

David J Vocadlo

List of Publications by Year in descending order

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170
papers

11,832
citations

23567

58
h-index

30087

103
g-index

178
all docs

178
docs citations

178
times ranked

9006
citing authors

#	ARTICLE	IF	CITATIONS
1	Catalysis by hen egg-white lysozyme proceeds via a covalent intermediate. <i>Nature</i> , 2001, 412, 835-838.	27.8	588
2	A potent mechanism-inspired O-GlcNAcase inhibitor that blocks phosphorylation of tau in vivo. <i>Nature Chemical Biology</i> , 2008, 4, 483-490.	8.0	576
3	A chemical approach for identifying O-GlcNAc-modified proteins in cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003, 100, 9116-9121.	7.1	496
4	Increasing O-GlcNAc slows neurodegeneration and stabilizes tau against aggregation. <i>Nature Chemical Biology</i> , 2012, 8, 393-399.	8.0	493
5	Mechanistic insights into glycosidase chemistry. <i>Current Opinion in Chemical Biology</i> , 2008, 12, 539-555.	6.1	363
6	O-GlcNAcase Uses Substrate-assisted Catalysis. <i>Journal of Biological Chemistry</i> , 2005, 280, 25313-25322.	3.4	333
7	O-GlcNAcylation Regulates Cancer Metabolism and Survival Stress Signaling via Regulation of the HIF-1 Pathway. <i>Molecular Cell</i> , 2014, 54, 820-831.	9.7	307
8	Hijacking a biosynthetic pathway yields a glycosyltransferase inhibitor within cells. <i>Nature Chemical Biology</i> , 2011, 7, 174-181.	8.0	291
9	NAG-thiazoline, An N-Acetyl- β -hexosaminidase Inhibitor That Implicates Acetamido Participation. <i>Journal of the American Chemical Society</i> , 1996, 118, 6804-6805.	13.7	248
10	Crystallographic Evidence for Substrate-assisted Catalysis in a Bacterial β -Hexosaminidase. <i>Journal of Biological Chemistry</i> , 2001, 276, 10330-10337.	3.4	239
11	<i>Drosophila</i> O-GlcNAc transferase (OGT) is encoded by the Polycomb group (PcG) gene, <i>super sex combs</i> (<i>sxc</i>). <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 13427-13432.	7.1	214
12	O-GlcNAc and neurodegeneration: biochemical mechanisms and potential roles in Alzheimer's disease and beyond. <i>Chemical Society Reviews</i> , 2014, 43, 6839-6858.	38.1	209
13	Hyper-O-GlcNAcylation Is Anti-apoptotic and Maintains Constitutive NF- κ B Activity in Pancreatic Cancer Cells. <i>Journal of Biological Chemistry</i> , 2013, 288, 15121-15130.	3.4	205
14	The Emerging Link between O-GlcNAc and Alzheimer Disease. <i>Journal of Biological Chemistry</i> , 2014, 289, 34472-34481.	3.4	205
15	Structure and mechanism of a bacterial β -glucosaminidase having O-GlcNAcase activity. <i>Nature Structural and Molecular Biology</i> , 2006, 13, 365-371.	8.2	182
16	HCF-1 Is Cleaved in the Active Site of O-GlcNAc Transferase. <i>Science</i> , 2013, 342, 1235-1239.	12.6	162
17	Developing inhibitors of glycan processing enzymes as tools for enabling glycobiology. <i>Nature Chemical Biology</i> , 2012, 8, 683-694.	8.0	159
18	Analysis of PUGNAc and NAG-thiazoline as Transition State Analogues for Human O-GlcNAcase: A Mechanistic and Structural Insights into Inhibitor Selectivity and Transition State Poise. <i>Journal of the American Chemical Society</i> , 2007, 129, 635-644.	13.7	155

