

Christina M Woo

List of Publications by Year in descending order

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Version: 2024-02-01

34
papers

1,292
citations

471371

17
h-index

377752

34
g-index

41
all docs

41
docs citations

41
times ranked

1192
citing authors

#	ARTICLE	IF	CITATIONS
1	Isotope-targeted glycoproteomics (IsoTaG): a mass-independent platform for intact N- and O-glycopeptide discovery and analysis. <i>Nature Methods</i> , 2015, 12, 561-567.	9.0	238
2	Mapping and Quantification of Over 2000 O-linked Glycopeptides in Activated Human T Cells with Isotope-Targeted Glycoproteomics (Isotag). <i>Molecular and Cellular Proteomics</i> , 2018, 17, 764-775.	2.5	138
3	Engineering a Proximity-Directed O-GlcNAc Transferase for Selective Protein O-GlcNAcylation in Cells. <i>ACS Chemical Biology</i> , 2020, 15, 1059-1066.	1.6	92
4	Labeling Preferences of Diazirines with Protein Biomolecules. <i>Journal of the American Chemical Society</i> , 2021, 143, 6691-6700.	6.6	92
5	Development of IsoTaG, a Chemical Glycoproteomics Technique for Profiling Intact N- and O-Glycopeptides from Whole Cell Proteomes. <i>Journal of Proteome Research</i> , 2017, 16, 1706-1718.	1.8	82
6	Community evaluation of glycoproteomics informatics solutions reveals high-performance search strategies for serum glycopeptide analysis. <i>Nature Methods</i> , 2021, 18, 1304-1316.	9.0	74
7	Small Molecule Interactome Mapping by Photoaffinity Labeling Reveals Binding Site Hotspots for the NSAIDs. <i>Journal of the American Chemical Society</i> , 2018, 140, 4259-4268.	6.6	67
8	Target protein deglycosylation in living cells by a nanobody-fused split O-GlcNAcase. <i>Nature Chemical Biology</i> , 2021, 17, 593-600.	3.9	62
9	Aspartate Residues Far from the Active Site Drive O-GlcNAc Transferase Substrate Selection. <i>Journal of the American Chemical Society</i> , 2019, 141, 12974-12978.	6.6	53
10	The Metabolic Chemical Reporter 6-Azido-6-deoxy-glucose Further Reveals the Substrate Promiscuity of <i>O</i> -GlcNAc Transferase and Catalyzes the Discovery of Intracellular Protein Modification by <i>O</i> -Glucose. <i>Journal of the American Chemical Society</i> , 2018, 140, 7092-7100.	6.6	47
11	A Binding Site Hotspot Map of the FKBP12-Rapamycin-FRB Ternary Complex by Photoaffinity Labeling and Mass Spectrometry-Based Proteomics. <i>Journal of the American Chemical Society</i> , 2019, 141, 11759-11764.	6.6	38
12	Isotope-targeted glycoproteomics (IsoTaG) analysis of sialylated N- and O-glycopeptides on an Orbitrap Fusion Tribrid using azido and alkynyl sugars. <i>Analytical and Bioanalytical Chemistry</i> , 2017, 409, 579-588.	1.9	26
13	The O-GlcNAc Modification on Kinases. <i>ACS Chemical Biology</i> , 2020, 15, 602-617.	1.6	24
14	A metabolic labeling approach for glycoproteomic analysis reveals altered glycoprotein expression upon GALNT3 knockdown in ovarian cancer cells. <i>Journal of Proteomics</i> , 2016, 145, 91-102.	1.2	21
15	Mapping the Small Molecule Interactome by Mass Spectrometry. <i>Biochemistry</i> , 2018, 57, 186-193.	1.2	21
16	Development of Photolenalidomide for Cellular Target Identification. <i>Journal of the American Chemical Society</i> , 2022, 144, 606-614.	6.6	20
17	Discovery of a Celecoxib Binding Site on Prostaglandin E Synthase (PTGES) with a Cleavable Chelation-Assisted Biotin Probe. <i>ACS Chemical Biology</i> , 2019, 14, 2527-2532.	1.6	19
18	O-Acetylated Chemical Reporters of Glycosylation Can Display Metabolism-Dependent Background Labeling of Proteins but Are Generally Reliable Tools for the Identification of Glycoproteins. <i>Frontiers in Chemistry</i> , 2020, 8, 318.	1.8	18

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19	Synthesis of an electronically-tuned minimally interfering alkynyl photo-affinity label to measure small molecule-protein interactions. <i>Tetrahedron</i> , 2018, 74, 3273-3277.	1.0	17
20	Isotope Targeted Glycoproteomics (IsoTaG) to Characterize Intact, Metabolically Labeled Glycopeptides from Complex Proteomes. <i>Current Protocols in Chemical Biology</i> , 2016, 8, 59-82.	1.7	15
21	Small Molecule Interactome Mapping by Photo-Affinity Labeling (SIM-PAL) to Identify Binding Sites of Small Molecules on a Proteome-Wide Scale. <i>Current Protocols in Chemical Biology</i> , 2019, 11, e75.	1.7	13
22	Enantioselective Synthesis and Biological Evaluation of Sanglifehrin-A and B and Analogs. <i>Angewandte Chemie - International Edition</i> , 2021, 60, 17045-17052.	7.2	13
23	O-GlcNAc Engineering on a Target Protein in Cells with Nanobody-OGT and Nanobody-splitOGA. <i>Current Protocols</i> , 2021, 1, e117.	1.3	12
24	Truncation of the TPR domain of OGT alters substrate and glycosite selection. <i>Analytical and Bioanalytical Chemistry</i> , 2021, 413, 7385-7399.	1.9	12
25	The schizophrenia-associated variant in SLC39A8 alters protein glycosylation in the mouse brain. <i>Molecular Psychiatry</i> , 2022, 27, 1405-1415.	4.1	11
26	Writing and Erasing O-GlcNAc on Casein Kinase 2 Alpha Alters the Phosphoproteome. <i>ACS Chemical Biology</i> , 2022, 17, 1111-1121.	1.6	10
27	A chiral trick to map protein ligandability. <i>Nature Chemistry</i> , 2019, 11, 1080-1082.	6.6	9
28	A Chemoproteomics Approach to Profile Phospholipase D-Derived Phosphatidyl Alcohol Interactions. <i>ACS Chemical Biology</i> , 2022, 17, 3276-3283.	1.6	9
29	Proteomic dataset for altered glycoprotein expression upon GALNT3 knockdown in ovarian cancer cells. <i>Data in Brief</i> , 2016, 8, 342-349.	0.5	7
30	4-Deoxy-4-fluoro-GalNAz (4FGalNAz) Is a Metabolic Chemical Reporter of O-GlcNAc Modifications, Highlighting the Notable Substrate Flexibility of O-GlcNAc Transferase. <i>ACS Chemical Biology</i> , 2022, 17, 159-170.	1.6	6
31	Ironing out New Antibiotic Mechanisms with Xanthocillin X. <i>ACS Central Science</i> , 2021, 7, 403-405.	5.3	4
32	The Crossroads of Glycoscience, Infection, and Immunology. <i>Frontiers in Microbiology</i> , 2021, 12, 731008.	1.5	3
33	Writing and erasing O-GlcNAc from target proteins in cells. <i>Biochemical Society Transactions</i> , 2021, 49, 2891-2901.	1.6	3
34	Enantioselective Synthesis and Biological Evaluation of Sanglifehrin-A and B and Analogs. <i>Angewandte Chemie</i> , 2021, 133, 17182-17189.	1.6	2