

# Magali Remaud-Simeon

## List of Publications by Year in descending order

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152  
papers

6,255  
citations

50276

46  
h-index

88630

70  
g-index

153  
all docs

153  
docs citations

153  
times ranked

3982  
citing authors

#	ARTICLE	IF	CITATIONS
1	Homopolysaccharides from lactic acid bacteria. <i>International Dairy Journal</i> , 2001, 11, 675-685.	3.0	284
2	Functional metagenomics to mine the human gut microbiome for dietary fiber catabolic enzymes. <i>Genome Research</i> , 2010, 20, 1605-1612.	5.5	228
3	Biodiversity of Exopolysaccharides Produced from Sucrose by Sourdough Lactic Acid Bacteria. <i>Journal of Agricultural and Food Chemistry</i> , 2009, 57, 10889-10897.	5.2	160
4	Amylosucrase, a Glucan-synthesizing Enzyme from the Î±-Amylase Family. <i>Journal of Biological Chemistry</i> , 2001, 276, 25273-25278.	3.4	135
5	Amylosucrase from <i>Neisseria polysaccharea</i> : novel catalytic properties. <i>FEBS Letters</i> , 2000, 471, 219-223.	2.8	133
6	Amylose Synthesized in Vitro by Amylosucrase: Morphology, Structure, and Properties. <i>Biomacromolecules</i> , 2005, 6, 1000-1011.	5.4	119
7	Understanding the Polymerization Mechanism of Glycoside-Hydrolase Family 70 Glucansucrases. <i>Journal of Biological Chemistry</i> , 2006, 281, 31254-31267.	3.4	119
8	Characterization of <i>Leuconostoc mesenteroides</i> NRRL B-512F dextranase (DSRS) and identification of amino-acid residues playing a key role in enzyme activity. <i>Applied Microbiology and Biotechnology</i> , 1997, 48, 465-472.	3.6	116
9	Sequence Analysis of the Gene Encoding Amylosucrase from <i>Neisseria polysaccharea</i> and Characterization of the Recombinant Enzyme. <i>Journal of Bacteriology</i> , 1999, 181, 375-381.	2.2	111
10	Glucansucrases: molecular engineering and oligosaccharide synthesis. <i>Journal of Molecular Catalysis B: Enzymatic</i> , 2000, 10, 117-128.	1.8	100
11	Molecular Basis of the Amylose-like Polymer Formation Catalyzed by <i>Neisseria polysaccharea</i> Amylosucrase. <i>Journal of Biological Chemistry</i> , 2004, 279, 726-734.	3.4	96
12	Characterization of dextran-producing <i>Weissella</i> strains isolated from sourdoughs and evidence of constitutive dextranase expression. <i>FEMS Microbiology Letters</i> , 2010, 311, 18-26.	1.8	89
13	Growth and energetics of <i>Leuconostoc mesenteroides</i> NRRL B-1299 during metabolism of various sugars and their consequences for dextranase production. <i>Applied and Environmental Microbiology</i> , 1997, 63, 2159-2165.	3.1	89
14	Insights into lid movements of <i>Burkholderia cepacia</i> lipase inferred from molecular dynamics simulations. <i>Proteins: Structure, Function and Bioinformatics</i> , 2009, 77, 509-523.	2.6	88
15	Molecular Characterization of DSR-E, an Î±-1,2 Linkage-Synthesizing Dextranase with Two Catalytic Domains. <i>Journal of Bacteriology</i> , 2002, 184, 5753-5761.	2.2	87
16	Geometric algorithms for the conformational analysis of long protein loops. <i>Journal of Computational Chemistry</i> , 2004, 25, 956-967.	3.3	86
17	Crystal Structures of Amylosucrase from <i>Neisseria polysaccharea</i> in Complex with d-Glucose and the Active Site Mutant Glu328Gln in Complex with the Natural Substrate Sucrose. <i>Biochemistry</i> , 2001, 40, 9032-9039.	2.5	85
18	In Vitro Fermentation of Linear and Î±-1,2-Branched Dextran by the Human Fecal Microbiota. <i>Applied and Environmental Microbiology</i> , 2011, 77, 5307-5315.	3.1	84

