## Flavia Vischi Winck

List of Publications by Year in descending order

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516710 477307 35 870 16 citations h-index papers

g-index 40 40 40 1720 docs citations times ranked citing authors all docs

29

#	Article	IF	CITATIONS
1	Characterization of the Nuclear Proteome of Chlamydomonas in Response to Salt Stress. Phycology, 2022, 2, 280-296.	3.6	2
2	Growth and saxitoxin production responses to copper (CuCl2) exposure by the cyanobacterium Raphidiopsis raciborskii. Journal of Applied Phycology, 2021, 33, 891-900.	2.8	1
3	A Reductionist Approach Using Primary and Metastatic Cell–Derived Extracellular Vesicles Reveals Hub Proteins Associated with Oral Cancer Prognosis. Molecular and Cellular Proteomics, 2021, 20, 100118.	3.8	12
4	Plant Proteomics and Systems Biology. Advances in Experimental Medicine and Biology, 2021, 1346, 51-66.	1.6	2
5	Introduction: Advances in Plant Omics and Systems Biology. Advances in Experimental Medicine and Biology, 2021, 1346, 1-9.	1.6	3
6	Analysis of autotrophic, mixotrophic and heterotrophic phenotypes in the microalgae Chlorella vulgaris using time-resolved proteomics and transcriptomics approaches. Algal Research, 2020, 51, 102060.	4.6	44
7	Gene regulatory networks on transfer entropy (GRNTE): a novel approach to reconstruct gene regulatory interactions applied to a case study for the plant pathogen Phytophthora infestans. Theoretical Biology and Medical Modelling, 2019, 16, 7.	2.1	20
8	Where do we aspire to publish? A position paper on scientific communication in biochemistry and molecular biology. Brazilian Journal of Medical and Biological Research, 2019, 52, e8935.	1.5	1
9	Extracellular vesicles from oral squamous carcinoma cells display pro―and antiâ€angiogenic properties. Oral Diseases, 2018, 24, 725-731.	3.0	15
10	Development of a <i>Chlamydomonas reinhardtii</i> metabolic network dynamic model to describe distinct phenotypes occurring at different CO <sub>2</sub> levels. PeerJ, 2018, 6, e5528.	2.0	19
11	Analysis of Sensitive CO2 Pathways and Genes Related to Carbon Uptake and Accumulation in Chlamydomonas reinhardtii through Genomic Scale Modeling and Experimental Validation. Frontiers in Plant Science, 2016, 7, 43.	3.6	16
12	Editorial: Advances in Microalgae Biology and Sustainable Applications. Frontiers in Plant Science, 2016, 7, 1385.	3.6	1
13	EEF1D modulates proliferation and epithelial–mesenchymal transition in oral squamous cell carcinoma. Clinical Science, 2016, 130, 785-799.	4.3	33
14	Insights into immune responses in oral cancer through proteomic analysis of saliva and salivary extracellular vesicles. Scientific Reports, 2015, 5, 16305.	3.3	109
15	Functional annotation and biological interpretation of proteomics data. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2015, 1854, 46-54.	2.3	45
16	Integrative analysis to select cancer candidate biomarkers to targeted validation. Oncotarget, 2015, 6, 43635-43652.	1.8	18
17	A stepâ€byâ€step protocol for formaldehydeâ€assisted isolation of regulatory elements from <i>Arabidopsis thaliana</i> . Journal of Integrative Plant Biology, 2014, 56, 527-538.	8.5	34
18	In Silico Analysis for Biomass Synthesis under Different CO2 Levels for Chlamydomonas reinhardtii Utilizing a Flux Balance Analysis Approach. Advances in Intelligent Systems and Computing, 2014, , 279-285.	0.6	3

#	Article	IF	Citations
19	Systems Analysis of the Response of Photosynthesis, Metabolism, and Growth to an Increase in Irradiance in the Photosynthetic Model Organism <i>Chlamydomonas reinhardtii</i> $\hat{A}$ $\hat{A}$ $\hat{A}$ . Plant Cell, 2014, 26, 2310-2350.	6.6	123
20	Phosphoproteome analysis reveals differences in phosphosite profiles between tumorigenic and non-tumorigenic epithelial cells. Journal of Proteomics, 2014, 96, 67-81.	2.4	11
21	Carbon acquisition and accumulation in microalgae Chlamydomonas: Insights from "omics― approaches. Journal of Proteomics, 2013, 94, 207-218.	2.4	24
22	Biochemical, Pharmacological, and Structural Characterization of New Basic Bbil-TX from <i>Bothriopsis bilineata</i> Snake Venom. BioMed Research International, 2013, 2013, 1-12.	1.9	24
23	Genome-Wide Identification of Regulatory Elements and Reconstruction of Gene Regulatory Networks of the Green Alga Chlamydomonas reinhardtii under Carbon Deprivation. PLoS ONE, 2013, 8, e79909.	2.5	14
24	The nuclear proteome of the green alga <i>Chlamydomonas reinhardtii</i> . Proteomics, 2012, 12, 95-100.	2.2	16
25	AN OPTIMIZED METHOD FOR THE ISOLATION OF NUCLEI FROM CHLAMYDOMONAS REINHARDTII (CHLOROPHYCEAE)1. Journal of Phycology, 2011, 47, 333-340.	2.3	13
26	Isolation and characterization of a new serine protease with thrombin-like activity (TLBm) from the venom of the snake Bothrops marajoensis. Toxicon, 2010, 55, 745-753.	1.6	33
27	The untiring search for the most complete proteome representation: reviewing the methods. Briefings in Functional Genomics & Proteomics, 2008, 7, 312-321.	3.8	18
28	Xylella fastidiosa disturbs nitrogen metabolism and causes a stress response in sweet orange Citrus sinensis cv. Pera. Journal of Experimental Botany, 2007, 58, 2733-2744.	4.8	17
29	LaRbp38: A Leishmania amazonensis protein that binds nuclear and kinetoplast DNAs. Biochemical and Biophysical Research Communications, 2007, 358, 854-860.	2.1	7
30	Absence of Classical Heat Shock Response in the Citrus Pathogen Xylella fastidiosa. Current Microbiology, 2007, 54, 119-123.	2.2	6
31	Biochemical, Pharmacological and Structural Characterization of Two PLA2 Isoforms Cdr-12 and Cdr-13 from Crotalus durissus ruruima Snake Venom. Protein Journal, 2007, 26, 39-49.	1.6	23
32	Isolation and Biochemical Characterization of a Galactoside Binding Lectin from Bauhinia variegata Candida (BvcL) Seeds. Protein Journal, 2007, 26, 193-201.	1.6	31
33	Comparative analysis of two-dimensional electrophoresis maps (2-DE) of Helicobacter pylori from Brazilian patients with chronic gastritis and duodenal ulcer: a preliminary report. Revista Do Instituto De Medicina Tropical De Sao Paulo, 2006, 48, 175-177.	1.1	8
34	Biochemical and enzymatic characterization of two basic Asp49 phospholipase A2 isoforms from Lachesis muta muta (Surucucu) venom. Biochimica Et Biophysica Acta - General Subjects, 2005, 1726, 75-86.	2.4	32
35	Proteome analysis of the plant pathogen Xylella fastidiosa reveals major cellular and extracellular proteins and a peculiar codon bias distribution. Proteomics, 2003, 3, 224-237.	2.2	87