

Vito Alessandro Lasorsa

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

26

papers

286

citations

9

h-index

16

g-index

31

ext. papers

475

ext. citations

5.6

avg, IF

3.23

L-index

#	Paper	IF	Citations
26	Exome and deep sequencing of clinically aggressive neuroblastoma reveal somatic mutations that affect key pathways involved in cancer progression. <i>Oncotarget</i> , 2016 , 7, 21840-52	3.3	53
25	An 18 gene expression-based score classifier predicts the clinical outcome in stage 4 neuroblastoma. <i>Journal of Translational Medicine</i> , 2016 , 14, 142	8.5	30
24	Genetic Analysis of the Coronavirus SARS-CoV-2 Host Protease in Different Populations. <i>Frontiers in Genetics</i> , 2020 , 11, 872	4.5	22
23	BAP1 induces cell death via interaction with 14-3-3 in neuroblastoma. <i>Cell Death and Disease</i> , 2018 , 9, 458	9.8	21
22	Common variants at 21q22.3 locus influence and gene expression and susceptibility to severe COVID-19. <i>IScience</i> , 2021 , 24, 102322	6.1	21
21	HIF-1 transcription activity: HIF1A driven response in normoxia and in hypoxia. <i>BMC Medical Genetics</i> , 2019 , 20, 37	2.1	20
20	Somatic mutations in specific and connected subpathways are associated with short neuroblastoma patients survival and indicate proteins targetable at onset of disease. <i>International Journal of Cancer</i> , 2018 , 143, 2525-2536	7.5	15
19	Neural crest-derived tumor neuroblastoma and melanoma share 1p13.2 as susceptibility locus that shows a long-range interaction with the SLC16A1 gene. <i>Carcinogenesis</i> , 2020 , 41, 284-295	4.6	11
18	miR-622 is a novel potential biomarker of breast carcinoma and impairs motility of breast cancer cells through targeting NUAK1 kinase. <i>British Journal of Cancer</i> , 2020 , 123, 426-437	8.7	10
17	Exploring Shared Susceptibility between Two Neural Crest Cells Originating Conditions: Neuroblastoma and Congenital Heart Disease. <i>Genes</i> , 2019 , 10,	4.2	8
16	Association of PARP1 polymorphisms with response to chemotherapy in patients with high-risk neuroblastoma. <i>Journal of Cellular and Molecular Medicine</i> , 2020 , 24, 4072-4081	5.6	8
15	A Targeted Gene Panel for Circulating Tumor DNA Sequencing in Neuroblastoma. <i>Frontiers in Oncology</i> , 2020 , 10, 596191	5.3	8
14	Genomic coamplification of CDK4/MDM2/FRS2 is associated with very poor prognosis and atypical clinical features in neuroblastoma patients. <i>Genes Chromosomes and Cancer</i> , 2020 , 59, 277-285	5	8
13	A comprehensive characterization of rare mitochondrial DNA variants in neuroblastoma. <i>Oncotarget</i> , 2016 , 7, 49246-49258	3.3	8
12	FKBP51s signature in peripheral blood mononuclear cells of melanoma patients as a possible predictive factor for immunotherapy. <i>Cancer Immunology, Immunotherapy</i> , 2017 , 66, 1143-1151	7.4	7
11	Functional characterization of full-length BARD1 strengthens its role as a tumor suppressor in neuroblastoma. <i>Journal of Cancer</i> , 2020 , 11, 1495-1504	4.5	7
10	Transcription Factors Involved in Tumorigenesis Are Over-Represented in Mutated Active DNA-Binding Sites in Neuroblastoma. <i>Cancer Research</i> , 2020 , 80, 382-393	10.1	7

9	Genetic analysis of the novel SARS-CoV-2 host receptor TMPRSS2 in different populations		5
8	Kinome multigenic panel identified novel druggable EPHB4-V871I somatic variant in high-risk neuroblastoma. <i>Journal of Cellular and Molecular Medicine</i> , 2020 , 24, 6459-6471	5.6	4
7	19p loss is significantly enriched in older age neuroblastoma patients and correlates with poor prognosis. <i>Npj Genomic Medicine</i> , 2020 , 5, 18	6.2	4
6	Gene Expression Signature of Acquired Chemoresistance in Neuroblastoma Cells. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	2
5	Discovery of -Enprioline as a Novel Drug for the Treatment of Vincristine Resistant Neuroblastoma. <i>International Journal of Molecular Sciences</i> , 2020 , 21,	6.3	2
4	Estimating asymptomatic SARS-CoV-2 infections in a geographic area of low disease incidence. <i>BMC Infectious Diseases</i> , 2021 , 21, 350	4	1
3	Regulatory Noncoding and Predicted Pathogenic Coding Variants of Predispose to Severe COVID-19. <i>International Journal of Molecular Sciences</i> , 2021 , 22,	6.3	1
2	The H159Y Variant Is Associated with Severe COVID-19: A Retrospective Study of 500 Patients from Southern Italy. <i>Genes</i> , 2021 , 12,	4.2	1
1	FGFR1 is a potential therapeutic target in neuroblastoma.. <i>Cancer Cell International</i> , 2022 , 22, 174	6.4	1