

Tracey M Gloster

List of Publications by Year in descending order

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Version: 2024-02-01

39
papers

2,714
citations

236912

25
h-index

289230

40
g-index

44
all docs

44
docs citations

44
times ranked

3216
citing authors

#	ARTICLE	IF	CITATIONS
1	Hijacking a biosynthetic pathway yields a glycosyltransferase inhibitor within cells. <i>Nature Chemical Biology</i> , 2011, 7, 174-181.	8.0	291
2	Recent structural insights into the expanding world of carbohydrate-active enzymes. <i>Current Opinion in Structural Biology</i> , 2005, 15, 637-645.	5.7	264
3	Glycosidase inhibition: assessing mimicry of the transition state. <i>Organic and Biomolecular Chemistry</i> , 2010, 8, 305-320.	2.8	217
4	Developing inhibitors of glycan processing enzymes as tools for enabling glycobiology. <i>Nature Chemical Biology</i> , 2012, 8, 683-694.	8.0	159
5	Iminosugar Glycosidase Inhibitors: A Structural and Thermodynamic Dissection of the Binding of Isofagomine and 1-Deoxynojirimycin to β -Glucosidases. <i>Journal of the American Chemical Society</i> , 2003, 125, 14313-14323.	13.7	154
6	An anti-CRISPR viral ring nuclease subverts type III CRISPR immunity. <i>Nature</i> , 2020, 577, 572-575.	27.8	139
7	Structural snapshots of the reaction coordinate for O-GlcNAc transferase. <i>Nature Chemical Biology</i> , 2012, 8, 966-968.	8.0	132
8	Advances in understanding glycosyltransferases from a structural perspective. <i>Current Opinion in Structural Biology</i> , 2014, 28, 131-141.	5.7	127
9	Glycosidase Inhibition: An Assessment of the Binding of 18 Putative Transition-State Mimics. <i>Journal of the American Chemical Society</i> , 2007, 129, 2345-2354.	13.7	124
10	Characterization and Three-dimensional Structures of Two Distinct Bacterial Xyloglucanases from Families GH5 and GH12. <i>Journal of Biological Chemistry</i> , 2007, 282, 19177-19189.	3.4	103
11	Structure and mechanism of a Type III CRISPR defence DNA nuclease activated by cyclic oligoadenylate. <i>Nature Communications</i> , 2020, 11, 500.	12.8	97
12	Divergence of Catalytic Mechanism within a Glycosidase Family Provides Insight into Evolution of Carbohydrate Metabolism by Human Gut Flora. <i>Chemistry and Biology</i> , 2008, 15, 1058-1067.	6.0	81
13	Structural and mechanistic insight into N-glycan processing by endo- β -mannosidase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 781-786.	7.1	74
14	Elevation of Global O-GlcNAc in Rodents Using a Selective O-GlcNAcase Inhibitor Does Not Cause Insulin Resistance or Perturb Glucohomeostasis. <i>Chemistry and Biology</i> , 2010, 17, 949-958.	6.0	71
15	Inhibition of O-GlcNAcase Using a Potent and Cell-Permeable Inhibitor Does Not Induce Insulin Resistance in 3T3-L1 Adipocytes. <i>Chemistry and Biology</i> , 2010, 17, 937-948.	6.0	67
16	Structural Studies of the β -Glycosidase from <i>Sulfolobus solfataricus</i> in Complex with Covalently and Noncovalently Bound Inhibitors. <i>Biochemistry</i> , 2004, 43, 6101-6109.	2.5	62
17	Metabolic Inhibitors of O-GlcNAc Transferase That Act In Vivo Implicate Decreased O-GlcNAc Levels in Leptin-Mediated Nutrient Sensing. <i>Angewandte Chemie - International Edition</i> , 2018, 57, 7644-7648.	13.8	56
18	Mechanism, Structure, and Inhibition of O-GlcNAc Processing Enzymes. <i>Current Signal Transduction Therapy</i> , 2010, 5, 74-91.	0.5	54

