## Yu-Ju Chen

List of Publications by Year in descending order

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ΥΠ-ΠΙ ΟΗΕΝ

#	Article	IF	CITATIONS
1	Kinase Sensing Based on Protein Interactions at the Catalytic Site. Chemistry - A European Journal, 2022, , .	3.3	3
2	Streamlined single-cell proteomics by an integrated microfluidic chip and data-independent acquisition mass spectrometry. Nature Communications, 2022, 13, 37.	12.8	85
3	Investigating a Boronateâ€Affinityâ€Guided Acylation Reaction for Labelling Native Antibodies. Chemistry - A European Journal, 2022, 28, .	3.3	4
4	Kinase Sensing Based on Protein Interactions at the Catalytic Site. Chemistry - A European Journal, 2022, 28, e202200655.	3.3	0
5	Electrochemical biosensors based on peptide-kinase interactions at the kinase docking site. Biosensors and Bioelectronics, 2022, 207, 114177.	10.1	10
6	Unveiling a novel serpinB2-tripeptidyl peptidase II signaling axis during senescence. Journal of Cell Science, 2022, 135, .	2.0	2
7	Phosphoproteomics Reveals the Role of Constitutive KAP1 Phosphorylation by B-cell Receptor Signaling in Chronic Lymphocytic Leukemia. Molecular Cancer Research, 2022, 20, 1222-1232.	3.4	1
8	Asia-Oceania HUPO: Past, Present, and Future. Molecular and Cellular Proteomics, 2021, 20, 100048.	3.8	1
9	An Enzyme-Mediated Aza-Michael Addition Is Involved in the Biosynthesis of an Imidazoyl Hybrid Product of Conidiogenone B. Organic Letters, 2021, 23, 1904-1909.	4.6	8
10	Integrating site-specific peptide reporters and targeted mass spectrometry enables rapid substrate-specific kinase assay at the nanogram cell level. Analytica Chimica Acta, 2021, 1155, 338341.	5.4	2
11	Toxic or Not Toxic, That Is the Carbon Quantum Dot's Question: A Comprehensive Evaluation with Zebrafish Embryo, Eleutheroembryo, and Adult Models. Polymers, 2021, 13, 1598.	4.5	24
12	A data-independent acquisition-based global phosphoproteomics system enables deep profiling. Nature Communications, 2021, 12, 2539.	12.8	44
13	ZIC-cHILIC Functionalized Magnetic Nanoparticle for Rapid and Sensitive Glycopeptide Enrichment from <1 µL Serum. Nanomaterials, 2021, 11, 2159.	4.1	1
14	Fracture risks in patients with atopic dermatitis. Annals of Allergy, Asthma and Immunology, 2021, 127, 667-673.e2.	1.0	8
15	MITF functions as a tumor suppressor in non-small cell lung cancer beyond the canonically oncogenic role. Aging, 2021, 13, 646-674.	3.1	5
16	Progress Identifying and Analyzing the Human Proteome: 2021ÂMetrics from the HUPO Human Proteome Project. Journal of Proteome Research, 2021, 20, 5227-5240.	3.7	30
17	Endothelial Yin Yang 1 Phosphorylation at S118 Induces Atherosclerosis Under Flow. Circulation Research, 2021, 129, 1158-1174.	4.5	10
18	ZIC-cHILIC-Based StageTip for Simultaneous Glycopeptide Enrichment and Fractionation toward Large-Scale N-Sialoglycoproteomics. Analytical Chemistry, 2021, 93, 15931-15940.	6.5	19

#	Article	IF	CITATIONS
19	Sample Size-Comparable Spectral Library Enhances Data-Independent Acquisition-Based Proteome Coverage of Low-Input Cells. Analytical Chemistry, 2021, 93, 17003-17011.	6.5	17
20	Transcriptomic and Proteomic Profiling of Human Mesenchymal Stem Cell Derived from Umbilical Cord in the Study of Preterm Birth. Proteomics - Clinical Applications, 2020, 14, e1900024.	1.6	6
21	Proteomic Analysis Reveals Antiâ€Fibrotic Effects of Blue Light Photobiomodulation on Fibroblasts. Lasers in Surgery and Medicine, 2020, 52, 358-372.	2.1	6