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19	Oligosaccharide and Sucrose Complexes of Amylosucrase. <i>Journal of Biological Chemistry</i> , 2002, 277, 47741-47747.	3.4	83
20	Production and use of glucosyltransferases from <i>Leuconostoc mesenteroides</i> NRRL B-1299 for the synthesis of oligosaccharides containing $\alpha$ -1,2 linkages. <i>Applied Biochemistry and Biotechnology</i> , 1994, 44, 101-117.	2.9	82
21	Prebiotic effects of oligosaccharides on selected vaginal lactobacilli and pathogenic microorganisms. <i>Anaerobe</i> , 2005, 11, 145-153.	2.1	80
22	Functional and Structural Characterization of $\alpha$ -1,2 Branching Sucrase Derived from DSR-E Glucansucrase. <i>Journal of Biological Chemistry</i> , 2012, 287, 7915-7924.	3.4	78
23	Functional Metagenomics Reveals Novel Pathways of Prebiotic Breakdown by Human Gut Bacteria. <i>PLoS ONE</i> , 2013, 8, e72766.	2.5	78
24	Cloning and sequencing of a gene coding for a novel dextransucrase from <i>Leuconostoc mesenteroides</i> NRRL B-1299 synthesizing only $\alpha$ -1,6 and $\alpha$ -1,3 linkages. <i>Gene</i> , 1996, 182, 23-32.	2.2	77
25	CAZyme discovery and design for sweet dreams. <i>Current Opinion in Chemical Biology</i> , 2014, 19, 17-24.	6.1	74
26	$\alpha$ -D-Glucan-Based Dendritic Nanoparticles Prepared by in Vitro Enzymatic Chain Extension of Glycogen. <i>Biomacromolecules</i> , 2006, 7, 1720-1728.	5.4	72
27	<i>Leuconostoc mesenteroides</i> glucansucrase synthesis of flavonoid glucosides by acceptor reactions in aqueous-organic solvents. <i>Carbohydrate Research</i> , 2006, 341, 855-863.	2.3	70
28	Characterization of the Different Dextransucrase Activities Excreted in Glucose, Fructose, or Sucrose Medium by <i>Leuconostoc mesenteroides</i> NRRL B-1299. <i>Applied and Environmental Microbiology</i> , 1998, 64, 1298-1302.	3.1	69
29	Combinatorial engineering to enhance amylosucrase performance: construction, selection, and screening of variant libraries for increased activity. <i>FEBS Letters</i> , 2004, 560, 91-97.	2.8	68
30	Crystal Structure of the Covalent Intermediate of Amylosucrase from <i>Neisseria polysaccharea</i> . <i>Biochemistry</i> , 2004, 43, 3104-3110.	2.5	67
31	Role of the Two Catalytic Domains of DSR-E Dextransucrase and Their Involvement in the Formation of Highly $\alpha$ -1,2 Branched Dextran. <i>Journal of Bacteriology</i> , 2005, 187, 296-303.	2.2	67
32	Sequence analysis of the gene encoding alternansucrase, a sucrose glucosyltransferase from <i>Leuconostoc mesenteroides</i> NRRL B-1355. <i>FEMS Microbiology Letters</i> , 2000, 182, 81-85.	1.8	65
33	Exopolysaccharide (EPS) Synthesis by <i>Oenococcus oeni</i> : From Genes to Phenotypes. <i>PLoS ONE</i> , 2014, 9, e98898.	2.5	65
34	Design of $\alpha$ -Transglucosidases of Controlled Specificity for Programmed Chemoenzymatic Synthesis of Antigenic Oligosaccharides. <i>Journal of the American Chemical Society</i> , 2009, 131, 7379-7389.	13.7	64
35	Characterization of glucan-producing <i>Leuconostoc</i> strains isolated from sourdough. <i>International Journal of Food Microbiology</i> , 2010, 144, 1-9.	4.7	62
36	Transglucosidases as efficient tools for oligosaccharide and glucoconjugate synthesis. <i>Current Opinion in Microbiology</i> , 2010, 13, 293-300.	5.1	61

