Roman Fischer

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7044550/publications.pdf

Version: 2024-02-01

155 papers 10,601 citations

51 h-index 93 g-index

183

183 docs citations

times ranked

183

19912 citing authors

#	Article	IF	CITATIONS
1	Interactome screening of (i) C9orf72 (i) dipeptide repeats reveals VCP sequestration and functional impairment by polyGA. Brain, 2022, 145, 684-699.	3.7	15
2	Integrated Plasma and Tissue Proteomics Reveals Attractin Release by Intraluminal Thrombus of Abdominal Aortic Aneurysms and Improves Aneurysm Growth Prediction in Humans. Annals of Surgery, 2022, 275, 1206-1211.	2.1	13
3	ABPP-HT*â€"Deep Meets Fast for Activity-Based Profiling of Deubiquitylating Enzymes Using Advanced DIA Mass Spectrometry Methods. International Journal of Molecular Sciences, 2022, 23, 3263.	1.8	7
4	Sequence and structural variations determining the recruitment of WNK kinases to the KLHL3 E3 ligase. Biochemical Journal, 2022, 479, 661-675.	1.7	4
5	Factor inhibiting HIF can catalyze two asparaginyl hydroxylations in VNVN motifs of ankyrin fold proteins. Journal of Biological Chemistry, 2022, 298, 102020.	1.6	4
6	Chronic inflammatory arthritis drives systemic changes in circadian energy metabolism. Proceedings of the National Academy of Sciences of the United States of America, 2022, 119, e2112781119.	3.3	11
7	The effects of endogenously―and exogenously―nduced hyperketonemia on exercise performance and adaptation. Physiological Reports, 2022, 10, .	0.7	8
8	T-cell trans-synaptic vesicles are distinct and carry greater effector content than constitutive extracellular vesicles. Nature Communications, $2022,13,.$	5. 8	18
9	Abstract 1247: Comprehensive molecular profiling to predict first-line immunochemotherapy outcomes in inoperable esophageal adenocarcinoma. Cancer Research, 2022, 82, 1247-1247.	0.4	0
10	LARP1 isoform expression in human cancer cell lines. RNA Biology, 2021, 18, 237-247.	1.5	11
11	Value of lipocalin 2 as a potential biomarker for bacterial meningitis. Clinical Microbiology and Infection, 2021, 27, 724-730.	2.8	9
12	Assessment of different screening methods for selecting palaeontological bone samples for peptide sequencing. Journal of Proteomics, 2021, 230, 103986.	1.2	3
13	Detection and quantification of novel Câ€ŧerminal TDPâ€43 fragments in ALSâ€₹DP. Brain Pathology, 2021, 31, e12923.	2.1	26
14	<i>Drosophila</i> Sex Peptide controls the assembly of lipid microcarriers in seminal fluid. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	16
15	Functional analysis of a gene-edited mouse model to gain insights into the disease mechanisms of a titin missense variant. Basic Research in Cardiology, 2021, 116, 14.	2.5	16
16	SPRTN protease-cleaved MRE11 decreases DNA repair and radiosensitises cancer cells. Cell Death and Disease, 2021, 12, 165.	2.7	8
17	ABPP-HT - High-Throughput Activity-Based Profiling of Deubiquitylating Enzyme Inhibitors in a Cellular Context. Frontiers in Chemistry, 2021, 9, 640105.	1.8	15
18	Global proteomic analysis of extracellular matrix in mouse and human brain highlights relevance to cerebrovascular disease. Journal of Cerebral Blood Flow and Metabolism, 2021, 41, 2423-2438.	2.4	14