22	GPERâ€induced signaling is essential for the survival of breast cancer stem cells. International Journal of Cancer, 2020, 146, 1674-1685.	5.1	37
23	Palmitoyl Acyltransferase Activity of ZDHHC13 Regulates Skin Barrier Development Partly by Controlling PADi3 and TGM1 Protein Stability. Journal of Investigative Dermatology, 2020, 140, 959-970.e3.	0.7	10
24	Fluorescence "Turn-on―Lectin Sensors Fabricated by Ligand-Assisted Labeling Probes for Detecting Protein–Glycoprotein Interactions. Biomacromolecules, 2020, 21, 815-824.	5.4	3
25	Standardization and harmonization of distributed multi-center proteotype analysis supporting precision medicine studies. Nature Communications, 2020, 11, 5248.	12.8	49
26	Identification of Conomarphin Variants in the Conus eburneus Venom and the Effect of Sequence and PTM Variations on Conomarphin Conformations. Marine Drugs, 2020, 18, 503.	4.6	3
27	Proteogenomics of Non-smoking Lung Cancer in East Asia Delineates Molecular Signatures of Pathogenesis and Progression. Cell, 2020, 182, 226-244.e17.	28.9	178
28	Direct Oligosaccharide Profiling Using Thin-Layer Chromatography Coupled with Ionic Liquid-Stabilized Nanomatrix-Assisted Laser Desorption-Ionization Mass Spectrometry. Analytical Chemistry, 2019, 91, 11544-11552.	6.5	14
29	Boronate affinity-based photoactivatable magnetic nanoparticles for the oriented and irreversible conjugation of Fc-fused lectins and antibodies. Chemical Science, 2019, 10, 8600-8609.	7.4	8
30	Regulation of miRNA Biogenesis and Histone Modification by K63-Polyubiquitinated DDX17 Controls Cancer Stem-like Features. Cancer Research, 2019, 79, 2549-2563.	0.9	45
31	High Glucose Triggers Nucleotide Imbalance through O-GlcNAcylation of Key Enzymes and Induces KRAS Mutation in Pancreatic Cells. Cell Metabolism, 2019, 29, 1334-1349.e10.	16.2	72
32	Thymidylate kinase is critical for DNA repair <i>via</i> ATMâ€dependent Tip60 complex formation. FASEB Journal, 2019, 33, 2017-2025.	0.5	6
33	Alteration of mesenchymal stem cells polarity by laminar shear stimulation promoting β-catenin nuclear localization. Biomaterials, 2019, 190-191, 1-10.	11.4	14
34	Internationalization and investment-cash flow sensitivity: Evidence from Taiwan. Asia Pacific Management Review, 2019, 24, 154-160.	4.4	1
35	Systematic Protein Prioritization for Targeted Proteomics Studies through Literature Mining. Journal of Proteome Research, 2018, 17, 1383-1396.	3.7	16
36	ROS-independent ER stress-mediated NRF2 activation promotes warburg effect to maintain stemness-associated properties of cancer-initiating cells. Cell Death and Disease, 2018, 9, 194.	6.3	73

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37	Chemotherapy Immunophenoprofiles in Nonâ€Smallâ€Cell Lung Cancer by Personalized Membrane Proteomics. Proteomics - Clinical Applications, 2018, 12, 1700040.	1.6	6
38	Inducing hair follicle neogenesis with secreted proteins enriched in embryonic skin. Biomaterials, 2018, 167, 121-131.	11.4	29
39	FAM198B Is Associated with Prolonged Survival and Inhibits Metastasis in Lung Adenocarcinoma via Blockage of ERK-Mediated MMP-1 Expression. Clinical Cancer Research, 2018, 24, 916-926.	7.0	23
40	Glycoproteomic Alterations in Drug-Resistant Nonsmall Cell Lung Cancer Cells Revealed by Lectin Magnetic Nanoprobe-Based Mass Spectrometry. Journal of Proteome Research, 2018, 17, 3761-3773.	3.7	26
