

Bernd Roschitzki

List of Publications by Year in descending order

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Version: 2024-02-01

56
papers

4,610
citations

172386

29
h-index

155592

55
g-index

58
all docs

58
docs citations

58
times ranked

8299
citing authors

#	ARTICLE	IF	CITATIONS
1	Reduced thrombogenicity of surface-treated Nitinol implants steered by altered protein adsorption. <i>Acta Biomaterialia</i> , 2022, 137, 331-345.	4.1	13
2	Grape ASR-Silencing Sways Nuclear Proteome, Histone Marks and Interplay of Intrinsically Disordered Proteins. <i>International Journal of Molecular Sciences</i> , 2022, 23, 1537.	1.8	2
3	BAF complexes drive proliferation and block myogenic differentiation in fusion-positive rhabdomyosarcoma. <i>Nature Communications</i> , 2021, 12, 6924.	5.8	25
4	In search of cerebrospinal fluid biomarkers of fatigue in multiple sclerosis: A proteomics study. <i>Journal of Sleep Research</i> , 2019, 28, e12721.	1.7	4
5	An Integrated Systems Approach Unveils New Aspects of Microoxia-Mediated Regulation in <i>Bradyrhizobium diazoefficiens</i> . <i>Frontiers in Microbiology</i> , 2019, 10, 924.	1.5	31
6	Engineered peptide barcodes for in-depth analyses of binding protein libraries. <i>Nature Methods</i> , 2019, 16, 421-428.	9.0	37
7	Brain Citrullination Patterns and T Cell Reactivity of Cerebrospinal Fluid-Derived CD4+ T Cells in Multiple Sclerosis. <i>Frontiers in Immunology</i> , 2019, 10, 540.	2.2	31
8	Diurnal changes in concerted plant protein phosphorylation and acetylation in <i>Arabidopsis</i> organs and seedlings. <i>Plant Journal</i> , 2019, 99, 176-194.	2.8	59
9	Isoform-specific localization of DNMT3A regulates DNA methylation fidelity at bivalent CpG islands. <i>EMBO Journal</i> , 2017, 36, 3421-3434.	3.5	99
10	<i>Mycobacterium smegmatis</i> PafBC is involved in regulation of DNA damage response. <i>Scientific Reports</i> , 2017, 7, 13987.	1.6	34
11	Differential regulation of germ line apoptosis and germ cell differentiation by CPEB family members in <i>C. elegans</i> . <i>PLoS ONE</i> , 2017, 12, e0182270.	1.1	5
12	Identification of Fungal T Cell Epitopes by Mass Spectrometry-Based Proteomics. <i>Methods in Molecular Biology</i> , 2017, 1625, 65-74.	0.4	0
13	High-throughput proteomics reveal alarmins as amplifiers of tissue pathology and inflammation after spinal cord injury. <i>Scientific Reports</i> , 2016, 6, 21607.	1.6	96
14	Natural Genetic Variation Differentially Affects the Proteome and Transcriptome in <i>Caenorhabditis elegans</i> . <i>Molecular and Cellular Proteomics</i> , 2016, 15, 1670-1680.	2.5	23
15	Natural Genetic Variation Influences Protein Abundances in <i>C. elegans</i> Developmental Signalling Pathways. <i>PLoS ONE</i> , 2016, 11, e0149418.	1.1	28
16	Independent Candidate Serum Protein Biomarkers of Response to Adalimumab and to Infliximab in Rheumatoid Arthritis: An Exploratory Study. <i>PLoS ONE</i> , 2016, 11, e0153140.	1.1	19
17	Proteomic Analysis of the Rat Canalicular Membrane Reveals Expression of a Complex System of P4-ATPases in Liver. <i>PLoS ONE</i> , 2016, 11, e0158033.	1.1	7
18	Inconsistencies in the red blood cell membrane proteome analysis: generation of a database for research and diagnostic applications. <i>Database: the Journal of Biological Databases and Curation</i> , 2015, 2015, bav056-bav056.	1.4	25