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19	Network Analysis of the CSF Proteome Characterizes Convergent Pathways of Cellular Dysfunction in ALS. Frontiers in Neuroscience, 2021, 15, 642324.	1.4	6
20	Thymosin \hat{l}^24 protects against aortic aneurysm via endocytic regulation of growth factor signaling. Journal of Clinical Investigation, 2021, 131, .	3.9	15
21	Assessing the degradation of ancient milk proteinsÂthrough site-specific deamidation patterns. Scientific Reports, 2021, 11, 7795.	1.6	22
22	Functional Genomic Analysis of a <i>RUNX3</i> Polymorphism Associated With Ankylosing Spondylitis. Arthritis and Rheumatology, 2021, 73, 980-990.	2.9	10
23	Kawasaki Disease Patient Stratification and Pathway Analysis Based on Host Transcriptomic and Proteomic Profiles. International Journal of Molecular Sciences, 2021, 22, 5655.	1.8	6
24	The chaperonin CCT8 controls proteostasis essential for T cell maturation, selection, and function. Communications Biology, 2021, 4, 681.	2.0	6
25	Phenotypic manifestation of α-synuclein strains derived from Parkinson's disease and multiple system atrophy in human dopaminergic neurons. Nature Communications, 2021, 12, 3817.	5.8	52
26	Diabetic mitochondria are resistant to palmitoyl CoA inhibition of respiration, which is detrimental during ischemia. FASEB Journal, 2021, 35, e21765.	0.2	4
27	Proteomic Signature of Dysfunctional Circulating Endothelial Colonyâ€Forming Cells of Young Adults. Journal of the American Heart Association, 2021, 10, e021119.	1.6	3
28	A modified density gradient proteomic-based method to analyze endolysosomal proteins in cardiac tissue. IScience, 2021, 24, 102949.	1.9	1
29	Age-dependent changes in protein incorporation into collagen-rich tissues of mice by in vivo pulsed SILAC labelling. ELife, 2021, 10, .	2.8	22
30	The Bloom syndrome complex senses RPA-coated single-stranded DNA to restart stalled replication forks. Nature Communications, 2021, 12, 585.	5.8	48
31	Regulation of CYLD activity and specificity by phosphorylation and ubiquitin-binding CAP-Gly domains. Cell Reports, 2021, 37, 109777.	2.9	20
32	Defactinib inhibits PYK2 phosphorylation of IRF5 and reduces intestinal inflammation. Nature Communications, 2021, 12, 6702.	5.8	13
33	FOXN1 forms higher-order nuclear condensates displaced by mutations causing immunodeficiency. Science Advances, 2021, 7, eabj9247.	4.7	10
34	Advancing mechanistic understanding and biomarker development in amyotrophic lateral sclerosis. Expert Review of Proteomics, 2021, 18, 977-994.	1.3	5
35	Lineage-Restricted Regulation of SCD and Fatty Acid Saturation by MITF Controls Melanoma Phenotypic Plasticity. Molecular Cell, 2020, 77, 120-137.e9.	4.5	87
36	An ALS-linked mutation in TDP-43 disrupts normal protein interactions in the motor neuron response to oxidative stress. Neurobiology of Disease, 2020, 144, 105050.	2.1	30