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19	Structural snapshots of the reaction coordinate for O-GlcNAc transferase. <i>Nature Chemical Biology</i> , 2012, 8, 966-968.	8.0	132
20	A Strategy for Functional Proteomic Analysis of Glycosidase Activity from Cell Lysates. <i>Angewandte Chemie - International Edition</i> , 2004, 43, 5338-5342.	13.8	131
21	Aspartate 313 in the <i>Streptomyces plicatus</i> Hexosaminidase Plays a Critical Role in Substrate-assisted Catalysis by Orienting the 2-Acetamido Group and Stabilizing the Transition State. <i>Journal of Biological Chemistry</i> , 2002, 277, 40055-40065.	3.4	126
22	O-GlcNAc processing enzymes: catalytic mechanisms, substrate specificity, and enzyme regulation. <i>Current Opinion in Chemical Biology</i> , 2012, 16, 488-497.	6.1	122
23	In Vivo Modulation of O-GlcNAc Levels Regulates Hippocampal Synaptic Plasticity through Interplay with Phosphorylation. <i>Journal of Biological Chemistry</i> , 2009, 284, 174-181.	3.4	115
24	Pharmacological inhibition of O-GlcNAcase (OGA) prevents cognitive decline and amyloid plaque formation in bigenic tau/APP mutant mice. <i>Molecular Neurodegeneration</i> , 2014, 9, 42.	10.8	114
25	O-GlcNAc occurs cotranslationally to stabilize nascent polypeptide chains. <i>Nature Chemical Biology</i> , 2015, 11, 319-325.	8.0	113
26	O-GlcNAc Modification of tau Directly Inhibits Its Aggregation without Perturbing the Conformational Properties of tau Monomers. <i>Journal of Molecular Biology</i> , 2014, 426, 1736-1752.	4.2	110
27	mTOR/MYC Axis Regulates O-GlcNAc Transferase Expression and O-GlcNAcylation in Breast Cancer. <i>Molecular Cancer Research</i> , 2015, 13, 923-933.	3.4	109
28	Identification of Asp174 and Asp175 as the Key Catalytic Residues of Human O-GlcNAcase by Functional Analysis of Site-Directed Mutants. <i>Biochemistry</i> , 2006, 45, 3835-3844.	2.5	107
29	Mechanism of Action and Identification of Asp242 as the Catalytic Nucleophile of <i>Vibrio furnisii</i> N-Acetyl- β -D-glucosaminidase Using 2-Acetamido-2-deoxy-5-fluoro- β -L-idopyranosyl Fluoride. <i>Biochemistry</i> , 2000, 39, 117-126.	2.5	106
30	Elevation of Global O-GlcNAc Levels in 3T3-L1 Adipocytes by Selective Inhibition of O-GlcNAcase Does Not Induce Insulin Resistance. <i>Journal of Biological Chemistry</i> , 2008, 283, 34687-34695.	3.4	106
31	Inhibition of O-GlcNAcase leads to elevation of O-GlcNAc tau and reduction of tauopathy and cerebrospinal fluid tau in rTg4510 mice. <i>Molecular Neurodegeneration</i> , 2017, 12, 39.	10.8	106
32	Increasing O-GlcNAc levels: An overview of small-molecule inhibitors of O-GlcNAcase. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2010, 1800, 107-121.	2.4	105
33	Small Molecule Inhibitors of a Glycoside Hydrolase Attenuate Inducible AmpC-mediated β -Lactam Resistance. <i>Journal of Biological Chemistry</i> , 2007, 282, 21382-21391.	3.4	103
34	Mapping O-GlcNAc modification sites on tau and generation of a site-specific O-GlcNAc tau antibody. <i>Amino Acids</i> , 2011, 40, 857-868.	2.7	103
35	Insights into O-Linked N-Acetylglucosamine ([O-9]O-GlcNAc) Processing and Dynamics through Kinetic Analysis of O-GlcNAc Transferase and O-GlcNAcase Activity on Protein Substrates. <i>Journal of Biological Chemistry</i> , 2012, 287, 15395-15408.	3.4	102
36	Detailed Comparative Analysis of the Catalytic Mechanisms of β -N-Acetylglucosaminidases from Families 3 and 20 of Glycoside Hydrolases. <i>Biochemistry</i> , 2005, 44, 12809-12818.	2.5	98

