Gilberto B Domont

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3031880/publications.pdf

Version: 2024-02-01

55	1,021	17 h-index	29
papers	citations		g-index
59	59	59	2125
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Proteomics of ZIKV infected amniotic fluids of microcephalic fetuses reveals extracellular matrix and immune system dysregulation. Proteomics - Clinical Applications, 2022, 16, e2100041.	0.8	5
2	Proteomic profiles of Zika virusâ€infected placentas bearing fetuses with microcephaly. Proteomics - Clinical Applications, 2022, 16, e2100042.	0.8	5
3	Improving hemocompatibility of decellularized liver scaffold using Custodiol solution. Materials Science and Engineering C, 2022, , 112642.	3 . 8	4
4	Sheltered in Stromal Tissue Cells, Trypanosoma cruzi Orchestrates Inflammatory Neovascularization via Activation of the Mast Cell Chymase Pathway. Pathogens, 2022, 11, 187.	1,2	2
5	Proteomic Analysis of Embryo Isolated From Mature Jatropha curcas L. Seeds. Frontiers in Plant Science, 2022, 13, 843764.	1.7	1
6	Scents and Flavors of Mass Spectrometry in Latin America. Journal of Proteome Research, 2022, 21, 1-1.	1.8	0
7	Cues from human atrial extracellular matrix enrich the atrial differentiation of human induced pluripotent stem cell-derived cardiomyocytes. Biomaterials Science, 2021, 9, 3737-3749.	2.6	8
8	Monitoring casbene synthase in Jatropha curcas tissues using targeted proteomics. Plant Methods, 2021, 17, 15.	1.9	1
9	Exploring the biological activities and proteome of Brazilian scorpion Rhopalurus agamemnon venom. Journal of Proteomics, 2021, 237, 104119.	1.2	7
10	Ancient enamel peptides recovered from the South American Pleistocene species Notiomastodon platensis and Myocastor cf. coypus. Journal of Proteomics, 2021, 240, 104187.	1.2	10
11	The human melanoma proteome atlasâ€"Defining the molecular pathology. Clinical and Translational Medicine, 2021, 11, e473.	1.7	14
12	The Human Melanoma Proteome Atlasâ€"Complementing the melanoma transcriptome. Clinical and Translational Medicine, 2021, 11, e451.	1.7	20
13	Quantitative profiling of axonal guidance proteins during the differentiation of human neurospheres. Biochimica Et Biophysica Acta - Proteins and Proteomics, 2021, 1869, 140656.	1.1	6
14	Topological Dissection of Proteomic Changes Linked to the Limbic Stage of Alzheimer's Disease. Frontiers in Immunology, 2021, 12, 750665.	2.2	5
15	Short-Term Effect of Induced Alterations in Testosterone Levels on Fasting Plasma Amino Acid Levels in Healthy Young Men. Life, 2021, 11, 1276.	1.1	2
16	Deep Proteomic Analysis on Biobanked Paraffine-Archived Melanoma with Prognostic/Predictive Biomarker Read-Out. Cancers, 2021, 13, 6105.	1.7	3
17	Mapping the Melanoma Plasma Proteome (MPP) Using Single-Shot Proteomics Interfaced with the WiMT Database. Cancers, 2021, 13, 6224.	1.7	4
18	Novel functional proteins coded by the human genome discovered in metastases of melanoma patients. Cell Biology and Toxicology, 2020, 36, 261-272.	2.4	9

#	Article	IF	CITATIONS
19	Proteome Dynamics of the Developing AçaÃ-Berry Pericarp (<i>Euterpe oleracea</i> Mart.). Journal of Proteome Research, 2020, 19, 437-445.	1.8	6
20	Proteomic analysis of whole saliva in chronic periodontitis. Journal of Proteomics, 2020, 213, 103602.	1.2	29
21	A high-stringency blueprint of the human proteome. Nature Communications, 2020, 11, 5301.	5 . 8	152
22	Quantitative Proteome Analysis of Jatropha curcas L. Genotypes with Contrasting Levels of Phorbol Esters. Proteomics, 2020, 20, 1900273.	1.3	1
23	Molecular alterations in the extracellular matrix in the brains of newborns with congenital Zika syndrome. Science Signaling, 2020, 13 , .	1.6	39
24	Back to the Future with Proteins and Proteomics. Brazilian Journal of Analytical Chemistry, 2020, 7, 3-10.	0.3	0
25	Metabolomic profiling suggests systemic signatures of premature aging induced by Hutchinson–Gilford progeria syndrome. Metabolomics, 2019, 15, 100.	1.4	4
26	Quantitative Subcellular Proteomics of the Orbitofrontal Cortex of Schizophrenia Patients. Journal of Proteome Research, 2019, 18, 4240-4253.	1.8	21
27	Proteomic signatures of brain regions affected by tau pathology in early and late stages of Alzheimer's disease. Neurobiology of Disease, 2019, 130, 104509.	2.1	46
28	Computational fluid dynamic analysis of physical forces playing a role in brain organoid cultures in two different multiplex platforms. BMC Developmental Biology, 2019, 19, 3.	2.1	31
29	Proteomic Analysis and Functional Validation of a Brassica oleracea Endochitinase Involved in Resistance to Xanthomonas campestris. Frontiers in Plant Science, 2019, 10, 414.	1.7	13
30	Inâ€Depth Proteome Analysis of Ricinus communis Pollens. Proteomics, 2019, 19, 1800347.	1.3	0
31	Launching the C-HPP neXt-CP50 Pilot Project for Functional Characterization of Identified Proteins with No Known Function. Journal of Proteome Research, 2018, 17, 4042-4050.	1.8	41
32	Common Features Between the Proteomes of Floral and Extrafloral Nectar From the Castor Plant (Ricinus Communis) and the Proteomes of Exudates From Carnivorous Plants. Frontiers in Plant Science, 2018, 9, 549.	1.7	8
33	Mass spectrometry evaluation of a neuroblastoma SH-SY5Y cell culture protocol. Analytical Biochemistry, 2018, 559, 51-54.	1.1	2
34	Aging-related compensated hypogonadism: Role of metabolomic analysis in physiopathological and therapeutic evaluation. Journal of Steroid Biochemistry and Molecular Biology, 2018, 183, 39-50.	1.2	30
35	7-Ketocholesterol overcomes drug resistance in chronic myeloid leukemia cell lines beyond MDR1 mechanism. Journal of Proteomics, 2017, 151, 12-23.	1.2	22
36	DiagnoProt: a tool for discovery of new molecules by mass spectrometry. Bioinformatics, 2017, 33, 1883-1885.	1.8	7

