Elke Hammer

List of Publications by Year in descending order

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141 4,272 37 56
papers citations h-index g-index

147 147 147 5983

times ranked

citing authors

docs citations

all docs

#	Article	IF	CITATIONS
1	Novel Activities of Glycolytic Enzymes in Bacillus subtilis. Molecular and Cellular Proteomics, 2009, 8, 1350-1360.	3.8	221
2	Characterization of the human neutrophil alloantigen-3a. Nature Medicine, 2010, 16, 45-48.	30.7	143
3	BLOOD COMPONENTS: A novel approach to pathogen reduction in platelet concentrates using shortâ€wave ultraviolet light. Transfusion, 2009, 49, 2612-2624.	1.6	138
4	Associations of circulating plasma microRNAs with age, body mass index and sex in a population-based study. BMC Medical Genomics, 2015, 8, 61.	1.5	133
5	Profiling of alterations in platelet proteins during storage of platelet concentrates. Transfusion, 2007, 47, 1221-1233.	1.6	103
6	Dehalogenation of Chlorinated Hydroxybiphenyls by Fungal Laccase. Applied and Environmental Microbiology, 2001, 67, 4377-4381.	3.1	100
7	Transformation of Triclosan by Trametes versicolor and Pycnoporus cinnabarinus. Applied and Environmental Microbiology, 2000, 66, 4157-4160.	3.1	96
8	Physical interactions between tricarboxylic acid cycle enzymes in Bacillus subtilis: Evidence for a metabolon. Metabolic Engineering, 2011, 13, 18-27.	7.0	94
9	Identification of periodontitis associated changes in the proteome of whole human saliva by mass spectrometric analysis. Journal of Clinical Periodontology, 2013, 40, 825-832.	4.9	88
10	Comparative evaluation of saliva collection methods for proteome analysis. Clinica Chimica Acta, 2013, 419, 42-46.	1.1	85
11	Genomewide metaâ€analysis identifies loci associated with <scp>IGF</scp> â€l and <scp>IGFBP</scp> â€3 levels with impact on ageâ€related traits. Aging Cell, 2016, 15, 811-824.	6.7	83
12	Reduced Degradation of the Chemokine MCP-3 by Matrix Metalloproteinase-2 Exacerbates Myocardial Inflammation in Experimental Viral Cardiomyopathy. Circulation, 2011, 124, 2082-2093.	1.6	81
13	Synthesis of 3-(3,4-dihydroxyphenyl)-propionic acid derivatives by N-coupling of amines using laccase. Tetrahedron, 2002, 58, 7589-7593.	1.9	80
14	Rapid Modulation of the Organic Anion Transporting Polypeptide 2B1 (OATP2B1, SLCO2B1) Function by Protein Kinase C-mediated Internalization. Journal of Biological Chemistry, 2010, 285, 11336-11347.	3.4	75
15	Identification, Characterization, and Structure Analysis of the Cyclic di-AMP-binding PII-like Signal Transduction Protein DarA. Journal of Biological Chemistry, 2015, 290, 3069-3080.	3.4	69
16	Characterization of the Human Myocardial Proteome in Inflammatory Dilated Cardiomyopathy by Label-free Quantitative Shotgun Proteomics of Heart Biopsies. Journal of Proteome Research, 2011, 10, 2161-2171.	3.7	66
17	Myocardial gene expression profiles and cardiodepressant autoantibodies predict response of patients with dilated cardiomyopathy to immunoadsorption therapy. European Heart Journal, 2013, 34, 666-675.	2.2	64
18	Novel Penicillins Synthesized by Biotransformation Using Laccase from Trametes spec Chemical and Pharmaceutical Bulletin, 2006, 54, 632-638.	1.3	60