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19	Structural, Kinetic, and Thermodynamic Analysis of Glucoimidazole-Derived Glycosidase Inhibitors. <i>Biochemistry</i> , 2006, 45, 11879-11884.	2.5	47
20	The CRISPR ancillary effector Can2 is a dual-specificity nuclease potentiating type III CRISPR defence. <i>Nucleic Acids Research</i> , 2021, 49, 2777-2789.	14.5	46
21	Structural basis for cyclophellitol inhibition of a Î²-glucosidase. <i>Organic and Biomolecular Chemistry</i> , 2007, 5, 444-446.	2.8	45
22	Structural, Thermodynamic, and Kinetic Analyses of Tetrahydrooxazine-derived Inhibitors Bound to Î²-Glucosidases. <i>Journal of Biological Chemistry</i> , 2004, 279, 49236-49242.	3.4	37
23	Dissection of Conformationally Restricted Inhibitors Binding to a Î²-Glucosidase. <i>ChemBioChem</i> , 2006, 7, 738-742.	2.6	34
24	Structural Snapshots for Mechanism-Based Inactivation of a Glycoside Hydrolase by Cyclopropyl Carbasugars. <i>Angewandte Chemie - International Edition</i> , 2016, 55, 14978-14982.	13.8	30
25	Revealing the mechanism for covalent inhibition of glycoside hydrolases by carbasugars at an atomic level. <i>Nature Communications</i> , 2018, 9, 3243.	12.8	28
26	Atomic resolution analyses of the binding of xylobiose-derived deoxynojirimycin and isofagomine to xylanase Xyn10A. Electronic supplementary information (ESI) available: kinetics and structural methods. See http://www.rsc.org/suppdata/cc/b4/b405152a/ . <i>Chemical Communications</i> , 2004, , 1794.	4.1	26
27	Tetramerisation of the CRISPR ring nuclease Crn3/Csx3 facilitates cyclic oligoadenylate cleavage. <i>ELife</i> , 2020, 9, .	6.0	22
28	Sialidase and Sialyltransferase Inhibitors: Targeting Pathogenicity and Disease. <i>Frontiers in Molecular Biosciences</i> , 2021, 8, 705133.	3.5	19
29	Linear Eyring Plots Conceal a Change in the Rate-Limiting Step in an Enzyme Reaction. <i>Biochemistry</i> , 2018, 57, 6757-6761.	2.5	16
30	Development of inhibitors as research tools for carbohydrate-processing enzymes. <i>Biochemical Society Transactions</i> , 2012, 40, 913-928.	3.4	15
31	A xylobiose-derived isofagomine lactam glycosidase inhibitor binds as its amide tautomer. Electronic supplementary information (ESI) available: details of data and structure quality for complex of 1 with Xyn10A. See http://www.rsc.org/suppdata/cc/b3/b301829f/ . <i>Chemical Communications</i> , 2003, , 944-945.	4.1	13
32	Exploitation of carbohydrate processing enzymes in biocatalysis. <i>Current Opinion in Chemical Biology</i> , 2020, 55, 180-188.	6.1	11
33	New Irreversible Î±-D-Glucuronidase Inhibitors and Activity-Based Probes. <i>Chemistry - A European Journal</i> , 2018, 24, 19081-19088.	3.3	9
34	Dissecting the Mechanism of (R)-3-Hydroxybutyrate Dehydrogenase by Kinetic Isotope Effects, Protein Crystallography, and Computational Chemistry. <i>ACS Catalysis</i> , 2020, 10, 15019-15032.	11.2	8
35	Analysis of the product streams obtained on butanosolv pretreatment of draff. <i>Biomass and Bioenergy</i> , 2020, 141, 105680.	5.7	8
36	Structural Snapshots for Mechanism-Based Inactivation of a Glycoside Hydrolase by Cyclopropyl Carbasugars. <i>Angewandte Chemie</i> , 2016, 128, 15202-15206.	2.0	7

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37	Metabolic Inhibitors of O ⁶ -GlcNAc Transferase That Act In ^{Vivo} Implicate Decreased O ⁶ -GlcNAc Levels in Leptin ⁻ Mediated Nutrient Sensing. <i>Angewandte Chemie</i> , 2018, 130, 7770-7774.	2.0	7
38	Structure, dynamics, and molecular inhibition of the <i>Staphylococcus aureus</i> m1A22-tRNA methyltransferase TrmK. <i>Journal of Biological Chemistry</i> , 2022, 298, 102040.	3.4	4
39	Kinetic and Structural Characterization of Sialidases (Kdnases) from Ascomycete Fungal Pathogens. <i>ACS Chemical Biology</i> , 2021, 16, 2632-2640.	3.4	1