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37	Structure of an O-GlcNAc transferase homolog provides insight into intracellular glycosylation. <i>Nature Structural and Molecular Biology</i> , 2008, 15, 764-765.	8.2	98
38	Synthesis and Use of Mechanism-Based Protein-Profiling Probes for Retaining β -Glucosaminidases Facilitate Identification of <i>Pseudomonas aeruginosa</i> NagZ. <i>Journal of the American Chemical Society</i> , 2008, 130, 327-335.	13.7	95
39	Analysis of Keystone Enzyme in Agar Hydrolysis Provides Insight into the Degradation (of a Tj ETQq1 1 0.784314 rgBT /Overlock 10 T	8.4	89
40	Structural and functional insight into human O-GlcNAcase. <i>Nature Chemical Biology</i> , 2017, 13, 610-612.	8.0	88
41	Discovery of MK-8719, a Potent O-GlcNAcase Inhibitor as a Potential Treatment for Tauopathies. <i>Journal of Medicinal Chemistry</i> , 2019, 62, 10062-10097.	6.4	87
42	<i>Streptococcus pneumoniae</i> Endohexosaminidase D, Structural and Mechanistic Insight into Substrate-assisted Catalysis in Family 85 Glycoside Hydrolases. <i>Journal of Biological Chemistry</i> , 2009, 284, 11676-11689.	3.4	85
43	Substrate-Guided Front-Face Reaction Revealed by Combined Structural Snapshots and Metadynamics for the Polypeptide <i>N</i> -Acetylgalactosaminyltransferase...2. <i>Angewandte Chemie - International Edition</i> , 2014, 53, 8206-8210.	13.8	80
44	Structural and mechanistic insight into the basis of mucopolysaccharidosis IIIB. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 6560-6565.	7.1	79
45	The β -Lactamase Gene Regulator AmpR Is a Tetramer That Recognizes and Binds the d-Ala-d-Ala Motif of Its Repressor UDP-N-acetylmuramic Acid (MurNAc)-pentapeptide. <i>Journal of Biological Chemistry</i> , 2015, 290, 2630-2643.	3.4	77
46	Differential Effects of an O-GlcNAcase Inhibitor on Tau Phosphorylation. <i>PLoS ONE</i> , 2012, 7, e35277.	2.5	76
47	Elevation of Global O-GlcNAc in Rodents Using a Selective O-GlcNAcase Inhibitor Does Not Cause Insulin Resistance or Perturb Gluconeogenesis. <i>Chemistry and Biology</i> , 2010, 17, 949-958.	6.0	71
48	Visualizing the Reaction Coordinate of an O-GlcNAc Hydrolase. <i>Journal of the American Chemical Society</i> , 2010, 132, 1807-1809.	13.7	70
49	Crystal Structure of β -D-Xylosidase from <i>Thermoanaerobacterium saccharolyticum</i> , a Family 39 Glycoside Hydrolase. <i>Journal of Molecular Biology</i> , 2004, 335, 155-165.	4.2	69
50	O-GlcNAcase Catalyzes Cleavage of Thioglycosides without General Acid Catalysis. <i>Journal of the American Chemical Society</i> , 2005, 127, 17202-17203.	13.7	69
51	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
52	Active Site Plasticity within the Glycoside Hydrolase NagZ Underlies a Dynamic Mechanism of Substrate Distortion. <i>Chemistry and Biology</i> , 2012, 19, 1471-1482.	6.0	67
53	A divergent synthesis of 2-acyl derivatives of PUGNAc yields selective inhibitors of O-GlcNAcase. <i>Organic and Biomolecular Chemistry</i> , 2006, 4, 839.	2.8	65
54	Inactivation of the Glycoside Hydrolase NagZ Attenuates Antipseudomonal β -Lactam Resistance in <i>Pseudomonas aeruginosa</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2009, 53, 2274-2282.	3.2	65