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19	Cometabolic Degradation of Dibenzofuran by Biphenyl-Cultivated <i>Ralstonia</i> sp. Strain SBUG 290. Applied and Environmental Microbiology, 2000, 66, 4528-4531.	3.1	55
20	Monitoring of changes in the membrane proteome during stationary phase adaptation of <i>Bacillus subtilis</i> using <i>in vivo</i> labeling techniques. Proteomics, 2008, 8, 2062-2076.	2.2	55
21	Characterization of the EGFR interactome reveals associated protein complex networks and intracellular receptor dynamics. Proteomics, 2013, 13, 3131-3144.	2.2	54
22	Isolation and Characterization of a Dibenzofuran-Degrading Yeast: Identification of Oxidation and Ring Cleavage Products. Applied and Environmental Microbiology, 1998, 64, 2215-2219.	3.1	54
23	Novel Cephalosporins Synthesized by Amination of 2,5-Dihydroxybenzoic Acid Derivatives Using Fungal Laccases II. Chemical and Pharmaceutical Bulletin, 2007, 55, 412-416.	1.3	53
24	Laccase-catalyzed carbon–carbon bond formation: oxidative dimerization of salicylic esters by air in aqueous solution. Tetrahedron, 2005, 61, 4615-4619.	1.9	51
25	Laccase-induced cross-coupling of 4-aminobenzoic acid with para-dihydroxylated compounds 2,5-dihydroxy-N-(2-hydroxyethyl)-benzamide and 2,5-dihydroxybenzoic acid methyl ester. Journal of Molecular Catalysis B: Enzymatic, 2005, 35, 86-92.	1.8	51
26	HDAC (Histone Deacetylase) Inhibitor Valproic Acid Attenuates Atrial Remodeling and Delays the Onset of Atrial Fibrillation in Mice. Circulation: Arrhythmia and Electrophysiology, 2019, 12, e007071.	4.8	49
27	Biotransformation of Biphenyl by Paecilomyces lilacinus and Characterization of Ring Cleavage Products. Applied and Environmental Microbiology, 2001, 67, 1551-1557.	3.1	48
28	Synthesis of New N-Analogous Corollosporine Derivatives with Antibacterial Activity by Laccase-Catalyzed Amination. Chemical and Pharmaceutical Bulletin, 2008, 56, 781-786.	1.3	47
29	Transformation of 2-hydroxydibenzofuran by laccases of the white rot fungi Trametes versicolor and Pycnoporus cinnabarinus and characterization of oligomerization products. Biodegradation, 1997, 8, 321-327.	3.0	45
30	Timeâ€resolved quantitative proteome profiling of host–pathogen interactions: The response of <i>Staphylococcus aureus</i> RN1HG to internalisation by human airway epithelial cells. Proteomics, 2010, 10, 2801-2811.	2.2	45
31	Proteomic identification of potential prognostic biomarkers in resectable pancreatic ductal adenocarcinoma. Proteomics, 2014, 14, 945-955.	2.2	44
32	Proteomic analysis of doxorubicinâ€induced changes in the proteome of HepG2cells combining 2â€D DIGE and LCâ€MS/MS approaches. Proteomics, 2010, 10, 99-114.	2.2	43
33	Cohort profile: Greifswald approach to individualized medicine (GANI_MED). Journal of Translational Medicine, 2014, 12, 144.	4.4	43
34	A global Staphylococcus aureus proteome resource applied to the in vivo characterization of host-pathogen interactions. Scientific Reports, 2017, 7, 9718.	3.3	42
35	Identification of the Components Involved in Cyclic Di-AMP Signaling in Mycoplasma pneumoniae. Frontiers in Microbiology, 2017, 8, 1328.	3.5	42
36	Proteomic characterization of freezeâ€dried human plasma: providing treatment of bleeding disorders without the need for a cold chain. Transfusion, 2008, 48, 2356-2363.	1.6	41