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37	Characterization of a novel dextransucrase from <i>Weissella confusa</i> isolated from sourdough. <i>Applied Microbiology and Biotechnology</i> , 2013, 97, 5413-5422.	3.6	60
38	Enzymatic synthesis of glycamide surfactants by amidification reaction. <i>Tetrahedron</i> , 1997, 53, 5185-5194.	1.9	56
39	Structural Investigation of the Thermostability and Product Specificity of Amylosucrase from the Bacterium <i>Deinococcus geothermalis</i> . <i>Journal of Biological Chemistry</i> , 2012, 287, 6642-6654.	3.4	55
40	Effect of <i>Leuconostoc mesenteroides</i> NRRL B-512F Dextransucrase Carboxy-Terminal Deletions on Dextran and Oligosaccharide Synthesis. <i>Applied and Environmental Microbiology</i> , 1998, 64, 1644-1649.	3.1	54
41	Characterisation of a novel amylosucrase from <i>Deinococcus radiodurans</i> . <i>FEBS Letters</i> , 2005, 579, 1405-1410.	2.8	53
42	Novel oligosaccharides synthesized from sucrose donor and cellobiose acceptor by alternansucrase. <i>Carbohydrate Research</i> , 2001, 331, 403-411.	2.3	52
43	Inventory of the GH70 enzymes encoded by <i>Leuconostoc citreum</i> NRRL 1299 identification of three novel transglucosylases. <i>FEBS Journal</i> , 2015, 282, 2115-2130.	4.7	49
44	Lipase-catalyzed chemoselective n-acylation of amino-sugar derivatives in hydrophobic solvent: Acid-amine ion-pair effects. <i>Tetrahedron</i> , 1997, 53, 7587-7594.	1.9	48
45	Sucrose-Utilizing Transglucosidases for Biocatalysis. <i>Topics in Current Chemistry</i> , 2010, 294, 25-48.	4.0	48
46	Applying Pairwise Combinations of Amino Acid Mutations for Sorting Out Highly Efficient Glucosylation Tools for Chemo-Enzymatic Synthesis of Bacterial Oligosaccharides. <i>Journal of the American Chemical Society</i> , 2012, 134, 18677-18688.	13.7	48
47	Induction and Transcription Studies of the Dextransucrase Gene in <i>Leuconostoc mesenteroides</i> NRRL B-512F. <i>Applied and Environmental Microbiology</i> , 1999, 65, 5504-5509.	3.1	47
48	Control of Lipase Enantioselectivity by Engineering the Substrate Binding Site and Access Channel. <i>ChemBioChem</i> , 2009, 10, 2760-2771.	2.6	46
49	Characterization of the First $\alpha$ -(1 $\rightarrow$ 3) Branching Sucrases of the GH70 Family. <i>Journal of Biological Chemistry</i> , 2016, 291, 7687-7702.	3.4	45
50	Cloning and sequencing of a gene coding for an extracellular dextransucrase (DSRB) from <i>Leuconostoc mesenteroides</i> NRRL B-1299 synthesizing only a $\beta$ -D-glucan. <i>FEMS Microbiology Letters</i> , 1998, 159, 307-315.	1.8	44
51	Conserved Repeat Motifs and Glucan Binding by Glucansucrases of Oral Streptococci and <i>Leuconostoc mesenteroides</i> . <i>Journal of Bacteriology</i> , 2004, 186, 8301-8308.	2.2	44
52	New Efficient Recombinant Expression System To Engineer <i>Candida antarctica</i> Lipase B. <i>Applied and Environmental Microbiology</i> , 2010, 76, 2684-2687.	3.1	44
53	GH13 amylosucrases and GH70 branching sucrases, atypical enzymes in their respective families. <i>Cellular and Molecular Life Sciences</i> , 2016, 73, 2661-2679.	5.4	44
54	Identification of key amino acid residues in <i>Neisseria polysaccharea</i> amylosucrase. <i>FEBS Letters</i> , 2000, 474, 33-37.	2.8	43

