Ute Distler

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6238918/publications.pdf Version: 2024-02-01



HTE DISTIED

#	Article	IF	CITATIONS
1	Drift time-specific collision energies enable deep-coverage data-independent acquisition proteomics. Nature Methods, 2014, 11, 167-170.	19.0	411
2	Exosomes released by chronic lymphocytic leukemia cells induce the transition of stromal cells into cancer-associated fibroblasts. Blood, 2015, 126, 1106-1117.	1.4	399
3	A multicenter study benchmarks software tools for label-free proteome quantification. Nature Biotechnology, 2016, 34, 1130-1136.	17.5	321
4	Label-free quantification in ion mobility–enhanced data-independent acquisition proteomics. Nature Protocols, 2016, 11, 795-812.	12.0	258
5	Evaluation of FASP, SP3, and iST Protocols for Proteomic Sample Preparation in the Low Microgram Range. Journal of Proteome Research, 2017, 16, 4060-4072.	3.7	227
6	Quantitative profiling of the protein coronas that form around nanoparticles. Nature Protocols, 2014, 9, 2030-2044.	12.0	200
7	MaxDIA enables library-based and library-free data-independent acquisition proteomics. Nature Biotechnology, 2021, 39, 1563-1573.	17.5	115
8	Inâ€depth protein profiling of the postsynaptic density from mouse hippocampus using dataâ€independent acquisition proteomics. Proteomics, 2014, 14, 2607-2613.	2.2	103
9	IR-MALDI-MS Analysis of HPTLC-Separated Phospholipid Mixtures Directly from the TLC Plate. Analytical Chemistry, 2007, 79, 5793-5808.	6.5	88
10	Advances on the compositional analysis of glycosphingolipids combining thin-layer chromatography with mass spectrometry. Mass Spectrometry Reviews, 2010, 29, 425-479.	5.4	74
11	Shiga Toxin Receptor Gb3Cer/CD77: Tumor-Association and Promising Therapeutic Target in Pancreas and Colon Cancer. PLoS ONE, 2009, 4, e6813.	2.5	70
12	Quantum Chemical-Based Protocol for the Rational Design of Covalent Inhibitors. Journal of the American Chemical Society, 2016, 138, 8332-8335.	13.7	69
13	Matching IR-MALDI-o-TOF Mass Spectrometry with the TLC Overlay Binding Assay and Its Clinical Application for Tracing Tumor-Associated Glycosphingolipids in Hepatocellular and Pancreatic Cancer. Analytical Chemistry, 2008, 80, 1835-1846.	6.5	67
14	Inâ€depth evaluation of software tools for dataâ€independent acquisition based labelâ€free quantification. Proteomics, 2015, 15, 3140-3151.	2.2	66
15	Proteomic Analysis of Post-synaptic Density Fractions from Shank3 Mutant Mice Reveals Brain Region Specific Changes Relevant to Autism Spectrum Disorder. Frontiers in Molecular Neuroscience, 2017, 10, 26.	2.9	66
16	Visualizing transfer of microbial biomolecules by outer membrane vesicles in microbeâ€hostâ€communication in vivo. Journal of Extracellular Vesicles, 2021, 10, e12159.	12.2	66
17	Direct Coupling of High-Performance Thin-Layer Chromatography with UV Spectroscopy and IR-MALDI Orthogonal TOF MS for the Analysis of Cyanobacterial Toxins. Analytical Chemistry, 2009, 81, 3858-3866.	6.5	47
18	Molecular cause and functional impact of altered synaptic lipid signaling due to a <i>prgâ€1</i> gene <scp>SNP</scp> . EMBO Molecular Medicine, 2016, 8, 25-38.	6.9	40