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37	Oxidative ring cleavage of low chlorinated biphenyl derivatives by fungi leads to the formation of chlorinated lactone derivatives. Chemosphere, 2006, 64, 672-685.	8.2	39
38	Novel .BETALactam Antibiotics Synthesized by Amination of Catechols Using Fungal Laccase. Chemical and Pharmaceutical Bulletin, 2008, 56, 902-907.	1.3	39
39	The KupA and KupB Proteins of <i>Lactococcus lactis</i> IL1403 Are Novel c-di-AMP Receptor Proteins Responsible for Potassium Uptake. Journal of Bacteriology, 2019, 201, .	2.2	38
40	Characterisation of coupling products formed by biotransformation of biphenyl and diphenyl ether by the white rot fungus Pycnoporus cinnabarinus. Archives of Microbiology, 2000, 174, 393-398.	2.2	36
41	Evidence for synergistic control of glutamate biosynthesis by glutamate dehydrogenases and glutamate in <scp><i>B</i></scp> <i>acillus subtilisEnvironmental Microbiology, 2015, 17, 3379-3390.</i>	3.8	35
42	Title is missing!. Biodegradation, 1999, 10, 279-286.	3.0	33
43	Sex-specific metabolic and functional differences in human umbilical vein endothelial cells from twin pairs. Atherosclerosis, 2019, 291, 99-106.	0.8	31
44	Synthesis of Imidazol-2-yl Amino Acids by Using Cells from Alkane-Oxidizing Bacteria. Applied and Environmental Microbiology, 2003, 69, 1670-1679.	3.1	30
45	Degradation of phenylalkanes and characterization of aromatic intermediates acting as growth inhibiting substances in hydrocarbon utilizing yeast Candida maltosa. International Biodeterioration and Biodegradation, 2008, 62, 408-414.	3.9	30
46	CcpA forms complexes with CodY and RpoA in <i>Bacillusâ€f subtilis</i> . FEBS Journal, 2012, 279, 2201-2214.	4.7	30
47	Plasma proteome and metabolome characterization of an experimental human thyrotoxicosis model. BMC Medicine, 2017, 15, 6.	5.5	30
48	Fungal hydroxylation of dibenzofuran. Mycological Research, 1997, 101, 433-436.	2.5	29
49	Effect of selected environmental factors on degradation and mineralization of biaryl compounds by the bacterium Ralstonia pickettii in soil and compost. Chemosphere, 1998, 36, 2321-2335.	8.2	29
50	Decolorization of synthetic dyes by the deuteromycetePestalotiopsis guepinii CLPS no. 786 strain. Journal of Basic Microbiology, 2006, 46, 28-33.	3.3	28
51	Cofilin Oligomer Formation Occurs In Vivo and Is Regulated by Cofilin Phosphorylation. PLoS ONE, 2013, 8, e71769.	2.5	26
52	Hydroxylation of biphenyl by the yeast Trichosporon mucoides. Archives of Microbiology, 2000, 174, 353-361.	2.2	25
53	Novel Ring Cleavage Products in the Biotransformation of Biphenyl by the Yeast Trichosporon mucoides. Applied and Environmental Microbiology, 2001, 67, 4158-4165.	3.1	25
54	Carbon-oxygen bond formation by fungal laccases: cross-coupling of 2,5-dihydroxy-N-(2-hydroxyethyl)-benzamide with the solvents water, methanol, and other alcohols. Applied Microbiology and Biotechnology, 2007, 76, 407-416.	3.6	25

