

Jong-Moon Park

List of Publications by Year in descending order

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Version: 2024-02-01

20
papers

306
citations

932766

10
h-index

887659

17
g-index

20
all docs

20
docs citations

20
times ranked

540
citing authors

#	ARTICLE	IF	CITATIONS
1	Review of Three-Dimensional Liquid Chromatography Platforms for Bottom-Up Proteomics. <i>International Journal of Molecular Sciences</i> , 2020, 21, 1524.	1.8	50
2	Proteomic Analysis of Serum from Patients with Major Depressive Disorder to Compare Their Depressive and Remission Statuses. <i>Psychiatry Investigation</i> , 2015, 12, 249.	0.7	44
3	Integrated analysis of global proteome, phosphoproteome and glycoproteome enables complementary interpretation of disease-related protein networks. <i>Scientific Reports</i> , 2015, 5, 18189.	1.6	34
4	Proteomics in Forensic Analysis: Applications for Human Samples. <i>Applied Sciences (Switzerland)</i> , 2021, 11, 3393.	1.3	21
5	Dihydropyruvate dehydrogenase as a potential UVB target in skin epidermis; using an integrated approach of label-free quantitative proteomics and targeted metabolite analysis. <i>Journal of Proteomics</i> , 2015, 117, 70-85.	1.2	20
6	Effects of FGF2-secreting adipose-derived stem cells in thioacetamide-induced hepatic fibrosis. <i>Journal of Cellular and Molecular Medicine</i> , 2018, 22, 5165-5169.	1.6	19
7	Neural Ganglia Transcriptome and Peptidome Associated with Sexual Maturation in Female Pacific Abalone (<i>Haliotis discus hannai</i>). <i>Genes</i> , 2019, 10, 268.	1.0	18
8	Establishment of functional epithelial organoids from human lacrimal glands. <i>Stem Cell Research and Therapy</i> , 2021, 12, 247.	2.4	18
9	UBA2 activates Wnt/ β -catenin signaling pathway during protection of R28 retinal precursor cells from hypoxia by extracellular vesicles derived from placental mesenchymal stem cells. <i>Stem Cell Research and Therapy</i> , 2020, 11, 428.	2.4	12
10	De novo transcriptome profile of coccolithophorid alga <i>Emiliana huxleyi</i> CCMP371 at different calcium concentrations with proteome analysis. <i>PLoS ONE</i> , 2019, 14, e0221938.	1.1	11
11	Development of an automated high-throughput sample preparation protocol for LC-MS/MS analysis of glycosylated peptides. <i>Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences</i> , 2018, 1092, 88-94.	1.2	10
12	Enrichment and analysis of glycosylated proteins. <i>Reviews in Analytical Chemistry</i> , 2022, 41, 83-97.	1.5	10
13	Label-free quantitative proteomic analysis of human periodontal ligament stem cells by high-resolution mass spectrometry. <i>Journal of Periodontal Research</i> , 2019, 54, 53-62.	1.4	9
14	An automated high-throughput sample preparation method using double-filtration for serum metabolite LC-MS analysis. <i>Analytical Methods</i> , 2019, 11, 4060-4065.	1.3	8
15	Proteomic Profiling of <i>Emiliana huxleyi</i> Using a Three-Dimensional Separation Method Combined with Tandem Mass Spectrometry. <i>Molecules</i> , 2020, 25, 3028.	1.7	5
16	Proteomic Analysis of the Vitreous Body in Proliferative and Non-Proliferative Diabetic Retinopathy. <i>Current Proteomics</i> , 2021, 18, 143-152.	0.1	5
17	Proteomic and Metabolomic Analyses of Maggots in Porcine Corpses for Post-Mortem Interval Estimation. <i>Applied Sciences (Switzerland)</i> , 2021, 11, 7885.	1.3	5
18	Comparative Proteomic Profiling of Marine and Freshwater <i>Synechocystis</i> Strains Using Liquid Chromatography-Tandem Mass Spectrometry. <i>Journal of Marine Science and Engineering</i> , 2020, 8, 790.	1.2	4

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19	Discovery of Post-Translational Modifications in <i>Emiliana huxleyi</i> . <i>Molecules</i> , 2021, 26, 2027.	1.7	2
20	A Quantitative Proteomic Analysis to Reveal Effects of N-acetylcysteine on H ₂ O ₂ -induced Cytotoxicity. <i>Current Proteomics</i> , 2021, 18, 403-414.	0.1	1