Valerie C Wasinger

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/161608/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Activation of the eIF2α/ATF4 axis drives triple-negative breast cancer radioresistance by promoting glutathione biosynthesis. Redox Biology, 2021, 43, 101993.	3.9	30
2	Current Trends in IBD—Development of Mucosal-Based Biomarkers and a Novel Minimally Invasive Recoverable Sampling System. Inflammatory Bowel Diseases, 2021, 27, S17-S24.	0.9	6
3	Spp24 is associated with endocytic signalling, lipid metabolism, and discrimination of tissue integrity for â€ ⁻ leaky-gut' in inflammatory bowel disease. Scientific Reports, 2020, 10, 12932.	1.6	13
4	The Molecular Floodgates of Stress-Induced Senescence Reveal Translation, Signalling and Protein Activity Central to the Post-Mortem Proteome. International Journal of Molecular Sciences, 2020, 21, 6422.	1.8	3
5	CHTOP in Chemoresistant Epithelial Ovarian Cancer: A Novel and Potential Therapeutic Target. Frontiers in Oncology, 2019, 9, 557.	1.3	11
6	Editorial: metabolomic biomarkers for colorectal adenocarcinoma and in the differentiation between irritable bowel syndrome and ulcerative colitis in clinical remission—confounded by the gut microbiome?. Alimentary Pharmacology and Therapeutics, 2019, 49, 1086-1087.	1.9	2
7	Analysis of the Preserved Amino Acid Bias in Peptide Profiles of Iron Age Teeth from a Tropical Environment Enable Sexing of Individuals Using Amelogenin MRM. Proteomics, 2019, 19, e1800341.	1.3	24
8	Molecular Pathophysiology of Epithelial Barrier Dysfunction in Inflammatory Bowel Diseases. Proteomes, 2018, 6, 17.	1.7	77
9	Identification of protein biomarkers and signaling pathways associated with prostate cancer radioresistance using label-free LC-MS/MS proteomic approach. Scientific Reports, 2017, 7, 41834.	1.6	59
10	Serological Epithelial Component Proteins Identify Intestinal Complications in Crohn's Disease. Molecular and Cellular Proteomics, 2017, 16, 1244-1257.	2.5	13
11	Amyotrophic lateral sclerosis-like superoxide dismutase 1 proteinopathy is associated with neuronal loss in Parkinson's disease brain. Acta Neuropathologica, 2017, 134, 113-127.	3.9	78
12	Impaired Intestinal Permeability Contributes to Ongoing Bowel Symptoms in Patients With Inflammatory Bowel Disease and Mucosal Healing. Gastroenterology, 2017, 153, 723-731.e1.	0.6	193
13	Urinary biomarkers in prostate cancer detection and monitoring progression. Critical Reviews in Oncology/Hematology, 2017, 118, 15-26.	2.0	64
14	Proteomics in Inflammatory Bowel Disease: Approach Using Animal Models. Digestive Diseases and Sciences, 2017, 62, 2266-2276.	1.1	6
15	Current application of proteomics in biomarker discovery for inflammatory bowel disease. World Journal of Gastrointestinal Pathophysiology, 2016, 7, 27.	0.5	36
16	Proteomics discovery of chemoresistant biomarkers for ovarian cancer therapy. Expert Review of Proteomics, 2016, 13, 905-915.	1.3	8
17	Impact of Perturbed Pancreatic β-Cell Cholesterol Homeostasis on Adipose Tissue and Skeletal Muscle Metabolism. Diabetes, 2016, 65, 3610-3620.	0.3	28
18	Low Mass Blood Peptides Discriminative of Inflammatory Bowel Disease (IBD) Severity: A Quantitative Proteomic Perspective, Molecular and Cellular Proteomics, 2016, 15, 256-265	2.5	30