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55	Comparative analyses of laccaseâ€eatalyzed amination reactions for production of novel βâ€lactam antibiotics. Biotechnology and Applied Biochemistry, 2012, 59, 295-306.	3.1	25
56	A proteomics workflow for quantitative and time-resolved analysis of adaptation reactions of internalized bacteria. Methods, 2013, 61, 244-250.	3.8	25
57	Molecular and proteome analyses highlight the importance of the Cpx envelope stress system for acid stress and cell wall stability in <i>Escherichia coli</i> i>NicrobiologyOpen, 2016, 5, 582-596.	3.0	25
58	Isolation and characterization of biarylic structure-degrading yeasts: hydroxylation potential of dibenzofuran. Environmental Pollution, 2002, 118, 379-382.	7.5	24
59	In-depth proteomic analysis of the human cerumen—A potential novel diagnostically relevant biofluid. Journal of Proteomics, 2013, 83, 119-129.	2.4	24
60	Proteome Analysis Reveals New Mechanisms of Bcl11b-loss Driven Apoptosis. Journal of Proteome Research, 2010, 9, 3799-3811.	3.7	23
61	The mitochondrial respiratory chain has a critical role in the antiviral process in Coxsackievirus B3-induced myocarditis. Laboratory Investigation, 2012, 92, 125-134.	3.7	23
62	Effect of Experimental Thyrotoxicosis onto Blood Coagulation: A Proteomics Study. European Thyroid Journal, 2015, 4, 119-124.	2.4	23
63	Protective effects of endothelin receptor A and B inhibitors against doxorubicin-induced cardiomyopathy. Biochemical Pharmacology, 2015, 94, 109-129.	4.4	23
64	Changes of myocardial gene expression and protein composition in patients with dilated cardiomyopathy after immunoadsorption with subsequent immunoglobulin substitution. Basic Research in Cardiology, 2016, 111, 53.	5.9	23
65	Selection of autochthonous yeast strains able to degrade biphenyl. World Journal of Microbiology and Biotechnology, 2001, 17, 591-594.	3.6	22
66	OMICS-based exploration of the molecular phenotype of resident cardiac progenitor cells from adult murine heart. Journal of Proteomics, 2012, 75, 5304-5315.	2.4	22
67	Utilising the EGFR interactome to identify mechanisms of drug resistance in non-small cell lung cancer â€" Proof of concept towards a systems pharmacology approach. European Journal of Pharmaceutical Sciences, 2016, 94, 20-32.	4.0	22
68	Sex-specific differences in the intracellular proteome of human endothelial cells from dizygotic twins. Journal of Proteomics, 2019, 201, 48-56.	2.4	22
69	Oxidation and ring cleavage of dibenzofuran by the filamentous fungus Paecilomyces lilacinus. Archives of Microbiology, 2004, 182, 51-59.	2.2	21
70	Production of aromatic compounds during methanogenic degradation of straw in rice field soil. FEMS Microbiology Ecology, 2005, 52, 43-48.	2.7	21
71	Biotransformation of biphenyl by the filamentous fungus Talaromyces helicus. World Journal of Microbiology and Biotechnology, 2005, 21, 101-106.	3.6	21
72	C–N coupling of 3-methylcatechol with primary amines using native and recombinant laccases from Trametes versicolor and Pycnoporus cinnabarinus. Tetrahedron, 2011, 67, 9311-9321.	1.9	21

