Maria Antonia O Caballol

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4564707/publications.pdf

Version: 2024-02-01

687363 794594 19 711 13 19 citations g-index h-index papers 19 19 19 1240 docs citations times ranked citing authors all docs

#	Article	lF	Citations
1	Multi-omics analysis reveals that co-exposure to phthalates and metals disturbs urea cycle and choline metabolism. Environmental Research, 2021, 192, 110041.	7.5	24
2	New Partners Identified by Mass Spectrometry Assay Reveal Functions of NCAM2 in Neural Cytoskeleton Organization. International Journal of Molecular Sciences, 2021, 22, 7404.	4.1	6
3	An Innovative Approach for The Integration of Proteomics and Metabolomics Data In Severe Septic Shock Patients Stratified for Mortality. Scientific Reports, 2018, 8, 6681.	3.3	28
4	Next-Generation Sequencing and Quantitative Proteomics of Hutchinson-Gilford progeria syndrome-derived cells point to a role of nucleotide metabolism in premature aging. PLoS ONE, 2018, 13, e0205878.	2.5	16
5	Proteolysis in septic shock patients: plasma peptidomic patterns are associated with mortality. British Journal of Anaesthesia, 2018, 121, 1065-1074.	3.4	37
6	Differential protein expression of hippocampal cells associated with heavy metals (Pb, As, and MeHg) neurotoxicity: Deepening into the molecular mechanism of neurodegenerative diseases. Journal of Proteomics, 2018, 187, 106-125.	2.4	38
7	A proteomic study of mesenchymal stem cells from equine umbilical cord. Theriogenology, 2017, 100, 8-15.	2.1	7
8	A multicentric study to evaluate the use of relative retention times in targeted proteomics. Journal of Proteomics, 2017, 152, 138-149.	2.4	9
9	Redox proteomic profiling of neuroketal-adducted proteins in human brain: Regional vulnerability at middle age increases in the elderly. Free Radical Biology and Medicine, 2016, 95, 1-15.	2.9	28
10	Multicenter experiment for quality control of peptide-centric LCâ€"MS/MS analysis â€" A longitudinal performance assessment with nLC coupled to orbitrap MS analyzers. Journal of Proteomics, 2015, 127, 264-274.	2.4	14
11	A Novel Phospholipase A2(D49) from the Venom of theCrotalus oreganus abyssus(North American) Tj ETQq1 1 C	0.784314	rgBT/Overloc
12	Isolation and Biochemical Characterization of a New Thrombin-Like Serine Protease from <i>Bothrops pirajai </i> Snake Venom. BioMed Research International, 2014, 2014, 1-13.	1.9	18
13	Proteomic analysis of outer membrane vesicles from the probiotic strain <i>Escherichia coli</i> Nissle 1917. Proteomics, 2014, 14, 222-229.	2.2	81
14	Insights into Maize LEA Proteins: From Proteomics to Functional Approaches. Plant and Cell Physiology, 2012, 53, 312-329.	3.1	76
15	Differential proteomic analysis of Acidithiobacillus ferrooxidans cells maintained in contact with bornite or chalcopyrite: Proteins involved with the early bacterial response. Process Biochemistry, 2011, 46, 770-776.	3.7	7
16	Acetylation of GAGA Factor Modulates Its Interaction with DNA. Biochemistry, 2010, 49, 9140-9151.	2,5	7
17	Modification of brain lipids but not phenotype in α-synucleinopathy transgenic mice by long-term dietary n-3 fatty acids. Neurochemistry International, 2010, 56, 318-328.	3.8	16
18	LCâ€MSMS identification of <i>Arabidopsis thaliana</i> heatâ€stable seed proteins: Enriching for LEAâ€type proteins by acid treatment. Journal of Mass Spectrometry, 2007, 42, 1485-1495.	1.6	37

#	÷	Article	IF	CITATIONS
1	9	Proteome, salicylic acid, and jasmonic acid changes in cucumber plants inoculated with <i>Trichoderma asperellum</i> strain T34. Proteomics, 2007, 7, 3943-3952.	2.2	252