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55	Laccases from Marine Organisms and Their Applications in the Biodegradation of Toxic and Environmental Pollutants: a Review. <i>Applied Biochemistry and Biotechnology</i> , 2019, 187, 583-611.	2.9	42
56	Dextranase production by <i>Leuconostoc mesenteroides</i> NRRL B-1299. Comparison with <i>L. mesenteroides</i> NRRL B-512F. <i>Enzyme and Microbial Technology</i> , 1997, 20, 523-530.	3.2	41
57	Characterisation of the activator effect of glycogen on amylosucrase from <i>Neisseria polysaccharea</i> . <i>FEMS Microbiology Letters</i> , 2000, 186, 103-108.	1.8	41
58	Towards a novel explanation of <i>Pseudomonas cepacia</i> lipase enantioselectivity via molecular modelling of the enantiomer trajectory into the active site. <i>Tetrahedron: Asymmetry</i> , 2003, 14, 1807-1817.	1.8	41
59	Construction of a fully active truncated alternansucrase partially deleted of its carboxy-terminal domain. <i>FEBS Letters</i> , 2006, 580, 763-768.	2.8	41
60	Cloning, purification and characterization of a thermostable amylosucrase from <i>Deinococcus geothermalis</i> . <i>FEMS Microbiology Letters</i> , 2008, 285, 25-32.	1.8	41
61	Sucrose analogs: an attractive (bio)source for glycodiversification. <i>Natural Product Reports</i> , 2012, 29, 945.	10.3	40
62	Polymeric Iminosugars Improve the Activity of Carbohydrate-Processing Enzymes. <i>Bioconjugate Chemistry</i> , 2015, 26, 766-772.	3.6	40
63	Glucosylation of $\beta$ -butyl- and $\beta$ -octyl-D-glucopyranosides by dextranase and alternansucrase from <i>Leuconostoc mesenteroides</i> . <i>Carbohydrate Research</i> , 2003, 338, 855-864.	2.3	38
64	Investigations on the Determinants Responsible for Low Molar Mass Dextran Formation by DSR-M Dextranase. <i>ACS Catalysis</i> , 2017, 7, 7106-7119.	11.2	37
65	Harnessing glycoenzyme engineering for synthesis of bioactive oligosaccharides. <i>Interface Focus</i> , 2019, 9, 20180069.	3.0	37
66	Increased amylosucrase activity and specificity, and identification of regions important for activity, specificity and stability through molecular evolution. <i>FEBS Journal</i> , 2006, 273, 673-681.	4.7	35
67	A Structure-Controlled Investigation of Lipase Enantioselectivity by a Path-Planning Approach. <i>ChemBioChem</i> , 2008, 9, 1308-1317.	2.6	35
68	Combinatorial engineering to enhance thermostability of amylosucrase. <i>Protein Science</i> , 2008, 17, 967-976.	7.6	33
69	Synthesis of dextrans with controlled amounts of $\beta$ -1,2 linkages using the transglucosidase GBD $\alpha$ -CD2. <i>Applied Microbiology and Biotechnology</i> , 2010, 86, 545-554.	3.6	33
70	Enzymatic amidification for the synthesis of biodegradable surfactants: Synthesis of N-acylated hydroxylated amines. <i>Journal of Molecular Catalysis B: Enzymatic</i> , 1998, 5, 13-17.	1.8	32
71	Lipase-catalyzed enantioselective transesterification toward esters of 2-bromo-tolylacetic acids. <i>Tetrahedron: Asymmetry</i> , 2003, 14, 317-323.	1.8	32
72	Encapsulation in Lentikats of Dextranase from <i>Leuconostoc mesenteroides</i> NRRL B-1299, and its Effect on Product Selectivity. <i>Biocatalysis and Biotransformation</i> , 2003, 21, 325-331.	2.0	32

