

Cãtia Santa

List of Publications by Year in descending order

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Version: 2024-02-01

27
papers

717
citations

758635

12
h-index

642321

23
g-index

27
all docs

27
docs citations

27
times ranked

1344
citing authors

#	ARTICLE	IF	CITATIONS
1	Proteome dataset of sea bass (<i>Dicentrarchus labrax</i>) skin-scales exposed to fluoxetine and estradiol. <i>Data in Brief</i> , 2022, 41, 107971.	0.5	0
2	Decoding the radiomic and proteomic phenotype of epicardial adipose tissue associated with adverse left atrial remodelling and post-operative atrial fibrillation in aortic stenosis. <i>European Heart Journal Cardiovascular Imaging</i> , 2022, 23, 1248-1259.	0.5	4
3	Systematic Review and Meta-Analysis of Mass Spectrometry Proteomics Applied to Human Peripheral Fluids to Assess Potential Biomarkers of Schizophrenia. <i>International Journal of Molecular Sciences</i> , 2022, 23, 4917.	1.8	10
4	Systematic Review and Meta-Analysis on MS-Based Proteomics Applied to Human Peripheral Fluids to Assess Potential Biomarkers of Bipolar Disorder. <i>International Journal of Molecular Sciences</i> , 2022, 23, 5460.	1.8	9
5	Disclosing proteins in the leaves of cork oak plants associated with the immune response to <i>Phytophthora cinnamomi</i> inoculation in the roots: A long-term proteomics approach. <i>PLoS ONE</i> , 2021, 16, e0245148.	1.1	9
6	Spotted Fever Group <i>Rickettsia</i> Trigger Species-Specific Alterations in Macrophage Proteome Signatures with Different Impacts in Host Innate Inflammatory Responses. <i>Microbiology Spectrum</i> , 2021, 9, e0081421.	1.2	4
7	Epicardial adipose tissue volume and annexin A2/fetuin-A signalling are linked to coronary calcification in advanced coronary artery disease: Computed tomography and proteomic biomarkers from the EPICHEART study. <i>Atherosclerosis</i> , 2020, 292, 75-83.	0.4	25
8	Influence of EPICardial adipose tissue in HEART diseases (EPICHEART) study: Protocol for a translational study in coronary atherosclerosis. <i>Revista Portuguesa De Cardiologia</i> , 2020, 39, 625-633.	0.2	2
9	Influence of EPICardial adipose tissue in HEART diseases (EPICHEART) study: Protocol for a translational study in coronary atherosclerosis. <i>Revista Portuguesa De Cardiologia (English Edition)</i> , 2020, 39, 625-633.	0.2	0
10	Meningeal T cell-derived IL-17 controls synaptic plasticity and short-term memory. <i>Science Immunology</i> , 2019, 4, .	5.6	184
11	Experimental data from flesh quality assessment and shelf life monitoring of high pressure processed European sea bass (<i>Dicentrarchus labrax</i>) fillets. <i>Data in Brief</i> , 2019, 26, 104451.	0.5	7
12	High pressure processing of European sea bass (<i>Dicentrarchus labrax</i>) fillets and tools for flesh quality and shelf life monitoring. <i>Journal of Food Engineering</i> , 2019, 262, 83-91.	2.7	39
13	A Pathogen and a Non-pathogen Spotted Fever Group <i>Rickettsia</i> Trigger Differential Proteome Signatures in Macrophages. <i>Frontiers in Cellular and Infection Microbiology</i> , 2019, 9, 43.	1.8	23
14	SWATH Mass Spectrometry Applied to Cerebrospinal Fluid Differential Proteomics: Establishment of a Sample-Specific Method. <i>Methods in Molecular Biology</i> , 2019, 2044, 169-189.	0.4	1
15	SWATH-MS as a tool for biomarker discovery: From basic research to clinical applications. <i>Proteomics</i> , 2017, 17, 1600278.	1.3	139
16	Matrisome Profiling During Intervertebral Disc Development And Ageing. <i>Scientific Reports</i> , 2017, 7, 11629.	1.6	35
17	Neuroproteomics Using Short GeLC-SWATH: From the Evaluation of Proteome Changes to the Clarification of Protein Function. <i>Neuromethods</i> , 2017, , 107-138.	0.2	7
18	Circulating biomarkers in schizophrenia: a proteomics perspective. <i>International Journal of Clinical Neurosciences and Mental Health</i> , 2017, , S05.	0.7	3

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19	Protein precipitation of diluted samples in SDS-containing buffer with acetone leads to higher protein recovery and reproducibility in comparison with TCA/acetone approach. <i>Proteomics</i> , 2016, 16, 1847-1851.	1.3	42
20	A reference library of peripheral blood mononuclear cells for SWATH-MS analysis. <i>Proteomics - Clinical Applications</i> , 2016, 10, 760-764.	0.8	11
21	Unraveling Mesenchymal Stem Cells™ Dynamic Secretome Through Nontargeted Proteomics Profiling. <i>Methods in Molecular Biology</i> , 2016, 1416, 521-549.	0.4	18
22	Intermittent cardiac overload results in adaptive hypertrophy and provides protection against left ventricular acute pressure overload insult. <i>Journal of Physiology</i> , 2015, 593, 3885-3897.	1.3	33
23	Short GeLC-SWATH: A fast and reliable quantitative approach for proteomic screenings. <i>Proteomics</i> , 2015, 15, 757-762.	1.3	79
24	Mitochondria proteome profiling: A comparative analysis between gel- and gel-free approaches. <i>Talanta</i> , 2013, 115, 277-283.	2.9	12
25	Characterization of mitochondrial proteome in a severe case of ETF-QO deficiency. <i>Journal of Proteomics</i> , 2011, 75, 221-228.	1.2	17
26	Neuroproteomics – LC-MS Quantitative Approaches. , 0, , .		4
27	Analysis of the quantitative proteomic signature in PBMCs of first-episode psychosis patients. <i>Frontiers in Cellular Neuroscience</i> , 0, 13, .	1.8	0