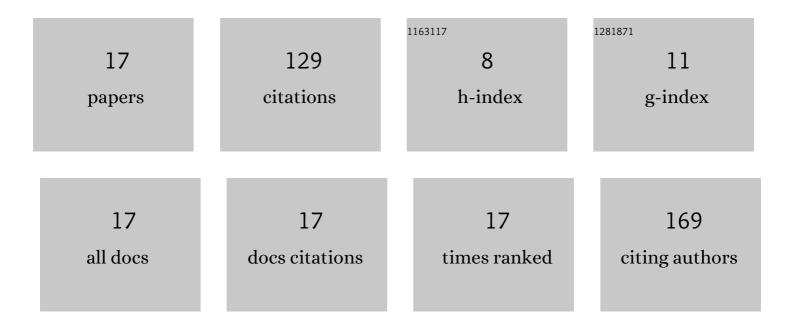
Han-Ju Chien

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Rapid determination of bioactive compounds in the different organs of Salvia Miltiorrhiza by UPLC-MS/MS. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2019, 1104, 81-88.	2.3	17
2	Honey proteomic signatures for the identification of honey adulterated with syrup, producing country, and nectar source using SWATH-MS approach. Food Chemistry, 2021, 354, 129590.	8.2	16
3	Graphene flakes enhance the detection of TiO2-enriched catechins by SALDI-MS after microwave-assisted enrichment. Talanta, 2016, 153, 347-352.	5.5	14
4	Proteomics for species authentication of cod and corresponding fishery products. Food Chemistry, 2022, 374, 131631.	8.2	14
5	Metabolic Disturbances in Adult-Onset Still's Disease Evaluated Using Liquid Chromatography/Mass Spectrometry-Based Metabolomic Analysis. PLoS ONE, 2016, 11, e0168147.	2.5	9
6	2-DE combined with two-layer feature selection accurately establishes the origin of oolong tea. Food Chemistry, 2016, 211, 392-399.	8.2	9
7	Rapid determination of isoflavones and other bioactive compounds in soybean using SWATH-MS. Analytica Chimica Acta, 2020, 1103, 122-133.	5.4	9
8	SWATH-MS-based quantitative proteomics reveals a uniquely intricate defense response in Cnaphalocrocis medinalis-resistant rice. Scientific Reports, 2020, 10, 6597.	3.3	9
9	A rapid, simple, and high-throughput UPLC-MS/MS method for simultaneous determination of bioactive constituents in Salvia miltiorrhiza with positive/negative ionization switching. Journal of Pharmaceutical and Biomedical Analysis, 2018, 161, 94-100.	2.8	8
10	Determination of adulteration, geographical origins, and species of food by mass spectrometry. Mass Spectrometry Reviews, 2023, 42, 2273-2323.	5.4	6
11	Proteomic analysis of "Oriental Beauty―oolong tea leaves with different degrees of leafhopper infestation. Rapid Communications in Mass Spectrometry, 2020, 34, e8825.	1.5	4
12	Preparation and comparison of Fe3O4@graphene oxide nanoclusters for analysis of glimepiride in urine by surface-assisted laser desorption/ionization time-of-flight mass spectrometry. Analytical and Bioanalytical Chemistry, 2020, 412, 4057-4065.	3.7	4
13	Comparative Proteomic Analysis of Rat Bronchoalveolar Lavage Fluid after Exposure to Zinc Oxide Nanoparticles. Mass Spectrometry, 2017, 6, S0066-S0066.	0.6	3
14	Inhibition of the clinical isolates of Acinetobacter baumannii by Pseudomonas aeruginosa: InÂvitro assessment of a case-based study. Journal of Microbiology, Immunology and Infection, 2020, 55, 60-60.	3.1	3
15	Metabolic disturbances in systemic lupus erythematosus evaluated with UPLC-MS/MS. Clinical and Experimental Rheumatology, 0, , .	0.8	3
16	Proteomic analysis of rat kidney under maleic acid treatment by SWATHâ€MS technology. Rapid Communications in Mass Spectrometry, 2020, 34, e8633.	1.5	1
17	Metabolic disturbances in systemic lupus erythematosus evaluated with UPLC-MS/MS. Clinical and Experimental Rheumatology, 2021, , .	0.8	0