

# Scott J Walmsley

## List of Publications by Year in descending order

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Version: 2024-02-01

19  
papers

441  
citations

758635

12  
h-index

839053

18  
g-index

19  
all docs

19  
docs citations

19  
times ranked

872  
citing authors

#	ARTICLE	IF	CITATIONS
1	LuciPHOR: Algorithm for Phosphorylation Site Localization with False Localization Rate Estimation Using Modified Target-Decoy Approach. <i>Molecular and Cellular Proteomics</i> , 2013, 12, 3409-3419.	2.5	69
2	Comprehensive Analysis of Protein Digestion Using Six Trypsins Reveals the Origin of Trypsin As a Significant Source of Variability in Proteomics. <i>Journal of Proteome Research</i> , 2013, 12, 5666-5680.	1.8	63
3	Customized Consensus Spectral Library Building for Untargeted Quantitative Metabolomics Analysis with Data Independent Acquisition Mass Spectrometry and MetaboDIA Workflow. <i>Analytical Chemistry</i> , 2017, 89, 4897-4906.	3.2	42
4	Cloning and characterization of five cDNAs encoding peritrophin-A domains from the cat flea, <i>Ctenocephalides felis</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2003, 33, 1061-1073.	1.2	41
5	Performance of a High-Pressure Liquid Chromatography-Ion Mobility-Mass Spectrometry System for Metabolic Profiling. <i>Analytical Chemistry</i> , 2017, 89, 6384-6391.	3.2	40
6	Proteomic analysis of brush-border membrane vesicles isolated from purified proximal convoluted tubules. <i>American Journal of Physiology - Renal Physiology</i> , 2010, 298, F1323-F1331.	1.3	32
7	Identification of serum biomarkers for canine B-cell lymphoma by use of surface-enhanced laser desorption-ionization time-of-flight mass spectrometry. <i>American Journal of Veterinary Research</i> , 2007, 68, 405-410.	0.3	25
8	Metabolism and biomarkers of heterocyclic aromatic amines in humans. <i>Genes and Environment</i> , 2021, 43, 29.	0.9	22
9	Methods and Challenges for Computational Data Analysis for DNA Adductomics. <i>Chemical Research in Toxicology</i> , 2019, 32, 2156-2168.	1.7	16
10	A Perspective and Framework for Developing Sample Type Specific Databases for LC/MS-Based Clinical Metabolomics. <i>Metabolites</i> , 2020, 10, 8.	1.3	16
11	Development of a DNA Adductome Mass Spectral Database. <i>Chemical Research in Toxicology</i> , 2020, 33, 852-854.	1.7	16
12	Proteomic profiling of the effect of metabolic acidosis on the apical membrane of the proximal convoluted tubule. <i>American Journal of Physiology - Renal Physiology</i> , 2012, 302, F1465-F1477.	1.3	14
13	Comprehensive Analysis of DNA Adducts Using Data-Independent wSIM/MS <sup>2</sup> Acquisition and wSIM-City. <i>Analytical Chemistry</i> , 2021, 93, 6491-6500.	3.2	11
14	A prototypic small molecule database for bronchoalveolar lavage-based metabolomics. <i>Scientific Data</i> , 2018, 5, 180060.	2.4	10
15	Effects of <i>GSTT1</i> Genotype on the Detoxification of 1,3-Butadiene Derived Diepoxide and Formation of Promutagenic DNA-DNA Cross-Links in Human Hapmap Cell Lines. <i>Chemical Research in Toxicology</i> , 2021, 34, 119-131.	1.7	10
16	Kinetics of DNA Adducts and Abasic Site Formation in Tissues of Mice Treated with a Nitrogen Mustard. <i>Chemical Research in Toxicology</i> , 2020, 33, 988-998.	1.7	7
17	The Cooked Meat Carcinogen 2-Amino-1-methyl-6-phenylimidazo[4,5- <i>b</i> ]pyridine Hair Dosimeter, DNA Adductomics Discovery, and Associations with Prostate Cancer Pathology Biomarkers. <i>Chemical Research in Toxicology</i> , 2022, 35, 703-730.	1.7	4
18	Identification of two cDNAs encoding synaptic vesicle protein 2 (SV2)-like proteins from epithelial tissues in the cat flea, <i>Ctenocephalides felis</i> . <i>Insect Molecular Biology</i> , 2004, 13, 225-230.	1.0	3

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19	Truncated rank correlation (TRC) as a robust measure of test-retest reliability in mass spectrometry data. <i>Statistical Applications in Genetics and Molecular Biology</i> , 2019, 18, .	0.2	0