## Wilfried Niessen

## List of Publications by Year in descending order

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430874 302126 1,837 39 18 39 citations h-index g-index papers 39 39 39 2088 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Matrix effects in quantitative pesticide analysis using liquid chromatography–mass spectrometry. Mass Spectrometry Reviews, 2006, 25, 881-899.	5.4	385
2	Bioanalytical LC–MS/MS of protein-based biopharmaceuticals. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2013, 929, 161-179.	2.3	204
3	Progress in liquid chromatography–mass spectrometry instrumentation and its impact on high-throughput screening. Journal of Chromatography A, 2003, 1000, 413-436.	3.7	178
4	Use of quadrupole time-of-flight mass spectrometry in the elucidation of unknown compounds present in environmental water. Rapid Communications in Mass Spectrometry, 2005, 19, 169-178.	1.5	132
5	Confirmation of organic micropollutants detected in environmental samples by liquid chromatography tandem mass spectrometry: Achievements and pitfalls. TrAC - Trends in Analytical Chemistry, 2006, 25, 1030-1042.	11.4	101
6	Fragmentation of toxicologically relevant drugs in positiveâ€ion liquid chromatography–tandem mass spectrometry. Mass Spectrometry Reviews, 2011, 30, 626-663.	5.4	87
7	High-performance liquid chromatography–mass spectrometry-based acetylcholinesterase assay for the screening of inhibitors in natural extracts. Journal of Chromatography A, 2006, 1112, 303-310.	3.7	84
8	Interstellar dust, chirality, comets and the origins of life: Life from dead stars?. Journal of Biological Physics, 1995, 20, 61-70.	1.5	68
9	Targeted LC–MS derivatization for aldehydes and carboxylic acids with a new derivatization agent 4-APEBA. Analytical and Bioanalytical Chemistry, 2010, 397, 665-675.	3.7	66
10	Mass spectrometry-based biochemical assays for enzyme-inhibitor screening. TrAC - Trends in Analytical Chemistry, 2007, 26, 867-883.	11.4	62
11	Group-specific fragmentation of pesticides and related compounds in liquid chromatography–tandem mass spectrometry. Journal of Chromatography A, 2010, 1217, 4061-4070.	3.7	59
12	Fragmentation of toxicologically relevant drugs in negativeâ€ion liquid chromatography–tandem mass spectrometry. Mass Spectrometry Reviews, 2012, 31, 626-665.	5.4	40
13	Determination and identification of estrogenic compounds generated with biosynthetic enzymes using hyphenated screening assays, high resolution mass spectrometry and off-line NMR. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2010, 878, 667-674.	2.3	38
14	Production and on-line acetylcholinesterase bioactivity profiling of chemical and biological degradation products of tacrine. Journal of Pharmaceutical and Biomedical Analysis, 2010, 53, 609-616.	2.8	27
15	Stability-indicating study of the anti-Alzheimer's drug galantamine hydrobromide. Journal of Pharmaceutical and Biomedical Analysis, 2011, 55, 85-92.	2.8	25
16	At-line nanofractionation with parallel mass spectrometry and bioactivity assessment for the rapid screening of thrombin and factor Xa inhibitors in snake venoms. Toxicon, 2016, 110, 79-89.	1.6	23
17	Neurotoxicity fingerprinting of venoms using on-line microfluidic AChBP profiling. Toxicon, 2018, 148, 213-222.	1.6	23
18	An efficient analytical platform for on-line microfluidic profiling of neuroactive snake venoms towards nicotinic receptor affinity. Toxicon, 2013, 61, 112-124.	1.6	22

