

Paula DÃ-ez

List of Publications by Year in descending order

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Version: 2024-02-01

37
papers

566
citations

777949

13
h-index

759306

22
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38
all docs

38
docs citations

38
times ranked

1249
citing authors

#	ARTICLE	IF	CITATIONS
1	Dynamic Intracellular Metabolic Cell Signaling Profiles During Ag-Dependent B-Cell Differentiation. <i>Frontiers in Immunology</i> , 2021, 12, 637832.	2.2	4
2	Tracking the Antibody Immunome in Sporadic Colorectal Cancer by Using Antigen Self-Assembled Protein Arrays. <i>Cancers</i> , 2021, 13, 2718.	1.7	9
3	Deepening into Intracellular Signaling Landscape through Integrative Spatial Proteomics and Transcriptomics in a Lymphoma Model. <i>Biomolecules</i> , 2021, 11, 1776.	1.8	8
4	A Novel Cytotoxic Conjugate Derived from the Natural Product Podophyllotoxin as a Direct-Target Protein Dual Inhibitor. <i>Molecules</i> , 2020, 25, 4258.	1.7	7
5	New Hybrids Derived from Podophyllic Aldehyde and Diterpenylhydroquinones with Selectivity toward Osteosarcoma Cells. <i>ACS Medicinal Chemistry Letters</i> , 2018, 9, 328-333.	1.3	9
6	Screening Phage-Display Antibody Libraries Using Protein Arrays. <i>Methods in Molecular Biology</i> , 2018, 1701, 365-380.	0.4	12
7	In-Depth Proteomic Characterization of Classical and Non-Classical Monocyte Subsets. <i>Proteomes</i> , 2018, 6, 8.	1.7	18
8	Understanding and utilizing the biomolecule/nanosystems interface. , 2018, , 207-297.		19
9	Functional proteomic insights in B-cell chronic lymphocytic leukemia. <i>Expert Review of Proteomics</i> , 2017, 14, 137-146.	1.3	8
10	Decoding the anticancer activity of VO-clioquinol compound: the mechanism of action and cell death pathways in human osteosarcoma cells. <i>Metallomics</i> , 2017, 9, 891-901.	1.0	27
11	Screening and Validation of Novel Biomarkers in Osteoarticular Pathologies by Comprehensive Combination of Protein Array Technologies. <i>Journal of Proteome Research</i> , 2017, 16, 1890-1899.	1.8	23
12	Proteomic Biomarker Identification in Cerebrospinal Fluid for Leptomeningeal Metastases with Neurological Complications. <i>Advances in Experimental Medicine and Biology</i> , 2017, 974, 85-96.	0.8	5
13	CSF analysis for protein biomarker identification in patients with leptomeningeal metastases from CNS lymphoma. <i>Expert Review of Proteomics</i> , 2017, 14, 363-372.	1.3	6
14	Comprehensive and Systematic Analysis of the Immunocompatibility of Polyelectrolyte Capsules. <i>Bioconjugate Chemistry</i> , 2017, 28, 556-564.	1.8	39
15	Functional insights into the cellular response triggered by a bile-acid platinum compound conjugated to biocompatible ferric nanoparticles using quantitative proteomic approaches. <i>Nanoscale</i> , 2017, 9, 9960-9972.	2.8	11
16	A systematic approach for peptide characterization of B-cell receptor in chronic lymphocytic leukemia cells. <i>Oncotarget</i> , 2017, 8, 42836-42846.	0.8	7
17	Methods for Selecting Phage Display Antibody Libraries. <i>Current Pharmaceutical Design</i> , 2017, 22, 6490-6499.	0.9	3
18	Multipronged functional proteomics approaches for global identification of altered cell signalling pathways in B-cell chronic lymphocytic leukaemia. <i>Proteomics</i> , 2016, 16, 1193-1203.	1.3	15

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19	Deciphering the effect of an oxovanadium(IV) complex with the flavonoid chrysin (VOChrys) on intracellular cell signalling pathways in an osteosarcoma cell line. <i>Metallomics</i> , 2016, 8, 739-749.	1.0	40
20	Proteogenomics for the Comprehensive Analysis of Human Cellular and Serum Antibody Repertoires. <i>Advances in Experimental Medicine and Biology</i> , 2016, 926, 153-162.	0.8	2
21	Outside Back Cover: Multipronged functional proteomics approaches for global identification of altered cell signalling pathways in B-cell chronic lymphocytic leukaemia. <i>Proteomics</i> , 2016, 16, NA-NA.	1.3	0
22	Nanotechnology in the Fabrication of Protein Microarrays. <i>Methods in Molecular Biology</i> , 2016, 1368, 197-208.	0.4	4
23	Sensing parasites: Proteomic and advanced bio-detection alternatives. <i>Journal of Proteomics</i> , 2016, 136, 145-156.	1.2	22
24	Evaluation Strategies of Nanomaterials Toxicity. , 2015, , .		4
25	NAPPA as a Real New Method for Protein Microarray Generation. <i>Microarrays (Basel, Switzerland)</i> , 2015, 4, 214-227.	1.4	24
26	High-throughput phage-display screening in array format. <i>Enzyme and Microbial Technology</i> , 2015, 79-80, 34-41.	1.6	1
27	Quest for Missing Proteins: Update 2015 on Chromosome-Centric Human Proteome Project. <i>Journal of Proteome Research</i> , 2015, 14, 3415-3431.	1.8	53
28	Integration of Proteomics and Transcriptomics Data Sets for the Analysis of a Lymphoma B-Cell Line in the Context of the Chromosome-Centric Human Proteome Project. <i>Journal of Proteome Research</i> , 2015, 14, 3530-3540.	1.8	16
29	In Vitro Transcription/Translation System: A Versatile Tool in the Search for Missing Proteins. <i>Journal of Proteome Research</i> , 2015, 14, 3441-3451.	1.8	11
30	Emerging Nanotechniques in Proteomics. <i>Comprehensive Analytical Chemistry</i> , 2014, 63, 137-157.	0.7	0
31	Surfing Transcriptomic Landscapes. A Step beyond the Annotation of Chromosome 16 Proteome. <i>Journal of Proteome Research</i> , 2014, 13, 158-172.	1.8	26
32	Protein Microarrays: Overview, Applications and Challenges. <i>Translational Bioinformatics</i> , 2014, , 147-173.	0.0	6
33	Chromosome 19 Annotations with Disease Speciation: A First Report from the Global Research Consortium. <i>Journal of Proteome Research</i> , 2013, 12, 135-150.	1.8	16
34	Protein arrays as tool for studies at the host-pathogen interface. <i>Journal of Proteomics</i> , 2013, 94, 387-400.	1.2	12
35	Protein Microarrays: Technological Aspects, Applications and Intellectual Property. <i>Recent Patents on Biotechnology</i> , 2013, 7, 142-152.	0.4	4
36	Biomarker Discovery by Novel Sensors Based on Nanoproteomics Approaches. <i>Sensors</i> , 2012, 12, 2284-2308.	2.1	59

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37	Data Analysis Strategies for Protein Microarrays. Microarrays (Basel, Switzerland), 2012, 1, 64-83.	1.4	34