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19	Identification and characterization of chloroplast casein kinase II from <i>Oryza sativa</i> (rice). <i>Journal of Experimental Botany</i> , 2015, 66, 175-187.	2.4	18
20	Characterization of recombinant human and bovine thyroid-stimulating hormone preparations by mass spectrometry and determination of their endotoxin content. <i>BMC Veterinary Research</i> , 2013, 9, 141.	0.7	2
21	Infusion of anti-Nogo-A antibodies in adult rats increases growth and synapse related proteins in the absence of behavioral alterations. <i>Experimental Neurology</i> , 2013, 250, 52-68.	2.0	10
22	Directed shotgun proteomics guided by saturated RNA-seq identifies a complete expressed prokaryotic proteome. <i>Genome Research</i> , 2013, 23, 1916-1927.	2.4	91
23	Altered Activation of Endothelial Anti- and Proapoptotic Pathways by High-Density Lipoprotein from Patients with Coronary Artery Disease. <i>Circulation</i> , 2013, 127, 891-904.	1.6	303
24	iTRAQ-Based and Label-Free Proteomics Approaches for Studies of Human Adenovirus Infections. <i>International Journal of Proteomics</i> , 2013, 2013, 1-16.	2.0	73
25	Secretome Analysis Defines the Major Role of SecDF in <i>Staphylococcus aureus</i> Virulence. <i>PLoS ONE</i> , 2013, 8, e63513.	1.1	31
26	A Novel Th Cell Epitope of <i>Candida albicans</i> Mediates Protection from Fungal Infection. <i>Journal of Immunology</i> , 2012, 188, 5636-5643.	0.4	95
27	Who is who in litter decomposition? Metaproteomics reveals major microbial players and their biogeochemical functions. <i>ISME Journal</i> , 2012, 6, 1749-1762.	4.4	537
28	Insights into the Gene Expression Profile of Uncultivable Hemotrophic <i>Mycoplasma suis</i> during Acute Infection, Obtained Using Proteome Analysis. <i>Journal of Bacteriology</i> , 2012, 194, 1505-1514.	1.0	12
29	Discovery of serum proteomic biomarkers for prediction of response to infliximab (a monoclonal) Tj ETQq1 1 0.784314 rgBT /Overlock 2012, 77, 372-382.	1.2	45
30	Identification of Proteins Associated with the <i>Pseudomonas aeruginosa</i> Biofilm Extracellular Matrix. <i>Journal of Proteome Research</i> , 2012, 11, 4906-4915.	1.8	198
31	Soil metaproteomics – Comparative evaluation of protein extraction protocols. <i>Soil Biology and Biochemistry</i> , 2012, 54, 14-24.	4.2	178
32	Characterization of the phosphoproteome of mature <i>Arabidopsis</i> pollen. <i>Plant Journal</i> , 2012, 72, 89-101.	2.8	73
33	Antibody Phage Display Assisted Identification of Junction Plakoglobin as a Potential Biomarker for Atherosclerosis. <i>PLoS ONE</i> , 2012, 7, e47985.	1.1	13
34	Extracellular Hemoglobin Polarizes the Macrophage Proteome toward Hb-Clearance, Enhanced Antioxidant Capacity and Suppressed HLA Class 2 Expression. <i>Journal of Proteome Research</i> , 2011, 10, 2397-2408.	1.8	36
35	Identification of Distinct Amino Acids as ADP-Ribose Acceptor Sites by Mass Spectrometry. <i>Methods in Molecular Biology</i> , 2011, 780, 57-66.	0.4	17
36	Plastid Proteome Assembly without Toc159: Photosynthetic Protein Import and Accumulation of N-Acetylated Plastid Precursor Proteins. <i>Plant Cell</i> , 2011, 23, 3911-3928.	3.1	77

