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## List of Publications by Year in descending order

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1170033 939365 32 454 9 18 citations h-index g-index papers 36 36 36 658 citing authors all docs docs citations times ranked

#	Article	IF	Citations
1	A comprehensive network map of IL-17A signaling pathway. Journal of Cell Communication and Signaling, 2023, 17, 209-215.	1.8	10
2	A network map of apelin-mediated signaling. Journal of Cell Communication and Signaling, 2022, 16, 137-143.	1.8	9
3	A multi-cellular molecular signaling and functional network map of C–C motif chemokine ligand 18 (CCL18): a chemokine with immunosuppressive and pro-tumor functions. Journal of Cell Communication and Signaling, 2022, 16, 293-300.	1.8	8
4	The network map of Elabela signaling pathway in physiological and pathological conditions. Journal of Cell Communication and Signaling, 2022, 16, 145-154.	1.8	8
5	A modular map of Bradykinin-mediated inflammatory signaling network. Journal of Cell Communication and Signaling, 2022, 16, 301-310.	1.8	14
6	Temporal Quantitative Phosphoproteomics Profiling of Interleukin-33 Signaling Network Reveals Unique Modulators of Monocyte Activation. Cells, 2022, 11, 138.	1.8	4
7	The network map of urotensin-II mediated signaling pathway in physiological and pathological conditions. Journal of Cell Communication and Signaling, 2022, 16, 601-608.	1.8	2
8	Dissecting <i>Plasmodium yoelii</i> Pathobiology: Proteomic Approaches for Decoding Novel Translational and Post-Translational Modifications. ACS Omega, 2022, 7, 8246-8257.	1.6	2
9	<i>Leptospira</i> and Leptospirosis: New Systems Science Insights on Proteome, Posttranslational Modifications, and Pathogen-Host Interaction. OMICS A Journal of Integrative Biology, 2022, 26, 280-289.	1.0	2
10	Competitive interaction of thymol with cviR inhibits quorum sensing and associated biofilm formation in Chromobacterium violaceum. International Microbiology, 2022, 25, 629-638.	1.1	5
11	Unique Posttranslational Modification Sites of Acetylation, Citrullination, Glutarylation, and Phosphorylation Are Found to Be Specific to the Proteins Partitioned in the Triton X-114 Fractions of <i>Leptospira</i> . ACS Omega, 2022, 7, 18569-18576.	1.6	1
12	A comprehensive review on current understanding of bradykinin in COVID-19 and inflammatory diseases. Molecular Biology Reports, 2022, 49, 9915-9927.	1.0	13
13	Revisiting Regulated Cell Death Responses in Viral Infections. International Journal of Molecular Sciences, 2022, 23, 7023.	1.8	11
14	A complete map of the Calcium/calmodulin-dependent protein kinase kinase 2 (CAMKK2) signaling pathway. Journal of Cell Communication and Signaling, 2021, 15, 283-290.	1.8	25
15	A pathway map of AXL receptor-mediated signaling network. Journal of Cell Communication and Signaling, 2021, 15, 143-148.	1.8	13
16	A network map of endothelin mediated signaling pathway. Journal of Cell Communication and Signaling, 2021, 15, 277-282.	1.8	15
17	The unique molecular targets associated antioxidant and antifibrotic activity of curcumin in in vitro model of acute lung injury: A proteomic approach. BioFactors, 2021, 47, 627-644.	2.6	3
18	SARS-CoV-2 signaling pathway map: A functional landscape of molecular mechanisms in COVID-19. Journal of Cell Communication and Signaling, 2021, 15, 601-608.	1.8	15

#	Article	IF	CITATIONS
19	Extracellular Proteome Analysis Shows the Abundance of Histidine Kinase Sensor Protein, DNA Helicase, Putative Lipoprotein Containing Peptidase M75 Domain and Peptidase C39 Domain Protein in Leptospira interrogans Grown in EMJH Medium. Pathogens, 2021, 10, 852.	1.2	4
20	Novel Post-Translational Modifications and Molecular Substrates in Glioma Identified by Bioinformatics. OMICS A Journal of Integrative Biology, 2021, 25, 463-473.	1.0	4
21	Broadening COVID-19 Interventions to Drug Innovation: Neprilysin Pathway as a Friend, Foe, or Promising Molecular Target?. OMICS A Journal of Integrative Biology, 2021, 25, 408-416.	1.0	5
22	Mapping Post-Translational Modifications in Brain Regions in Alzheimer's Disease Using Proteomics Data Mining. OMICS A Journal of Integrative Biology, 2021, 25, 525-536.	1.0	7
23	Unraveling Toxoplasma gondii GT1 Strain Virulence and New Protein-Coding Genes with Proteogenomic Analyses. OMICS A Journal of Integrative Biology, 2021, 25, 591-604.	1.0	6
24	COVID19 Disease Map, a computational knowledge repository of virus–host interaction mechanisms. Molecular Systems Biology, 2021, 17, e10387.	3.2	53
25	A comprehensive pathway map of IL-18-mediated signalling. Journal of Cell Communication and Signaling, 2020, 14, 257-266.	1.8	66
26	Triton Xâ€114 Fractionated Subcellular Proteome of <i>Leptospira interrogans</i> Shows Selective Enrichment of Pathogenic and Outer Membrane Proteins in the Detergent Fraction. Proteomics, 2020, 20, e2000170.	1.3	13
27	Proteogenomic examination of esophageal squamous cell carcinoma (ESCC): new lines of inquiry. Expert Review of Proteomics, 2020, 17, 649-662.	1.3	1
28	Proteomics Analysis Revealed the Importance of Inflammation-Mediated Downstream Pathways and the Protective Role of Curcumin in Bleomycin-Induced Pulmonary Fibrosis in C57BL/6 Mice. Journal of Proteome Research, 2020, 19, 2950-2963.	1.8	12
29	Rise of Clinical Microbial Proteogenomics: A Multiomics Approach to Nontuberculous Mycobacteriumâ€"The Case of <i>Mycobacterium abscessus</i> UC22. OMICS A Journal of Integrative Biology, 2019, 23, 1-16.	1.0	8
30	Omics data-driven analysis identifies laminin-integrin-mediated signaling pathway as a determinant for cell differentiation in oral squamous cell carcinoma. Indian Journal of Pathology and Microbiology, 2019, 62, 529.	0.1	3
31	A network map of IL-33 signaling pathway. Journal of Cell Communication and Signaling, 2018, 12, 615-624.	1.8	90
32	The Role of Nanotechnology in Targeting Carcinogens—A Comparison on Synthesis of Gold and Zinc Nanoparticles. Journal of Biomaterials and Tissue Engineering, 2014, 4, 591-599.	0.0	O