## Jun Han

## List of Publications by Year in descending order

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126907 128289 3,956 71 33 60 citations h-index g-index papers 73 73 73 6505 docs citations times ranked citing authors all docs

| #  | Article   | IF           | Citations |
|----|---|--------------|-----------|
| 1  | An isotope-labeled chemical derivatization method for the quantitation of short-chain fatty acids in human feces by liquid chromatography–tandem mass spectrometry. Analytica Chimica Acta, 2015, 854, 86-94.                       | 5.4          | 380       |
| 2  | Adora2b-elicited Per2 stabilization promotes a HIF-dependent metabolic switch crucial for myocardial adaptation to ischemia. Nature Medicine, 2012, 18, 774-782.  | 30.7         | 278       |
| 3  | Effect of Antibiotic Treatment on the Intestinal Metabolome. Antimicrobial Agents and Chemotherapy, 2011, 55, 1494-1503.  | 3.2          | 258       |
| 4  | Hydrogen/Deuterium Exchange Mass Spectrometry with Top-Down Electron Capture Dissociation for Characterizing Structural Transitions of a 17 kDa Protein. Journal of the American Chemical Society, 2009, 131, 12801-12808.          | 13.7         | 174       |
| 5  | Diet and specific microbial exposure trigger features of environmental enteropathy in a novel murine model. Nature Communications, 2015, 6, 7806.   | 12.8         | 172       |
| 6  | HIF1A Reduces Acute Lung Injury by Optimizing Carbohydrate Metabolism in the Alveolar Epithelium. PLoS Biology, 2013, 11, e1001665.   | 5 <b>.</b> 6 | 138       |
| 7  | Towards high-throughput metabolomics using ultrahigh-field Fourier transform ion cyclotron resonance mass spectrometry. Metabolomics, 2008, 4, 128-140.   | 3.0          | 136       |
| 8  | Metabolic Profiling of Bile Acids in Human and Mouse Blood by LC–MS/MS in Combination with Phospholipid-Depletion Solid-Phase Extraction. Analytical Chemistry, 2015, 87, 1127-1136.  | <b>6.</b> 5  | 134       |
| 9  | Electron Capture Dissociation of Electrosprayed Protein Ions for Spatially Resolved Hydrogen Exchange Measurements. Journal of the American Chemical Society, 2008, 130, 11574-11575.   | 13.7         | 111       |
| 10 | Defects in myosin VB are associated with a spectrum of previously undiagnosed low $\hat{l}^3 \hat{a} \in g$ lutamyltransferase cholestasis. Hepatology, 2017, 65, 1655-1669.  | 7.3          | 107       |
| 11 | Recent advancements in matrix-assisted laser desorption/ionization mass spectrometry imaging. Current Opinion in Biotechnology, 2017, 43, 62-69.  | 6.6          | 107       |
| 12 | Impact of <i>Salmonella</i> Infection on Host Hormone Metabolism Revealed by Metabolomics. Infection and Immunity, 2011, 79, 1759-1769.   | 2.2          | 104       |
| 13 | Metabolomic analysis of key central carbon metabolism carboxylic acids as their 3â€nitrophenylhydrazones by <scp>UPLC</scp> / <scp>ESI</scp> â€ <scp>MS</scp> . Electrophoresis, 2013, 34, 2891-2900.                               | 2.4          | 100       |
| 14 | <scp><i>M</i></scp> <i>ycobacterium leprae</i> intracellular survival relies on cholesterol accumulation in infected macrophages: a potential target for new drugs for leprosy treatment. Cellular Microbiology, 2014, 16, 797-815. | 2.1          | 83        |
| 15 | Characterizing Short-Lived Protein Folding Intermediates by Top-Down Hydrogen Exchange Mass<br>Spectrometry. Analytical Chemistry, 2010, 82, 8591-8597.   | 6.5          | 78        |
| 16 | Hydroxyflavones as a New Family of Matrices for MALDI Tissue Imaging. Analytical Chemistry, 2013, 85, 7566-7573.  | 6.5          | 72        |
