

Lyann Sim

List of Publications by Year in descending order

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30
papers

1,369
citations

471509

17
h-index

454955

30
g-index

30
all docs

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docs citations

30
times ranked

1397
citing authors

#	ARTICLE	IF	CITATIONS
1	Mammalian sialyltransferases allow efficient <i>Escherichia coli</i> -based production of mucin-type O-glycoproteins but can also transfer Kdo. <i>Glycobiology</i> , 2022, 32, 429-440.	2.5	2
2	7-Fluorosialyl Glycosides Are Hydrolysis Resistant but Readily Assembled by Sialyltransferases Providing Easy Access to More Metabolically Stable Glycoproteins. <i>ACS Central Science</i> , 2021, 7, 345-354.	11.3	16
3	Discovery and Development of Promiscuous O-Glycan Hydrolases for Removal of Intact Sialyl T-Antigen. <i>ACS Chemical Biology</i> , 2021, 16, 2004-2015.	3.4	7
4	Prevention of vascular-allograft rejection by protecting the endothelial glycocalyx with immunosuppressive polymers. <i>Nature Biomedical Engineering</i> , 2021, 5, 1202-1216.	22.5	12
5	Comparison of α 2,6-sialyltransferases for sialylation of therapeutic proteins. <i>Glycobiology</i> , 2019, 29, 735-747.	2.5	7
6	An enzymatic pathway in the human gut microbiome that converts A to universal O type blood. <i>Nature Microbiology</i> , 2019, 4, 1475-1485.	13.3	56
7	Directed evolution of bacterial polysialyltransferases. <i>Glycobiology</i> , 2019, 29, 588-598.	2.5	8
8	A Bacterial Expression Platform for Production of Therapeutic Proteins Containing Human-like O-Linked Glycans. <i>Cell Chemical Biology</i> , 2019, 26, 203-212.e5.	5.2	35
9	Characterization of a thermostable endoglucanase from <i>Cellulomonas fimi</i> ATCC484. <i>Biochemistry and Cell Biology</i> , 2018, 96, 68-76.	2.0	10
10	X-ray crystallographic structure of a bacterial polysialyltransferase provides insight into the biosynthesis of capsular polysialic acid. <i>Scientific Reports</i> , 2017, 7, 5842.	3.3	13
11	Structural and biochemical characterization of the N-terminal domain of flocculin LgFlo1p from <i>Saccharomyces pastorianus</i> reveals a unique specificity for phosphorylated mannose. <i>FEBS Journal</i> , 2013, 280, 1073-1083.	4.7	18
12	Unexpected High Digestion Rate of Cooked Starch by the Ct-Maltase-Glucoamylase Small Intestine Mucosal α -Glucosidase Subunit. <i>PLoS ONE</i> , 2012, 7, e35473.	2.5	43
13	Mapping the intestinal alpha-glucogenic enzyme specificities of starch digesting maltase-glucoamylase and sucrase-isomaltase. <i>Bioorganic and Medicinal Chemistry</i> , 2011, 19, 3929-3934.	3.0	69
14	Structural Basis for Substrate Selectivity in Human Maltase-Glucoamylase and Sucrase-Isomaltase N-terminal Domains. <i>Journal of Biological Chemistry</i> , 2010, 285, 17763-17770.	3.4	173
15	New Glucosidase Inhibitors from an Ayurvedic Herbal Treatment for Type 2 Diabetes: Structures and Inhibition of Human Intestinal Maltase-Glucoamylase with Compounds from <i>Salacia reticulata</i> . <i>Biochemistry</i> , 2010, 49, 443-451.	2.5	134
16	Specific starch digestion of maize alpha-limit dextrins by recombinant mucosal glucosidase enzymes. <i>FASEB Journal</i> , 2010, 24, 231.6.	0.5	1
17	Total Syntheses of Casuarine and Its α -Glucoside: Complementary Inhibition towards Glycoside Hydrolases of the GH31 and GH37 Families. <i>Chemistry - A European Journal</i> , 2009, 15, 1627-1636.	3.3	92
18	Synthesis of 2-deoxy-2-fluoro and 1,2-ene derivatives of the naturally occurring glycosidase inhibitor, salacinol, and their inhibitory activities against recombinant human maltase glucoamylase. <i>Carbohydrate Research</i> , 2008, 343, 951-956.	2.3	5

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19	Human Intestinal Maltaseâ€“Glucoamylase: Crystal Structure of the N-Terminal Catalytic Subunit and Basis of Inhibition and Substrate Specificity. <i>Journal of Molecular Biology</i> , 2008, 375, 782-792.	4.2	248
20	Studies Directed toward the Stereochemical Structure Determination of the Naturally Occurring Glucosidase Inhibitor, Kotalanol: Synthesis and Inhibitory Activities against Human Maltase Glucoamylase of Seven-Carbon, Chain-Extended Homologues of Salacinol. <i>Journal of Organic Chemistry</i> , 2008, 73, 6172-6181.	3.2	43
21	Luminal Starch Substrate â€œBrakeâ€•on Maltase-Glucoamylase Activity Is Located within the Glucoamylase Subunit3. <i>Journal of Nutrition</i> , 2008, 138, 685-692.	2.9	81
22	New Synthetic Routes to Chain-Extended Selenium, Sulfur, and Nitrogen Analogues of the Naturally Occurring Glucosidase Inhibitor Salacinol and their Inhibitory Activities against Recombinant Human Maltase Glucoamylase. <i>Journal of Organic Chemistry</i> , 2007, 72, 6562-6572.	3.2	39
23	New Chain-Extended Analogues of Salacinol and Blintol and Their Glycosidase Inhibitory Activities. Mapping the Active-Site Requirements of Human Maltase Glucoamylase. <i>Journal of Organic Chemistry</i> , 2007, 72, 180-186.	3.2	32
24	Synthesis of S-alkylated sulfonium-ions and their glucosidase inhibitory activities against recombinant human maltase glucoamylase. <i>Carbohydrate Research</i> , 2007, 342, 901-912.	2.3	24
25	Synthesis and glycosidase inhibitory activities of chain-modified analogues of the glycosidase inhibitors salacinol and blintol. <i>Carbohydrate Research</i> , 2007, 342, 1888-1894.	2.3	8
26	Evidence of native starch degradation with human small intestinal maltaseâ€“glucoamylase (recombinant). <i>FEBS Letters</i> , 2007, 581, 2381-2388.	2.8	58
27	Synthesis of analogues of salacinol containing a carboxylate inner salt and their inhibitory activities against human maltase glucoamylase. <i>Carbohydrate Research</i> , 2007, 342, 1661-1667.	2.3	10
28	A New Class of Glucosidase Inhibitor:â€• Analogues of the Naturally Occurring Glucosidase Inhibitor Salacinol with Different Ring Heteroatom Substituents and Acyclic Chain Extension. <i>Journal of Organic Chemistry</i> , 2006, 71, 3007-3013.	3.2	37
29	Inhibition of recombinant human maltase glucoamylase by salacinol and derivatives. <i>FEBS Journal</i> , 2006, 273, 2673-2683.	4.7	74
30	Synthesis, enzymatic activity, and X-ray crystallography of an unusual class of amino acids. <i>Bioorganic and Medicinal Chemistry</i> , 2006, 14, 8332-8340.	3.0	14