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37	The Immunomodulatory Metabolite Itaconate Modifies NLRP3 and Inhibits Inflammasome Activation. Cell Metabolism, 2020, 32, 468-478.e7.	7.2	283
38	The role of birds at Çatalhöyük revealed by the analysis of eggshell. Quaternary International, 2020, 543, 50-60.	0.7	10
39	Quality Control of ER Membrane Proteins by the RNF185/Membralin Ubiquitin Ligase Complex. Molecular Cell, 2020, 79, 768-781.e7.	4.5	41
40	Identifying collagen VI as a target of fibrotic diseases regulated by CREBBP/EP300. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 20753-20763.	3.3	45
41	Living off the land: Terrestrial-based diet and dairying in the farming communities of the Neolithic Balkans. PLoS ONE, 2020, 15, e0237608.	1.1	21
42	CSF extracellular vesicle proteomics demonstrates altered protein homeostasis in amyotrophic lateral sclerosis. Clinical Proteomics, 2020, 17, 31.	1.1	27
43	Supramolecular attack particles are autonomous killing entities released from cytotoxic T cells. Science, 2020, 368, 897-901.	6.0	98
44	Phosphoproteomics of CD2 signaling reveals AMPK-dependent regulation of lytic granule polarization in cytotoxic T cells. Science Signaling, 2020, 13, .	1.6	18
45	TEX264 coordinates p97- and SPRTN-mediated resolution of topoisomerase 1-DNA adducts. Nature Communications, 2020, 11, 1274.	5.8	64
46	What's the catch? Archaeological application of rapid collagen-based species identification for Pacific Salmon. Journal of Archaeological Science, 2020, 116, 105116.	1.2	19
47	MaxQuant Software for Ion Mobility Enhanced Shotgun Proteomics. Molecular and Cellular Proteomics, 2020, 19, 1058-1069.	2.5	128
48	Male reproductive aging arises via multifaceted mating-dependent sperm and seminal proteome declines, but is postponable in <i>Drosophila</i> . Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 17094-17103.	3.3	39
49	DeamiDATE 1.0: Site-specific deamidation as a tool to assess authenticity of members of ancient proteomes. Journal of Archaeological Science, 2020, 115, 105080.	1.2	36
50	Characterization of exosomes in peritoneal fluid of endometriosis patients. Fertility and Sterility, 2020, 113, 364-373.e2.	0.5	35
51	Glycosylated Siglec-6 expression in syncytiotrophoblast-derived extracellular vesicles from preeclampsia placentas. Biochemical and Biophysical Research Communications, 2020, 533, 838-844.	1.0	19
52	Scottish soldiers from the Battle of Dunbar 1650: A prosopographical approach to a skeletal assemblage. PLoS ONE, 2020, 15, e0243369.	1.1	7
53	Interaction mapping of endoplasmic reticulum ubiquitin ligases identifies modulators of innate immune signalling. ELife, 2020, 9, .	2.8	61
54	Ataxin-3 Links NOD2 and TLR2 Mediated Innate Immune Sensing and Metabolism in Myeloid Cells. Frontiers in Immunology, 2019, 10, 1495.	2.2	11

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55	SPRTN protease and checkpoint kinase 1 cross-activation loop safeguards DNA replication. Nature Communications, 2019, 10, 3142.	5.8	36
56	Nitric Oxide Modulates Metabolic Remodeling in Inflammatory Macrophages through TCA Cycle Regulation and Itaconate Accumulation. Cell Reports, 2019, 28, 218-230.e7.	2.9	149
57	Mechanism of Signalling and Adaptation through the Rhodobacter sphaeroides Cytoplasmic Chemoreceptor Cluster. International Journal of Molecular Sciences, 2019, 20, 5095.	1.8	9
58	Extracellular vesicle integrins act as a nexus for platelet adhesion in cerebral microvessels. Scientific Reports, 2019, 9, 15847.	1.6	9
59	Divergent allocation of sperm and the seminal proteome along a competition gradient in $\langle i \rangle$ Drosophila melanogaster $\langle i \rangle$. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 17925-17933.	3.3	76
60	New insights into Neolithic milk consumption through proteomic analysis of dental calculus. Archaeological and Anthropological Sciences, 2019, 11, 6183-6196.	0.7	45
61	Proteomics analysis of the matrisome from MC38 experimental mouse liver metastases. American Journal of Physiology - Renal Physiology, 2019, 317, G625-G639.	1.6	7
62	Comprehensive Landscape of Active Deubiquitinating Enzymes Profiled by Advanced Chemoproteomics. Frontiers in Chemistry, 2019, 7, 592.	1.8	41
63	Birds of prey and humans in prehistoric Europe: A view from El Mirón Cave, Cantabria (Spain). Journal of Archaeological Science: Reports, 2019, 24, 244-252.	0.2	8
64	Diabetes causes marked inhibition of mitochondrial metabolism in pancreatic \hat{l}^2 -cells. Nature Communications, 2019, 10, 2474.	5.8	223
65	<scp>RASSF</scp> 1A is required for the maintenance of nuclear actin levels. EMBO Journal, 2019, 38, e101168.	3.5	37
66	Palaeoproteomics resolves sloth relationships. Nature Ecology and Evolution, 2019, 3, 1121-1130.	3.4	91
67	Molecular insights into an ancient form of Paget's disease of bone. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 10463-10472.	3.3	24
68	A late Middle Pleistocene Denisovan mandible from the Tibetan Plateau. Nature, 2019, 569, 409-412.	13.7	302
69	MDC1 Interacts with TOPBP1 to Maintain Chromosomal Stability during Mitosis. Molecular Cell, 2019, 74, 571-583.e8.	4.5	97
70	Adaptation to HIF1 $\hat{1}$ ± Deletion in Hypoxic Cancer Cells by Upregulation of GLUT14 and Creatine Metabolism. Molecular Cancer Research, 2019, 17, 1531-1544.	1.5	22
71	Tumor Imaging Using Radiolabeled Matrix Metalloproteinase–Activated Anthrax Proteins. Journal of Nuclear Medicine, 2019, 60, 1474-1482.	2.8	6
72	Development of a Sensitive, Scalable Method for Spatial, Cell-Type-Resolved Proteomics of the Human Brain. Journal of Proteome Research, 2019, 18, 1787-1795.	1.8	39