| 17 | MALDI Tissue Imaging of Ocular Lens α-Crystallin. , 2006, 47, 2990.   |              | 70        |
| 18 | Metabolomic profiling of a modified alcohol liquid diet model for liver injury in the mouse uncovers new markers of disease. Toxicology and Applied Pharmacology, 2008, 232, 236-243.   | 2.8          | 67        |

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|----|--|------|-----------|
| 19 | Structure and Dynamics of Small Soluble Aβ(1–40) Oligomers Studied by Top-Down Hydrogen Exchange Mass Spectrometry. Biochemistry, 2012, 51, 3694-3703.   | 2.5  | 64        |
| 20 | Conformer-Specific Hydrogen Exchange Analysis of Aβ(1–42) Oligomers by Top-Down Electron Capture Dissociation Mass Spectrometry. Analytical Chemistry, 2011, 83, 5386-5393.  | 6.5  | 62        |
| 21 | Proteomics and Phosphoproteomics Analysis of Human Lens Fiber Cell Membranes., 2013, 54, 1135.   |      | 61        |
| 22 | Mass spectrometry-based technologies for high-throughput metabolomics. Bioanalysis, 2009, 1, 1665-1684.  | 1.5  | 60        |
| 23 | Comprehensive Imaging of Porcine Adrenal Gland Lipids by MALDI-FTMS Using Quercetin as a Matrix.<br>Analytical Chemistry, 2014, 86, 638-646.   | 6.5  | 56        |
| 24 | Comprehensive Analysis of Oil Sands Processed Water by Direct-Infusion Fourier-Transform Ion Cyclotron Resonance Mass Spectrometry with and without Offline UHPLC Sample Prefractionation. Environmental Science & Environment | 10.0 | 49        |
| 25 | Metabolomics: towards understanding host–microbe interactions. Future Microbiology, 2010, 5, 153-161.  | 2.0  | 48        |
| 26 | Analysis of Selected Sugars and Sugar Phosphates in Mouse Heart Tissue by Reductive Amination and Liquid Chromatography-Electrospray Ionization Mass Spectrometry. Analytical Chemistry, 2013, 85, 5965-5973.  | 6.5  | 45        |
| 27 | Proteolysis and Mass Spectrometric Analysis of an Integral Membrane:Â Aquaporin 0. Journal of Proteome Research, 2004, 3, 807-812.   | 3.7  | 44        |
| 28 | Mass Spectrometry-Based Structural Proteomics. European Journal of Mass Spectrometry, 2012, 18, 251-267.   | 1.0  | 44        |
| 29 | Dithranol as a MALDI Matrix for Tissue Imaging of Lipids by Fourier Transform Ion Cyclotron<br>Resonance Mass Spectrometry. Analytical Chemistry, 2012, 84, 8391-8398.   | 6.5  | 44        |
| 30 | Metabolic Signatures of Triatomine Vectors of Trypanosoma cruzi Unveiled by Metabolomics. PLoS ONE, 2013, 8, e77283.   | 2.5  | 43        |
| 31 | Metabonomics Reveals Drastic Changes in Anti-Inflammatory/Pro-Resolving Polyunsaturated Fatty<br>Acids-Derived Lipid Mediators in Leprosy Disease. PLoS Neglected Tropical Diseases, 2013, 7, e2381.   | 3.0  | 41        |
| 32 | The Deubiquitinase Activity of the Salmonella Pathogenicity Island 2 Effector, SseL, Prevents Accumulation of Cellular Lipid Droplets. Infection and Immunity, 2011, 79, 4392-4400.  | 2.2  | 40        |
| 33 | Verification and spatial localization of aquaporin-5 in the ocular lens. Experimental Eye Research, 2013, 108, 94-102.   | 2.6  | 40        |
| 34 | Spatial Differences in an Integral Membrane Proteome Detected in Laser Capture Microdissected Samples. Journal of Proteome Research, 2008, 7, 2696-2702.   | 3.7  | 38        |
| 35 | Matrix coating assisted by an electric field (MCAEF) for enhanced tissue imaging by MALDI-MS. Chemical Science, 2015, 6, 729-738.  | 7.4  | 36        |
| 36 | Metabolomic profiling of prostate cancer by matrix assisted laser desorption/ionization-Fourier transform ion cyclotron resonance mass spectrometry imaging using Matrix Coating Assisted by an Electric Field (MCAEF). Biochimica Et Biophysica Acta - Proteins and Proteomics, 2017, 1865, 755-767.  | 2.3  | 35        |

