## Hongliang Jiang

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Network pharmacology combined with metabolomics and lipidomics to reveal the hypolipidemic mechanism of <i>Alismatis rhizoma</i> in hyperlipidemic mice. Food and Function, 2022, 13, 4714-4733.	4.6	16
2	Simultaneous determination of abiraterone and its five metabolites in human plasma by LC-MS/MS: Application to pharmacokinetic study in healthy Chinese subjects. Journal of Pharmaceutical and Biomedical Analysis, 2022, 217, 114826.	2.8	3
3	Comprehensive metabolic profiling of Alismatis Rhizoma triterpenes in rats based on characteristic ions and a triterpene database. Journal of Pharmaceutical Analysis, 2021, 11, 96-107.	5.3	8
4	Discrimination of turmeric from different origins in China by MRM-based curcuminoid profiling and multivariate analysis. Food Chemistry, 2021, 338, 127794.	8.2	15
5	Preparation of bovine serum albumin restricted access octadecyl/phenyl-mixed-functionalized magnetic silica nanoparticles for magnetic solid phase extraction of antidepressants in aquatic products followed by UHPLC-MS/MS. Talanta, 2021, 221, 121458.	5.5	9
6	The Protein-Binding Behavior of Platinum Anticancer Drugs in Blood Revealed by Mass Spectrometry. Pharmaceuticals, 2021, 14, 104.	3.8	26
7	Qualitative distribution of endogenous sphingolipids in plasma of human and rodent species by UPLC-Q-Exactive-MS. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2021, 1173, 122684.	2.3	3
8	Interaction of Natural Compounds in Licorice and Turmeric with HIV-NCp7 Zinc Finger Domain: Potential Relevance to the Mechanism of Antiviral Activity. Molecules, 2021, 26, 3563.	3.8	3
9	Systematic Identification of Proteins Binding with Cisplatin in Blood by Affinity Chromatography and a Four-Dimensional Proteomic Method. Journal of Proteome Research, 2021, 20, 4553-4565.	3.7	6
10	Quantitative proteomics of plasma and liver reveals the mechanism of turmeric in preventing hyperlipidemia in mice. Food and Function, 2021, 12, 10484-10499.	4.6	8
11	Improved structural annotation of triterpene metabolites of traditional Chinese medicine in vivo based on quantitative structure-retention relationships combined with characteristic ions: Alismatis Rhizoma as an example. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2021, 1187, 123012.	2.3	5
12	Novel PEP-PAN@PSF rods extraction of EDCs in environmental water, sediment, and fish homogenate followed by pre-column derivatization and UHPLC-MS/MS detection. Talanta, 2020, 210, 120661.	5.5	16
13	Spectrum-effect relationship for anti-tumor activity of shikonins and shikonofurans in medicinal Zicao by UHPLC-MS/MS and chemometric approaches. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2020, 1136, 121924.	2.3	23
14	Qualitative distribution of endogenous phosphatidylcholine and sphingomyelin in serum using LC-MS/MS based profiling. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2020, 1155, 122289.	2.3	8
15	Magnetic solid-phase extraction modified Quick, Easy, Cheap, Effective, Rugged and Safe method combined with pre-column derivatization and ultra-high performance liquid chromatography-tandem mass spectrometry for determination of estrogens and estrogen mimics in pork and chicken samples. Journal of Chromatography A, 2020, 1622, 461137.	3.7	21
16	Identification of the lipid-lowering component of triterpenes from Alismatis rhizoma based on the MRM-based characteristic chemical profiles and support vector machine model. Analytical and Bioanalytical Chemistry, 2019, 411, 3257-3268.	3.7	26
17	The Influence of Proinflammatory Cytokines on Voriconazole Trough Concentration in Patients With Different Forms of Hematologic Disorders. Journal of Clinical Pharmacology, 2019, 59, 1340-1350.	2.0	13
18	Lysophospholipid profiles of apolipoprotein E-deficient mice reveal potential lipid biomarkers associated with atherosclerosis progression using validated UPLC-QTRAP-MS/MS-based lipidomics approach. Journal of Pharmaceutical and Biomedical Analysis, 2019, 171, 148-157.	2.8	8

