Yadira Palacios

List of Publications by Year in descending order

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1040056 1199594 14 241 9 12 citations h-index g-index papers 14 14 14 425 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	<i>TNFRSF1B</i> and <i>TNF</i> Variants Are Associated With Differences in Levels of Soluble Tumor Necrosis Factor Receptors in Patients With Severe COVID-19. Journal of Infectious Diseases, 2022, 226, 778-787.	4.0	6
2	Diverse Cell Death Mechanisms Are Simultaneously Activated in Macrophages Infected by Virulent Mycobacterium tuberculosis. Pathogens, 2022, 11, 492.	2.8	7
3	Immunosuppressant Therapies in COVID-19: Is the TNF Axis an Alternative?. Pharmaceuticals, 2022, 15, 616.	3.8	7
4	Transmembrane TNF and Its Receptors TNFR1 and TNFR2 in Mycobacterial Infections. International Journal of Molecular Sciences, 2021, 22, 5461.	4.1	16
5	Severe COVID-19 Patients Show an Increase in Soluble TNFR1 and ADAM17, with a Relationship to Mortality. International Journal of Molecular Sciences, 2021, 22, 8423.	4.1	32
6	Comparative proteomic profiling of triple-negative breast cancer reveals that up-regulation of RhoGDI-2 is associated to the inhibition of caspase 3 and caspase 9. Journal of Proteomics, 2014, 111, 198-211.	2.4	19
7	HIV-1 V3 loop crown epitope-focused mimotope selection by patient serum from random phage display libraries: Implications for the epitope structural features. Molecular Immunology, 2013, 54, 148-156.	2.2	18
8	Proteomic Profiling Reveals That Resveratrol Inhibits HSP27 Expression and Sensitizes Breast Cancer Cells to Doxorubicin Therapy. PLoS ONE, 2013, 8, e64378.	2.5	66
9	Short constrained peptides derived from phage display libraries as epitope models: the case of mAb 2F5. Retrovirology, 2012, 9, .	2.0	0
10	Polypeptide Modulators of Caspase Recruitment Domain (CARD)-CARD-mediated Protein-Protein Interactions. Journal of Biological Chemistry, 2011, 286, 44457-44466.	3.4	15
11	Characterization of dequalinium as a XIAP antagonist that targets the BIR2 domain. Apoptosis: an International Journal on Programmed Cell Death, 2011, 16, 460-467.	4.9	10
12	Constrained peptide models from phage display libraries highlighting the cognate epitope-specific potential of the anti-HIV-1 mAb 2F5. Immunology Letters, 2011, 136, 80-89.	2.5	14
13	P12-05. Using combinatorial phage-display libraries as a source of mimetic peptides of three HIV-1 envelope epitopes. Retrovirology, 2009, 6, .	2.0	0
14	Collection of phage–peptide probes for HIV-1 immunodominant loop-epitope. Journal of Microbiological Methods, 2007, 68, 225-235.	1.6	31