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73	Colonic epithelial cell diversity in health and inflammatory bowel disease. Nature, 2019, 567, 49-55.	13.7	486
74	BMP signaling inhibition in $\langle i \rangle$ Drosophila $\langle i \rangle$ secondary cells remodels the seminal proteome and self and rival ejaculate functions. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 24719-24728.	3.3	29
75	Integrated Physiological and Biochemical Assessments for the Prediction of Growth of Abdominal Aortic Aneurysms in Humans. Annals of Surgery, 2019, 270, e1-e3.	2.1	16
76	Quantitative Proteomics Identification of Seminal Fluid Proteins in Male Drosophila melanogaster. Molecular and Cellular Proteomics, 2019, 18, S46-S58.	2.5	66
77	Lack of activity of recombinant HIF prolyl hydroxylases (PHDs) on reported non-HIF substrates. ELife, 2019, 8, .	2.8	70
78	Composition and structure of synaptic ectosomes exporting antigen receptor linked to functional CD40 ligand from helper T cells. ELife, $2019, 8, .$	2.8	57
79	Cardiac Dysfunction and Metabolic Inflexibility in a Mouse Model of Diabetes Without Dyslipidemia. Diabetes, 2018, 67, 1057-1067.	0.3	28
80	< scp>FBXL $<$ /scp> 13 directs the proteolysis of $<$ scp>CEP $<$ /scp> 192 to regulate centrosome homeostasis and cell migration. EMBO Reports, 2018, 19, .	2.0	18
81	Cerebrospinal fluid macrophage biomarkers in amyotrophic lateral sclerosis. Annals of Neurology, 2018, 83, 258-268.	2.8	107
82	A robust mass spectrometry method for rapid profiling of erythrocyte ghost membrane proteomes. Clinical Proteomics, 2018, 15, 14.	1.1	28
83	Itaconate is an anti-inflammatory metabolite that activates Nrf2 via alkylation of KEAP1. Nature, 2018, 556, 113-117.	13.7	1,115
84	UFLCâ€Derived CSF Extracellular Vesicle Origin and Proteome. Proteomics, 2018, 18, e1800257.	1.3	36
85	Colorectal cancer liver metastatic growth depends on PAD4-driven citrullination of the extracellular matrix. Nature Communications, 2018, 9, 4783.	5.8	134
86	Bi-directional signaling by membrane-bound KitL induces proliferation and coordinates thymic endothelial cell and thymocyte expansion. Nature Communications, 2018, 9, 4685.	5.8	9
87	Ancient proteins from ceramic vessels at \tilde{A} ‡atalh \tilde{A} ¶y \tilde{A} ½k West reveal the hidden cuisine of early farmers. Nature Communications, 2018, 9, 4064.	5.8	105
88	\hat{l}^2 -TrCP- and Casein Kinase II-Mediated Degradation of Cyclin F Controls Timely Mitotic Progression. Cell Reports, 2018, 24, 3404-3412.	2.9	37
89	Tetrahydrobiopterin modulates ubiquitin conjugation to UBC13/UBE2N and proteasome activity by S-nitrosation. Scientific Reports, 2018, 8, 14310.	1.6	5
90	Proteomic profiling of the plasma of Gambian children with cerebral malaria. Malaria Journal, 2018, 17, 337.	0.8	16

