

Florence Roux-Dalvai

List of Publications by Year in descending order

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Version: 2024-02-01

19
papers

725
citations

686830

13
h-index

839053

18
g-index

21
all docs

21
docs citations

21
times ranked

1262
citing authors

#	ARTICLE	IF	CITATIONS
1	Extensive Analysis of the Cytoplasmic Proteome of Human Erythrocytes Using the Peptide Ligand Library Technology and Advanced Mass Spectrometry. <i>Molecular and Cellular Proteomics</i> , 2008, 7, 2254-2269.	2.5	208
2	In-depth Exploration of Cerebrospinal Fluid by Combining Peptide Ligand Library Treatment and Label-free Protein Quantification. <i>Molecular and Cellular Proteomics</i> , 2010, 9, 1006-1021.	2.5	116
3	Neutrophil-Derived Exosomes: A New Mechanism Contributing to Airway Smooth Muscle Remodeling. <i>American Journal of Respiratory Cell and Molecular Biology</i> , 2016, 55, 450-461.	1.4	101
4	Extensive and Accurate Benchmarking of DIA Acquisition Methods and Software Tools Using a Complex Proteomic Standard. <i>Journal of Proteome Research</i> , 2021, 20, 4801-4814.	1.8	47
5	Fast and Accurate Bacterial Species Identification in Urine Specimens Using LC-MS/MS Mass Spectrometry and Machine Learning*. <i>Molecular and Cellular Proteomics</i> , 2019, 18, 2492-2505.	2.5	42
6	Portrait of blood-derived extracellular vesicles in patients with Parkinson's disease. <i>Neurobiology of Disease</i> , 2019, 124, 163-175.	2.1	33
7	Mitochondrial Inhibitory Factor 1 (IFI1) Is Present in Human Serum and Is Positively Correlated with HDL-Cholesterol. <i>PLoS ONE</i> , 2011, 6, e23949.	1.1	29
8	Oxidized DNA Precursors Cleanup by NUDT1 Contributes to Vascular Remodeling in Pulmonary Arterial Hypertension. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2021, 203, 614-627.	2.5	22
9	Mapping the Extracellular and Membrane Proteome Associated with the Vasculature and the Stroma in the Embryo. <i>Molecular and Cellular Proteomics</i> , 2013, 12, 2293-2312.	2.5	21
10	A metabolic labeling approach for glycoproteomic analysis reveals altered glycoprotein expression upon GALNT3 knockdown in ovarian cancer cells. <i>Journal of Proteomics</i> , 2016, 145, 91-102.	1.2	21
11	Interactions with plant pathogens influence lipopeptides production and antimicrobial activity of <i>Bacillus subtilis</i> strain PTB185. <i>Biological Control</i> , 2021, 154, 104497.	1.4	18
12	Mass Spectrometry-based Absolute Quantification of 20S Proteasome Status for Controlled Ex-vivo Expansion of Human Adipose-derived Mesenchymal Stromal/Stem Cells. <i>Molecular and Cellular Proteomics</i> , 2019, 18, 744-759.	2.5	17
13	Mass spectrometry-based proteomics identify novel serum osteoarthritis biomarkers. <i>Arthritis Research and Therapy</i> , 2022, 24, .	1.6	17
14	Vitamin C alters the amount of specific endoplasmic reticulum associated proteins involved in lipid metabolism in the liver of mice synthesizing a nonfunctional Werner syndrome (Wrn) mutant protein. <i>PLoS ONE</i> , 2018, 13, e0193170.	1.1	9
15	Proteomic dataset for altered glycoprotein expression upon GALNT3 knockdown in ovarian cancer cells. <i>Data in Brief</i> , 2016, 8, 342-349.	0.5	7
16	Accessing to the minor proteome of red blood cells through the influence of the nanoparticle surface properties on the corona composition. <i>International Journal of Nanomedicine</i> , 2015, 10, 1869.	3.3	6
17	DIA proteomics data from a UPS1-spiked E.coli protein mixture processed with six software tools. <i>Data in Brief</i> , 2022, 41, 107829.	0.5	5
18	Proteomic Analysis of Maternal Urine for the Early Detection of Preeclampsia and Fetal Growth Restriction. <i>Journal of Clinical Medicine</i> , 2021, 10, 4679.	1.0	3

#	ARTICLE	IF	CITATIONS
19	Abstract A42: Identification of glutamate and aspartate ADP-ribosylation sites onto histones by mass mass spectrometry. , 2017, , .		0