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55	Analysis of a New Family of Widely Distributed Metal-independent β -Mannosidases Provides Unique Insight into the Processing of N-Linked Glycans. <i>Journal of Biological Chemistry</i> , 2011, 286, 15586-15596.	3.4	65
56	Mislocalization of TDP-43 in the G93A mutant SOD1 transgenic mouse model of ALS. <i>Neuroscience Letters</i> , 2009, 458, 70-74.	2.1	64
57	Molecular Basis for Inhibition of GH84 Glycoside Hydrolases by Substituted Azepanes: Conformational Flexibility Enables Probing of Substrate Distortion. <i>Journal of the American Chemical Society</i> , 2009, 131, 5390-5392.	13.7	62
58	NagZ Inactivation Prevents and Reverts β -Lactam Resistance, Driven by AmpD and PBP 4 Mutations, in <i>Pseudomonas aeruginosa</i> . <i>Antimicrobial Agents and Chemotherapy</i> , 2010, 54, 3557-3563.	3.2	61
59	Providing β -lactams a helping hand: targeting the AmpC β -lactamase induction pathway. <i>Future Microbiology</i> , 2011, 6, 1415-1427.	2.0	61
60	Inhibition of O-GlcNAcase by a gluco-configured nagstatin and a PUGNAc-imidazole hybrid inhibitor. <i>Chemical Communications</i> , 2006, , 4372-4374.	4.1	60
61	Characterization of a beta-N-acetylhexosaminidase and a beta-N-acetylglucosaminidase/beta-glucosidase from <i>Cellulomonas fimi</i> . <i>FEBS Journal</i> , 2006, 273, 2929-2941.	4.7	60
62	Fluorescence-Quenched Substrates for Live Cell Imaging of Human Glucocerebrosidase Activity. <i>Journal of the American Chemical Society</i> , 2015, 137, 1181-1189.	13.7	59
63	Post-translational O-GlcNAcylation is essential for nuclear pore integrity and maintenance of the pore selectivity filter. <i>Journal of Molecular Cell Biology</i> , 2016, 8, 2-16.	3.3	57
64	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
65	Mechanism, Structure, and Inhibition of O-GlcNAc Processing Enzymes. <i>Current Signal Transduction Therapy</i> , 2010, 5, 74-91.	0.5	54
66	A Case for Reverse Protonation: Identification of Glu160 as an Acid/Base Catalyst in <i>Thermoanaerobacterium saccharolyticum</i> β -Xylosidase and Detailed Kinetic Analysis of a Site-Directed Mutant. <i>Biochemistry</i> , 2002, 41, 9736-9746.	2.5	50
67	Structures of lactate dehydrogenase A (LDHA) in apo, ternary and inhibitor-bound forms. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2015, 71, 185-195.	2.5	49
68	Molecular Basis for G Protein Control of the Prokaryotic ATP Sulfurylase. <i>Molecular Cell</i> , 2006, 21, 109-122.	9.7	48
69	Crystal Structure of the AmpR Effector Binding Domain Provides Insight into the Molecular Regulation of Inducible AmpC β -Lactamase. <i>Journal of Molecular Biology</i> , 2010, 400, 998-1010.	4.2	48
70	Mechanism of <i>Thermoanaerobacterium saccharolyticum</i> β -Xylosidase: Kinetic Studies. <i>Biochemistry</i> , 2002, 41, 9727-9735.	2.5	47
71	AmpG Inactivation Restores Susceptibility of Pan- β -Lactam-Resistant <i>Pseudomonas aeruginosa</i> Clinical Strains. <i>Antimicrobial Agents and Chemotherapy</i> , 2011, 55, 1990-1996.	3.2	47
72	Pharmacological Inhibition of O-GlcNAcase Enhances Autophagy in Brain through an mTOR-Independent Pathway. <i>ACS Chemical Neuroscience</i> , 2018, 9, 1366-1379.	3.5	47

