Andreas Pich

List of Publications by Year in descending order

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414034 430442 1,313 79 18 32 citations h-index g-index papers 81 81 81 2216 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Unravelling the Proteomics of HLA-B*57:01+ Antigen Presenting Cells during Abacavir Medication. Journal of Personalized Medicine, 2022, 12, 40.	1.1	2
2	AQUA Mutant Protein Quantification of Endomyocardial Biopsy-Sized Samples From a Patient With Hypertrophic Cardiomyopathy. Frontiers in Cardiovascular Medicine, 2022, 9, 816330.	1.1	0
3	Proteome profile of patients with excellent and poor speech intelligibility after cochlear implantation: Can perilymph proteins predict performance?. PLoS ONE, 2022, 17, e0263765.	1.1	1
4	Natural antibodies and CRP drive anaphylatoxin production by urate crystals. Scientific Reports, 2022, 12, 4483.	1.6	2
5	Proteomic Profiling and T Cell Receptor Usage of Abacavir Susceptible Subjects. Biomedicines, 2022, 10, 693.	1.4	1
6	<i>Pseudomonas aeruginosa</i> postâ€translational responses to elevated <scp>câ€diâ€GMP</scp> levels. Molecular Microbiology, 2022, 117, 1213-1226.	1.2	6
7	TcdB of Clostridioides difficile Mediates RAS-Dependent Necrosis in Epithelial Cells. International Journal of Molecular Sciences, 2022, 23, 4258.	1.8	5
8	Bioinformatic Analysis of the Perilymph Proteome to Generate a Human Protein Atlas. Frontiers in Cell and Developmental Biology, 2022, 10, 847157.	1.8	2
9	Meteorin-like promotes heart repair through endothelial KIT receptor tyrosine kinase. Science, 2022, 376, 1343-1347.	6.0	34
10	Possibilities of Molecular Perilymph Diagnostics in Patients with Cochlea Implant Surgeries. Laryngo-Rhino- Otologie, 2022, , .	0.2	0
11	Alpha-1 antitrypsin deficiency impairs lung antibacterial immunity in mice. JCI Insight, 2021, 6, .	2.3	12
12	Profilin2 regulates actin rod assembly in neuronal cells. Scientific Reports, 2021, 11, 10287.	1.6	7
13	The barrier functions of crude cervical mucus plugs against HIV-1 infection in the context of cell-free and cell-to-cell transmission. Aids, 2021, 35, 2105-2117.	1.0	4
14	Reconstruction of the miR-506-Quaking axis in Idiopathic Pulmonary Fibrosis using integrative multi-source bioinformatics. Scientific Reports, 2021, 11, 12456.	1.6	3
15	Myeloid-Derived Growth Factor Protects Against Pressure Overload–Induced Heart Failure by Preserving Sarco/Endoplasmic Reticulum Ca ²⁺ -ATPase Expression in Cardiomyocytes. Circulation, 2021, 144, 1227-1240.	1.6	27
16	The Coâ€mutational Spectrum Determines the Therapeutic Response in Murine FGFR2 Fusionâ€Driven Cholangiocarcinoma. Hepatology, 2021, 74, 1357-1370.	3.6	13
17	Personalized Proteomics for Precision Diagnostics in Hearing Loss: Disease-Specific Analysis of Human Perilymph by Mass Spectrometry. ACS Omega, 2021, 6, 21241-21254.	1.6	7
18	A mouse model of cardiogenic shock. Cardiovascular Research, 2021, 117, 2414-2415.	1.8	2

