Kristin E Burnum

List of Publications by Year in descending order

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117625 133252 3,856 67 34 h-index citations papers

g-index 71 71 71 6373 docs citations times ranked citing authors all docs

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#	Article	IF	Citations
1	Solvent-free matrix dry-coating for MALDI imaging of phospholipids. Journal of the American Society for Mass Spectrometry, 2008, 19, 882-886.	2.8	211
2	Uncovering biologically significant lipid isomers with liquid chromatography, ion mobility spectrometry and mass spectrometry. Analyst, The, 2016, 141, 1649-1659.	3.5	196
3	Proteomic Profiling of Exosomes Leads to the Identification of Novel Biomarkers for Prostate Cancer. PLoS ONE, 2013, 8, e82589.	2.5	179
4	Automated mass spectrometry imaging of over 2000 proteins from tissue sections at 100- $\hat{1}\frac{1}{4}$ m spatial resolution. Nature Communications, 2020, 11, 8.	12.8	178
5	MPLEx: a Robust and Universal Protocol for Single-Sample Integrative Proteomic, Metabolomic, and Lipidomic Analyses. MSystems, 2016, 1 , .	3.8	166
6	MERS-CoV and H5N1 influenza virus antagonize antigen presentation by altering the epigenetic landscape. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E1012-E1021.	7.1	142
7	Spatial and temporal alterations of phospholipids determined by mass spectrometry during mouse embryo implantation. Journal of Lipid Research, 2009, 50, 2290-2298.	4.2	136
8	Quantitative proteomics analysis of adsorbed plasma proteins classifies nanoparticles with different surface properties and size. Proteomics, 2011, 11, 4569-4577.	2.2	135
9	Membrane Stresses Induced by Overproduction of Free Fatty Acids in Escherichia coli. Applied and Environmental Microbiology, 2011, 77, 8114-8128.	3.1	135
10	Metagenomic and metaproteomic insights into bacterial communities in leaf-cutter ant fungus gardens. ISME Journal, 2012, 6, 1688-1701.	9.8	126
11	High spatial resolution imaging of biological tissues using nanospray desorption electrospray ionization mass spectrometry. Nature Protocols, 2019, 14, 3445-3470.	12.0	125
12	Leucoagaricus gongylophorus Produces Diverse Enzymes for the Degradation of Recalcitrant Plant Polymers in Leaf-Cutter Ant Fungus Gardens. Applied and Environmental Microbiology, 2013, 79, 3770-3778.	3.1	98
13	Multi-platform 'Omics Analysis of Human Ebola Virus Disease Pathogenesis. Cell Host and Microbe, 2017, 22, 817-829.e8.	11.0	88
14	Matrix-Assisted Laser Desorption/Ionization Imaging Mass Spectrometry for the Investigation of Proteins and Peptides. Annual Review of Analytical Chemistry, 2008, 1, 689-705.	5.4	86
15	Coupling Front-End Separations, Ion Mobility Spectrometry, and Mass Spectrometry For Enhanced Multidimensional Biological and Environmental Analyses. Annual Review of Analytical Chemistry, 2017, 10, 71-92.	5 . 4	84
16	High Spatial Resolution Imaging of Mouse Pancreatic Islets Using Nanospray Desorption Electrospray Ionization Mass Spectrometry. Analytical Chemistry, 2018, 90, 6548-6555.	6.5	76
17	Metabolic Reprogramming during Purine Stress in the Protozoan Pathogen Leishmania donovani. PLoS Pathogens, 2014, 10, e1003938.	4.7	74
18	Mass spectrometry for translational proteomics: progress and clinical implications. Genome Medicine, 2012, 4, 63.	8.2	71