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73	Isolation and chemoenzymatic treatment of glycoalkaloids from green, sprouting and rotting <i>Solanum tuberosum</i> potatoes for solanidine recovery. <i>Food Chemistry</i> , 2017, 220, 257-265.	8.2	32
74	Title is missing!. <i>Biotechnology Letters</i> , 1999, 13, 749-755.	0.5	30
75	Characterization of dextransucrases from <i>Leuconostoc mesenteroides</i> NRRL B-1299. <i>Applied Biochemistry and Biotechnology</i> , 1997, 62, 47-59.	2.9	28
76	High-level production and purification of a fully active recombinant dextransucrase from <i>Leuconostoc mesenteroides</i> NRRL B-512F. <i>FEMS Microbiology Letters</i> , 2006, 261, 203-210.	1.8	28
77	NMR-Based Structural Glycomics for High-Throughput Screening of Carbohydrate-Active Enzyme Specificity. <i>Analytical Chemistry</i> , 2011, 83, 1202-1206.	6.5	28
78	Extending the Structural Diversity of Flavonoid Glycosides with Engineered Glucansucrases. <i>ChemCatChem</i> , 2014, 6, 2282-2291.	3.7	28
79	Lipase-catalysed synthesis of biosurfactants by transacylation of N-methyl-glucamine and fatty-acid methyl esters. <i>Tetrahedron</i> , 1997, 53, 7629-7634.	1.9	27
80	Deciphering an Undecided Enzyme: Investigations of the Structural Determinants Involved in the Linkage Specificity of Alternansucrase. <i>ACS Catalysis</i> , 2019, 9, 2222-2237.	11.2	27
81	Structure and Property Engineering of $\alpha$ -D-Glucans Synthesized by Dextransucrase Mutants. <i>Biomacromolecules</i> , 2012, 13, 187-195.	5.4	26
82	Computer-Aided Engineering of a Transglycosylase for the Glucosylation of an Unnatural Disaccharide of Relevance for Bacterial Antigen Synthesis. <i>ACS Catalysis</i> , 2015, 5, 1186-1198.	11.2	26
83	A dextran with unique rheological properties produced by the dextransucrase from <i>Oenococcus oeni</i> DSM 17330. <i>Carbohydrate Polymers</i> , 2018, 179, 10-18.	10.2	26
84	Maltooligosaccharide disproportionation reaction: an intrinsic property of amylosucrase from <i>Neisseria polysaccharea</i> . <i>FEBS Letters</i> , 2002, 527, 67-70.	2.8	25
85	Genome Sequence of <i>Weissella confusa</i> LBAE C39-2, Isolated from a Wheat Sourdough. <i>Journal of Bacteriology</i> , 2012, 194, 1608-1609.	2.2	25
86	Crystallization and preliminary X-ray studies of recombinant amylosucrase from <i>Neisseria polysaccharea</i> . <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2000, 56, 203-205.	2.5	24
87	A novel family of glucosyl 1,5-anhydro-d-fructose derivatives synthesised by transglucosylation with dextransucrase from <i>Leuconostoc mesenteroides</i> NRRL B-512F. <i>Carbohydrate Research</i> , 2005, 340, 395-401.	2.3	24
88	Factors affecting $\alpha$ -1,2 glucooligosaccharide synthesis by <i>Leuconostoc mesenteroides</i> NRRL B-1299 dextransucrase. <i>Biotechnology and Bioengineering</i> , 2001, 74, 498-504.	3.3	23
89	The structure of amylosucrase from <i>Deinococcus radiodurans</i> has an unusual open active-site topology. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2013, 69, 973-978.	0.7	23
90	Probing impact of active site residue mutations on stability and activity of <i>Neisseria polysaccharea</i> amylosucrase. <i>Protein Science</i> , 2013, 22, 1754-1765.	7.6	23