UTE DISTLER

#	Article	IF	CITATIONS
19	Mast Cell–deficient <i>KitW-sh</i> "Sash―Mutant Mice Display Aberrant Myelopoiesis Leading to the Accumulation of Splenocytes That Act as Myeloid-Derived Suppressor Cells. Journal of Immunology, 2013, 190, 5534-5544.	0.8	36
20	Synaptic phospholipids as a new target for cortical hyperexcitability and E/I balance in psychiatric disorders. Molecular Psychiatry, 2018, 23, 1699-1710.	7.9	33
21	Differences in CD75s- and iso-CD75s-ganglioside content and altered mRNA expression of sialyltransferases ST6CAL1 and ST3CAL6 in human hepatocellular carcinomas and nontumoral liver tissues. Glycobiology, 2011, 21, 584-594.	2.5	30
22	Biomedical applications of ion mobility-enhanced data-independent acquisition-based label-free quantitative proteomics. Expert Review of Proteomics, 2014, 11, 675-684.	3.0	29
23	Tumor-associated CD75s- and iso-CD75s-gangliosides are potential targets for adjuvant therapy in pancreatic cancer. Molecular Cancer Therapeutics, 2008, 7, 2464-2475.	4.1	28
24	CMTM6 expressed on the adaxonal Schwann cell surface restricts axonal diameters in peripheral nerves. Nature Communications, 2020, 11, 4514.	12.8	27
25	Rapid Antigen Processing and Presentation of a Protective and Immunodominant HLA-B*27-restricted Hepatitis C Virus-specific CD8+ T-cell Epitope. PLoS Pathogens, 2012, 8, e1003042.	4.7	25
26	Asymmetric Disulfanylbenzamides as Irreversible and Selective Inhibitors of <i>Staphylococcus aureus</i> Sortase A. ChemMedChem, 2020, 15, 839-850.	3.2	24
27	Purification and Properties of Yeast Proteases Secreted by Wickerhamomyces anomalus 227 and Metschnikovia pulcherrima 446 during Growth in a White Grape Juice. Fermentation, 2017, 3, 2.	3.0	23
28	Enhancing Sensitivity of Microflow-Based Bottom-Up Proteomics through Postcolumn Solvent Addition. Analytical Chemistry, 2019, 91, 7510-7515.	6.5	22
29	Application of thinâ€layer chromatography/infrared matrixâ€assisted laser desorption/ionization orthogonal timeâ€ofâ€flight mass spectrometry to structural analysis of bacteriaâ€binding glycosphingolipids selected by affinity detection. Rapid Communications in Mass Spectrometry, 2010, 24, 1032-1038.	1.5	21
30	Fungicide resistance towards fludioxonil conferred by overexpression of the phosphatase gene Mo PTP 2 in Magnaporthe oryzae. Molecular Microbiology, 2018, 111, 662-677.	2.5	21
31	Proteomic Analysis of Brain Region and Sex-Specific Synaptic Protein Expression in the Adult Mouse Brain. Cells, 2020, 9, 313.	4.1	20
32	Fluorovinylsulfones and -Sulfonates as Potent Covalent Reversible Inhibitors of the Trypanosomal Cysteine Protease Rhodesain: Structure–Activity Relationship, Inhibition Mechanism, Metabolism, and In Vivo Studies. Journal of Medicinal Chemistry, 2021, 64, 12322-12358.	6.4	20
33	Naphthoquinones as Covalent Reversible Inhibitors of Cysteine Proteases—Studies on Inhibition Mechanism and Kinetics. Molecules, 2020, 25, 2064.	3.8	20
34	Proteogenomics analysis unveils a TFG-RET gene fusion and druggable targets in papillary thyroid carcinomas. Nature Communications, 2020, 11, 2056.	12.8	19
35	New Cysteine Protease Inhibitors: Electrophilic (Het)arenes and Unexpected Prodrug Identification for the Trypanosoma Protease Rhodesain. Molecules, 2020, 25, 1451.	3.8	16
36	OpenTIMS, TimsPy, and TimsR: Open and Easy Access to timsTOF Raw Data. Journal of Proteome Research, 2021, 20, 2122-2129.	3.7	15