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|----|---|-----|-----------|
| 37 | Intense Light-Mediated Circadian Cardioprotection via Transcriptional Reprogramming of the Endothelium. Cell Reports, 2019, 28, 1471-1484.e11.  | 6.4 | 35        |
| 38 | Metabolomics Reveals Phospholipids as Important Nutrient Sources during Salmonella Growth in Bile In Vitro and <i>In Vivo</i> . Journal of Bacteriology, 2011, 193, 4719-4725.  | 2.2 | 32        |
| 39 | Metabolomic insights into system-wide coordination of vertebrate metamorphosis. BMC Developmental Biology, 2014, 14, 5.   | 2.1 | 32        |
| 40 | Repression of Salmonella Host Cell Invasion by Aromatic Small Molecules from the Human Fecal Metabolome. Applied and Environmental Microbiology, 2017, 83, .  | 3.1 | 31        |
| 41 | Freeâ€flow electrophoresis for topâ€down proteomics by Fourier transform ion cyclotron resonance mass spectrometry. Proteomics, 2008, 8, 2798-2808.   | 2.2 | 30        |
| 42 | Isotope-labeling derivatization with 3-nitrophenylhydrazine for LC/multiple-reaction monitoring-mass-spectrometry-based quantitation of carnitines in dried blood spots. Analytica Chimica Acta, 2018, 1037, 177-187. | 5.4 | 29        |
| 43 | Hydrophilic bile acids prevent liver damage caused by lack of biliary phospholipid in Mdr2 mice.<br>Journal of Lipid Research, 2019, 60, 85-97.   | 4.2 | 28        |
| 44 | MALDI tissue profiling of integral membrane proteins from ocular tissues. Journal of the American Society for Mass Spectrometry, 2008, 19, 814-822.   | 2.8 | 25        |
| 45 | Quantitation of low molecular weight sugars by chemical derivatizationâ€liquid chromatography/multiple reaction monitoring/mass spectrometry. Electrophoresis, 2016, 37, 1851-1860.                                   | 2.4 | 25        |
| 46 | Cardiac Ryanodine Receptor (Ryr2)-mediated Calcium Signals Specifically Promote Glucose Oxidation via Pyruvate Dehydrogenase. Journal of Biological Chemistry, 2016, 291, 23490-23505.                                | 3.4 | 23        |
| 47 | Increased sulfation of bile acids in mice and human subjects with sodium taurocholate cotransporting polypeptide deficiency. Journal of Biological Chemistry, 2019, 294, 11853-11862.                                 | 3.4 | 22        |
| 48 | Repression of Salmonella enterica <i>phoP</i> Expression by Small Molecules from Physiological Bile. Journal of Bacteriology, 2012, 194, 2286-2296.   | 2.2 | 19        |
| 49 | The use of matrix coating assisted by an electric field (MCAEF) to enhance mass spectrometric imaging of human prostate cancer biomarkers. Journal of Mass Spectrometry, 2016, 51, 86-95.                             | 1.6 | 19        |
| 50 | Using multiple structural proteomics approaches for the characterization of prion proteins. Journal of Proteomics, 2013, 81, 31-42.   | 2.4 | 18        |
| 51 | Accurate molecular weight analysis of histones using FFE and RPâ€HPLC on monolithic capillary columns. Journal of Separation Science, 2009, 32, 2691-2698.  | 2.5 | 17        |
| 52 | Peering into molecular mechanisms of action with frogSCOPE. General and Comparative Endocrinology, 2010, 168, 190-198.  | 1.8 | 17        |
| 53 | Comprehensive bile acid profiling in hereditary intrahepatic cholestasis: Genetic and clinical correlations. Liver International, 2018, 38, 1676-1685.  | 3.9 | 14        |
| 54 | Top-down hydrogen/deuterium exchange and ECD-stitched FTICR-MS for probing structural dynamics of a 29-kDa enzyme. International Journal of Mass Spectrometry, 2012, 325-327, 130-138.                                | 1.5 | 12        |