41	Launching the C-HPP neXt-CP50 Pilot Project for Functional Characterization of Identified Proteins with No Known Function. Journal of Proteome Research, 2018, 17, 4042-4050.	3.7	41
42	DNA Demethylation by DNMT3A and DNMT3B in vitro and of Methylated Episomal DNA in Transiently Transfected Cells. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2018, 1861, 1048-1061.	1.9	6
43	Surface markers of human embryonic stem cells: a meta analysis of membrane proteomics reports. Expert Review of Proteomics, 2018, 15, 911-922.	3.0	8
44	Subcellular Proteome Landscape of Human Embryonic Stem Cells Revealed Missing Membrane Proteins. Journal of Proteome Research, 2018, 17, 4138-4151.	3.7	19
45	β-Amyloid Induces Pathology-Related Patterns of Tau Hyperphosphorylation at Synaptic Terminals. Journal of Neuropathology and Experimental Neurology, 2018, 77, 814-826.	1.7	46
46	Directed strain evolution restructures metabolism for 1-butanol production in minimal media. Metabolic Engineering, 2018, 49, 153-163.	7.0	22
47	A Cloud-Based Metabolite and Chemical Prioritization System for the Biology/Disease-Driven Human Proteome Project. Journal of Proteome Research, 2018, 17, 4345-4357.	3.7	7
48	Evaluating the Possibility of Detecting Variants in Shotgun Proteomics via LeTE-Fusion Analysis Pipeline. Journal of Proteome Research, 2018, 17, 2937-2952.	3.7	5
49	Glucose intake hampers PKA-regulated HSP90 chaperone activity. ELife, 2018, 7, .	6.0	16
50	Identification of Potential Plasma Biomarkers for Nonalcoholic Fatty Liver Disease by Integrating Transcriptomics and Proteomics in Laying Hens. Journal of Nutrition, 2017, 147, 293-303.	2.9	28
51	Proteome analysis of human embryonic stem cells organelles. Journal of Proteomics, 2017, 162, 108-118.	2.4	12
52	Exploring the expression bar code of SAA variants for gastric cancer detection. Proteomics, 2017, 17, 1600356.	2.2	3
53	Data for whole and mitochondrial proteome of human embryonic stem cells. Data in Brief, 2017, 13, 371-376.	1.0	2
54	One-Pot Two-Nanoprobe Assay Uncovers Targeted Glycoprotein Biosignature. Analytical Chemistry, 2017, 89, 3973-3980.	6.5	30

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55	Phosphoproteomics Reveals HMGA1, a CK2 Substrate, as a Drug-Resistant Target in Non-Small Cell Lung Cancer. Scientific Reports, 2017, 7, 44021.	3.3	31
56	An Integrated System to Remotely Trigger Intracellular Signal Transduction by Upconversion Nanoparticle-mediated Kinase Photoactivation. Journal of Visualized Experiments, 2017, , .	0.3	1
57	Identification of Siglec Ligands Using a Proximity Labeling Method. Journal of Proteome Research, 2017, 16, 3929-3941.	3.7	73
58	Decoding the Effect of Isobaric Substitutions on Identifying Missing Proteins and Variant Peptides in Human Proteome. Journal of Proteome Research, 2017, 16, 4415-4424.	3.7	8
59	Role of S-Palmitoylation by ZDHHC13 in Mitochondrial function and Metabolism in Liver. Scientific Reports, 2017, 7, 2182.	3.3	66
60	iTop-Q: an Intelligent Tool for Top-down Proteomics Quantitation Using DYAMOND Algorithm. Analytical Chemistry, 2017, 89, 13128-13136.	6.5	2
61	Combining membrane proteomics and computational three-way pathway analysis revealed signalling pathways preferentially regulated in human iPSCs and human ESCs. Scientific Reports, 2017, 7, 15055.	3.3	3
62	A Role Model with Endless Enthusiasm for Science: In Memory of Tsutomu Masujima. Journal of the Mass Spectrometry Society of Japan, 2017, 65, 150-153.	0.1	0