VALERIE C WASINGER

#	Article	IF	CITATIONS
19	Absolute quantification of human tear lactoferrin using multiple reaction monitoring technique with stable-isotopic labeling. Analytical Biochemistry, 2016, 496, 30-34.	1.1	9
20	Proteomic identification of the lactate dehydrogenase A in a radioresistant prostate cancer xenograft mouse model for improving radiotherapy. Oncotarget, 2016, 7, 74269-74285.	0.8	24
21	Reverse-Polynomial Dilution Calibration Methodology Extends Lower Limit of Quantification and Reduces Relative Residual Error in Targeted Peptide Measurements in Blood Plasma. Molecular and Cellular Proteomics, 2015, 14, 441-454.	2.5	3
22	Proteomic Analysis of Urine to Identify Breast Cancer Biomarker Candidates Using a Label-Free LC-MS/MS Approach. PLoS ONE, 2015, 10, e0141876.	1.1	87
23	Abstract 2001: Identification of lactate dehydrogenase A (LDHA) as a potential therapeutic target for prostate cancer radiotherapy. , 2015, , .		0
24	Proteomics for Breast Cancer Urine Biomarkers. Advances in Clinical Chemistry, 2014, 63, 123-167.	1.8	30
25	Profilin-1 Overexpression in MDA-MB-231 Breast Cancer Cells Is Associated with Alterations in Proteomics Biomarkers of Cell Proliferation, Survival, and Motility as Revealed by Global Proteomics Analyses. OMICS A Journal of Integrative Biology, 2014, 18, 778-791.	1.0	29
26	Green fluorescent protein expression triggers proteome changes in breast cancer cells. Experimental Cell Research, 2014, 320, 33-45.	1.2	26
27	Identification of plasma Complement C3 as a potential biomarker for neuroblastoma using a quantitative proteomic approach. Journal of Proteomics, 2014, 96, 1-12.	1.2	19
28	A Standardized and Reproducible Urine Preparation Protocol for Cancer Biomarkers Discovery. Biomarkers in Cancer, 2014, 6, BIC.S17991.	3.6	15
29	In vivolocalization of antibodies raised againstEimeria maximawall forming bodies during sexual intracellular development. Parasitology, 2014, 141, 1726-1735.	0.7	6
30	Proteomics and metabolomics in inflammatory bowel disease. Journal of Gastroenterology and Hepatology (Australia), 2013, 28, 1076-1086.	1.4	32
31	Proteome profiles of vaginal fluids from women affected by bacterial vaginosis and healthy controls: outcomes of rifaximin treatment. Journal of Antimicrobial Chemotherapy, 2013, 68, 2648-2659.	1.3	19
32	Evaluation of chemical cleaning of UF membranes fouled with whey protein isolates via analysis of residual protein components on membranes surface. Separation and Purification Technology, 2013, 103, 241-250.	3.9	39
33	Tear Fluid Protein Biomarkers. Advances in Clinical Chemistry, 2013, 62, 151-196.	1.8	41
34	Current Status and Advances in Quantitative Proteomic Mass Spectrometry. International Journal of Proteomics, 2013, 2013, 1-12.	2.0	128
35	Advances in Quantitative Mass Spectrometry. International Journal of Proteomics, 2013, 2013, 1-2.	2.0	0
36	Glycosylation in a Mammalian Expression System Is Critical for the Production of Functionally Active Leukocyte Immunoglobulin-like Receptor A3 Protein. Journal of Biological Chemistry, 2013, 288, 32873-32885.	1.6	14

VALERIE C WASINGER

#	Article	IF	CITATIONS
37	Preliminary identification of differentially expressed tear proteins in keratoconus. Molecular Vision, 2013, 19, 2124-34.	1.1	39
38	Tryptophan Metabolome Signature Differences in Inflammatory Bowel Diseases. Gastroenterology, 2011, 140, S-840.	0.6	0
39	Proteomic Identification Of Factors That Regulate Mast Cell Localization To The Airway Smooth Muscle Cells In Asthma. , 2011, , .		Ο
40	Protein and peptide fractionation, enrichment and depletion: Tools for the complex proteome. Proteomics, 2011, 11, 513-534.	1.3	88
41	Postâ€ŧranslation modification of proteins in tears. Electrophoresis, 2010, 31, 1853-1861.	1.3	49
42	Mass and charge selective protein fractionation for the differential analysis of T-cell and CD34+ stem cell proteins from cord blood. Journal of Proteomics, 2010, 73, 571-578.	1.2	3
43	Tear lipocalin is the predominant phosphoprotein in human tear fluid. Experimental Eye Research, 2010, 90, 344-349.	1.2	18
44	Peptide enrichment and protein fractionation using selective electrophoresis. Proteomics, 2008, 8, 4197-4208.	1.3	22
45	Prefractionation, Enrichment, Desalting and Depleting of Low Volume and Low Abundance Proteins and Peptides Using the MF10. Methods in Molecular Biology, 2008, 424, 257-275.	0.4	7
46	Genetic and proteomic characterization of rifaximin resistance in Bifidobacterium infantis BI07. Research in Microbiology, 2007, 158, 355-362.	1.0	17
47	Identification of Vascular Surface Proteins by in Vivo Biotinylation:  A Method Sufficiently Sensitive To Detect Changes in Rat Liver 2 Weeks after Partial Hepatectomy. Journal of Proteome Research, 2007, 6, 3108-3113.	1.8	3
48	Overview of the HUPO Plasma Proteome Project: Results from the pilot phase with 35 collaborating laboratories and multiple analytical groups, generating a core dataset of 3020 proteins and a publicly-available database. , 2006, , 1-35.		4
49	A functional annotation of subproteomes in human plasma. , 2006, , 329-351.		Ο
50	Holistic Biology of Microorganisms: Genomics, Transcriptomics, and Proteomics. Methods of Biochemical Analysis, 2005, , 1-14.	0.2	1
51	Proteomics: An Overview. Inflammatory Bowel Diseases, 2005, 11, 927-936.	0.9	11
52	A proteomic view ofBifidobacterium infantis generated by multi-dimensional chromatography coupled with tandem mass spectrometry. Proteomics, 2005, 5, 1859-1867.	1.3	37
53	Two-dimensional liquid chromatography/tandem mass spectrometry analysis of Gradiflowâ,,¢ fractionated native human plasma. Proteomics, 2005, 5, 3397-3401.	1.3	29
54	A functional annotation of subproteomes in human plasma. Proteomics, 2005, 5, 3506-3519.	1.3	82