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19	Elucidation of the Mechanism of Action for Metal Based Anticancer Drugs by Mass Spectrometry-Based Quantitative Proteomics. Molecules, 2019, 24, 581.	3.8	19
20	Discovery and characterisation of lycorineâ€type alkaloids in <i>Lycoris</i> spp. (Amaryllidaceae) using UHPLCâ€QTOFâ€MS. Phytochemical Analysis, 2019, 30, 268-277.	2.4	10
21	Metabolic Stability and Metabolite Characterization of Capilliposide B and Capilliposide C by LC–QTRAP–MS/MS. Pharmaceutics, 2018, 10, 178.	4.5	4
22	A targeted strategy to identify untargeted metabolites from in vitro to in vivo: Rapid and sensitive metabolites profiling of licorice in rats using ultra-high performance liquid chromatography coupled with triple quadrupole-linear ion trap mass spectrometry. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2018, 1092, 40-50.	2.3	31
23	Impact of curcumin on the pharmacokinetics of rosuvastatin in rats and dogs based on the conjugated metabolites. Xenobiotica, 2017, 47, 267-275.	1.1	23
24	Dose optimisation of voriconazole with therapeutic drug monitoring in children: a single-centre experience in China. International Journal of Antimicrobial Agents, 2017, 49, 483-487.	2.5	26
25	The strategy for establishment of the multiple reaction monitoring based characteristic chemical profile of triterpenes in Alismatis rhizoma using two combined tandem mass spectrometers. Journal of Chromatography A, 2017, 1524, 121-134.	3.7	20
26	Identification and characterization of curcuminoids in turmeric using ultra-high performance liquid chromatography-quadrupole time of flight tandem mass spectrometry. Journal of Chromatography A, 2017, 1521, 110-122.	3.7	36
27	Tissue distribution of capilliposide B, capilliposide C and their bioactive metabolite in mice using liquid â€ŧandem mass spectrometry. Biomedical Chromatography, 2017, 31, e3895.	1.7	4
28	An integrated strategy for establishment of metabolite profile of endogenous lysoglycerophospholipids by two LC-MS/MS platforms. Talanta, 2017, 162, 530-539.	5.5	14
29	An integrated strategy for establishment of curcuminoid profile in turmeric using two LC–MS/MS platforms. Journal of Pharmaceutical and Biomedical Analysis, 2017, 132, 93-102.	2.8	29
30	HRMS for the quantification of xenobiotics in biological samples. Bioanalysis, 2017, 9, 1939-1941.	1.5	1
31	Untargeted screening of unknown xenobiotics and potential toxins in plasma of poisoned patients using high-resolution mass spectrometry: Generation of xenobiotic fingerprint using background subtraction. Analytica Chimica Acta, 2016, 944, 37-43.	5.4	18
32	Bile acid profiles in diabetic (db/db) mice and their wild type littermates. Journal of Pharmaceutical and Biomedical Analysis, 2016, 131, 473-481.	2.8	27
33	An integrated approach for profiling oxidative metabolites and glutathione adducts using liquid chromatography coupled with ultraviolet detection and triple quadrupole-linear ion trap mass spectrometry. Journal of Pharmaceutical and Biomedical Analysis, 2016, 129, 482-491.	2.8	5
34	Optimization of solid-phase extraction and liquid chromatography-tandem mass spectrometry for simultaneous determination of capilliposide B and its active metabolite in rat urine and feces: Overcoming nonspecific binding. Journal of Pharmaceutical and Biomedical Analysis, 2016, 131, 6-12.	2.8	9
35	Metabolism and bioactivation of the tricyclic antidepressant amitriptyline in human liver microsomes and human urine. Bioanalysis, 2016, 8, 1365-1381.	1.5	5
36	The Hypolipidemic Effect of Total Saponins from Kuding Tea in Highâ€Fat Dietâ€Induced Hyperlipidemic Mice and Its Composition Characterized by UPLCâ€QTOFâ€MS/MS. Journal of Food Science, 2016, 81, H1313-9.	3.1	21

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37	The metabolic change of serum lysophosphatidylcholines involved in the lipid lowering effect of triterpenes from Alismatis rhizoma on high-fat diet induced hyperlipidemia mice. Journal of Ethnopharmacology, 2016, 177, 10-18.	4.1	55
38	Cell-based assays in combination with ultra-high performance liquid chromatography-quadrupole time of flight tandem mass spectrometry for screening bioactive capilliposide C metabolites generated by rat intestinal microflora. Journal of Pharmaceutical and Biomedical Analysis, 2016, 119, 130-138.	2.8	10
39	A novel fourâ€step approach for systematic identification of naphthoquinones in <i>Juglans cathayensis</i> dode using various scan functions of liquid chromatographyâ€ŧandem mass spectrometry along with data mining strategies. Phytochemical Analysis, 2015, 26, 413-422.	2.4	7
40	Systematic identification of shikonins and shikonofurans in medicinal Zicao species using ultra-high performance liquid chromatography quadrupole time of flight tandem mass spectrometry combined with a data mining strategy. Journal of Chromatography A, 2015, 1425, 158-172.	3.7	23
41	A novel dereplication strategy for the identification of two new trace compounds in the extract of Gastrodia elata using UHPLC/Q-TOF-MS/MS. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2015, 988, 45-52.	2.3	60
42	Determination of thailandepsin B in rat plasma with a LC–MS/MS method. Bioanalysis, 2015, 7, 39-52.	1.5	4
43	Preventive effects of turmeric on the high-fat diet-induced hyperlipidaemia in mice associated with a targeted metabolomic approach for the analysis of serum lysophosphatidylcholine using LC-MS/MS. Journal of Functional Foods, 2014, 11, 130-141.	3.4	19
44	The rapid discovery and identification of physalins in the calyx of Physalis alkekengi L.var.franchetii (Mast.) Makino using ultra-high performance liquid chromatography–quadrupole time of flight tandem mass spectrometry together with a novel three-step data mining strategy. Journal of Chromatography A, 2014, 1361, 139-152.	3.7	39
45	Simultaneous determination of capilliposide B and capilliposide C in rat plasma by LC–MS/MS and its application to a PK study. Bioanalysis, 2014, 6, 935-945.	1.5	17
46	Systematic evaluation of supported liquid extraction in reducing matrix effect and improving extraction efficiency in LC–MS/MS based bioanalysis for 10 model pharmaceutical compounds. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2012, 891-892, 71-80.	2.3	73
47	Development and validation of a sensitive LC/MS/MS method for the simultaneous determination of naloxone and its metabolites in mouse plasma. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2011, 879, 2663-2668.	2.3	21
48	Determination of carboplatin in human plasma using HybridSPE-precipitation along with liquid chromatography–tandem mass spectrometry. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2011, 879, 2162-2170.	2.3	31
49	Use of liquid chromatography–electrospray ionization tandem mass spectrometry to identify diarylheptanoids in turmeric (Curcuma longa L.) rhizome. Journal of Chromatography A, 2006, 1111, 21-31.	3.7	108
50	Metabolic profiling and phylogenetic analysis of medicinal Zingiber species: Tools for authentication of ginger (Zingiber officinale Rosc.). Phytochemistry, 2006, 67, 1673-1685.	2.9	138