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91	A variant NuRD complex containing PWWP2A/B excludes MBD2/3 to regulate transcription at active genes. Nature Communications, 2018, 9, 3798.	5.8	40
92	Glutamate receptor $\hat{l}'2$ serum antibodies in pediatric opsoclonus myoclonus ataxia syndrome. Neurology, 2018, 91, e714-e723.	1.5	43
93	53BP1 cooperation with the REV7–shieldin complex underpins DNA structure-specific NHEJ. Nature, 2018, 560, 122-127.	13.7	222
94	A TFEB nuclear export signal integrates amino acid supply and glucose availability. Nature Communications, 2018, 9, 2685.	5.8	84
95	Proteomic evidence of dietary sources in ancient dental calculus. Proceedings of the Royal Society B: Biological Sciences, 2018, 285, 20180977.	1.2	97
96	BRAF/MAPK and GSK3 signaling converges to control MITF nuclear export. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E8668-E8677.	3.3	50
97	Proteo-metabolomics reveals compensation between ischemic and non-injured contralateral kidneys after reperfusion. Scientific Reports, 2018, 8, 8539.	1.6	39
98	Development and validation of response markers to predict survival and pleurodesis success in patients with malignant pleural effusion (PROMISE): a multicohort analysis. Lancet Oncology, The, 2018, 19, 930-939.	5.1	92
99	The Jumonji-C oxygenase JMJD7 catalyzes (3S)-lysyl hydroxylation of TRAFAC GTPases. Nature Chemical Biology, 2018, 14, 688-695.	3.9	31
100	Proteomic profiling of the brain of mice with experimental cerebral malaria. Journal of Proteomics, 2018, 180, 61-69.	1.2	5
101	PCNA dependent cellular activities tolerate dramatic perturbations in PCNA client interactions. DNA Repair, 2017, 50, 22-35.	1.3	12
102	A Ribosomopathy Reveals Decoding Defective Ribosomes Driving Human Dysmorphism. American Journal of Human Genetics, 2017, 100, 506-522.	2.6	69
103	Expanding Proteome Coverage with CHarge Ordered Parallel Ion aNalysis (CHOPIN) Combined with Broad Specificity Proteolysis. Journal of Proteome Research, 2017, 16, 1288-1299.	1.8	92
104	Inflammatory Stroke Extracellular Vesicles Induce Macrophage Activation. Stroke, 2017, 48, 2292-2296.	1.0	49
105	A novel role for endothelial tetrahydrobiopterin in mitochondrial redox balance. Free Radical Biology and Medicine, 2017, 104, 214-225.	1.3	49
106	New criteria for the molecular identification of cereal grains associated with archaeological artefacts. Scientific Reports, 2017, 7, 6633.	1.6	63
107	Association of Distinct Fine Specificities of Antiâ ^{**} Citrullinated Peptide Antibodies With Elevated Immune Responses to <i>Prevotella intermedia</i> in a Subgroup of Patients With Rheumatoid Arthritis and Periodontitis. Arthritis and Rheumatology, 2017, 69, 2303-2313.	2.9	37
108	A novel workflow combining plaque imaging, plaque and plasma proteomics identifies biomarkers of human coronary atherosclerotic plaque disruption. Clinical Proteomics, 2017, 14, 22.	1.1	16