#	Article	IF	CITATIONS
55	Overview of the HUPO Plasma Proteome Project: Results from the pilot phase with 35 collaborating laboratories and multiple analytical groups, generating a core dataset of 3020 proteins and a publicly-available database. Proteomics, 2005, 5, 3226-3245.	1.3	766
56	Characterization of the Role of the Rab GTPase-activating Protein AS160 in Insulin-regulated GLUT4 Trafficking. Journal of Biological Chemistry, 2005, 280, 37803-37813.	1.6	330
57	Changes in Gene Expression Associated with Stable Drug and Radiation Resistance in Small Cell Lung Cancer Cells are Similar to those Caused by a Single X-Ray Dose. Radiation Research, 2004, 161, 495-503.	0.7	11
58	Akt Mediates Insulin-stimulated Phosphorylation of Ndrg2. Journal of Biological Chemistry, 2004, 279, 18623-18632.	1.6	76
59	Studies on NADH oxidase and alkyl hydroperoxide reductase produced by Porphyromonas gingivalis. Oral Microbiology and Immunology, 2004, 19, 137-143.	2.8	27
60	Cross-species identification of proteins from proteome profiles of the marine oligotrophic ultramicrobacterium, Sphingopyxis alaskensis. Proteomics, 2004, 4, 1779-1788.	1.3	23
61	Depletion of the highly abundant protein albumin from human plasma using the Gradiflow. Proteomics, 2003, 3, 279-287.	1.3	100
62	A Cortactin-CD2-associated Protein (CD2AP) Complex Provides a Novel Link between Epidermal Growth Factor Receptor Endocytosis and the Actin Cytoskeleton. Journal of Biological Chemistry, 2003, 278, 21805-21813.	1.6	192
63	Proteomic tools for biomedicine. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2002, 771, 33-48.	1.2	33
64	The dynamic range of protein expression: A challenge for proteomic research. Electrophoresis, 2000, 21, 1104-1115.	1.3	603
65	The proteome of Mycoplasma genitalium. FEBS Journal, 2000, 267, 1571-1582.	0.2	84
66	Cross-species characterisation of abundantly expressedOchrobactrum anthropi gene products. Electrophoresis, 1999, 20, 2196-2203.	1.3	15
67	Small genes/gene-products inEscherichia coliK-12. FEMS Microbiology Letters, 1998, 169, 375-382.	0.7	44
68	Low molecular weight proteins: A challenge for post-genomic research. Electrophoresis, 1998, 19, 536-544.	1.3	58
69	Proteomic â€~contigs' ofOchrobactrum anthropi, application of extensive pH gradients. Electrophoresis, 1997, 18, 1373-1383.	1.3	59
70	Conserved Motifs as the Basis for Recognition of Homologous Proteins Across Species Boundaries Using Peptide-mass Fingerprinting. , 1997, 32, 370-378.		26
71	Progress with gene-product mapping of the Mollicutes:Mycoplasma genitalium. Electrophoresis, 1995, 16, 1090-1094.	1.3	892