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19	Ion mobility spectrometry and the omics: Distinguishing isomers, molecular classes and contaminant ions in complex samples. TrAC - Trends in Analytical Chemistry, 2019, 116, 292-299.	11.4	71
20	High-Speed Tandem Mass Spectrometric in Situ Imaging by Nanospray Desorption Electrospray Ionization Mass Spectrometry. Analytical Chemistry, 2013, 85, 9596-9603.	6.5	69
21	SPE-IMS-MS: An automated platform for sub-sixty second surveillance of endogenous metabolites and xenobiotics in biofluids. Clinical Mass Spectrometry, 2016, 2, 1-10.	1.9	63
22	Uterine FK506-binding protein 52 (FKBP52)–peroxiredoxin-6 (PRDX6) signaling protects pregnancy from overt oxidative stress. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 15577-15582.	7.1	62
23	Plasma lipidome reveals critical illness and recovery from human Ebola virus disease. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 3919-3928.	7.1	62
24	Imaging Mass Spectrometry Reveals Unique Protein Profiles during Embryo Implantation. Endocrinology, 2008, 149, 3274-3278.	2.8	61
25	Quantitative Mass Spectrometry Imaging of Prostaglandins as Silver Ion Adducts with Nanospray Desorption Electrospray Ionization. Analytical Chemistry, 2018, 90, 7246-7252.	6.5	61
26	Proteome insights into the symbiotic relationship between a captive colony of <i>Nasutitermes corniger</i> and its hindgut microbiome. ISME Journal, 2011, 5, 161-164.	9.8	57
27	Proteomic and Transcriptomic Analyses of " <i>Candidatus</i> Pelagibacter ubique―Describe the First P _{II} -Independent Response to Nitrogen Limitation in a Free-Living Alphaproteobacterium. MBio, 2013, 4, e00133-12.	4.1	54
28	Enhancing bottomâ€up and topâ€down proteomic measurements with ion mobility separations. Proteomics, 2015, 15, 2766-2776.	2.2	54
29	Advancing the High Throughput Identification of Liver Fibrosis Protein Signatures Using Multiplexed Ion Mobility Spectrometry. Molecular and Cellular Proteomics, 2014, 13, 1119-1127.	3.8	51
30	SNaPP: Simplified Nanoproteomics Platform for Reproducible Global Proteomic Analysis of Nanogram Protein Quantities. Endocrinology, 2016, 157, 1307-1314.	2.8	48
31	Three-dimensional imaging of lipids and metabolites in tissues by nanospray desorption electrospray ionization mass spectrometry. Analytical and Bioanalytical Chemistry, 2015, 407, 2063-2071.	3.7	47
32	Surfactant-assisted one-pot sample preparation for label-free single-cell proteomics. Communications Biology, 2021, 4, 265.	4.4	46
33	The Long Noncoding RNA Paupar Modulates PAX6 Regulatory Activities to Promote Alpha Cell Development and Function. Cell Metabolism, 2019, 30, 1091-1106.e8.	16.2	45
34	MPLEx: a method for simultaneous pathogen inactivation and extraction of samples for multi-omics profiling. Analyst, The, 2017, 142, 442-448.	3. 5	43
35	Proteome and computational analyses reveal new insights into the mechanisms of hepatitis C virus-mediated liver disease posttransplantation. Hepatology, 2012, 56, 28-38.	7. 3	39
36	<i>pmartR</i> : Quality Control and Statistics for Mass Spectrometry-Based Biological Data. Journal of Proteome Research, 2019, 18, 1418-1425.	3.7	39