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19	The Binary Toxin of Clostridioides difficile Alters the Proteome and Phosphoproteome of HEp-2 Cells. Frontiers in Microbiology, 2021, 12, 725612.	1.5	11
20	Reversible Growth-Arrest of a Spontaneously-Derived Human MSC-Like Cell Line. International Journal of Molecular Sciences, 2020, 21, 4752.	1.8	18
21	Overexpression of Macrophage-Inducible C-Type Lectin Mincle Aggravates Proinflammatory Responses to <i>Streptococcus pneumoniae</i> with Fatal Outcome in Mice. Journal of Immunology, 2020, 205, 3390-3399.	0.4	7
22	The Proteome and Secretome of Cortical Brain Cells Infected With Herpes Simplex Virus. Frontiers in Neurology, 2020, 11, 844.	1.1	7
23	Binding of Macrophage Receptor MARCO, LDL, and LDLR to Disease-Associated Crystalline Structures. Frontiers in Immunology, 2020, 11, 596103.	2.2	10
24	Identification of the Cleavage Domain within Glycoprotein G of Herpes Simplex Virus Type 2. Viruses, 2020, 12, 1428.	1.5	4
25	The Loss of HLA-F/KIR3DS1 Ligation Is Mediated by Hemoglobin Peptides. International Journal of Molecular Sciences, 2020, 21, 8012.	1.8	4
26	Comprehensive Bioinformatics Identifies Key microRNA Players in ATG7-Deficient Lung Fibroblasts. International Journal of Molecular Sciences, 2020, 21, 4126.	1.8	11
27	Inflammatory Drivers of Cardiovascular Disease: Molecular Characterization of Senescent Coronary Vascular Smooth Muscle Cells. Frontiers in Physiology, 2020, 11, 520.	1.3	23
28	Pleiotropic cardiac functions controlled by ischemia-induced lncRNA H19. Journal of Molecular and Cellular Cardiology, 2020, 146, 43-59.	0.9	12
29	GC-MS and LC-MS/MS pilot studies on the guanidine (NG)-dimethylation in native, asymmetrically and symmetrically NG-dimethylated arginine-vasopressin peptides and proteins in human red blood cells. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2020, 1141, 122024.	1.2	5
30	C-reactive protein (CRP) recognizes uric acid crystals and recruits proteases C1 and MASP1. Scientific Reports, 2020, 10, 6391.	1.6	19
31	Pharmacokinetic Studies of Antisense Oligonucleotides Using MALDI-TOF Mass Spectrometry. Frontiers in Pharmacology, 2020, 11, 220.	1.6	2
32	Genetic information from discordant sibling pairs points to ESRP2 as a candidate trans-acting regulator of the CF modifier gene SCNN1B. Scientific Reports, 2020, 10, 22447.	1.6	4
33	miR-21-KO Alleviates Alveolar Structural Remodeling and Inflammatory Signaling in Acute Lung Injury. International Journal of Molecular Sciences, 2020, 21, 822.	1.8	9
34	The Mechanistic Differences in HLA-Associated Carbamazepine Hypersensitivity. Pharmaceutics, 2019, 11, 536.	2.0	12
35	E. coli primase and DNA polymerase III holoenzyme are able to bind concurrently to a primed template during DNA replication. Scientific Reports, 2019, 9, 14460.	1.6	4
36	<scp>TIP</scp> 30 counteracts cardiac hypertrophy and failure by inhibiting translational elongation. EMBO Molecular Medicine, 2019, 11, e10018.	3.3	17

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37	The Core Proteome of Biofilm-Grown Clinical Pseudomonas aeruginosa Isolates. Cells, 2019, 8, 1129.	1.8	26
38	Therapeutic modulation of RNA-binding protein Rbm38 facilitates re-endothelialization after arterial injury. Cardiovascular Research, 2019, 115 , 1804 - 1810 .	1.8	12
39	Proteome and Phosphoproteome Analysis in TNF Long Term-Exposed Primary Human Monocytes. International Journal of Molecular Sciences, 2019, 20, 1241.	1.8	8
40	HLA-F Allele-Specific Peptide Restriction Represents an Exceptional Proteomic Footprint. International Journal of Molecular Sciences, 2019, 20, 5572.	1.8	5
41	Stable depletion of RUNX1-ETO in Kasumi-1 cells induces expression and enhanced proteolytic activity of Cathepsin G and Neutrophil Elastase. PLoS ONE, 2019, 14, e0225977.	1.1	4
42	Difference in Mono-O-Glucosylation of Ras Subtype GTPases Between Toxin A and Toxin B From Clostridioides difficile Strain 10463 and Lethal Toxin From Clostridium sordellii Strain 6018. Frontiers in Microbiology, 2018, 9, 3078.	1.5	19
43	Quantitative Phosphoproteome Analysis of Clostridioides difficile Toxin B Treated Human Epithelial Cells. Frontiers in Microbiology, 2018, 9, 3083.	1.5	5
44	Identification of targets of monoclonal antibodies that inhibit adhesion and growth in Mycoplasma mycoides subspecies mycoides. Veterinary Immunology and Immunopathology, 2018, 204, 11-18.	0.5	6
45	Environmentâ€driven changes of mRNA and protein levels in <i>Pseudomonas aeruginosa</i> . Environmental Microbiology, 2018, 20, 3952-3963.	1.8	19
46	Kaposi's Sarcoma-Associated Herpesvirus Nonstructural Membrane Protein pK15 Recruits the Class II Phosphatidylinositol 3-Kinase PI3K-C2 $\hat{l}\pm$ To Activate Productive Viral Replication. Journal of Virology, 2018, 92, .	1.5	18
47	Burst-Like Transcription of Mutant and Wildtype MYH7-Alleles as Possible Origin of Cell-to-Cell Contractile Imbalance in Hypertrophic Cardiomyopathy. Frontiers in Physiology, 2018, 9, 359.	1.3	39
48	Quantitative Secretomics Reveals Extrinsic Signals Involved in Human Pluripotent Stem Cell Cardiomyogenesis. Proteomics, 2018, 18, e1800102.	1.3	23
49	MS-based quantification of RhoA/B and RhoC ADP-ribosylation. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2018, 1092, 268-271.	1.2	3
50	Proteomic Analysis of Human Pluripotent Stem Cell Cardiomyogenesis Revealed Altered Expression of Metabolic Enzymes and PDLIM5 Isoforms. Journal of Proteome Research, 2017, 16, 1133-1149.	1.8	32
51	Proteome Analysis of Human Perilymph Using an Intraoperative Sampling Method. Journal of Proteome Research, 2017, 16, 1911-1923.	1.8	59
52	Quantification of small GTPase glucosylation by clostridial glucosylating toxins using multiplexed MRM analysis. Proteomics, 2017, 17, 1700016.	1.3	10
53	Quantitative Assessment of Sialoâ€Glycoproteins and Nâ€Glycans during Cardiomyogenic Differentiation of Human Induced Pluripotent Stem Cells. ChemBioChem, 2017, 18, 1317-1331.	1.3	44
54	Glucosyltransferase-dependent and -independent effects of TcdB on the proteome of HEp-2 cells. Proteomics, 2017, 17, 1600435.	1.3	15

