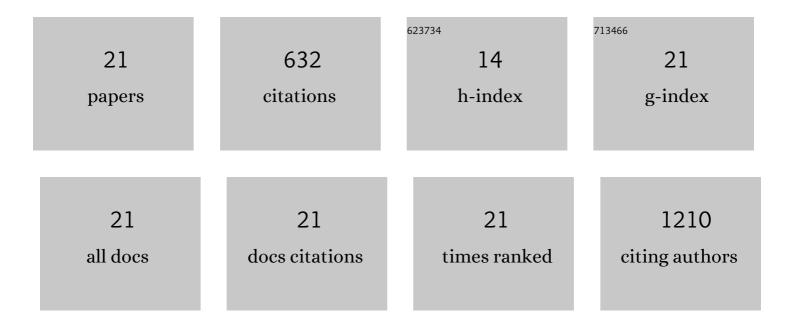
Isabella Venza

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3665489/publications.pdf

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#	Article	IF	CITATIONS
1	Cellular Mechanisms of Oxidative Stress and Action in Melanoma. Oxidative Medicine and Cellular Longevity, 2015, 2015, 1-11.	4.0	88
2	Epigenetic regulation of p14 and p16 expression in cutaneous and uveal melanoma. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2015, 1849, 247-256.	1.9	58
3	PGE2 induces <i><i>i>interleukin-8</i></i> derepression in human astrocytoma through coordinated DNA demethylation and histone hyperacetylation. Epigenetics, 2012, 7, 1315-1330.	2.7	50
4	Heavy Metals and Epigenetic Alterations in Brain Tumors. Current Genomics, 2015, 15, 457-463.	1.6	50
5	Class I-specific histone deacetylase inhibitor MS-275 overrides TRAIL-resistance in melanoma cells by downregulating c-FLIP. International Immunopharmacology, 2014, 21, 439-446.	3.8	41
6	Impact of DNA methyltransferases on the epigenetic regulation of tumor necrosis factor-related apoptosis-inducing ligand (TRAIL) receptor expression in malignant melanoma. Biochemical and Biophysical Research Communications, 2013, 441, 743-750.	2.1	39
7	Neutrophils Directly Recognize Group B Streptococci and Contribute to Interleukin-1Î ² Production during Infection. PLoS ONE, 2016, 11, e0160249.	2.5	39
8	DNA methylation-induced E-cadherin silencing is correlated with the clinicopathological features of melanoma. Oncology Reports, 2016, 35, 2451-2460.	2.6	38
9	Epigenetic marks responsible for cadmium-induced melanoma cell overgrowth. Toxicology in Vitro, 2015, 29, 242-250.	2.4	34
10	DSS1 promoter hypomethylation and overexpression predict poor prognosis in melanoma and squamous cell carcinoma patients. Human Pathology, 2017, 60, 137-146.	2.0	28
11	Possible protective role of the ABCA4 gene c.1268A>G missense variant in Stargardt disease and syndromic retinitis pigmentosa in a Sicilian family: Preliminary data. International Journal of Molecular Medicine, 2017, 39, 1011-1020.	4.0	27
12	Epigenetic Effects of Cadmium in Cancer: Focus on Melanoma. Current Genomics, 2015, 15, 420-435.	1.6	27
13	miR-92a-3p and MYCBP2 are involved in MS-275-induced and c-myc-mediated TRAIL-sensitivity in melanoma cells. International Immunopharmacology, 2016, 40, 235-243.	3.8	24
14	Phage display revisited: Epitope mapping of a monoclonal antibody directed against <i>Neisseria meningitidis</i> adhesin A using the PROFILER technology. MAbs, 2016, 8, 741-750.	5.2	19
15	Identification of microRNA Expression Patterns in Cutaneous and Uveal Melanoma Cell Lines. Tumori, 2014, 100, e4-e7.	1.1	13
16	The overriding of TRAIL resistance by the histone deacetylase inhibitor MS-275 involves c-myc up-regulation in cutaneous, uveal, and mucosal melanoma. International Immunopharmacology, 2015, 28, 313-321.	3.8	13
17	Functional characterization of a monoclonal antibody epitope using a lambda phage display-deep sequencing platform. Scientific Reports, 2016, 6, 31458.	3.3	12
18	Epitope Mapping of a Monoclonal Antibody Directed against Neisserial Heparin Binding Antigen Using Next Generation Sequencing of Antigen-Specific Libraries. PLoS ONE, 2016, 11, e0160702.	2.5	11

ISABELLA VENZA

#	Article	IF	CITATIONS
19	NOD2 triggers PGE2 synthesis leading to IL-8 activation in Staphylococcus aureus-infected human conjunctival epithelial cells. Biochemical and Biophysical Research Communications, 2013, 440, 551-557.	2.1	9
20	Epidrugs in the Immunotherapy of Cutaneous and Uveal Melanoma. Anti-Cancer Agents in Medicinal Chemistry, 2017, 17, 190-205.	1.7	7
21	Association of the DSS1 c.143G>A Polymorphism with Skin Squamous Cell Carcinoma. Journal of Investigative Dermatology, 2010, 130, 1719-1725.	0.7	5