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91	Structural Insights into the Carbohydrate Binding Ability of an Î±-(1â†’2) Branching Sucrase from Glycoside Hydrolase Family 70. <i>Journal of Biological Chemistry</i> , 2016, 291, 7527-7540.	3.4	23
92	Optimized and Automated Protocols for High-Throughput Screening of Amylosucrase Libraries. <i>Journal of Biomolecular Screening</i> , 2007, 12, 715-723.	2.6	22
93	Genome Sequences of Three <i>Leuconostoc citreum</i> Strains, LBAE C10, LBAE C11, and LBAE E16, Isolated from Wheat Sourdoughs. <i>Journal of Bacteriology</i> , 2012, 194, 1610-1611.	2.2	22
94	Evaluation of dough rheological properties and bread texture of pearl millet-wheat flour mix. <i>Journal of Food Science and Technology</i> , 2016, 53, 2061-2066.	2.8	22
95	Towards the molecular understanding of glycogen elongation by amylosucrase. <i>Proteins: Structure, Function and Bioinformatics</i> , 2006, 66, 118-126.	2.6	21
96	Capillary electrophoresis analysis of glucooligosaccharide regioisomers. <i>Electrophoresis</i> , 2004, 25, 861-869.	2.4	20
97	Programmed chemo-enzymatic synthesis of the oligosaccharide component of a carbohydrate-based antibacterial vaccine candidate. <i>Chemical Communications</i> , 2015, 51, 2581-2584.	4.1	20
98	Isolation of a Gene from <i>Leuconostoc citreum</i> B/110-1-2 Encoding a Novel Dextranucrase Enzyme. <i>Current Microbiology</i> , 2011, 62, 1260-1266.	2.2	19
99	Probing Substrate Promiscuity of Amylosucrase from <i>Neisseria polysaccharea</i> . <i>ChemCatChem</i> , 2013, 5, 2288-2295.	3.7	19
100	Natural and engineered transglycosylases: Green tools for the enzyme-based synthesis of glycoproducts. <i>Current Opinion in Chemical Biology</i> , 2021, 61, 96-106.	6.1	19
101	Bacterial Î±-Glucan and Branching Sucasres from GH70 Family: Discovery, Structureâ€“Function Relationship Studies and Engineering. <i>Microorganisms</i> , 2021, 9, 1607.	3.6	19
102	Understanding the Polymerization Mechanism of Glycoside-Hydrolase Family 70 Glucansucasres. <i>Journal of Biological Chemistry</i> , 2006, 281, 31254-31267.	3.4	19
103	One-step synthesis of isomalto-oligosaccharide syrups and dextrans of controlled size using engineered dextranucrase. <i>Biocatalysis and Biotransformation</i> , 2008, 26, 141-151.	2.0	18
104	Combinatorial Engineering of Dextranucrase Specificity. <i>PLoS ONE</i> , 2013, 8, e77837.	2.5	18
105	A mixed molecular modelingâ€“robotics approach to investigate lipase large molecular motions. <i>Proteins: Structure, Function and Bioinformatics</i> , 2011, 79, 2517-2529.	2.6	17
106	Engineering of <i>Candida antarctica</i> lipase B for poly(Î¼-caprolactone) synthesis. <i>European Polymer Journal</i> , 2017, 95, 809-819.	5.4	17
107	Fructosylation of phenolic compounds by levansucrase from <i>Gluconacetobacter diazotrophicus</i> . <i>Enzyme and Microbial Technology</i> , 2019, 122, 19-25.	3.2	17
108	Enzymatic synthesis of polysaccharide-based copolymers. <i>Green Chemistry</i> , 2018, 20, 4012-4022.	9.0	16

