Valerie C Wasinger

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Progress with gene-product mapping of the Mollicutes:Mycoplasma genitalium. Electrophoresis, 1995, 16, 1090-1094.	1.3	892
2	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
3	The dynamic range of protein expression: A challenge for proteomic research. Electrophoresis, 2000, 21, 1104-1115.	1.3	603
4	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
5	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
6	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
7	Current Status and Advances in Quantitative Proteomic Mass Spectrometry. International Journal of Proteomics, 2013, 2013, 1-12.	2.0	128
8	Depletion of the highly abundant protein albumin from human plasma using the Gradiflow. Proteomics, 2003, 3, 279-287.	1.3	100
9	Protein and peptide fractionation, enrichment and depletion: Tools for the complex proteome. Proteomics, 2011, 11, 513-534.	1.3	88
10	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
11	The proteome of Mycoplasma genitalium. FEBS Journal, 2000, 267, 1571-1582.	0.2	84
12	A functional annotation of subproteomes in human plasma. Proteomics, 2005, 5, 3506-3519.	1.3	82
13	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
14	Molecular Pathophysiology of Epithelial Barrier Dysfunction in Inflammatory Bowel Diseases. Proteomes, 2018, 6, 17.	1.7	77
15	Akt Mediates Insulin-stimulated Phosphorylation of Ndrg2. Journal of Biological Chemistry, 2004, 279, 18623-18632.	1.6	76
16	Urinary biomarkers in prostate cancer detection and monitoring progression. Critical Reviews in Oncology/Hematology, 2017, 118, 15-26.	2.0	64
17	Proteomic â€~contigs' ofOchrobactrum anthropi, application of extensive pH gradients. Electrophoresis, 1997, 18, 1373-1383.	1.3	59
18	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

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19	Low molecular weight proteins: A challenge for post-genomic research. Electrophoresis, 1998, 19, 536-544.	1.3	58
20	Postâ€ŧranslation modification of proteins in tears. Electrophoresis, 2010, 31, 1853-1861.	1.3	49
21	Small genes/gene-products inEscherichia coliK-12. FEMS Microbiology Letters, 1998, 169, 375-382.	0.7	44
22	Tear Fluid Protein Biomarkers. Advances in Clinical Chemistry, 2013, 62, 151-196.	1.8	41
23	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
24	Preliminary identification of differentially expressed tear proteins in keratoconus. Molecular Vision, 2013, 19, 2124-34.	1.1	39
25	A proteomic view ofBifidobacterium infantis generated by multi-dimensional chromatography coupled with tandem mass spectrometry. Proteomics, 2005, 5, 1859-1867.	1.3	37
26	Current application of proteomics in biomarker discovery for inflammatory bowel disease. World Journal of Gastrointestinal Pathophysiology, 2016, 7, 27.	0.5	36
27	Proteomic tools for biomedicine. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2002, 771, 33-48.	1.2	33
28	Proteomics and metabolomics in inflammatory bowel disease. Journal of Gastroenterology and Hepatology (Australia), 2013, 28, 1076-1086.	1.4	32
29	Proteomics for Breast Cancer Urine Biomarkers. Advances in Clinical Chemistry, 2014, 63, 123-167.	1.8	30
30	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
31	Activation of the elF2α/ATF4 axis drives triple-negative breast cancer radioresistance by promoting glutathione biosynthesis. Redox Biology, 2021, 43, 101993.	3.9	30
32	Two-dimensional liquid chromatography/tandem mass spectrometry analysis of Gradiflowâ,,¢ fractionated native human plasma. Proteomics, 2005, 5, 3397-3401.	1.3	29
33	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
34	Impact of Perturbed Pancreatic β-Cell Cholesterol Homeostasis on Adipose Tissue and Skeletal Muscle Metabolism. Diabetes, 2016, 65, 3610-3620.	0.3	28
35	Studies on NADH oxidase and alkyl hydroperoxide reductase produced by Porphyromonas gingivalis. Oral Microbiology and Immunology, 2004, 19, 137-143.	2.8	27
36	Conserved Motifs as the Basis for Recognition of Homologous Proteins Across Species Boundaries Using Peptide-mass Fingerprinting. , 1997, 32, 370-378.		26