63	Biomarker identification of hepatocellular carcinoma using a methodical literature mining strategy. Database: the Journal of Biological Databases and Curation, 2017, 2017, .	3.0	6
64	Absolute Phosphorylation Stoichiometry Analysis by Motif-Targeting Quantitative Mass Spectrometry. Methods in Molecular Biology, 2017, 1636, 313-325.	0.9	1
65	Untargeted, spectral libraryâ€free analysis of dataâ€independent acquisition proteomics data generated using Orbitrap mass spectrometers. Proteomics, 2016, 16, 2257-2271.	2.2	56
66	K63-polyubiquitinated HAUSP deubiquitinates HIF-11̂± and dictates H3K56 acetylation promoting hypoxia-induced tumour progression. Nature Communications, 2016, 7, 13644.	12.8	99
67	The Impact of dUTPase on Ribonucleotide Reductase-Induced Genome Instability in Cancer Cells. Cell Reports, 2016, 16, 1287-1299.	6.4	22
68	Functionalized HgTe nanoparticles promote laser-induced solid phase ionization/dissociation for comprehensive glycan sequencing. Analyst, The, 2016, 141, 6093-6103.	3.5	10
69	Atomic force microscopy characterization of kinase-mediated phosphorylation of a peptide monolayer. Scientific Reports, 2016, 6, 36793.	3.3	10
70	Temporal regulation of Lsp1 O-GlcNAcylation and phosphorylation during apoptosis of activated B cells. Nature Communications, 2016, 7, 12526.	12.8	28
71	Chemical Inhibition of Human Thymidylate Kinase and Structural Insights into the Phosphate Binding Loop and Ligand-Induced Degradation. Journal of Medicinal Chemistry, 2016, 59, 9906-9918.	6.4	15
72	The Shp2-induced epithelial disorganization defect is reversed by HDAC6 inhibition independent of Cdc42. Nature Communications, 2016, 7, 10420.	12.8	6

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73	A photo-cleavable biotin affinity tag for the facile release of a photo-crosslinked carbohydrate-binding protein. Bioorganic and Medicinal Chemistry, 2016, 24, 1216-1224.	3.0	14
74	The Influence of Phase Separation on Bioactivity of Spray Pyrolyzed Bioactive Glass. Journal of Nanoscience and Nanotechnology, 2015, 15, 4688-4696.	0.9	3
75	Mitochondrial translocation of EGFR regulates mitochondria dynamics and promotes metastasis in NSCLC. Oncotarget, 2015, 6, 37349-37366.	1.8	74
76	Analysis of Protein Stability by the Cycloheximide Chase Assay. Bio-protocol, 2015, 5, .	0.4	74
77	DcR3 suppresses influenza virus-induced macrophage activation and attenuates pulmonary inflammation and lethality. Journal of Molecular Medicine, 2015, 93, 1131-1143.	3.9	12
78	UV-activated multilayer nanomatrix provides one-step tunable carbohydrate structural characterization in MALDI-MS. Chemical Science, 2015, 6, 4790-4800.	7.4	14
79	MAGIC: An Automated N-Linked Clycoprotein Identification Tool Using a Y1-Ion Pattern Matching Algorithm and <i>in Silico</i> MS <sup>2</sup> Approach. Analytical Chemistry, 2015, 87, 2466-2473.	6.5	67
80	Mining Missing Membrane Proteins by High-pH Reverse-Phase StageTip Fractionation and Multiple Reaction Monitoring Mass Spectrometry. Journal of Proteome Research, 2015, 14, 3658-3669.	3.7	24
81	Qualification and Verification of Serological Biomarker Candidates for Lung Adenocarcinoma by Targeted Mass Spectrometry. Journal of Proteome Research, 2015, 14, 3039-3050.	3.7	18
82	Imaging Endogenous Bilirubins with Two-Photon Fluorescence of Bilirubin Dimers. Analytical Chemistry, 2015, 87, 7575-7582.	6.5	25
83	Quest for Missing Proteins: Update 2015 on Chromosome-Centric Human Proteome Project. Journal of Proteome Research, 2015, 14, 3415-3431.	3.7	53