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109	The identification of archaeological eggshell using peptide markers. Science and Technology of Archaeological Research, 2017, 3, 89-99.	2.4	23
110	Modulation of proteostasis counteracts oxidative stress and affects DNA base excision repair capacity in ATM-deficient cells. Nucleic Acids Research, 2017, 45, 10042-10055.	6.5	13
111	RYBP stimulates PRC1 to shape chromatin-based communication between Polycomb repressive complexes. ELife, 2016, 5, .	2.8	111
112	<scp>SCF</scp> (Fbxl17) ubiquitylation of Sufu regulates Hedgehog signaling and medulloblastoma development. EMBO Journal, 2016, 35, 1400-1416.	3.5	50
113	Integrative Phosphoproteomics Links IL-23R Signaling with Metabolic Adaptation in Lymphocytes. Scientific Reports, 2016, 6, 24491.	1.6	24
114	Proteomic changes in response to crystal formation in Drosophila Malpighian tubules. Fly, 2016, 10, 91-100.	0.9	12
115	Defining the HLA class lâ€associated viral antigen repertoire from HIVâ€1â€infected human cells. European Journal of Immunology, 2016, 46, 60-69.	1.6	57
116	Salt-Inducible Kinase 2 Couples Ovarian Cancer Cell Metabolism with Survival at the Adipocyte-Rich Metastatic Niche. Cancer Cell, 2016, 30, 273-289.	7.7	143
117	Abrogation of collagen-induced arthritis by a peptidyl arginine deiminase inhibitor is associated with modulation of T cell-mediated immune responses. Scientific Reports, 2016, 6, 26430.	1.6	76
118	Palaeoproteomic evidence identifies archaic hominins associated with the $Ch\tilde{A}^{\ddagger}$ telperronian at the Grotte du Renne. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 11162-11167.	3.3	251
119	Metalloprotease SPRTN/DVC1 Orchestrates Replication-Coupled DNA-Protein Crosslink Repair. Molecular Cell, 2016, 64, 704-719.	4.5	193
120	Plasma degradome affected by variable storage of human blood. Clinical Proteomics, 2016, 13, 26.	1.1	27
121	A7.01â€Abrogation of collagen-induced arthritis by a second generation peptidyl arginine deiminase inhibitor is associated with a shift from TH1/TH17 to TH2-mediated immune responses. Annals of the Rheumatic Diseases, 2016, 75, A56.1-A56.	0.5	0
122	Identification of an immunodominant peptide from citrullinated tenascin-C as a major target for autoantibodies in rheumatoid arthritis. Annals of the Rheumatic Diseases, 2016, 75, 1876-1883.	0.5	58
123	Crystal structure of i>Porphyromonas gingivalis ii>peptidylarginine deiminase: implications for autoimmunity in rheumatoid arthritis. Annals of the Rheumatic Diseases, 2016, 75, 1255-1261.	0.5	66
124	Structure of the Human Protein Kinase ZAK in Complex with Vemurafenib. ACS Chemical Biology, 2016, 11, 1595-1602.	1.6	19
125	Protein sequences bound to mineral surfaces persist into deep time. ELife, 2016, 5, .	2.8	176
126	Gelâ€aided sample preparation (GASP)â€"A simplified method for gelâ€assisted proteomic sample generation from protein extracts and intact cells. Proteomics, 2015, 15, 1224-1229.	1.3	104