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37	Green fluorescent protein expression triggers proteome changes in breast cancer cells. Experimental Cell Research, 2014, 320, 33-45.	1.2	26
38	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
39	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
40	Cross-species identification of proteins from proteome profiles of the marine oligotrophic ultramicrobacterium, Sphingopyxis alaskensis. Proteomics, 2004, 4, 1779-1788.	1.3	23
41	Peptide enrichment and protein fractionation using selective electrophoresis. Proteomics, 2008, 8, 4197-4208.	1.3	22
42	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
43	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
44	Tear lipocalin is the predominant phosphoprotein in human tear fluid. Experimental Eye Research, 2010, 90, 344-349.	1.2	18
45	Genetic and proteomic characterization of rifaximin resistance in Bifidobacterium infantis BI07. Research in Microbiology, 2007, 158, 355-362.	1.0	17
46	Cross-species characterisation of abundantly expressedOchrobactrum anthropi gene products. Electrophoresis, 1999, 20, 2196-2203.	1.3	15
47	A Standardized and Reproducible Urine Preparation Protocol for Cancer Biomarkers Discovery. Biomarkers in Cancer, 2014, 6, BIC.S17991.	3.6	15
48	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
49	Serological Epithelial Component Proteins Identify Intestinal Complications in Crohn's Disease. Molecular and Cellular Proteomics, 2017, 16, 1244-1257.	2.5	13
50	Spp24 is associated with endocytic signalling, lipid metabolism, and discrimination of tissue integrity for †leaky-gut' in inflammatory bowel disease. Scientific Reports, 2020, 10, 12932.	1.6	13
51	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
52	Proteomics: An Overview. Inflammatory Bowel Diseases, 2005, 11, 927-936.	0.9	11
53	CHTOP in Chemoresistant Epithelial Ovarian Cancer: A Novel and Potential Therapeutic Target. Frontiers in Oncology, 2019, 9, 557.	1.3	11
54	Absolute quantification of human tear lactoferrin using multiple reaction monitoring technique with stable-isotopic labeling. Analytical Biochemistry, 2016, 496, 30-34.	1.1	9

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#	Article	IF	CITATIONS
55	Proteomics discovery of chemoresistant biomarkers for ovarian cancer therapy. Expert Review of Proteomics, 2016, 13, 905-915.	1.3	8
56	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
57	In vivolocalization of antibodies raised againstEimeria maximawall forming bodies during sexual intracellular development. Parasitology, 2014, 141, 1726-1735.	0.7	6
58	Proteomics in Inflammatory Bowel Disease: Approach Using Animal Models. Digestive Diseases and Sciences, 2017, 62, 2266-2276.	1.1	6
59	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
60	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
61	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
62	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
63	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
64	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
65	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
66	Holistic Biology of Microorganisms: Genomics, Transcriptomics, and Proteomics. Methods of Biochemical Analysis, 2005, , 1-14.	0.2	1
67	A functional annotation of subproteomes in human plasma. , 2006, , 329-351.		0
68	Tryptophan Metabolome Signature Differences in Inflammatory Bowel Diseases. Gastroenterology, 2011, 140, S-840.	0.6	0
69	Proteomic Identification Of Factors That Regulate Mast Cell Localization To The Airway Smooth Muscle Cells In Asthma. , 2011, , .		0
70	Advances in Quantitative Mass Spectrometry. International Journal of Proteomics, 2013, 2013, 1-2.	2.0	0
71	Abstract 2001: Identification of lactate dehydrogenase A (LDHA) as a potential therapeutic target for prostate cancer radiotherapy. , 2015, , .		0