84	Glutathionylspermidine in the Modification of Protein SH Groups: The Enzymology and Its Application to Study Protein Glutathionylation. Molecules, 2015, 20, 1452-1474.	3.8	10
85	Effect of sialylation on EGFR phosphorylation and resistance to tyrosine kinase inhibition. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 6955-6960.	7.1	102
86	dbSNO 2.0: a resource for exploring structural environment, functional and disease association and regulatory network of protein S-nitrosylation. Nucleic Acids Research, 2015, 43, D503-D511.	14.5	65
87	Large-scale determination of absolute phosphorylation stoichiometries in human cells by motif-targeting quantitative proteomics. Nature Communications, 2015, 6, 6622.	12.8	139
88	Cyclic Alopecia and Abnormal Epidermal Cornification in Zdhhc13 -Deficient Mice Reveal the Importance of Palmitoylation in Hair and Skin Differentiation. Journal of Investigative Dermatology, 2015, 135, 2603-2610.	0.7	15
89	Integrating proteomics with electrochemistry for identifying kinase biomarkers. Chemical Science, 2015, 6, 4756-4766.	7.4	30
90	Monodispersity of magnetic immuno-nanoprobes enhances the detection sensitivity of low abundance biomarkers in one drop of serum. Analyst, The, 2015, 140, 7678-7686.	3.5	5

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91	Rapid High-pH Reverse Phase StageTip for Sensitive Small-Scale Membrane Proteomic Profiling. Analytical Chemistry, 2015, 87, 12016-12023.	6.5	47
92	Informatics View on the Challenges of Identifying Missing Proteins from Shotgun Proteomics. Journal of Proteome Research, 2015, 14, 5396-5407.	3.7	14
93	Slug is temporally regulated by cyclin E in cell cycle and controls genome stability. Oncogene, 2015, 34, 1116-1125.	5.9	32
94	GSHSite: Exploiting an Iteratively Statistical Method to Identify S-Glutathionylation Sites with Substrate Specificity. PLoS ONE, 2015, 10, e0118752.	2.5	26
95	An Intelligent System for Identifying Acetylated Lysine on Histones and Nonhistone Proteins. BioMed Research International, 2014, 2014, 1-11.	1.9	20
96	Global Analysis of Cdc14 Dephosphorylation Sites Reveals Essential Regulatory Role in Mitosis and Cytokinesis. Molecular and Cellular Proteomics, 2014, 13, 594-605.	3.8	25
97	Distinct Subpopulations of Head and Neck Cancer Cells with Different Levels of Intracellular Reactive Oxygen Species Exhibit Diverse Stemness, Proliferation, and Chemosensitivity. Cancer Research, 2014, 74, 6291-6305.	0.9	104
98	Construction of Nitrogenâ€Containing 9â€Membered Ring Epoxy Vinyl Ethers via Gold(I)â€Catalyzed Intramolecular Cyclization Reactions of Acyclic 5â€Azaâ€2,3â€epoxyâ€7â€ynâ€1â€ols. Journal of the Chinese C Society, 2014, 61, 1281-1288.	hemical	2
99	RegPhos 2.0: an updated resource to explore protein kinase–substrate phosphorylation networks in mammals. Database: the Journal of Biological Databases and Curation, 2014, 2014, bau034.	3.0	38
100	Incorporating Amino Acids Composition and Functional Domains for Identifying Bacterial Toxin Proteins. BioMed Research International, 2014, 2014, 1-7.	1.9	3
101	dbGSH: a database of <i>S</i> -glutathionylation. Bioinformatics, 2014, 30, 2386-2388.	4.1	50
102	Phosphoproteomic analyses reveal that galectin-1 augments the dynamics of B-cell receptor signaling. Journal of Proteomics, 2014, 103, 241-253.	2.4	12
103	Cancer-associated fibroblasts regulate the plasticity of lung cancer stemness via paracrine signalling. Nature Communications, 2014, 5, 3472.	12.8	317
