## Iain D G Campuzano

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9245520/publications.pdf

Version: 2024-02-01

41 papers 3,787 citations

26 h-index

218677

289244 40 g-index

41 all docs

41 docs citations

41 times ranked

3243 citing authors

| #  | Article  | IF   | Citations  |
|----|--|------|------------|
| 1  | Evidence for Macromolecular Protein Rings in the Absence of Bulk Water. Science, 2005, 310, 1658-1661.   | 12.6 | 551        |
| 2  | Structural Characterization of Drug-like Compounds by Ion Mobility Mass Spectrometry: Comparison of Theoretical and Experimentally Derived Nitrogen Collision Cross Sections. Analytical Chemistry, 2012, 84, 1026-1033.                           | 6.5  | 340        |
| 3  | Enhancements in travelling wave ion mobility resolution. Rapid Communications in Mass Spectrometry, 2011, 25, 1559-1566.   | 1.5  | 334        |
| 4  | Recommendations for reporting ion mobility Mass Spectrometry measurements. Mass Spectrometry Reviews, 2019, 38, 291-320.   | 5.4  | 315        |
| 5  | Deciphering Drift Time Measurements from Travelling Wave Ion Mobility Spectrometry-Mass<br>Spectrometry Studies. European Journal of Mass Spectrometry, 2009, 15, 113-130.   | 1.0  | 312        |
| 6  | Ion Mobility Mass Spectrometry of Peptide Ions: Effects of Drift Gas and Calibration Strategies. Analytical Chemistry, 2012, 84, 7124-7130.  | 6.5  | 281        |
| 7  | Best practices and benchmarks for intact protein analysis for top-down mass spectrometry. Nature<br>Methods, 2019, 16, 587-594.  | 19.0 | 241        |
| 8  | An integrated native mass spectrometry and top-down proteomics method that connects sequence to structure and function of macromolecular complexes. Nature Chemistry, 2018, 10, 139-148.   | 13.6 | 170        |
| 9  | Travelingâ€wave ion mobility mass spectrometry of protein complexes: accurate calibrated collision crossâ€sections of human insulin oligomers. Rapid Communications in Mass Spectrometry, 2012, 26, 1181-1193.                                     | 1.5  | 138        |
| 10 | Baseline resolution of isomers by traveling wave ion mobility mass spectrometry: investigating the effects of polarizable drift gases and ionic charge distribution. Journal of Mass Spectrometry, 2013, 48, 989-997.                              | 1.6  | 77         |
| 11 | Use of ion mobility mass spectrometry and a collision crossâ€section algorithm to study an organometallic ruthenium anticancer complex and its adducts with a DNA oligonucleotide. Rapid Communications in Mass Spectrometry, 2009, 23, 3563-3569. | 1.5  | <b>7</b> 5 |
| 12 | A method for direct measurement of ion mobilities using a travelling wave ion guide. International Journal of Mass Spectrometry, 2010, 298, 10-16.   | 1.5  | 74         |
| 13 | Isomer separation and gas-phase configurations of organoruthenium anticancer complexes: Ion mobility mass spectrometry and modeling. Journal of the American Society for Mass Spectrometry, 2009, 20, 1119-1122.                                   | 2.8  | <b>7</b> 3 |
| 14 | Sites of metabolic substitution: investigating metabolite structures utilising ion mobility and molecular modelling. Rapid Communications in Mass Spectrometry, 2010, 24, 3157-3162.   | 1.5  | 70         |
| 15 | Discovery of <i>N</i> -(1-Acryloylazetidin-3-yl)-2-(1 <i>H</i> -indol-1-yl)acetamides as Covalent Inhibitors of KRAS <sup>G12C</sup> . ACS Medicinal Chemistry Letters, 2019, 10, 1302-1308.   | 2.8  | 66         |
| 16 | Shape changes induced by N-terminal platination of ubiquitin by cisplatin. Journal of the American Society for Mass Spectrometry, 2010, 21, 1097-1106.   | 2.8  | 50         |
| 17 | Product ion mobility as a promising tool for assignment of positional isomers of drug metabolites.<br>Rapid Communications in Mass Spectrometry, 2011, 25, 3497-3503.  | 1.5  | 50         |
| 18 | Effects of Drift Gas on Collision Cross Sections of a Protein Standard in Linear Drift Tube and Traveling Wave Ion Mobility Mass Spectrometry. Analytical Chemistry, 2012, 84, 8524-8531.  | 6.5  | 47         |

