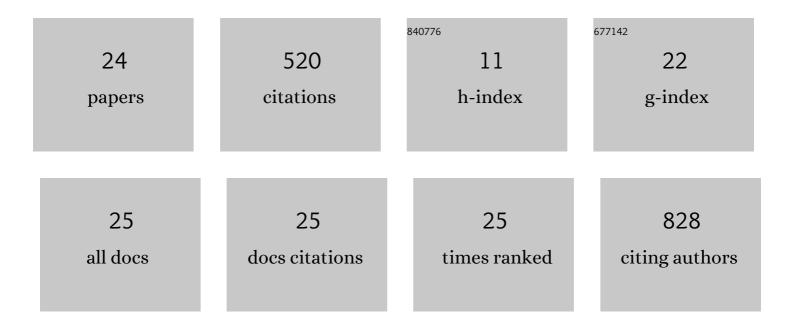
Ana Sofia Carvalho

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	LAMP2A regulates the loading of proteins into exosomes. Science Advances, 2022, 8, eabm1140.	10.3	69
2	Comparative analysis of the bronchoalveolar microbiome in Portuguese patients with different chronic lung disorders. Scientific Reports, 2021, 11, 15042.	3.3	5
3	Proteomic Landscape of Extracellular Vesicles for Diffuse Large B-Cell Lymphoma Subtyping. International Journal of Molecular Sciences, 2021, 22, 11004.	4.1	9
4	Extracellular Vesicle Proteome in Prostate Cancer: A Comparative Analysis of Mass Spectrometry Studies. International Journal of Molecular Sciences, 2021, 22, 13605.	4.1	3
5	WNK1 phosphorylation sites in TBC1D1 and TBC1D4 modulate cell surface expression of GLUT1. Archives of Biochemistry and Biophysics, 2020, 679, 108223.	3.0	12
6	Is the Proteome of Bronchoalveolar Lavage Extracellular Vesicles a Marker of Advanced Lung Cancer?. Cancers, 2020, 12, 3450.	3.7	14
7	Transcriptome Reprogramming of CD11b+ Bone Marrow Cells by Pancreatic Cancer Extracellular Vesicles. Frontiers in Cell and Developmental Biology, 2020, 8, 592518.	3.7	10
8	Extra-cellular vesicles carry proteome of cancer hallmarks. Frontiers in Bioscience - Landmark, 2020, 25, 398-436.	3.0	14
9	Methods and Algorithms for Quantitative Proteomics by Mass Spectrometry. Methods in Molecular Biology, 2020, 2051, 161-197.	0.9	10
10	Interplay between SUMOylation and NEDDylation regulates RPL11 localization and function. FASEB Journal, 2019, 33, 643-651.	0.5	20
11	Profiling of lung microbiota discloses differences in adenocarcinoma and squamous cell carcinoma. Scientific Reports, 2019, 9, 12838.	3.3	64
12	Bronchoalveolar Lavage Proteomics in Patients with Suspected Lung Cancer. Scientific Reports, 2017, 7, 42190.	3.3	46
13	Quantitative proteome analysis of an antibiotic resistant Escherichia coli exposed to tetracycline reveals multiple affected metabolic and peptidoglycan processes. Journal of Proteomics, 2017, 156, 20-28.	2.4	20
14	Red Blood Cells in Clinical Proteomics. Methods in Molecular Biology, 2017, 1619, 173-181.	0.9	5
15	Bronchoalveolar Lavage: Quantitative Mass Spectrometry-Based Proteomics Analysis in Lung Diseases. Methods in Molecular Biology, 2017, 1619, 487-494.	0.9	6
16	New insights into functional regulation in MS-based drug profiling. Scientific Reports, 2016, 6, 18826.	3.3	13
17	Review and Literature Mining on Proteostasis Factors and Cancer. Methods in Molecular Biology, 2016, 1449, 71-84.	0.9	15
18	Global MS-Based Proteomics Drug Profiling. Methods in Molecular Biology, 2016, 1449, 469-479.	0.9	2

2

#	Article	IF	CITATIONS
19	Sequence variation at <i>KLK</i> and <i>WFDC</i> clusters and its association to semen hyperviscosity and other male infertility phenotypes. Human Reproduction, 2016, 31, 2881-2891.	0.9	11
20	CHAPTER 6. Identification and Localization of Post-Translational Modifications by High-Resolution Mass Spectrometry. New Developments in Mass Spectrometry, 2016, , 116-132.	0.2	0
21	Clobal Mass Spectrometry and Transcriptomics Array Based Drug Profiling Provides Novel Insight into Glucosamine Induced Endoplasmic Reticulum Stress. Molecular and Cellular Proteomics, 2014, 13, 3294-3307.	3.8	42
22	Discussion on common data analysis strategies used in MSâ€based proteomics. Proteomics, 2011, 11, 604-619.	2.2	31
23	MUC2 mucin is a major carrier of the cancer-associated sialyl-Tn antigen in intestinal metaplasia and gastric carcinomas. Glycobiology, 2010, 20, 199-206.	2.5	93
24	Multiple Myeloma-Derived Extracellular Vesicles Modulate the Bone Marrow Immune Microenvironment. Frontiers in Immunology, 0, 13, .	4.8	6