3

#	Article	IF	CITATIONS
37	Application of iTRAQ Shotgun Proteomics for Measurement of Brain Proteins in Studies of Psychiatric Disorders. Advances in Experimental Medicine and Biology, 2017, 974, 219-227.	0.8	5
38	Proteomic assessment of colorectal cancers and respective resection margins from patients of the Amazon state of Brazil. Journal of Proteomics, 2017, 154, 59-68.	1.2	4
39	Synaptosomal Proteome of the Orbitofrontal Cortex from Schizophrenia Patients Using Quantitative Label-Free and iTRAQ-Based Shotgun Proteomics. Journal of Proteome Research, 2017, 16, 4481-4494.	1.8	44
40	Quantitative proteomic analysis identifies proteins and pathways related to neuronal development in differentiated SH-SY5Y neuroblastoma cells. EuPA Open Proteomics, 2017, 16, 1-11.	2.5	48
41	Quantitative proteomic analysis of the Saccharomyces cerevisiae industrial strains CAT-1 and PE-2. Journal of Proteomics, 2017, 151, 114-121.	1.2	18
42	It is time for top-down venomics. Journal of Venomous Animals and Toxins Including Tropical Diseases, 2017, 23, 44.	0.8	34
43	iTRAQ-Based Shotgun Proteomics Approach for Relative Protein Quantification. Methods in Molecular Biology, 2017, 1546, 267-274.	0.4	10
44	A Time-Based and Intratumoral Proteomic Assessment of a Recurrent Glioblastoma Multiforme. Frontiers in Oncology, 2016, 6, 183.	1.3	13
45	Time-course proteome analysis of developing extrafloral nectaries of <i>Ricinus communis</i> Proteomics, 2016, 16, 629-633.	1.3	17
46	Comparative proteome analysis reveals that blood and sugar meals induce differential protein expression in <i>Aedes aegypti</i> female heads. Proteomics, 2016, 16, 2582-2586.	1.3	10
47	CN-GELFrEE - Clear Native Gel-eluted Liquid Fraction Entrapment Electrophoresis. Journal of Visualized Experiments, 2016, , 53597.	0.2	9
48	Deep proteome analysis of gerontoplasts from the inner integument of developing seeds of Jatropha curcas. Journal of Proteomics, 2016, 143, 346-352.	1.2	12
49	Quest for Missing Proteins: Update 2015 on Chromosome-Centric Human Proteome Project. Journal of Proteome Research, 2015, 14, 3415-3431.	1.8	53
50	Proteomic Analysis of the Endosperm Ontogeny of <i>Jatropha curcas</i> L. Seeds. Journal of Proteome Research, 2015, 14, 2557-2568.	1.8	21
51	Oligomerization and Membrane-binding Properties of Covalent Adducts Formed by the Interaction of \hat{l}_{\pm} -Synuclein with the Toxic Dopamine Metabolite 3,4-Dihydroxyphenylacetaldehyde (DOPAL). Journal of Biological Chemistry, 2015, 290, 27660-27679.	1.6	100
52	Two decades of proteomics in Latin America: A personal view. Journal of Proteomics, 2014, 107, 83-92.	1.2	7
53	Microbial Proteomics. Journal of Proteome Research, 2012, 11, 1-1.	1.8	4
54	Proteome analysis of castor bean seeds. Pure and Applied Chemistry, 2010, 82, 259-267.	0.9	15

4	#	Article	lF	CITATIONS
	55	GO Explorer: A gene-ontology tool to aid in the interpretation of shotgun proteomics data. Proteome Science, 2009, 7, 6.	0.7	35