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37	The fungal cultivar of leafâ€cutter ants produces specific enzymes in response to different plant substrates. Molecular Ecology, 2016, 25, 5795-5805.	3.9	37
38	Moisture modulates soil reservoirs of active DNA and RNA viruses. Communications Biology, 2021, 4, 992.	4.4	33
39	Uterine Deletion of Trp53 Compromises Antioxidant Responses in the Mouse Decidua. Endocrinology, 2012, 153, 4568-4579.	2.8	32
40	Comparing identified and statistically significant lipids and polar metabolites in 15â€year old serum and dried blood spot samples for longitudinal studies. Rapid Communications in Mass Spectrometry, 2017, 31, 447-456.	1.5	31
41	Production of ent-kaurene from lignocellulosic hydrolysate in Rhodosporidium toruloides. Microbial Cell Factories, 2020, 19, 24.	4.0	30
42	Simultaneous Proteomic Discovery and Targeted Monitoring using Liquid Chromatography, Ion Mobility Spectrometry, and Mass Spectrometry. Molecular and Cellular Proteomics, 2016, 15, 3694-3705.	3.8	29
43	Enrichment and Broad Representation of Plant Biomass-Degrading Enzymes in the Specialized Hyphal Swellings of Leucoagaricus gongylophorus, the Fungal Symbiont of Leaf-Cutter Ants. PLoS ONE, 2015, 10, e0134752.	2.5	28
44	Broad Substrate-Specific Phosphorylation Events Are Associated With the Initial Stage of Plant Cell Wall Recognition in Neurospora crassa. Frontiers in Microbiology, 2019, 10, 2317.	3. 5	25
45	Evaluating lipid mediator structural complexity using ion mobility spectrometry combined with mass spectrometry. Bioanalysis, 2018, 10, 279-289.	1.5	22
46	The fungus gardens of leafâ€cutter ants undergo a distinct physiological transition during biomass degradation. Environmental Microbiology Reports, 2014, 6, 389-395.	2.4	21
47	Multiplexed Activity-based Protein Profiling of the Human Pathogen Aspergillus fumigatus Reveals Large Functional Changes upon Exposure to Human Serum. Journal of Biological Chemistry, 2012, 287, 33447-33459.	3.4	20
48	Characterizing the lipid and metabolite changes associated with placental function and pregnancy complications using ion mobility spectrometry-mass spectrometry and mass spectrometry imaging. Placenta, 2017, 60, S67-S72.	1.5	20
49	Application of multiplexed ion mobility spectrometry towards the identification of host protein signatures of treatment effect in pulmonary tuberculosis. Tuberculosis, 2018, 112, 52-61.	1.9	20
50	Galectin-1 Markedly Reduces the Incidence of Resorptions in Mice Missing Immunophilin FKBP52. Endocrinology, 2012, 153, 2486-2493.	2.8	19
51	The MPLEx Protocol for Multi-omic Analyses of Soil Samples. Journal of Visualized Experiments, 2018, ,	0.3	19
52	STEPS: A grid search methodology for optimized peptide identification filtering of MS/MS database search results. Proteomics, 2013, 13, 766-770.	2.2	18
53	Development of an ecophysiological model for <i>Diplosphaera colotermitum</i> TAV2, a termite hindgut Verrucomicrobium. ISME Journal, 2013, 7, 1803-1813.	9.8	18
54	Muscle Segment Homeobox Genes Direct Embryonic Diapause by Limiting Inflammation in the Uterus*. Journal of Biological Chemistry, 2015, 290, 15337-15349.	3.4	18

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55	Trp53 deficient mice predisposed to preterm birth display region-specific lipid alterations at the embryo implantation site. Scientific Reports, 2016, 6, 33023.	3.3	17
56	Unveiling molecular signatures of preeclampsia and gestational diabetes mellitus with multi-omics and innovative cheminformatics visualization tools. Molecular Omics, 2020, 16, 521-532.	2.8	16
57	Unfolded Protein Response Inhibition Reduces Middle East Respiratory Syndrome Coronavirus-Induced Acute Lung Injury. MBio, 2021, 12, e0157221.	4.1	16
58	New Views of Old Proteins: Clarifying the Enigmatic Proteome. Molecular and Cellular Proteomics, 2022, 21, 100254.	3.8	16
59	Urinary Virome Perturbations in Kidney Transplantation. Frontiers in Medicine, 2018, 5, 72.	2.6	12
60	Hanging drop sample preparation improves sensitivity of spatial proteomics. Lab on A Chip, 2022, 22, 2869-2877.	6.0	12
61	From Plants to Ants: Fungal Modification of Leaf Lipids for Nutrition and Communication in the Leaf-Cutter Ant Fungal Garden Ecosystem. MSystems, 2021, 6, .	3.8	11
62	High-Throughput Large-Scale Targeted Proteomics Assays for Quantifying Pathway Proteins in Pseudomonas putida KT2440. Frontiers in Bioengineering and Biotechnology, 2020, 8, 603488.	4.1	10
63	From Prevention to Disease Perturbations: A Multi-Omic Assessment of Exercise and Myocardial Infarctions. Biomolecules, 2021, 11, 40.	4.0	8
64	Uncovering Hidden Members and Functions of the Soil Microbiome Using <i>De Novo</i> Metaproteomics. Journal of Proteome Research, 2022, 21, 2023-2035.	3.7	6
65	Preserved and variable spatialâ€chemical changes of lipids across tomato leaves in response to central vein wounding reveals potential origin of linolenic acid in signal transduction cascade. Plant-Environment Interactions, 2021, 2, 28-35.	1.5	4
66	Uniformly ¹⁵ N-Labeled Recombinant Ricin A-Chain as an Internal Retention Time Standard for Increased Confidence in Forensic Identification of Ricin by Untargeted Nanoflow Liquid Chromatography-Tandem Mass Spectrometry. Analytical Chemistry, 2019, 91, 13372-13376.	6.5	3
67	Proteomic Sample Preparation Techniques: Toward Forensic Proteomic Applications. ACS Symposium Series, 2019, , 29-46.	0.5	O