104	Sequential Phosphoproteomic Enrichment through Complementary Metal-Directed Immobilized Metal Ion Affinity Chromatography. Analytical Chemistry, 2014, 86, 685-693.	6.5	100
105	Decoding the S-Nitrosoproteomic Atlas in Individualized Human Colorectal Cancer Tissues Using a Label-Free Quantitation Strategy. Journal of Proteome Research, 2014, 13, 4942-4958.	3.7	19
106	Sequential one-pot enzymatic synthesis of oligo-N-acetyllactosamine and its multi-sialylated extensions. Chemical Communications, 2014, 50, 5786-5789.	4.1	30
107	Interaction modes and approaches to glycopeptide and glycoprotein enrichment. Analyst, The, 2014, 139, 688-704.	3.5	111
108	HSP70 colocalizes with PLK1 at the centrosome and disturbs spindle dynamics in cells arrested in mitosis by arsenic trioxide. Archives of Toxicology, 2014, 88, 1711-1723.	4.2	31

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109	Chromosome-centric Human Proteome Project (C-HPP): Chromosome 12. Journal of Proteome Research, 2014, 13, 3160-3165.	3.7	4
110	GSK3β controls epithelial–mesenchymal transition and tumor metastasis by CHIP-mediated degradation of Slug. Oncogene, 2014, 33, 3172-3182.	5.9	118
111	Quantitative proteomics analysis highlights the role of redox hemostasis and energy metabolism in human embryonic stem cell differentiation to neural cells. Journal of Proteomics, 2014, 101, 1-16.	2.4	25
112	Characterization and identification of protein O-GlcNAcylation sites with substrate specificity. BMC Bioinformatics, 2014, 15, S1.	2.6	27
113	Palmitoyl Acyltransferase, Zdhhc13, Facilitates Bone Mass Acquisition by Regulating Postnatal Epiphyseal Development and Endochondral Ossification: A Mouse Model. PLoS ONE, 2014, 9, e92194.	2.5	24
114	Recent Development of Mass Spectrometric Technologies in Asia and Oceania. Mass Spectrometry, 2014, 3, K0006-K0006.	0.6	0
115	Decoding the Disease-Associated Proteins Encoded in the Human Chromosome 4. Journal of Proteome Research, 2013, 12, 33-44.	3.7	9
116	An iTRAQ Proteomic Study Reveals an Association between Diet-Induced Enhanced Fatty Acid Metabolism and the Development of Glucose Intolerance in Prediabetic Mice. Journal of Proteome Research, 2013, 12, 1120-1133.	3.7	14
117	Methods for detection and characterization of protein S-nitrosylation. Methods, 2013, 62, 138-150.	3.8	21
118	Quantitative Proteomics Reveals Diverse Roles of miR-148a from Gastric Cancer Progression to Neurological Development. Journal of Proteome Research, 2013, 12, 3993-4004.	3.7	20
119	A Chemically Functionalized Magnetic Nanoplatform for Rapid and Specific Biomolecular Recognition and Separation. Biomacromolecules, 2013, 14, 160-168.	5.4	33
120	BAD-Lectins: Boronic Acid-Decorated Lectins with Enhanced Binding Affinity for the Selective Enrichment of Glycoproteins. Analytical Chemistry, 2013, 85, 8268-8276.	6.5	33
121	Spectrum-based Method to Generate Good Decoy Libraries for Spectral Library Searching in Peptide Identifications. Journal of Proteome Research, 2013, 12, 2305-2310.	3.7	24
122	Quantitative Phosphoproteomic Study of Pressure-Overloaded Mouse Heart Reveals Dynamin-Related Protein 1 as a Modulator of Cardiac Hypertrophy. Molecular and Cellular Proteomics, 2013, 12, 3094-3107.	3.8	57
123	Synthesis and Evaluation of a Photoactive Probe with a Multivalent Carbohydrate for Capturing Carbohydrate–Lectin Interactions. Bioconjugate Chemistry, 2013, 24, 1895-1906.	3.6	18