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|----|--|------|-----------|
| 55 | Metabolomic insights into the effects of thyroid hormone on Rana [Lithobates] catesbeiana metamorphosis using whole-body Matrix Assisted Laser Desorption/Ionization-Mass Spectrometry Imaging (MALDI-MSI). General and Comparative Endocrinology, 2018, 265, 237-245. | 1.8  | 12        |
| 56 | Depletion of essential isoprenoids and ER stress induction following acute liver-specific deletion of HMG-CoA reductase. Journal of Lipid Research, 2020, 61, 1675-1686.   | 4.2  | 12        |
| 57 | Mapping of protein phosphorylation by dual enzyme digestion and matrix-assisted laser desorption ionization–quadrupole orthogonal time-of-flight mass spectrometry. Analytical Biochemistry, 2002, 310, 215-218.   | 2.4  | 10        |
| 58 | Comment on "Profiling Oil Sands Mixtures from Industrial Developments and Natural Groundwaters for Source Identification― Environmental Science & Environmental Science & 11013-11014.   | 10.0 | 10        |
| 59 | The Gut Microbiome and Metabolome of Two Riparian Communities in the Amazon. Frontiers in Microbiology, 2019, 10, 2003.  | 3.5  | 10        |
| 60 | Topâ€down analysis of recombinant histone H3 and its methylated analogs by ESI/FTâ€ICR mass spectrometry. Proteomics, 2010, 10, 3621-3630.   | 2.2  | 8         |
| 61 | NTCP Deficiency Causes Gallbladder Abnormalities in Mice and Human Beings. Cellular and Molecular Gastroenterology and Hepatology, 2021, 11, 831-839.  | 4.5  | 7         |
| 62 | Profiling of dissolved organic compounds in the oil sands region using complimentary liquid–liquid extraction and ultrahigh resolution Fourier transform mass spectrometry. Environmental Earth Sciences, 2017, 76, 1.   | 2.7  | 6         |
| 63 | Changes in plasma bile acid profiles after partial internal biliary diversion in PFIC2 patients. Annals of Translational Medicine, 2020, 8, 185-185.   | 1.7  | 5         |
| 64 | Acceleration of age-induced proteolysis in the guinea pig lens nucleus by in vivo exposure to hyperbaric oxygen: A mass spectrometry analysis. Experimental Eye Research, 2021, 210, 108697.   | 2.6  | 5         |
| 65 | Sequence and peptide map of guinea pig aquaporin 0. Molecular Vision, 2004, 10, 215-22.  | 1.1  | 5         |
| 66 | Molecular profiling of naphthenic acids in technical mixtures and oil sands processâ€affected water using polar reversedâ€phase liquid chromatography–mass spectrometry. Electrophoresis, 2016, 37, 3089-3100.   | 2.4  | 4         |
| 67 | Emerging Mass Spectrometry-Based Technologies for Analyses of Chromatin Changes: Analysis of Histones and Histone Modifications. Methods in Molecular Biology, 2011, 773, 259-303.   | 0.9  | 4         |
| 68 | Circulating Isovalerylcarnitine and Lung Cancer Risk: Evidence from Mendelian Randomization and Prediagnostic Blood Measurements. Cancer Epidemiology Biomarkers and Prevention, 2022, 31, 1966-1974.  | 2.5  | 4         |
| 69 | Dithranol as a Matrix for Matrix Assisted Laser Desorption/Ionization Imaging on a Fourier Transform Ion Cyclotron Resonance Mass Spectrometer. Journal of Visualized Experiments, 2013, , e50733.   | 0.3  | 3         |
| 70 | Supermolecule-assisted imaging of low-molecular-weight quaternary-ammonium compounds by MALDI-MS of their non-covalent complexes with cucurbit[7]uril. RSC Advances, 2020, 10, 34261-34265.  | 3.6  | 2         |
| 71 | Intestinal Deletion of 3-Hydroxy-3-Methylglutaryl-Coenzyme A Reductase Promotes Expansion of the Resident Stem Cell Compartment. Arteriosclerosis, Thrombosis, and Vascular Biology, 2022, 42, 381-394.  | 2.4  | 1         |