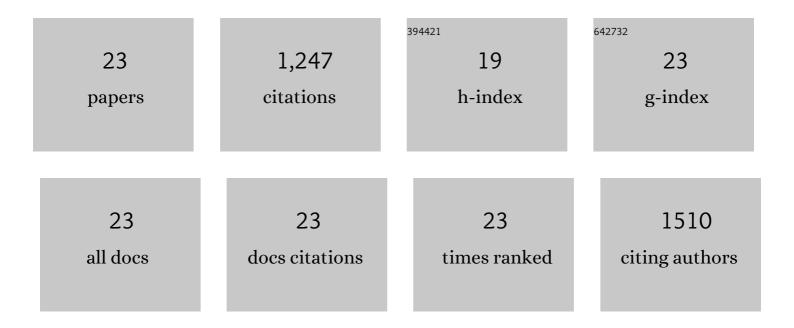
Shiaw-Lin Wu

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Mass Spectrometric Determination of Disulfide Linkages in Recombinant Therapeutic Proteins Using Online LCâ^'MS with Electron-Transfer Dissociation. Analytical Chemistry, 2009, 81, 112-122.	6.5	128
2	On-Line LCâ [~] 'MS Approach Combining Collision-Induced Dissociation (CID), Electron-Transfer Dissociation (ETD), and CID of an Isolated Charge-Reduced Species for the Trace-Level Characterization of Proteins with Post-Translational Modifications. Journal of Proteome Research, 2007, 6, 4230-4244.	3.7	126
3	Targeted Proteomics of Low-Level Proteins in Human Plasma by LC/MSn:  Using Human Growth Hormone as a Model System. Journal of Proteome Research, 2002, 1, 459-465.	3.7	123
4	Extended Range Proteomic Analysis (ERPA):Â A New and Sensitive LCâ^'MS Platform for High Sequence Coverage of Complex Proteins with Extensive Post-translational ModificationsComprehensive Analysis of Beta-Casein and Epidermal Growth Factor Receptor (EGFR)â€. Journal of Proteome Research, 2005, 4, 1155-1170.	3.7	121
5	Characterization and Comparison of Disulfide Linkages and Scrambling Patterns in Therapeutic Monoclonal Antibodies: Using LC-MS with Electron Transfer Dissociation. Analytical Chemistry, 2011, 83, 3133-3140.	6.5	96
6	Dynamic Profiling of the Post-translational Modifications and Interaction Partners of Epidermal Growth Factor Receptor Signaling after Stimulation by Epidermal Growth Factor Using Extended Range Proteomic Analysis (ERPA). Molecular and Cellular Proteomics, 2006, 5, 1610-1627.	3.8	84
7	Pim-2 Kinase Influences Regulatory T Cell Function and Stability by Mediating Foxp3 Protein N-terminal Phosphorylation. Journal of Biological Chemistry, 2015, 290, 20211-20220.	3.4	74
8	Identification of the Unpaired Cysteine Status and Complete Mapping of the 17 Disulfides of Recombinant Tissue Plasminogen Activator Using LCâ^'MS with Electron Transfer Dissociation/Collision Induced Dissociation. Analytical Chemistry, 2010, 82, 5296-5303.	6.5	67
9	A New Algorithm Using Cross-Assignment for Label-Free Quantitation with LCâ^'LTQ-FT MS. Journal of Proteome Research, 2007, 6, 2186-2194.	3.7	49
10	Characterization of the Glycosylation Occupancy and the Active Site in the Follow-on Protein Therapeutic: TNK-Tissue Plasminogen Activator. Analytical Chemistry, 2010, 82, 6154-6162.	6.5	45
11	Proteomic Analysis of High-Grade Dysplastic Cervical Cells Obtained from ThinPrep Slides Using Laser Capture Microdissection and Mass Spectrometry. Journal of Proteome Research, 2007, 6, 4256-4268.	3.7	42
12	A new and sensitive on-line liquid chromatography/mass spectrometric approach for top-down protein analysis: the comprehensive analysis of human growth hormone in anE. coli lysate using a hybrid linear ion trap/Fourier transform ion cyclotron resonance mass spectrometer. Rapid Communications in Mass Spectrometry, 2004, 18, 2201-2207.	1.5	41
13	Analysis of mouse brain microvascular endothelium using immunoâ€laser capture microdissection coupled to a hybrid linear ion trap with Fourier transformâ€mass spectrometry proteomics platform. Electrophoresis, 2008, 29, 2689-2695.	2.4	41
14	Aberrant Serum Immunoglobulin G Glycosylation in Chronic Hepatitis B Is Associated With Histological Liver Damage and Reversible by Antiviral Therapy. Journal of Infectious Diseases, 2015, 211, 115-124.	4.0	38
15	Complete Mapping of a Cystine Knot and Nested Disulfides of Recombinant Human Arylsulfatase A by Multi-Enzyme Digestion and LC-MS Analysis Using CID and ETD. Journal of the American Society for Mass Spectrometry, 2013, 24, 125-133.	2.8	36
16	A global comparability approach for biosimilar monoclonal antibodies using LC–tandem MS based proteomics. Journal of Pharmaceutical and Biomedical Analysis, 2013, 80, 126-135.	2.8	29
17	Glycation of interferon-beta-1b and human serum albumin in a lyophilized glucose formulation. International Journal of Pharmaceutics, 2006, 322, 136-145.	5.2	25
18	Mass spectrometric analysis of innovator, counterfeit, and followâ€on recombinant human growth hormone. Biotechnology Progress, 2009, 25, 207-218.	2.6	25

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#	Article	IF	CITATIONS
19	Analysis of Mouse Brain Microvascular Endothelium Using Laser Capture Microdissection Coupled with Proteomics. Methods in Molecular Biology, 2011, 686, 297-311.	0.9	19
20	Characterization of Site-Specific Glycosylation in Influenza A Virus Hemagglutinin Produced by <i>Spodoptera frugiperda</i> Insect Cell Line. Analytical Chemistry, 2017, 89, 11036-11043.	6.5	16
21	Development of LC-MS methods for quantitation of hepcidin and demonstration of siRNA-mediated hepcidin suppression in serum. Journal of Pharmacological and Toxicological Methods, 2015, 71, 110-119.	0.7	9
22	Integrated Proteomic and Genomic Analysis of Gastric Cancer Patient Tissues. Journal of Proteome Research, 2015, 14, 4995-5006.	3.7	7
23	Integrated Bottom-Up and Top-Down Liquid Chromatography–Mass Spectrometry for Characterization of Recombinant Human Growth Hormone Degradation Products. Analytical Chemistry, 2017, 89, 12771-12777.	6.5	6