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73	Molecular mechanisms regulating O-linked N-acetylglucosamine (O-GlcNAc) processing enzymes. <i>Current Opinion in Chemical Biology</i> , 2019, 53, 131-144.	6.1	46
74	MK-8719, a Novel and Selective <i>O</i> -GlcNAcase Inhibitor That Reduces the Formation of Pathological Tau and Ameliorates Neurodegeneration in a Mouse Model of Tauopathy. <i>Journal of Pharmacology and Experimental Therapeutics</i> , 2020, 374, 252-263.	2.5	45
75	O-GlcNAc posttranslational modifications regulate the entry of neurons into an axon branching program. <i>Developmental Neurobiology</i> , 2009, 69, 162-173.	3.0	43
76	Insight into a strategy for attenuating AmpC-mediated β -lactam resistance: Structural basis for selective inhibition of the glycoside hydrolase NagZ. <i>Protein Science</i> , 2009, 18, 1541-1551.	7.6	43
77	Quinolinic Acid Amyloid-like Fibrillar Assemblies Seed β -Synuclein Aggregation. <i>Journal of Molecular Biology</i> , 2018, 430, 3847-3862.	4.2	43
78	Biochemical and Structural Assessment of the 1-N-Azasugar GalNAc-isofagomine as a Potent Family 20 β -N-Acetylhexosaminidase Inhibitor. <i>Journal of Biological Chemistry</i> , 2001, 276, 42131-42137.	3.4	42
79	The Conformation and Function of a Multimodular Glycogen-Degrading Pneumococcal Virulence Factor. <i>Structure</i> , 2011, 19, 640-651.	3.3	42
80	Metabolic Inhibition of Sialyl-Lewis X Biosynthesis by 5-Thiofucose Remodels the Cell Surface and Impairs Selectin-Mediated Cell Adhesion*. <i>Journal of Biological Chemistry</i> , 2012, 287, 40021-40030.	3.4	42
81	Identification of Glu-277 as the catalytic nucleophile of <i>Thermoanaerobacterium saccharolyticum</i> β -xylosidase using electrospray MS. <i>Biochemical Journal</i> , 1998, 335, 449-455.	3.7	41
82	Differential Recognition and Hydrolysis of Host Carbohydrate Antigens by <i>Streptococcus pneumoniae</i> Family 98 Glycoside Hydrolases. <i>Journal of Biological Chemistry</i> , 2009, 284, 26161-26173.	3.4	41
83	A Convenient Approach to Stereoisomeric Iminocyclitols: Generation of Potent Brain-Permeable OGA Inhibitors. <i>Angewandte Chemie - International Edition</i> , 2015, 54, 15429-15433.	13.8	41
84	A Selective Inhibitor GalPUGNAc of Human Lysosomal β -Hexosaminidases Modulates Levels of the Ganglioside GM2 in Neuroblastoma Cells. <i>Angewandte Chemie - International Edition</i> , 2009, 48, 1300-1303.	13.8	39
85	Reduced protein O-glycosylation in the nervous system of the mutant SOD1 transgenic mouse model of amyotrophic lateral sclerosis. <i>Neuroscience Letters</i> , 2012, 516, 296-301.	2.1	39
86	Direct One-Step Fluorescent Labeling of <i>O</i> -GlcNAc-Modified Proteins in Live Cells Using Metabolic Intermediates. <i>Journal of the American Chemical Society</i> , 2018, 140, 15300-15308.	13.7	39
87	Inhibition of the Pneumococcal Virulence Factor StrH and Molecular Insights into N-Glycan Recognition and Hydrolysis. <i>Structure</i> , 2011, 19, 1603-1614.	3.3	38
88	Tools for probing and perturbing O-GlcNAc in cells and in vivo. <i>Current Opinion in Chemical Biology</i> , 2013, 17, 719-728.	6.1	38
89	The Development of Selective Inhibitors of NagZ: Increased Susceptibility of Gram-Negative Bacteria to β -Lactams. <i>ChemBioChem</i> , 2013, 14, 1973-1981.	2.6	38
90	A 1-acetamido derivative of 6-epi-valienamine: an inhibitor of a diverse group of β -N-acetylglucosaminidases. <i>Organic and Biomolecular Chemistry</i> , 2007, 5, 3013.	2.8	37

