

Hyeyoon Kim

List of Publications by Year in descending order

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Version: 2024-02-01

16
papers

290
citations

1163117

8
h-index

940533

16
g-index

17
all docs

17
docs citations

17
times ranked

512
citing authors

#	ARTICLE	IF	CITATIONS
1	Alterations in blood proteins in the prodromal stage of bipolar II disorders. <i>Scientific Reports</i> , 2022, 12, 3174.	3.3	4
2	Discovery of Proteins Responsible for Resistance to Three Chemotherapy Drugs in Breast Cancer Cells Using Proteomics and Bioinformatics Analysis. <i>Molecules</i> , 2022, 27, 1762.	3.8	9
3	In-depth proteome of perilymph in guinea pig model. <i>Proteomics</i> , 2021, 21, 2000138.	2.2	5
4	Identification of altered protein expression in major depressive disorder and bipolar disorder patients using liquid chromatography-tandem mass spectrometry. <i>Psychiatry Research</i> , 2021, 299, 113850.	3.3	10
5	Quantitative Proteomics Reveals Knockdown of CD44 Promotes Proliferation and Migration in Claudin-Low MDA-MB-231 and Hs 578T Breast Cancer Cell Lines. <i>Journal of Proteome Research</i> , 2021, 20, 3720-3733.	3.7	4
6	Mass Spectrometry-Based Proteomic Discovery of Prognostic Biomarkers in Adrenal Cortical Carcinoma. <i>Cancers</i> , 2021, 13, 3890.	3.7	12
7	Predictive protein markers for depression severity in mood disorders: A preliminary trans-diagnostic approach study. <i>Journal of Psychiatric Research</i> , 2021, 142, 63-72.	3.1	5
8	Plasma proteomic data in bipolar II disorders and major depressive disorders. <i>Data in Brief</i> , 2021, 39, 107495.	1.0	1
9	In-depth proteomic profiling captures subtype-specific features of craniopharyngiomas. <i>Scientific Reports</i> , 2021, 11, 21206.	3.3	4
10	Bioinformatic analysis of proteomic data for iron, inflammation, and hypoxic pathways in restless legs syndrome. <i>Sleep Medicine</i> , 2020, 75, 448-455.	1.6	13
11	Comparison of serum protein profiles between major depressive disorder and bipolar disorder. <i>BMC Psychiatry</i> , 2020, 20, 145.	2.6	29
12	In-depth blood proteome profiling analysis revealed distinct functional characteristics of plasma proteins between severe and non-severe COVID-19 patients. <i>Scientific Reports</i> , 2020, 10, 22418.	3.3	80
13	An efficient method for high-pH peptide fractionation based on C18 StageTips for in-depth proteome profiling. <i>Analytical Methods</i> , 2019, 11, 4693-4698.	2.7	13
14	Quantitative Proteome Analysis of Brain Subregions and Spinal Cord from Experimental Autoimmune Encephalomyelitis Mice by TMT-Based Mass Spectrometry. <i>Proteomics</i> , 2019, 19, e1800355.	2.2	18
15	Comparative proteomic analysis of human malignant ascitic fluids for the development of gastric cancer biomarkers. <i>Clinical Biochemistry</i> , 2018, 56, 55-61.	1.9	17
16	Quantitative Proteomic Analysis Identifies AHNAK (Neuroblast Differentiation-associated Protein) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 Cytology. <i>Molecular and Cellular Proteomics</i> , 2018, 17, 1788-1802.	3.8	66