UTE DISTLER

#	Article	IF	CITATIONS
37	Transmembrane BAX Inhibitor-1 Motif Containing Protein 5 (TMBIM5) Sustains Mitochondrial Structure, Shape, and Function by Impacting the Mitochondrial Protein Synthesis Machinery. Cells, 2020, 9, 2147.	4.1	14
38	Hybrid QconCAT-Based Targeted Absolute and Data-Independent Acquisition-Based Label-Free Quantification Enables In-Depth Proteomic Characterization of Wheat Amylase/Trypsin Inhibitor Extracts. Journal of Proteome Research, 2021, 20, 1544-1557.	3.7	13
39	NF-κB inducing kinase (NIK) is an essential post-transcriptional regulator of T-cell activation affecting F-actin dynamics and TCR signaling. Journal of Autoimmunity, 2018, 94, 110-121.	6.5	12
40	Tools for Pathogen Proteomics: Fishing with Biomimetic Nanosponges. ACS Nano, 2017, 11, 11768-11772.	14.6	10
41	Proteomic profiling of German Dornfelder grape berries using data-independent acquisition. Plant Physiology and Biochemistry, 2017, 118, 64-70.	5.8	9
42	Structural and mechanistic insights into the interaction of the circadian transcription factor BMAL1 with the KIX domain of the CREB-binding protein. Journal of Biological Chemistry, 2019, 294, 16604-16619.	3.4	9
43	Plasmodium falciparum S-Adenosylmethionine Synthetase Is Essential for Parasite Survival through a Complex Interaction Network with Cytoplasmic and Nuclear Proteins. Microorganisms, 2022, 10, 1419.	3.6	9
44	REGGAE: a novel approach for the identification of key transcriptional regulators. Bioinformatics, 2018, 34, 3503-3510.	4.1	8
45	Chronic intestinal inflammation in mice expressing viral Flip in epithelial cells. Mucosal Immunology, 2018, 11, 1621-1629.	6.0	8
46	The role of TCF3 as potential master regulator in blastemal Wilms tumors. International Journal of Cancer, 2019, 144, 1432-1443.	5.1	4
47	Label-Free Proteomics of Quantity-Limited Samples Using Ion Mobility-Assisted Data-Independent Acquisition Mass Spectrometry. Methods in Molecular Biology, 2021, 2228, 327-339.	0.9	4
48	Adaptive Mechanisms of Somatostatin-Positive Interneurons after Traumatic Brain Injury through a Switch of α Subunits in L-Type Voltage-Gated Calcium Channels. Cerebral Cortex, 2022, 32, 1093-1109.	2.9	4
49	The caspase-2 substrate p54nrb exhibits a multifaceted role in tumor cell death susceptibility via gene regulatory functions. Cell Death and Disease, 2022, 13, 386.	6.3	4
50	Quantitative proteomics analysis reveals core and variable tick salivary proteins at the tickâ€vertebrate host interface. Molecular Ecology, 2022, 31, 4162-4175.	3.9	4
51	Friend virus limits adaptive cellular immune responses by imprinting a maturation-resistant and T helper type 2-biased immunophenotype in dendritic cells. PLoS ONE, 2018, 13, e0192541.	2.5	3
52	Astrocytic ATX fuels synaptic phospholipid signaling involved in psychiatric disorders. Molecular Psychiatry, 2018, 23, 1685-1686.	7.9	1
53	GABAA Receptor-Stabilizing Protein Ubqln1 Affects Hyperexcitability and Epileptogenesis after Traumatic Brain Injury and in a Model of In Vitro Epilepsy in Mice. International Journal of Molecular Sciences, 2022, 23, 3902.	4.1	1
54	Gamma Irradiation Triggers Immune Escape in Glioma-Propagating Cells. Cancers, 2022, 14, 2728.	3.7	1