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91	Mammalian Notch is modified by d-Xyl-1-3-d-Xyl-1-3-d-Glc-1-O-Ser: Implementation of a method to study O-glycosylation. <i>Glycobiology</i> , 2010, 20, 287-299.	2.5	37
92	Metabolism of Vertebrate Amino Sugars with N-Glycolyl Groups. <i>Journal of Biological Chemistry</i> , 2012, 287, 28898-28916.	3.4	37
93	Characterization of the Glu and Asp Residues in the Active Site of Human β -Hexosaminidase B. <i>Biochemistry</i> , 2001, 40, 2201-2209.	2.5	36
94	Probing Synergy between Two Catalytic Strategies in the Glycoside Hydrolase <i>O</i> -GlcNAcase Using Multiple Linear Free Energy Relationships. <i>Journal of the American Chemical Society</i> , 2009, 131, 13415-13422.	13.7	36
95	Selective trihydroxyazepane NagZ inhibitors increase sensitivity of <i>Pseudomonas aeruginosa</i> to β -lactams. <i>Chemical Communications</i> , 2013, 49, 10983.	4.1	36
96	Functional analysis of a group A streptococcal glycoside hydrolase Spy1600 from family 84 reveals it is a β -N-acetylglucosaminidase and not a hyaluronidase. <i>Biochemical Journal</i> , 2006, 399, 241-247.	3.7	35
97	Enzymatic characterization and inhibition of the nuclear variant of human <i>O</i> -GlcNAcase. <i>Carbohydrate Research</i> , 2009, 344, 1079-1084.	2.3	34
98	Catalytic Promiscuity of <i>O</i> -GlcNAc Transferase Enables Unexpected Metabolic Engineering of Cytoplasmic Proteins with 2-Azido-2-deoxy-glucose. <i>ACS Chemical Biology</i> , 2017, 12, 206-213.	3.4	34
99	Analysis of transition state mimicry by tight binding aminothiazoline inhibitors provides insight into catalysis by human <i>O</i> -GlcNAcase. <i>Chemical Science</i> , 2016, 7, 3742-3750.	7.4	33
100	Genome-wide chemical mapping of <i>O</i> -GlcNAcylated proteins in <i>Drosophila melanogaster</i> . <i>Nature Chemical Biology</i> , 2017, 13, 161-167.	8.0	33
101	Carbohydrate Bis-acetal-Based Substrates as Tunable Fluorescence-Quenched Probes for Monitoring <i>exo</i> -Glycosidase Activity. <i>Journal of the American Chemical Society</i> , 2017, 139, 8392-8395.	13.7	31
102	Monitoring and modulating <i>O</i> -GlcNAcylation: assays and inhibitors of <i>O</i> -GlcNAc processing enzymes. <i>Current Opinion in Structural Biology</i> , 2021, 68, 157-165.	5.7	30
103	Multivalency To Inhibit and Discriminate Hexosaminidases. <i>Chemistry - A European Journal</i> , 2017, 23, 9022-9025.	3.3	28
104	Precision Mapping of <i>O</i> -Linked <i>N</i> -Acetylglucosamine Sites in Proteins Using Ultraviolet Photodissociation Mass Spectrometry. <i>Journal of the American Chemical Society</i> , 2020, 142, 11569-11577.	13.7	28
105	Pharmacological inhibition and knockdown of <i>O</i> -GlcNAcase reduces cellular internalization of β -synuclein preformed fibrils. <i>FEBS Journal</i> , 2021, 288, 452-470.	4.7	28
106	Tandem Bioorthogonal Labeling Uncovers Endogenous Cotranslationally <i>O</i> -GlcNAc Modified Nascent Proteins. <i>Journal of the American Chemical Society</i> , 2020, 142, 15729-15739.	13.7	27
107	<i>O</i> -GlcNAc Modification and the Tauopathies: Insights from Chemical Biology. <i>Current Alzheimer Research</i> , 2009, 6, 451-454.	1.4	25
108	Production of β -L-iduronidase in maize for the potential treatment of a human lysosomal storage disease. <i>Nature Communications</i> , 2012, 3, 1062.	12.8	25

