 5, 839-850.	2.3	25
103	A cultivation technique for <i>E. coli</i> fed-batch cultivations operating close to the maximum oxygen transfer capacity of the reactor. <i>Biotechnology Letters</i> , 2005, 27, 983-990.	2.2	20
104	Purification and characterisation of acidocin D20079, a bacteriocin produced by <i>Lactobacillus acidophilus</i> DSM 20079. <i>Journal of Biotechnology</i> , 2005, 117, 343-354.	3.8	98
105	Effect of postinduction nutrient feed composition and use of lactose as inducer during production of thermostable xylanase in <i>Escherichia coli</i> glucose-limited fed-batch cultivations. <i>Journal of Bioscience and Bioengineering</i> , 2005, 99, 477-484.	2.2	34
106	Optimized expression of soluble cyclomaltodextrinase of thermophilic origin in <i>Escherichia coli</i> by using a soluble fusion-tag and by tuning of inducer concentration. <i>Protein Expression and Purification</i> , 2005, 39, 54-60.	1.3	52
107	Two novel cyclodextrin-degrading enzymes isolated from thermophilic bacteria have similar domain structures but differ in oligomeric state and activity profile. <i>Journal of Bioscience and Bioengineering</i> , 2005, 100, 380-390.	2.2	30
108	A carbohydrate binding module as a diversity-carrying scaffold. <i>Protein Engineering, Design and Selection</i> , 2004, 17, 213-221.	2.1	51

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109	The modular xylanase Xyn10A from <i>Rhodothermus marinus</i> cell-attached, and its C-terminal domain has several putative homologues among cell-attached proteins within the phylum Bacteroidetes. <i>FEMS Microbiology Letters</i> , 2004, 241, 233-242.	1.8	27
110	Citrate synthase from <i>Thermus aquaticus</i> : a thermostable bacterial enzyme with a five-membered inter-subunit ionic network. <i>Extremophiles</i> , 2003, 7, 9-16.	2.3	14
111	Probing the stability of the modular family 10 xylanase from <i>Rhodothermus marinus</i> . <i>Extremophiles</i> , 2003, 7, 483-491.	2.3	38
112	The Modular Organisation and Stability of a Thermostable Family 10 Xylanase. <i>Biocatalysis and Biotransformation</i> , 2003, 21, 253-260.	2.0	10
113	Calcium Binding and Thermostability of Carbohydrate Binding Module CBM4-2 of Xyn10A from <i>Rhodothermus marinus</i> . <i>Biochemistry</i> , 2002, 41, 5720-5729.	2.5	41
114	The Solution Structure of the CBM4-2 Carbohydrate Binding Module from a Thermostable <i>Rhodothermus marinus</i> Xylanase. <i>Biochemistry</i> , 2002, 41, 5712-5719.	2.5	68
115	The Structure of <i>Rhodothermus marinus</i> Cel12A, A Highly Thermostable Family 12 Endoglucanase, at 1.8Å... Resolution. <i>Journal of Molecular Biology</i> , 2002, 320, 883-897.	4.2	48
116	Production of heterologous thermostable glycoside hydrolases and the presence of host-cell proteases in substrate limited fed-batch cultures of <i>Escherichia coli</i> BL21(DE3). <i>Applied Microbiology and Biotechnology</i> , 2002, 60, 408-416.	3.6	30
117	<i>Rhodothermus marinus</i> : a thermophilic bacterium producing dimeric and hexameric citrate synthase isoenzymes. <i>Extremophiles</i> , 2002, 6, 51-56.	2.3	11
118	Virtually complete ¹ H, ¹³ C and ¹⁵ N resonance assignments of the second family 4 xylan binding module of <i>Rhodothermus marinus</i> xylanase 10A. <i>Journal of Biomolecular NMR</i> , 2002, 22, 187-188.	2.8	4
119	Title is missing!. <i>Biotechnology Letters</i> , 2002, 24, 1191-1197.	2.2	4
120	Deletion of a cytotoxic, N-terminal putative signal peptide results in a significant increase in production yields in <i>Escherichia coli</i> and improved specific activity of Cel12A from <i>Rhodothermus marinus</i> . <i>Applied Microbiology and Biotechnology</i> , 2001, 55, 578-584.	3.6	27
121	Integrated flow-injection processing for on-line quantification of plasmid DNA during cultivation of <i>E. coli</i> . <i>Biotechnology and Bioengineering</i> , 2001, 73, 406-411.	3.3	10
122	Title is missing!. <i>Biotechnology Letters</i> , 2001, 23, 1135-1140.	2.2	5
123	Carbohydrate-binding modules from a thermostable <i>Rhodothermus marinus</i> xylanase: cloning, expression and binding studies. <i>Biochemical Journal</i> , 2000, 345, 53.	3.7	77
124	Carbohydrate-binding modules from a thermostable <i>Rhodothermus marinus</i> xylanase: cloning, expression and binding studies. <i>Biochemical Journal</i> , 2000, 345, 53-60.	3.7	89
125	Title is missing!. <i>Biotechnology Letters</i> , 2000, 22, 663-669.	2.2	17
126	Carbohydrate-binding modules from a thermostable <i>Rhodothermus marinus</i> xylanase: cloning, expression and binding studies. <i>Biochemical Journal</i> , 2000, 345 Pt 1, 53-60.	3.7	28

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127	Efficient production of truncated thermostable xylanases from <i>Rhodothermus marinus</i> in <i>Escherichia coli</i> fed-batch cultures. <i>Journal of Bioscience and Bioengineering</i> , 1999, 87, 598-606.	2.2	30
128	On-line detection of acetate formation in <i>Escherichia coli</i> cultures using dissolved oxygen responses to feed transients. , 1999, 64, 590-598.		84
129	Evidence for substrate binding of a recombinant thermostable xylanase originating from <i>Rhodothermus marinus</i> . <i>FEMS Microbiology Letters</i> , 1998, 168, 1-7.	1.8	23
130	Enzymatic specificity and hydrolysis pattern of the catalytic domain of the xylanase Xyn1 from <i>Rhodothermus marinus</i> . <i>Journal of Biotechnology</i> , 1998, 60, 23-35.	3.8	41
131	Evidence for substrate binding of a recombinant thermostable xylanase originating from <i>Rhodothermus marinus</i> . <i>FEMS Microbiology Letters</i> , 1998, 168, 1-7.	1.8	1
132	Cloning and sequence of a thermostable multidomain xylanase from the bacterium <i>Rhodothermus marinus</i> . <i>Biochimica Et Biophysica Acta Gene Regulatory Mechanisms</i> , 1997, 1353, 118-124.	2.4	51