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127	Identification of distinct circulating exosomes in Parkinson's disease. Annals of Clinical and Translational Neurology, 2015, 2, 353-361.	1.7	111
128	Animal origin of 13th-century uterine vellum revealed using noninvasive peptide fingerprinting. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 15066-15071.	3.3	140
129	Expanding the yeast protein arginine methylome. Proteomics, 2015, 15, 3232-3243.	1.3	21
130	Cells deficient in base-excision repair reveal cancer hallmarks originating from adjustments to genetic instability. Nucleic Acids Research, 2015, 43, 3667-3679.	6.5	39
131	Expression of citrulline and homocitrulline residues in the lungs of non-smokers and smokers: implications for autoimmunity in rheumatoid arthritis. Arthritis Research and Therapy, 2015, 17, 9.	1.6	102
132	Ancient proteins resolve the evolutionary history of Darwin's South American ungulates. Nature, 2015, 522, 81-84.	13.7	273
133	Striking Oxygen Sensitivity of the Peptidylglycine \hat{l}_{\pm} -Amidating Monooxygenase (PAM) in Neuroendocrine Cells. Journal of Biological Chemistry, 2015, 290, 24891-24901.	1.6	25
134	Hypoxia induces a lipogenic cancer cell phenotype via HIF1 \hat{l} ±-dependent and -independent pathways. Oncotarget, 2015, 6, 1920-1941.	0.8	72
135	Integration of Proteomics and Metabolomics to Unravel the Mechanism of Ischemia Reperfusion Injury in Donor Kidneys for Transplantation Transplantation, 2014, 98, 76.	0.5	0
136	RNA and Imidazoquinolines Are Sensed by Distinct TLR7/8 Ectodomain Sites Resulting in Functionally Disparate Signaling Events. Journal of Immunology, 2014, 192, 5963-5973.	0.4	38
137	Optimal Translational Termination Requires C4 Lysyl Hydroxylation of eRF1. Molecular Cell, 2014, 53, 645-654.	4.5	99
138	OGFOD1 catalyzes prolyl hydroxylation of RPS23 and is involved in translation control and stress granule formation. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 4031-4036.	3.3	105
139	Critical Role of Endoplasmic Reticulum Aminopeptidase 1 in Determining the Length and Sequence of Peptides Bound and Presented by HLA–B27. Arthritis and Rheumatology, 2014, 66, 284-294.	2.9	71
140	<scp>GOAT</scp> – A simple <scp>LC</scp> â€ <scp>MS</scp> / <scp>MS</scp> gradient optimization tool. Proteomics, 2014, 14, 1467-1471.	1.3	17
141	Direct evidence of milk consumption from ancient human dental calculus. Scientific Reports, 2014, 4, 7104.	1.6	184
142	Inhibition of Mitochondrial Aconitase by Succination in Fumarate Hydratase Deficiency. Cell Reports, 2013, 3, 689-700.	2.9	137
143	PfHPRT: A New Biomarker Candidate of Acute <i>Plasmodium falciparum</i> Infection. Journal of Proteome Research, 2013, 12, 1211-1222.	1.8	19
144	Two birds with one stone: Doing metabolomics with your proteomics kit. Proteomics, 2013, 13, 3371-3386.	1.3	24

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145	Detection of Multiple Autoantibodies in Patients with Ankylosing Spondylitis Using Nucleic Acid Programmable Protein Arrays. Molecular and Cellular Proteomics, 2012, 11, M9.00384.	2.5	77
146	Discovery of Candidate Serum Proteomic and Metabolomic Biomarkers in Ankylosing Spondylitis. Molecular and Cellular Proteomics, 2012, 11, M111.013904.	2.5	92
147	Detection of BK virus in urine from renal transplant subjects by mass spectrometry. Clinical Proteomics, 2012, 9, 4.	1.1	20
148	Renal Cyst Formation in Fh1-Deficient Mice Is Independent of the Hif/Phd Pathway: Roles for Fumarate in KEAP1 Succination and Nrf2 Signaling. Cancer Cell, 2011, 20, 524-537.	7.7	494
149	Activity-Based Chemical Proteomics Accelerates Inhibitor Development for Deubiquitylating Enzymes. Chemistry and Biology, 2011, 18, 1401-1412.	6.2	348
150	Comparative evaluation of labelâ€free SINQ normalized spectral index quantitation in the central proteomics facilities pipeline. Proteomics, 2011, 11, 2790-2797.	1.3	120
151	A Photoreactive Small-Molecule Probe for 2-Oxoglutarate Oxygenases. Chemistry and Biology, 2011, 18, 642-654.	6.2	46
152	Crystal structures of the endoplasmic reticulum aminopeptidase-1 (ERAP1) reveal the molecular basis for N-terminal peptide trimming. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 7745-7750.	3.3	216
153	Quantitative Mass Spectrometry Reveals Dynamics of Factor-inhibiting Hypoxia-inducible Factor-catalyzed Hydroxylation*. Journal of Biological Chemistry, 2011, 286, 33784-33794.	1.6	22
154	Detection of multiple autoantibodies in patients with ankylosing spondylitis using nucleic acid programmable protein arrays. Molecular and Cellular Proteomics, 2010, , .	2.5	7
155	Phylogeny by a polyphasic approach of the order Caulobacterales, proposal of Caulobacter mirabilis sp. nov., Phenylobacterium haematophilum sp. nov. and Phenylobacterium conjunctum sp. nov., and emendation of the genus Phenylobacterium. International Journal of Systematic and Evolutionary Microbiology. 2008, 58, 1939-1949.	0.8	57