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19	Analysis of acetylcholinesterase inhibitors: bioanalysis, degradation and metabolism. Biomedical Chromatography, 2011, 25, 278-299.	1.7	20
20	Developing strategies for isolation of minor impurities with mass spectrometry-directed fractionation. Journal of Chromatography A, 2002, 970, 131-140.	3.7	18
21	Identification of very long-chain (>C24) fatty acid methyl esters using gas chromatography coupled to quadrupole/time-of-flight mass spectrometry with atmospheric pressure chemical ionization source. Analytica Chimica Acta, 2019, 1051, 103-109.	5.4	18
22	High Throughput Screening Methodologies Classified for Major Drug Target Classes According to Target Signaling Pathways. Combinatorial Chemistry and High Throughput Screening, 2010, 13, 548-561.	1.1	17
23	Analytical workflow for rapid screening and purification of bioactives from venom proteomes. Toxicon, 2013, 76, 270-281.	1.6	16
24	Miniaturized Bioaffinity Assessment Coupled to Mass Spectrometry for Guided Purification of Bioactives from Toad and Cone Snail. Biology, 2014, 3, 139-156.	2.8	16
25	Solution-phase electrochemistry-nuclear magnetic resonance of small organic molecules. TrAC - Trends in Analytical Chemistry, 2015, 70, 31-39.	11.4	16
26	Photon-Independent Gas-Phase-Ion Formation in Capillary Electrophoresisâ^'Mass Spectrometry Using Atmospheric Pressure Photoionization. Analytical Chemistry, 2007, 79, 5351-5357.	6.5	15
27	Identification and quantification of drug–albumin adducts in serum samples from a drug exposure study in mice. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2013, 917-918, 53-61.	2.3	14
28	High-resolution metabolic profiling towards G protein-coupled receptors: Rapid and comprehensive screening of histamine H4 receptor ligands. Journal of Chromatography A, 2012, 1259, 213-220.	3.7	11
29	EC–SPE–stripline-NMR analysis of reactive products: a feasibility study. Analytical and Bioanalytical Chemistry, 2013, 405, 6711-6720.	3.7	11
30	At-line coupling of LC–MS to bioaffinity and selectivity assessment for metabolic profiling of ligands towards chemokine receptors CXCR1 and CXCR2. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2015, 1002, 42-53.	2.3	9
31	Tandem mass spectrometry study of p38α kinase inhibitors and related substances. Journal of Mass Spectrometry, 2013, 48, 718-731.	1.6	7
32	On-line coupling of surface plasmon resonance optical sensing to size-exclusion chromatography for affinity assessment of antibody samples. Journal of Chromatography A, 2016, 1452, 81-88.	3.7	6
33	Photohuperzine Aâ€"A new photoisomer of huperzine A: Structure elucidation, formation kinetics and activity assessment. Journal of Pharmaceutical and Biomedical Analysis, 2010, 52, 190-194.	2.8	5
34	Tandem mass spectrometry of organic nitro and halogen compounds: Competition between losses of molecules and of radicals. International Journal of Mass Spectrometry, 2021, 460, 116496.	1.5	5
35	Development of On-line Liquid Chromatography-Biochemical Detection for Soluble Epoxide Hydrolase Inhibitors in Mixtures. Chromatographia, 2013, 76, 13-21.	1.3	3
36	Development of Plate Reader and On-Line Microfluidic Screening to Identify Ligands of the 5-Hydroxytryptamine Binding Protein in Venoms. Toxins, 2015, 7, 2336-2353.	3.4	2

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#	Article	IF	CITATIONS
37	Interpretation of MS–MS spectra of small-molecule signal transduction inhibitors using accurate-m/z data and m/z-shifts with stable-isotope-labeled analogues and metabolites. International Journal of Mass Spectrometry, 2021, 464, 116559.	1.5	2
38	Optimization of information content in a mass spectrometry based flow-chemistry system by investigating different ionization approaches. Talanta, 2011, 84, 623-631.	5.5	1
39	Tandem mass spectrometry of small-molecule signal transduction inhibitors: Accurate-m/z data to adapt structure proposals of product ions. Journal of Pharmaceutical and Biomedical Analysis, 2021, 195, 113864.	2.8	1