

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	DIA proteomics data from a UPS1-spiked E.coli protein mixture processed with six software tools. Data in Brief, 2022, 41, 107829.	0.5	5
2	Mass spectrometry-based proteomics identify novel serum osteoarthritis biomarkers. Arthritis Research and Therapy, 2022, 24, .	1.6	17
3	Oxidized DNA Precursors Cleanup by NUDT1 Contributes to Vascular Remodeling in Pulmonary Arterial Hypertension. American Journal of Respiratory and Critical Care Medicine, 2021, 203, 614-627.	2.5	22
4	Interactions with plant pathogens influence lipopeptides production and antimicrobial activity of Bacillus subtilis strain PTB185. Biological Control, 2021, 154, 104497.	1.4	18
5	Extensive and Accurate Benchmarking of DIA Acquisition Methods and Software Tools Using a Complex Proteomic Standard. Journal of Proteome Research, 2021, 20, 4801-4814.	1.8	47
6	Proteomic Analysis of Maternal Urine for the Early Detection of Preeclampsia and Fetal Growth Restriction. Journal of Clinical Medicine, 2021, 10, 4679.	1.0	3
7	Fast and Accurate Bacterial Species Identification in Urine Specimens Using LC-MS/MS Mass Spectrometry and Machine Learning*. Molecular and Cellular Proteomics, 2019, 18, 2492-2505.	2.5	42
8	Mass Spectrometry-based Absolute Quantification of 20S Proteasome Status for Controlled Ex-vivo Expansion of Human Adipose-derived Mesenchymal Stromal/Stem Cells. Molecular and Cellular Proteomics, 2019, 18, 744-759.	2.5	17
9	Portrait of blood-derived extracellular vesicles in patients with Parkinson's disease. Neurobiology of Disease, 2019, 124, 163-175.	2.1	33
10	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. PLoS ONE, 2018, 13, e0193170.	1.1	9
11	Abstract A42: Identification of glutamate and aspartate ADP-ribosylation sites onto histones by mass spectrometry. , 2017, , .		0
12	A metabolic labeling approach for glycoproteomic analysis reveals altered glycoprotein expression upon GALNT3 knockdown in ovarian cancer cells. Journal of Proteomics, 2016, 145, 91-102.	1.2	21
13	Neutrophil-Derived Exosomes: A New Mechanism Contributing to Airway Smooth Muscle Remodeling. American Journal of Respiratory Cell and Molecular Biology, 2016, 55, 450-461.	1.4	101
14	Proteomic dataset for altered glycoprotein expression upon GALNT3 knockdown in ovarian cancer cells. Data in Brief, 2016, 8, 342-349.	0.5	7
15	Accessing to the minor proteome of red blood cells through the influence of the nanoparticle surface properties on the corona composition. International Journal of Nanomedicine, 2015, 10, 1869.	3.3	6
16	Mapping the Extracellular and Membrane Proteome Associated with the Vasculature and the Stroma in the Embryo. Molecular and Cellular Proteomics, 2013, 12, 2293-2312.	2.5	21
17	Mitochondrial Inhibitory Factor 1 (IF1) Is Present in Human Serum and Is Positively Correlated with HDL-Cholesterol. PLoS ONE, 2011, 6, e23949.	1.1	29
18	In-depth Exploration of Cerebrospinal Fluid by Combining Peptide Ligand Library Treatment and Label-free Protein Quantification. Molecular and Cellular Proteomics, 2010, 9, 1006-1021.	2.5	116

#	ARTICLE	IF	CITATIONS
19	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