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109	Optimizing the production of an $\alpha$ -1,2 branching sucrose in <i>Escherichia coli</i> using statistical design. <i>Applied Microbiology and Biotechnology</i> , 2014, 98, 5173-5184.	3.6	15
110	Essential role of amino acid position 226 in oligosaccharide elongation by amylosucrase from <i>Neisseria polysaccharea</i> . <i>Biotechnology and Bioengineering</i> , 2014, 111, 1719-1728.	3.3	15
111	Engineering a branching sucrose for flavonoid glucoside diversification. <i>Scientific Reports</i> , 2018, 8, 15153.	3.3	15
112	Futile Encounter Engineering of the DSR-M Dextran Sucrase Modifies the Resulting Polymer Length. <i>Biochemistry</i> , 2019, 58, 2853-2859.	2.5	15
113	Neutral Genetic Drift-Based Engineering of a Sucrose-Utilizing Enzyme toward Glycodiversification. <i>ACS Catalysis</i> , 2019, 9, 1241-1252.	11.2	15
114	Kinetic modeling of oligosaccharide synthesis catalyzed by <i>Leuconostoc mesenteroides</i> NRRL B-1299 dextran sucrose. , 1999, 63, 308-315.		14
115	Resolution of 2-bromo-o-tolyl-carboxylic acid by transesterification using lipases from <i>Rhizomucor miehei</i> and <i>Pseudomonas cepacia</i> . <i>Tetrahedron: Asymmetry</i> , 2001, 12, 2473-2480.	1.8	14
116	Synthesis of L-Rhamnose and N -Acetyl-D-Glucosamine Derivatives Entering in the Composition of Bacterial Polysaccharides by Use of Glucansucrases. <i>Journal of Carbohydrate Chemistry</i> , 2009, 28, 142-160.	1.1	14
117	Branching pattern of gluco-oligosaccharides and 1.5kDa dextran grafted by the $\alpha$ -1,2 branching sucrose GBD-CD2. <i>Carbohydrate Polymers</i> , 2013, 94, 567-576.	10.2	14
118	Reactor optimization for $\alpha$ -1,2 glucooligosaccharide synthesis by immobilized dextran sucrose. <i>Biotechnology and Bioengineering</i> , 2001, 75, 276-284.	3.3	13
119	A novel dextran sucrose is produced by <i>Leuconostoc citreum</i> strain B/110-1-2: an isolate used for the industrial production of dextran and dextran derivatives. <i>Journal of Industrial Microbiology and Biotechnology</i> , 2011, 38, 1499-1506.	3.0	13
120	Functionalization of natural compounds by enzymatic fructosylation. <i>Applied Microbiology and Biotechnology</i> , 2017, 101, 5223-5234.	3.6	13
121	Convergent Chemoenzymatic Strategy to Deliver a Diversity of <i>Shigella flexneri</i> Serotype-Specific O-Antigen Segments from a Unique Lightly Protected Tetrasaccharide Core. <i>Journal of Organic Chemistry</i> , 2021, 86, 2058-2075.	3.2	13
122	Crystal structure of the Glu328Gln mutant of <i>Neisseria polysaccharea</i> amylosucrase in complex with sucrose and maltoheptaose. <i>Biocatalysis and Biotransformation</i> , 2006, 24, 99-105.	2.0	12
123	Search for a dextran sucrose minimal motif involved in dextran binding. <i>FEBS Letters</i> , 2007, 581, 4675-4680.	2.8	12
124	Macromolecular structure and film properties of enzymatically-engineered high molar mass dextrans. <i>Carbohydrate Polymers</i> , 2018, 181, 337-344.	10.2	12
125	Chapter 28. Successes in engineering glucansucrases to enhance glycodiversification. <i>Carbohydrate Chemistry</i> , 2014, , 624-645.	0.3	12
126	A laundry detergent compatible lichenase: Statistical optimization for production under solid state fermentation on crude millet. <i>Industrial Crops and Products</i> , 2013, 43, 349-354.	5.2	11



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127	Overview of the glucansucrase equipment of <i>Leuconostoc citreum</i> LBAE-E16 and LBAE-C11, two strains isolated from sourdough. <i>FEMS Microbiology Letters</i> , 2015, 362, 1-8.	1.8	11
128	Synthetic Derivatives of (+)- <i>epi</i> - $\alpha$ -Bisabolol Are Formed by Mammalian Cytochromes P450 Expressed in a Yeast Reconstituted Pathway. <i>ACS Synthetic Biology</i> , 2020, 9, 368-380.	3.8	10
129	A specific oligosaccharide-binding site in the alternansucrase catalytic domain mediates alternan elongation. <i>Journal of Biological Chemistry</i> , 2020, 295, 9474-9489.	3.4	9
130	Processivity of dextransucrases synthesizing very-high-molar-mass dextran is mediated by sugar-binding pockets in domain V. <i>Journal of Biological Chemistry</i> , 2020, 295, 5602-5613.	3.4	9
131	One-pot bi-enzymatic cascade synthesis of puerarin polyfructosides. <i>Carbohydrate Polymers</i> , 2020, 247, 116710.	10.2	9
132	A pH-Based High-Throughput Screening of Sucrose-Utilizing Transglucosidases for the Development of Enzymatic Glucosylation Tools. <i>ChemCatChem</i> , 2010, 2, 969-975.	3.7	8
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