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73	Virusâ€induced dilated cardiomyopathy is characterized by increased levels of fibrotic extracellular matrix proteins and reduced amounts of energyâ€producing enzymes. Proteomics, 2011, 11, 4310-4320.	2.2	21
74	Targeted synthesis of novel \hat{l}^2 -lactam antibiotics by laccase-catalyzed reaction of aromatic substrates selected by pre-testing for their antimicrobial and cytotoxic activity. Applied Microbiology and Biotechnology, 2016, 100, 4885-4899.	3.6	21
75	Impact of blood sample collection methods on blood protein profiling studies. Clinica Chimica Acta, 2017, 471, 128-134.	1.1	21
76	A novel assay to assess the effect of pharmaceutical compounds on the differentiation of podocytes. British Journal of Pharmacology, 2017, 174, 163-176.	5.4	21
77	A Novel Artificial MicroRNA Expressing AAV Vector for Phospholamban Silencing in Cardiomyocytes Improves Ca2+ Uptake into the Sarcoplasmic Reticulum. PLoS ONE, 2014, 9, e92188.	2.5	19
78	Characterization of the Genetic Program Linked to the Development of Atrial Fibrillation in CREM-Ibl°C-X Mice. Circulation: Arrhythmia and Electrophysiology, 2017, 10, .	4.8	19
79	Biotransformation of Biarylic Compounds by Yeasts of the Genus Trichosporon. Systematic and Applied Microbiology, 2002, 25, 332-339.	2.8	18
80	Kidney protein profiling of Wilms' tumor patients by analysis of formalin-fixed paraffin-embedded tissue samples. Clinica Chimica Acta, 2014, 433, 235-241.	1.1	18
81	Correlation of gene expression and clinical parameters identifies a set of genes reflecting LV systolic dysfunction and morphological alterations. Physiological Genomics, 2019, 51, 356-367.	2.3	18
82	Cross-Sectional Association of Salivary Proteins with Age, Sex, Body Mass Index, Smoking, and Education. Journal of Proteome Research, 2017, 16, 2273-2281.	3.7	17
83	Endomyocardial proteomic signature corresponding to the response of patients with dilated cardiomyopathy to immunoadsorption therapy. Journal of Proteomics, 2017, 150, 121-129.	2.4	17
84	Using a Label Free Quantitative Proteomics Approach to Identify Changes in Protein Abundance in Multidrug-Resistant Mycobacterium tuberculosis. Indian Journal of Microbiology, 2015, 55, 219-230.	2.7	16
85	Biotransformation and reduction of estrogenicity of bisphenol A by the biphenyl-degrading Cupriavidus basilensis. Applied Microbiology and Biotechnology, 2017, 101, 3743-3758.	3.6	16
86	The role of the two-component systems Cpx and Arc in protein alterations upon gentamicin treatment in Escherichia coli. BMC Microbiology, 2017, 17, 197.	3. 3	16
87	Dynamic adaptation of myocardial proteome during heart failure development. PLoS ONE, 2017, 12, e0185915.	2.5	16
88	Laccase-catalyzed cross-linking of amino acids and peptides with dihydroxylated aromatic compounds. Amino Acids, 2010, 39, 671-683.	2.7	15
89	Viral myocarditis induced by Coxsackievirus B3 in <i>A.BY</i> / <i>SnJ</i> mice: Analysis of changes in the myocardial proteome. Proteomics, 2010, 10, 1802-1818.	2.2	15
90	Comparative evaluation of peptide desalting methods for salivary proteome analysis. Clinica Chimica Acta, 2014, 434, 16-20.	1.1	15

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91	The N-Terminal CCHC Zinc Finger Motif Mediates Homodimerization of Transcription Factor BCL11B. Molecular and Cellular Biology, 2018, 38, .	2.3	15
92	Global secretome analysis of resident cardiac progenitor cells from wildâ€type and transgenic heart failure mice: Why ambience matters. Journal of Cellular Physiology, 2019, 234, 10111-10122.	4.1	15
93	Adenine Nucleotide Translocase 1 Expression Is Coupled to the HSP27-Mediated TLR4 Signaling in Cardiomyocytes. Cells, 2019, 8, 1588.	4.1	14
94	Fibronectin modulates formation of PF4/heparin complexes and is a potential factor for reducing risk of developing HIT. Blood, 2019, 133, 978-989.	1.4	14
95	Angiotensin II-dependent hypertension causes reversible changes in the platelet proteome. Journal of Hypertension, 2011, 29, 2126-2137.	0.5	13
96	Absolute quantification of the Kdp subunits of <i>Escherichia coli</i> by multiple reaction monitoring. Proteomics, 2014, 14, 1630-1638.	2.2	13
97	Fumarate dependent protein composition under aerobic and anaerobic growth conditions in Escherichia coli. Journal of Proteomics, 2020, 212, 103583.	2.4	13
98	Brain Derived Neurotrophic Factor Contributes to the Cardiogenic Potential of Adult Resident Progenitor Cells in Failing Murine Heart. PLoS ONE, 2015, 10, e0120360.	2.5	12
99	Several adaptor proteins promote intracellular localisation of the transporter MRP4/ABCC4 in platelets and haematopoietic cells. Thrombosis and Haemostasis, 2017, 117, 105-115.	3.4	12
100	Mono-allelic expression of the IGF-I receptor does not affect IGF responses in human fibroblasts. European Journal of Endocrinology, 2004, 151, 521-529.	3.7	11
101	Proteome analysis of heart biopsies using a TRIzol-based protein extraction. Clinica Chimica Acta, 2015, 438, 246-247.	1.1	11
102	Circulating proteomic patterns in AF related left atrial remodeling indicate involvement of coagulation and complement cascade. PLoS ONE, 2018, 13, e0198461.	2.5	11
103	Low Dose Proteasome Inhibition Affects Alternative Splicing. Journal of Proteome Research, 2012, 11, 3947-3954.	3.7	10
104	Oxidation of triphenylarsine to triphenylarsineoxide by Trichoderma harzianum and other fungi. Chemosphere, 2001, 44, 697-700.	8.2	9
105	Proteomic analyses of age related changes in A.BY/SnJ mouse hearts. Proteome Science, 2013, 11, 29.	1.7	9
106	Plasma protein profiling of patients with intraductal papillary mucinous neoplasm of the pancreas as potential precursor lesions of pancreatic cancer. Clinica Chimica Acta, 2018, 477, 127-134.	1.1	9
107	Anaerobic formation and degradation of toxic aromatic compounds in agricultural and communal sewage deposits. Chemosphere, 1999, 38, 2561-2568.	8.2	8
108	A novel engineering tool in the Bacillus subtilis toolbox: inducer-free activation of gene expression by selection-driven promoter decryptification. Microbiology (United Kingdom), 2015, 161, 354-361.	1.8	8