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37	Recognition of Host Proteins by Helicobacter Cysteine-Rich Protein C. <i>Current Microbiology</i> , 2011, 63, 239-249.	1.0	9
38	Implementation and evaluation of relative and absolute quantification in shotgun proteomics with label-free methods. <i>Journal of Proteomics</i> , 2010, 73, 1740-1746.	1.2	139
39	A proteomics approach to study synergistic and antagonistic interactions of the fungal-bacterial consortium <i>Fusarium oxysporum</i> wild-type MSA 35. <i>Proteomics</i> , 2010, 10, 3292-3320.	1.3	17
40	Phosphoproteomic Analysis Reveals Interconnected System-Wide Responses to Perturbations of Kinases and Phosphatases in Yeast. <i>Science Signaling</i> , 2010, 3, rs4.	1.6	277
41	PARP1 ADP-ribosylates lysine residues of the core histone tails. <i>Nucleic Acids Research</i> , 2010, 38, 6350-6362.	6.5	226
42	Tyrosine phosphorylation by Src within the cavity of the adenine nucleotide translocase 1 regulates ADP/ATP exchange in mitochondria. <i>American Journal of Physiology - Cell Physiology</i> , 2010, 298, C740-C748.	2.1	55
43	Deterministic protein inference for shotgun proteomics data provides new insights into Arabidopsis pollen development and function. <i>Genome Research</i> , 2009, 19, 1786-1800.	2.4	151
44	Identification and Functional Characterization of N-Terminally Acetylated Proteins in <i>Drosophila melanogaster</i> . <i>PLoS Biology</i> , 2009, 7, e1000236.	2.6	149
45	Polystyrene beads as an alternative support material for epitope identification of a prion-antibody interaction using proteolytic excision-mass spectrometry. <i>Analytical and Bioanalytical Chemistry</i> , 2009, 395, 1395-1401.	1.9	7
46	Proteomic profiling of <i>Cronobacter turicensis</i> 3032, a food-borne opportunistic pathogen. <i>Proteomics</i> , 2009, 9, 3564-3579.	1.3	15
47	Phosphorylated serine and threonine residues promote site-specific fragmentation of singly charged, arginine-containing peptide ions. <i>Rapid Communications in Mass Spectrometry</i> , 2009, 23, 1435-1445.	0.7	10
48	Analysis of Ewing Sarcoma (EWS)-Binding Proteins: Interaction with hnRNP M, U, and RNA-Helicases p68/72 within Protein-RNA Complexes. <i>Journal of Proteome Research</i> , 2009, 8, 4455-4465.	1.8	34
49	Community proteogenomics reveals insights into the physiology of phyllosphere bacteria. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 16428-16433.	3.3	774
50	Epitope mapping on bovine prion protein using chemical cross-linking and mass spectrometry. <i>Journal of Mass Spectrometry</i> , 2008, 43, 185-195.	0.7	55
51	Mass Spectrometry: A Tool for Enhanced Detection of Hemoglobin Variants. <i>Clinical Chemistry</i> , 2008, 54, 69-76.	1.5	73
52	Phosphoproteome analysis of isoflurane-protected heart mitochondria: phosphorylation of adenine nucleotide translocator-1 on Tyr194 regulates mitochondrial function. <i>Cardiovascular Research</i> , 2008, 80, 20-29.	1.8	94
53	The Mass Distance Fingerprint: A statistical framework for de novo detection of predominant modifications using high-accuracy mass spectrometry. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2007, 854, 173-182.	1.2	13
54	Zn7Metallothionein-3 and the Synaptic Vesicle Cycle: Interaction of Metallothionein-3 with the Small GTPase Rab3A. <i>Biochemistry</i> , 2005, 44, 3159-3165.	1.2	54

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55	Redox Labile Site in a Zn ₄ Cluster of Cu ₄ Zn ₄ ~Metallothionein-3. Biochemistry, 2003, 42, 9822-9828.	1.2	58
56	A distinct Cu ₄ -thiolate cluster of human metallothionein-3 is located in the N-terminal domain. Journal of Biological Inorganic Chemistry, 2002, 7, 611-616.	1.1	52