124	Identification of Downstream Components of Ubiquitin-Conjugating Enzyme PHOSPHATE2 by Quantitative Membrane Proteomics in <i>Arabidopsis</i> Roots. Plant Cell, 2013, 25, 4044-4060.	6.6	242
125	Androgen modulates cardiac fibrosis contributing to gender differences on heart failure. Aging Male, 2013, 16, 22-27.	1.9	10
126	dbPTM 3.0: an informative resource for investigating substrate site specificity and functional association of protein post-translational modifications. Nucleic Acids Research, 2013, 41, D295-D305.	14.5	179

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127	Quantitative Proteomic Analysis of Human Lung Tumor Xenografts Treated with the Ectopic ATP Synthase Inhibitor Citreoviridin. PLoS ONE, 2013, 8, e70642.	2.5	26
128	Prioritization of Cancer Marker Candidates Based on the Immunohistochemistry Staining Images Deposited in the Human Protein Atlas. PLoS ONE, 2013, 8, e81079.	2.5	9
129	Thyroid Storm and Incidental Anterior Mediastinal Teratoma: Coincidence or Correlation?. Acta Cardiologica Sinica, 2013, 29, 467-70.	0.2	0
130	dbSNO: a database of cysteine <i>S</i> -nitrosylation. Bioinformatics, 2012, 28, 2293-2295.	4.1	71
131	Cancer Phosphoproteomics: Tools and Emerging Applications for Mining the Phosphoproteome in Cancer Biology. , 2012, , 161-188.		0
132	Membrane Proteomics for the Opportunity of Cancer Biomarker and Drug Target Discovery. , 2012, , 259-286.		0
133	Phosphoproteomic Analysis of Human Mesenchymal Stromal Cells during Osteogenic Differentiation. Journal of Proteome Research, 2012, 11, 586-598.	3.7	14
134	Complementary Fe <sup>3+</sup> ―and Ti <sup>4+</sup> â€immobilized metal ion affinity chromatography for purification of acidic and basic phosphopeptides. Rapid Communications in Mass Spectrometry, 2012, 26, 2186-2194.	1.5	43
135	In-depth Identification of Pathways Related to Cisplatin-induced Hepatotoxicity through an Integrative Method Based on an Informatics-assisted Label-free Protein Quantitation and Microarray Gene Expression Approach. Molecular and Cellular Proteomics, 2012, 11, M111.010884.	3.8	58
136	Tumor Cells Require Thymidylate Kinase to Prevent dUTP Incorporation during DNA Repair. Cancer Cell, 2012, 22, 36-50.	16.8	56
137	A new type of donor–acceptor dye bridged by the bidentate moiety; metal ion complexation enhancing DSSC performance. Journal of Materials Chemistry, 2011, 21, 4090.	6.7	16
138	Iron Oxide Nanomatrix Facilitating Metal Ionization in Matrix-Assisted Laser Desorption/Ionization Mass Spectrometry. Analytical Chemistry, 2011, 83, 9337-9343.	6.5	18
139	A Genuine Intramolecular Proton Relay System Undergoing Excited-State Double Proton Transfer Reaction. Journal of Physical Chemistry Letters, 2011, 2, 3063-3068.	4.6	94
140	Identification of SEC61Î <sup>2</sup> and its autoantibody as biomarkers for colorectal cancer. Clinica Chimica Acta, 2011, 412, 887-893.	1.1	18
141	Interplay between SIN3A and STAT3 Mediates Chromatin Conformational Changes and GFAP Expression during Cellular Differentiation. PLoS ONE, 2011, 6, e22018.	2.5	48
142	Surface Marker Epithelial Cell Adhesion Molecule and E-cadherin Facilitate the Identification and Selection of Induced Pluripotent Stem Cells. Stem Cell Reviews and Reports, 2011, 7, 722-735.	5.6	55
143	Rapid and specific influenza virus detection by functionalized magnetic nanoparticles and mass spectrometry. Journal of Nanobiotechnology, 2011, 9, 52.	9.1	71