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109	Cytokine-Mediated Alterations of Human Cardiac Fibroblast's Secretome. International Journal of Molecular Sciences, 2021, 22, 12262.	4.1	8
110	Direct mass spectrometric identification of ABCB1 (P-glycoprotein/MDR1) from the apical membrane fraction of human placenta using fourier transform ion cyclotron mass spectrometry. Pharmacogenetics and Genomics, 2006, 16 , 385 - 389 .	1.5	7
111	Antineoplastic agent busulfan regulates a network of genes related to coagulation and fibrinolysis. European Journal of Clinical Pharmacology, 2012, 68, 923-935.	1.9	7
112	Novel insights into the fungal oxidation of monoaromatic and biarylic environmental pollutants by characterization of two new ring cleavage enzymes. Applied Microbiology and Biotechnology, 2013, 97, 5043-5053.	3.6	7
113	Mass spectrometric phosphoproteome analysis of small-sized samples of human neutrophils. Clinica Chimica Acta, 2015, 451, 199-207.	1.1	7
114	Cellular Concentrations of the Transporters DctA and DcuB and the Sensor DcuS of Escherichia coli and the Contributions of Free and Complexed DcuS to Transcriptional Regulation by DcuR. Journal of Bacteriology, 2018, 200, .	2.2	7
115	Chronic \hat{l}^2 -adrenergic stimulation reverses depressed Ca handling in mice overexpressing inhibitor-2 of protein phosphatase 1. Journal of Molecular and Cellular Cardiology, 2018, 125, 195-204.	1.9	7
116	Impact of Storage Conditions on the Breast Milk Peptidome. Nutrients, 2020, 12, 2733.	4.1	7
117	Nup133 and ERα mediate the differential effects of hyperoxia-induced damage in male and female OPCs. Molecular and Cellular Pediatrics, 2020, 7, 10.	1.8	7
118	Purification and biochemical characterization of a lysosomal \hat{l}_{\pm} -fucosidase from the deuterostomia Asterias rubens. Biochimie, 2012, 94, 1199-1205.	2.6	6
119	Proteomic profile of platelets during reconstitution of platelet counts after apheresis. Proteomics - Clinical Applications, 2016, 10, 831-838.	1.6	6
120	From Proteomics to Personalized Medicine: The Importance of Isoflavone Dose and Estrogen Receptor Status in Breast Cancer Cells. Journal of Personalized Medicine, 2020, 10, 292.	2.5	6
121	Enhanced excretion of intermediates of aromatic amino acid catabolism during chlorophenol degradation due to nutrient limitation in the yeastCandida maltosa. Journal of Basic Microbiology, 1996, 36, 239-243.	3.3	5
122	The Other Side of the RAAS: Aldosterone Improves Migration of Cardiac Progenitor Cells. Journal of Cellular Physiology, 2015, 230, 2829-2836.	4.1	4
123	Data on the impact of the blood sample collection methods on blood protein profiling studies. Data in Brief, 2017, 14, 313-319.	1.0	4
124	Fetal Zone Steroids and Estrogen Show Sex Specific Effects on Oligodendrocyte Precursor Cells in Response to Oxidative Damage. International Journal of Molecular Sciences, 2021, 22, 6586.	4.1	4
125	Laccase-catalyzed derivatization of 6-aminopenicillanic, 7-aminocephalosporanic and 7-aminodesacetoxycephalosporanic acid. AMB Express, 2020, 10, 177.	3.0	4
126	Differences in Cell-Intrinsic Inflammatory Programs of Yolk Sac and Bone Marrow Macrophages. Cells, 2021, 10, 3564.	4.1	4

