

# Hsin-Yi Wu

## List of Publications by Year in descending order

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14  
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1307594

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#	ARTICLE	IF	CITATIONS
1	Post-Deconvolution MS/MS Spectra Extraction with Data-Independent Acquisition for Comprehensive Profiling of Urinary Glucuronide-Conjugated Metabolome. <i>Analytical Chemistry</i> , 2022, 94, 2740-2748.	6.5	5
2	Phosphoproteomics Reveals the Role of Constitutive KAP1 Phosphorylation by B-cell Receptor Signaling in Chronic Lymphocytic Leukemia. <i>Molecular Cancer Research</i> , 2022, 20, 1222-1232.	3.4	1
3	Quantitative Proteome Analysis Reveals <i>Melissa officinalis</i> Extract Targets Mitochondrial Respiration in Colon Cancer Cells. <i>Molecules</i> , 2022, 27, 4533.	3.8	3
4	Analyzing BMP2, FGFR, and TGF Beta Expressions in High-Grade Osteosarcoma Untreated and Treated Autografts Using Proteomic Analysis. <i>International Journal of Molecular Sciences</i> , 2022, 23, 7409.	4.1	1
5	Discovery of Spoilage Markers for Chicken Eggs Using Liquid Chromatography-High Resolution Mass Spectrometry-Based Untargeted and Targeted Foodomics. <i>Journal of Agricultural and Food Chemistry</i> , 2021, 69, 4331-4341.	5.2	18
6	Untargeted foodomics strategy using high-resolution mass spectrometry reveals potential indicators for fish freshness. <i>Analytica Chimica Acta</i> , 2020, 1127, 98-105.	5.4	33
7	Utilizing Experimental Mouse Model to Identify Effectors of Hepatocellular Carcinoma Induced by HBx Antigen. <i>Cancers</i> , 2020, 12, 409.	3.7	13
8	Assessment of Serological Early Biomarker Candidates for Lung Adenocarcinoma by using Multiple Reaction Monitoring-Mass Spectrometry. <i>Proteomics - Clinical Applications</i> , 2020, 14, e1900095.	1.6	7
9	Assessing the Similarity between Random Copolymer Drug Glatiramer Acetate by Using LC-MS Data Coupling with Hypothesis Testing. <i>Analytical Chemistry</i> , 2019, 91, 14281-14289.	6.5	2
10	Combining Alkaline Phosphatase Treatment and Hybrid Linear Ion Trap/Orbitrap High Mass Accuracy Liquid Chromatography-Mass Spectrometry Data for the Efficient and Confident Identification of Protein Phosphorylation. <i>Analytical Chemistry</i> , 2009, 81, 7778-7787.	6.5	19
11	Proteomics Analysis of Nasopharyngeal Carcinoma Cell Secretome Using a Hollow Fiber Culture System and Mass Spectrometry. <i>Journal of Proteome Research</i> , 2009, 8, 380-389.	3.7	32
12	Strategy for Determination of in Vitro Protein Acetylation Sites by Using Isotope-Labeled Acetyl Coenzyme A and Liquid Chromatography-Mass Spectrometry. <i>Analytical Chemistry</i> , 2008, 80, 6178-6189.	6.5	8
13	Analysis of protein phosphorylation using mass spectrometry. <i>Chang Gung Medical Journal</i> , 2008, 31, 217-27.	0.7	5
14	Mining Phosphopeptide Signals in Liquid Chromatography-Mass Spectrometry Data for Protein Phosphorylation Analysis. <i>Journal of Proteome Research</i> , 2007, 6, 1812-1821.	3.7	25