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|----|--|-----|-----------|
| 19 | Assigning Structures to Gas-Phase Peptide Cations and Cation-Radicals. An Infrared Multiphoton Dissociation, Ion Mobility, Electron Transfer, and Computational Study of a Histidine Peptide Ion. Journal of Physical Chemistry B, 2012, 116, 3445-3456. | 2.6 | 47        |
| 20 | Native MS Analysis of Bacteriorhodopsin and an Empty Nanodisc by Orthogonal Acceleration Time-of-Flight, Orbitrap and Ion Cyclotron Resonance. Analytical Chemistry, 2016, 88, 12427-12436.  | 6.5 | 44        |
| 21 | High-Throughput Mass Spectrometric Analysis of Covalent Protein-Inhibitor Adducts for the Discovery of Irreversible Inhibitors: A Complete Workflow. Journal of Biomolecular Screening, 2016, 21, 136-144.   | 2.6 | 42        |
| 22 | Submicrometer Emitter ESI Tips for Native Mass Spectrometry of Membrane Proteins in Ionic and Nonionic Detergents. Journal of the American Society for Mass Spectrometry, 2018, 29, 203-206.   | 2.8 | 41        |
| 23 | Native-MS Analysis of Monoclonal Antibody Conjugates by Fourier Transform Ion Cyclotron<br>Resonance Mass Spectrometry. Analytical Chemistry, 2018, 90, 745-751.   | 6.5 | 36        |
| 24 | Structural Resolution of 4-Substituted Proline Diastereomers with Ion Mobility Spectrometry via Alkali Metal Ion Cationization. Analytical Chemistry, 2015, 87, 3300-3307.   | 6.5 | 35        |
| 25 | Native and Denaturing MS Protein Deconvolution for Biopharma: Monoclonal Antibodies and Antibody–Drug Conjugates to Polydisperse Membrane Proteins and Beyond. Analytical Chemistry, 2019, 91, 9472-9480.  | 6.5 | 32        |
| 26 | Fourier Transform-Ion Cyclotron Resonance Mass Spectrometry as a Platform for Characterizing Multimeric Membrane Protein Complexes. Journal of the American Society for Mass Spectrometry, 2018, 29, 183-193.  | 2.8 | 29        |
| 27 | Denaturing and Native Mass Spectrometric Analytics for Biotherapeutic Drug Discovery Research: Historical, Current, and Future Personal Perspectives. Journal of the American Society for Mass Spectrometry, 2021, 32, 1861-1885.                        | 2.8 | 27        |
| 28 | Quantitative collisionâ€induced unfolding differentiates model antibody–drug conjugates. Protein Science, 2019, 28, 598-608.   | 7.6 | 26        |
| 29 | Extracting Charge and Mass Information from Highly Congested Mass Spectra Using Fourier-Domain Harmonics. Journal of the American Society for Mass Spectrometry, 2018, 29, 2067-2080.  | 2.8 | 23        |
| 30 | Coupling electrospray corona discharge, charge reduction and ion mobility mass spectrometry: From peptides to large macromolecular protein complexes. International Journal for Ion Mobility Spectrometry, 2013, 16, 51-60.                              | 1.4 | 19        |
| 31 | Ion Mobility and Mass Spectrometry Measurements of the Humanized IgGk NIST Monoclonal Antibody. ACS Symposium Series, 2015, , 75-112.  | 0.5 | 19        |
| 32 | Nanospray Ion Mobility Mass Spectrometry of Selected High Mass Species. Methods in Molecular Biology, 2011, 790, 57-70.  | 0.9 | 19        |
| 33 | High Mass Analysis with a Fourier Transform Ion Cyclotron Resonance Mass Spectrometer: From Inorganic Salt Clusters to Antibody Conjugates and Beyond. Journal of the American Society for Mass Spectrometry, 2020, 31, 1155-1162.                       | 2.8 | 16        |
| 34 | Intrinsic Mobility of Gaseous Cationic and Anionic Aggregates of Ionic Liquids. ChemPhysChem, 2011, 12, 1444-1447.   | 2.1 | 14        |
| 35 | Quantification of siRNA-Antibody Conjugates in Biological Matrices by Triplex-Forming<br>Oligonucleotide ELISA. Nucleic Acid Therapeutics, 2019, 29, 161-166.  | 3.6 | 13        |
| 36 | Rapid LC–MS Method for Accurate Molecular Weight Determination of Membrane and Hydrophobic Proteins. Analytical Chemistry, 2018, 90, 13616-13623.  | 6.5 | 12        |

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|----|--|-----|-----------|
| 37 | High-Resolution Demultiplexing (HRdm) Ion Mobility Spectrometry–Mass Spectrometry for Aspartic and Isoaspartic Acid Determination and Screening. Analytical Chemistry, 2022, 94, 6191-6199.  | 6.5 | 12        |
| 38 | Purification of guanine-quadruplex using monolithic stationary phase under ion-exchange conditions. Journal of Chromatography A, 2020, 1634, 461633.   | 3.7 | 10        |
| 39 | Unequivocal Identification of Aspartic Acid and <i>iso</i> Aspartic Acid by MALDI-TOF/TOF: From Peptide Standards to a Therapeutic Antibody. Journal of the American Society for Mass Spectrometry, 2021, 32, 1901-1909.   | 2.8 | 6         |
| 40 | Editorial and Review: 30th ASMS Sanibel Conference on Mass Spectrometryâ€"Computational Modelling in Mass Spectrometry and Ion Mobility: Methods for Ion Structure and Reactivity Determination. Journal of the American Society for Mass Spectrometry, 2018, 29, 2283-2286. | 2.8 | 1         |
| 41 | Editorial: Special JASMS Focus on Mass Spectrometry in Industry. Journal of the American Society for Mass Spectrometry, 2021, 32, 1850-1851.   | 2.8 | 0         |