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127	2D DIGE proteomic analysis of multidrug resistant and susceptible clinical Mycobacterium tuberculosis isolates. Journal of Integrated OMICS, 2014, 4, .	0.5	3
128	Pathophysiological aldosterone levels modify the secretory activity of cardiac progenitor cells. Molecular and Cellular Endocrinology, 2017, 439, 16-25.	3.2	3
129	Global plasma protein profiling reveals DCM characteristic protein signatures. Journal of Proteomics, 2019, 209, 103508.	2.4	3
130	Specific domain ν reduction of beta-2-glycoprotein I induces protein flexibility and alters pathogenic antibody binding. Scientific Reports, 2021, 11, 4542.	3.3	3
131	Riociguat attenuates the changes in left ventricular proteome and microRNA profile after experimental aortic stenosis in mice. British Journal of Pharmacology, 2022, 179, 4575-4592.	5.4	3
132	Plasma protein absolute quantification by nano-LC Q-TOF UDMSE for clinical biomarker verification. Medicine and Pharmacy Reports, 2017, 90, 425-430.	0.4	2
133	Serum starvation induces sexual dimorphisms in secreted proteins of human umbilical vein endothelial cells (HUVECs) from twin pairs. Proteomics, 2022, 22, e2100168.	2.2	2
134	Tight Complex Formation of the Fumarate Sensing DcuS-DcuR Two-Component System at the Membrane and Target Promoter Search by Free DcuR Diffusion. MSphere, 2022, 7, .	2.9	2
135	Deficiency in FTSJ1 Affects Neuronal Plasticity in the Hippocampal Formation of Mice. Biology, 2022, 11, 1011.	2.8	2
136	Characterization of the Human Myocardial Proteome in Dilated Cardiomyopathy by Label-Free Quantitative Shotgun Proteomics of Heart Biopsies. Methods in Molecular Biology, 2013, 1005, 67-76.	0.9	1
137	Analysis of DCM associated protein alterations of human right and left ventricles. Journal of Proteomics, 2021, 231, 104018.	2.4	1
138	Improved reconstitution of Trizol derived protein extracts provides high quality samples for comprehensive proteomic characterization of cell cultures. Journal of Integrated OMICS, 2015, 5, .	0.5	0
139	Oxidation of the Fungicide Biphenyl by Yeasts of the Genus Trichosporon. , 2003, , 293-296.		0
140	Comparative immunoproteome analysis of the response of susceptible A.BY/SnJ and resistant C57BL/6 mice to Coxsackievirus B3-infection. Journal of Integrated OMICS, 2012, 2, .	0.5	0
141	Fetal Zone Steroids Show Discrete Effects on Hyperoxia-Induced Attenuation of Migration in Cultured Oligodendrocyte Progenitor Cells. Oxidative Medicine and Cellular Longevity, 2022, 2022, 1-19.	4.0	0