

# Jong-Moon Park

## List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

19  
papers

173  
citations

7  
h-index

12  
g-index

20  
ext. papers

245  
ext. citations

3.9  
avg, IF

3.28  
L-index

#	Paper	IF	Citations
19	Proteomic analysis of serum from patients with major depressive disorder to compare their depressive and remission statuses. <i>Psychiatry Investigation</i> , <b>2015</b> , 12, 249-59	3.1	33
18	Review of Three-Dimensional Liquid Chromatography Platforms for Bottom-Up Proteomics. <i>International Journal of Molecular Sciences</i> , <b>2020</b> , 21,	6.3	31
17	Integrated analysis of global proteome, phosphoproteome, and glycoproteome enables complementary interpretation of disease-related protein networks. <i>Scientific Reports</i> , <b>2015</b> , 5, 18189	4.9	23
16	Dihydrolipoyl 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> , <b>2015</b> , 117, 70-85	3.9	15
15	Neural Ganglia Transcriptome and Peptidome Associated with Sexual Maturation in Female Pacific Abalone (). <i>Genes</i> , <b>2019</b> , 10,	4.2	11
14	Effects of FGF21-secreting adipose-derived stem cells in thioacetamide-induced hepatic fibrosis. <i>Journal of Cellular and Molecular Medicine</i> , <b>2018</b> , 22, 5165-5169	5.6	11
13	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> , <b>2018</b> , 1092, 88-94	3.2	8
12	De novo transcriptome profile of coccolithophorid alga <i>Emiliania huxleyi</i> CCMP371 at different calcium concentrations with proteome analysis. <i>PLoS ONE</i> , <b>2019</b> , 14, e0221938	3.7	7
11	An automated high-throughput sample preparation method using double-filtration for serum metabolite LC-MS analysis. <i>Analytical Methods</i> , <b>2019</b> , 11, 4060-4065	3.2	6
10	UBA2 activates Wnt/ $\beta$ -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> , <b>2020</b> , 11, 428	8.3	6
9	Label-free quantitative proteomic analysis of human periodontal ligament stem cells by high-resolution mass spectrometry. <i>Journal of Periodontal Research</i> , <b>2019</b> , 54, 53-62	4.3	5
8	Proteomic Profiling of Using a Three-Dimensional Separation Method Combined with Tandem Mass Spectrometry. <i>Molecules</i> , <b>2020</b> , 25,	4.8	3
7	Establishment of functional epithelial organoids from human lacrimal glands. <i>Stem Cell Research and Therapy</i> , <b>2021</b> , 12, 247	8.3	3
6	Proteomics in Forensic Analysis: Applications for Human Samples. <i>Applied Sciences (Switzerland)</i> , <b>2021</b> , 11, 3393	2.6	3
5	Proteomic Analysis of the Vitreous Body in Proliferative and Non-Proliferative Diabetic Retinopathy. <i>Current Proteomics</i> , <b>2021</b> , 18, 143-152	0.7	3
4	Comparative Proteomic Profiling of Marine and Freshwater Synechocystis Strains Using Liquid Chromatography-Tandem Mass Spectrometry. <i>Journal of Marine Science and Engineering</i> , <b>2020</b> , 8, 790	2.4	2
3	Enrichment and analysis of glycosylated proteins. <i>Reviews in Analytical Chemistry</i> , <b>2022</b> , 41, 83-97	2.3	1

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|---|---|-----|---|
| 2 | A Quantitative Proteomic Analysis to Reveal Effects of N-acetylcysteine on H <sub>2</sub> O <sub>2</sub> -induced Cytotoxicity. <i>Current Proteomics</i> , <b>2021</b> , 18, 403-414 | 0.7 | 1 |
| 1 | Proteomic and Metabolomic Analyses of Maggots in Porcine Corpses for Post-Mortem Interval Estimation. <i>Applied Sciences (Switzerland)</i> , <b>2021</b> , 11, 7885                  | 2.6 |   |