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55	Toxin A of the nosocomial pathogen <i>Clostridium difficile</i> induces primary effects in the proteome of HEpâ€2 cells. Proteomics - Clinical Applications, 2017, 11, 1600031.	0.8	14
56	The use of urinary proteomics in the assessment of suitability of mouse models for ageing. PLoS ONE, 2017, 12, e0166875.	1.1	17
57	Effect of hyperbaric oxygen on BDNF-release and neuroprotection: Investigations with human mesenchymal stem cells and genetically modified NIH3T3 fibroblasts as putative cell therapeutics. PLoS ONE, 2017, 12, e0178182.	1.1	20
58	Detection and Quantification of ADP-Ribosylated RhoA/B by Monoclonal Antibody. Toxins, 2016, 8, 100.	1.5	9
59	Unravelling postâ€ŧranscriptional PrmCâ€dependent regulatory mechanisms in <i>Pseudomonas aeruginosa</i> . Environmental Microbiology, 2016, 18, 3583-3592.	1.8	6
60	Evidence by chromatography and mass spectrometry that inorganic nitrite induces S -glutathionylation of hemoglobin in human red blood cells. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2016, 1019, 72-82.	1.2	16
61	Surface-bound bovine serum albumin carrier protein as present in recombinant cytokine preparations amplifies T helper 17 cell polarization. Scientific Reports, 2016, 6, 36598.	1.6	1
62	Cytoplasmic isoforms of Kaposi sarcoma herpesvirus LANA recruit and antagonize the innate immune DNA sensor cGAS. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E1034-43.	3.3	128
63	Proteomic characterization of pleural effusion, a specific host niche of Mycoplasma mycoides subsp. mycoides from cattle with contagious bovine pleuropneumonia (CBPP). Journal of Proteomics, 2016, 131, 93-103.	1.2	12
64	Proteome Alterations of Hippocampal Cells Caused by Clostridium botulinum C3 Exoenzyme. Journal of Proteome Research, 2015, 14, 4721-4733.	1.8	10
65	Human neutrophils are activated by a peptide fragment of Clostridium difficiletoxin B presumably via formyl peptide receptor. Cellular Microbiology, 2015, 17, 893-909.	1.1	17
66	C3-induced release of neurotrophic factors from Schwann cells $\hat{a} \in \text{``}$ potential mechanism behind its regeneration promoting activity. Neurochemistry International, 2015, 90, 232-245.	1.9	3
67	Myeloid-derived growth factor (C19orf10) mediates cardiac repair following myocardial infarction. Nature Medicine, 2015, 21, 140-149.	15.2	168
68	DXD Motif-Dependent and -Independent Effects of the Chlamydia trachomatis Cytotoxin CT166. Toxins, 2015, 7, 621-637.	1.5	15
69	SUMO modification of TBK1 at the adaptor-binding C-terminal coiled-coil domain contributes to its antiviral activity. Biochimica Et Biophysica Acta - Molecular Cell Research, 2015, 1853, 136-143.	1.9	29
70	Identification of ageing-associated naturally occurring peptides in human urine. Oncotarget, 2015, 6, 34106-34117.	0.8	31
71	Verification of a canine PSMA (FolH1) antibody. Anticancer Research, 2015, 35, 145-8.	0.5	6
72	Vimentin Mediates Uptake of C3 Exoenzyme. PLoS ONE, 2014, 9, e101071.	1.1	31

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73	Time-resolved cellular effects induced by TcdA from <i>Clostridium difficile</i> . Rapid Communications in Mass Spectrometry, 2014, 28, 1089-1100.	0.7	11
74	Proteome analysis in the assessment of ageing. Ageing Research Reviews, 2014, 18, 74-85.	5.0	18
75	Substrate Specificity of Clostridial Glucosylating Toxins and Their Function on Colonocytes Analyzed by Proteomics Techniques. Journal of Proteome Research, 2013, 12, 1604-1618.	1.8	50
76	Mir $\hat{a}^{-1}/417$ -92 Identifies BCL2 As a Therapeutic Target In BCR-ABL Positive B-Lineage Acute Lymphoblastic Leukemia. Blood, 2013, 122, 835-835.	0.6	0
77	Impact of clostridial glucosylating toxins on the proteome of colonic cells determined by isotope-coded protein labeling and LC-MALDI. Proteome Science, 2011, 9, 48.	0.7	38
78	Mass spectrometry-based methods for biomarker detection and analysis. Drug Discovery Today: Technologies, 2005, 2, 361-367.	4.0	6
79	Small Molecule/HLA Complexes Alter the Cellular Proteomic Content. , 0, , .		1