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109	A mechanism-based inactivator of glycoside hydrolases involving formation of a transient non-classical carbocation. <i>Nature Communications</i> , 2014, 5, 5590.	12.8	25
110	Structural, Mechanistic, and Computational Analysis of the Effects of Anomeric Fluorines on Anomeric Fluoride Departure in 5-Fluoroxylsyl Fluorides. <i>Journal of the American Chemical Society</i> , 2011, 133, 15826-15829.	13.7	24
111	Molecular Basis of 1,6-Anhydro Bond Cleavage and Phosphoryl Transfer by <i>Pseudomonas aeruginosa</i> 1,6-Anhydro-N-acetylmuramic Acid Kinase. <i>Journal of Biological Chemistry</i> , 2011, 286, 12283-12291.	3.4	24
112	Cryo-EM structure provides insights into the dimer arrangement of the O-linked β -N-acetylglucosamine transferase OGT. <i>Nature Communications</i> , 2021, 12, 6508.	12.8	24
113	Metabolism of Vertebrate Amino Sugars with N-Glycolyl Groups. <i>Journal of Biological Chemistry</i> , 2012, 287, 28882-28897.	3.4	23
114	Structural Analysis of a Family 101 Glycoside Hydrolase in Complex with Carbohydrates Reveals Insights into Its Mechanism. <i>Journal of Biological Chemistry</i> , 2015, 290, 25657-25669.	3.4	23
115	<i>Streptococcus pneumoniae</i> endohexosaminidase D; feasibility of using N-glycan oxazoline donors for synthetic glycosylation of a GlcNAc-asparagine acceptor. <i>Organic and Biomolecular Chemistry</i> , 2010, 8, 1861.	2.8	22
116	P4036: Pharmacokinetics and Pharmacodynamics to Support Clinical Studies of MK8719: an O-GlcNAcase Inhibitor for Progressive Supranuclear Palsy. <i>Alzheimer's and Dementia</i> , 2016, 12, P1028.	0.8	20
117	A Direct Fluorescent Activity Assay for Glycosyltransferases Enables Convenient High-Throughput Screening: Application to β -GlcNAc Transferase. <i>Angewandte Chemie - International Edition</i> , 2020, 59, 9601-9609.	13.8	19
118	The nutrient sensor OGT regulates Hipk stability and tumorigenic-like activities in <i>Drosophila</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 2004-2013.	7.1	19
119	Diverse perspectives on interdisciplinarity from Members of the College of the Royal Society of Canada. <i>Facets</i> , 2020, 5, 138-165.	2.4	19
120	Thermal Proteome Profiling Reveals the O-GlcNAc-Dependent Meltome. <i>Journal of the American Chemical Society</i> , 2022, 144, 3833-3842.	13.7	19
121	Characterization and downstream mannose phosphorylation of human recombinant β -glucuronidase produced in <i>Arabidopsis thaliana</i> complex glycan-deficient (<i>cg1</i>) seeds. <i>Plant Biotechnology Journal</i> , 2013, 11, 1034-1043.	8.3	18
122	Conformational flexibility of the glycosidase NagZ allows it to bind structurally diverse inhibitors to suppress β -lactam antibiotic resistance. <i>Protein Science</i> , 2017, 26, 1161-1170.	7.6	18
123	Chemoproteomic identification of CO ₂ -dependent lysine carboxylation in proteins. <i>Nature Chemical Biology</i> , 2022, 18, 782-791.	8.0	18
124	Role of β -Arg211 in the Active Site of Human β -Hexosaminidase B. <i>Biochemistry</i> , 2000, 39, 6219-6227.	2.5	17
125	6- ³ Azido-6- ³ -deoxy-UDP-N-acetylglucosamine as a glycosyltransferase substrate. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2011, 21, 1199-1201.	2.2	17
126	Design of glycosyltransferase inhibitors targeting human β -GlcNAc transferase (OGT). <i>MedChemComm</i> , 2014, 5, 1172-1178.	3.4	17

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127	Modifying the phenyl group of PUGNAc: reactivity tuning to deliver selective inhibitors for N-acetyl-d-glucosaminidases. <i>Organic and Biomolecular Chemistry</i> , 2016, 14, 3193-3197.	2.8	16
128	OZâ€13â€04: Early Clinical Results and Preclinical Validation of the Oâ€GlcNacase (OGA) Inhibitor Mkâ€8719 as a Novel Therapeutic for the Treatment of Tauopathies. <i>Alzheimer's and Dementia</i> , 2016, 12, P261.	0.8	15
129	Mechanism of Human Nucleocytoplasmic Hexosaminidase D. <i>Biochemistry</i> , 2016, 55, 2735-2747.	2.5	15
130	Bicyclic Picomolar OGA Inhibitors Enable Chemoproteomic Mapping of Its Endogenous Post-translational Modifications. <i>Journal of the American Chemical Society</i> , 2022, 144, 832-844.	13.7	15
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