144	Nanoprobeâ€based immobilized metal affinity chromatography for sensitive and complementary enrichment of multiply phosphorylated peptides. Proteomics, 2011, 11, 2639-2653.	2.2	20

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145	Nanoparticle-assisted MALDI-TOF MS combined with seed-layer surface preparation for quantification of small molecules. Analytica Chimica Acta, 2011, 697, 1-7.	5.4	30
146	Phosphoproteomics by Highly Selective IMAC Protocol. Neuromethods, 2011, , 181-196.	0.3	3
147	An Informatics-assisted Label-free Approach for Personalized Tissue Membrane Proteomics: Case Study on Colorectal Cancer. Molecular and Cellular Proteomics, 2011, 10, M110.003087.	3.8	50
148	Phosphoproteomics Identifies Oncogenic Ras Signaling Targets and Their Involvement in Lung Adenocarcinomas. PLoS ONE, 2011, 6, e20199.	2.5	35
149	SNOSite: Exploiting Maximal Dependence Decomposition to Identify Cysteine S-Nitrosylation with Substrate Site Specificity. PLoS ONE, 2011, 6, e21849.	2.5	77
150	Phosphoproteomics characterization of novel phosphorylated sites of lens proteins from normal and cataractous human eye lenses. Molecular Vision, 2011, 17, 186-98.	1.1	20
151	Dihydrobenzoic acid modified nanoparticle as a MALDI-TOF MS matrix for soft ionization and structure determination of small molecules with diverse structures. Journal of the American Society for Mass Spectrometry, 2010, 21, 1930-1939.	2.8	40
152	The Asia Oceania Human Proteome Organisation Membrane Proteomics Initiative. Preparation and characterisation of the carbonateâ€washed membrane standard. Proteomics, 2010, 10, 4142-4148.	2.2	26
153	Comparison of membrane fraction proteomic profiles of normal and cancerous human colorectal tissues with gelâ€assisted digestion and iTRAQ labeling mass spectrometry. FEBS Journal, 2010, 277, 3028-3038.	4.7	63
154	IDEAL-Q, an Automated Tool for Label-free Quantitation Analysis Using an Efficient Peptide Alignment Approach and Spectral Data Validation. Molecular and Cellular Proteomics, 2010, 9, 131-144.	3.8	114
155	<i>S</i> -Alkylating Labeling Strategy for Site-Specific Identification of the <i>S</i> -Nitrosoproteome. Journal of Proteome Research, 2010, 9, 6417-6439.	3.7	64
156	An Informatics-assisted Label-free Quantitation Strategy that Depicts Phosphoproteomic Profiles in Lung Cancer Cell Invasion. Journal of Proteome Research, 2010, 9, 5582-5597.	3.7	57
157	Temporal Proteomics Profiling of Lipid Rafts in CCR6-Activated T Cells Reveals the Integration of Actin Cytoskeleton Dynamics. Journal of Proteome Research, 2010, 9, 283-297.	3.7	36
158	Phosphorescent Ir(iii) complexes bearing double benzyldiphenylphosphine cyclometalates; strategic synthesis, fundamental and integration for white OLED fabrication. Journal of Materials Chemistry, 2010, 20, 7682.	6.7	67
159	Identification of in vivo phosphorylation sites of lens proteins from porcine eye lenses by a gel-free phosphoproteomics approach. Molecular Vision, 2010, 16, 294-302.	1.1	15
160	Role for α- <scp> </scp> -fucosidase in the control of <i>Helicobacter pylori</i> -infected gastric cancer cells. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 14581-14586.	7.1	69
161	Complementary Quantitative Proteomics Reveals that Transcription Factor AP-4 Mediates E-box-dependent Complex Formation for Transcriptional Repression of HDM2>. Molecular and Cellular Proteomics, 2009, 8, 2034-2050.	3.8	21
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