Wenying Jian

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Recent advances in application of hydrophilic interaction chromatography for quantitative bioanalysis. Journal of Separation Science, 2010, 33, 681-697.	2.5	139
2	Validation and application of an LC–MS/MS method for quantitation of three fatty acid ethanolamides as biomarkers for fatty acid hydrolase inhibition in human plasma. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2010, 878, 1687-1699.	2.3	63
3	LC–MS bioanalysis of intact proteins and peptides. Biomedical Chromatography, 2020, 34, e4633.	1.7	62
4	A workflow for absolute quantitation of large therapeutic proteins in biological samples at intact level using LC-HRMS. Bioanalysis, 2016, 8, 1679-1691.	1.5	57
5	Relative Quantitation of Glycoisoforms of Intact Apolipoprotein C3 in Human Plasma by Liquid Chromatography–High-Resolution Mass Spectrometry. Analytical Chemistry, 2013, 85, 2867-2874.	6.5	54
6	Rapid Detection and Characterization of in Vitro and Urinary <i>N</i> -Acetyl- <scp> </scp> -cysteine Conjugates Using Quadrupole-Linear Ion Trap Mass Spectrometry and Polarity Switching. Chemical Research in Toxicology, 2009, 22, 1246-1255.	3.3	52
7	Development and validation of an LC–MS/MS based method for quantification of 25 hydroxyvitamin D2 and 25 hydroxyvitamin D3 in human serum and plasma. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2014, 961, 62-70.	2.3	42
8	Simultaneous Catabolite Identification and Quantitation of Large Therapeutic Protein at the Intact Level by Immunoaffinity Capture Liquid Chromatography–High-Resolution Mass Spectrometry. Analytical Chemistry, 2017, 89, 6065-6075.	6.5	40
9	Evaluation of a High-Throughput Online Solid Phase Extraction–Tandem Mass Spectrometry System for In Vivo Bioanalytical Studies. Analytical Chemistry, 2011, 83, 8259-8266.	6.5	39
10	Important considerations for quantitation of small-molecule biomarkers using LC–MS. Bioanalysis, 2012, 4, 2431-2434.	1.5	36
11	Potential bias and mitigations when using stable isotope labeled parent drug as internal standard for LC–MS/MS quantitation of metabolites. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2010, 878, 3267-3276.	2.3	29
12	Analysis of polar metabolites by hydrophilic interaction chromatography–MS/MS. Bioanalysis, 2011, 3, 899-912.	1.5	27
13	Quantitation of intact monoclonal antibody in biological samples: comparison of different data processing strategies. Bioanalysis, 2018, 10, 1055-1067.	1.5	24
14	2019 White Paper on Recent Issues in Bioanalysis: Chromatographic Assays (Part 1 – Innovation in Small) Tj ET	⁻ Qq0 0 0 r 1.5	gBT /Overloc 24
15	2020 White Paper on Recent issues in Bioanalysis: BMV of Hybrid Assays, Acoustic MS, HRMS, Data Integrity, Endogenous Compounds, Microsampling and Microbiome (<u>Part 1</u> – Recommendations) Tj ET	Qq1 1 0.7 1.5	784314 rgBT 24
16	Bioanalysis, 2021, 19, 2097236. Quantitation of leukotriene B4 in human sputum as a biomarker using UPLC–MS/MS. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2013, 932, 59-65.	2.3	22
17	LC–MS Challenges in Characterizing and Quantifying Monoclonal Antibodies (mAb) and Antibody-Drug Conjugates (ADC) in Biological Samples. Current Pharmacology Reports, 2018, 4, 45-63.	3.0	21
18	Recent advances in absolute quantification of peptides and proteins using LC-MS. Reviews in Analytical Chemistry. 2014, 33.	3.2	20

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19	Intact Protein Mass Spectrometry for Therapeutic Protein Quantitation, Pharmacokinetics, and Biotransformation in Preclinical and Clinical Studies: An Industry Perspective. Journal of the American Society for Mass Spectrometry, 2021, 32, 1886-1900.	2.8	19
20	LC/MS/MS Bioanalysis of Protein–Drug Conjugates—The Importance of Incorporating Succinimide Hydrolysis Products. Analytical Chemistry, 2018, 90, 5314-5321. 2021 White Paper on Recent Issues in Bioanalysis: Mass Spec of Proteins, Extracellular Vesicles, CRISPR.	6.5	15
21	Chiral Assays, Óligos; Nanomedicines Bioanalýsis; ICH M1Ó Section 7.1; Non-Liquid & Rare Matrices; Regulatory InputsÂ(<u>Part 1A</u> – Recommendations on Endogenous Compounds, Small Molecules,) Tj ET(Qq1_1 0.7	'84314 rgBT
22	Bio-generation of stable isotope labeled internal standards for absolute and relative quantitation of drug metabolites in plasma samples by LC–MS/MS. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2013, 926, 92-100.	2.3	13
23	Bioanalytical workflow for novel scaffold protein–drug conjugates: quantitation of total Centyrin protein, conjugated Centyrin and free payload for Centyrin–drug conjugate in plasma and tissue samples using liquid chromatography–tandem mass spectrometry. Bioanalysis, 2018, 10, 1651-1665.	1.5	12
24	A Cross Company Perspective on the Assessment of Therapeutic Protein Biotransformation. Drug Metabolism and Disposition, 2022, 50, 846-857.	3.3	8
25	Conjugation of a peptide to an antibody engineered with free cysteines dramatically improves half-life and activity. MAbs, 2020, 12, 1794687.	5.2	7
26	LC–MS/MS quantification of 7α-hydroxy-4-cholesten-3-one (C4) in rat and monkey plasma. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2017, 1064, 49-55.	2.3	6
27	Novel advances in biotransformation and bioactivation research – 2020 year in review. Drug Metabolism Reviews, 2021, 53, 384-433.	3.6	4
28	Selectivity for quantitation of biomarkers using liquid chromatography and mass spectrometry. Bioanalysis, 2018, 10, 1461-1465.	1.5	3
29	Design, synthesis and preclinical evaluation of bio-conjugated amylinomimetic peptides as long-acting amylin receptor agonists. European Journal of Medicinal Chemistry, 2022, 236, 114330.	5.5	2
30	Bioanalysis of small and large molecule drugs, metabolites, and biomarkers by LC-MS. , 2020, , 3-38.		1
31	Application of middle-down approach in quantitation and catabolite identification of protein by LC–high-resolution mass spectrometry. Bioanalysis, 2021, 13, 465-479.	1.5	1