

Veronica Zelli

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

531
citations

840776

11
h-index

677142

22
g-index

23
all docs

23
docs citations

23
times ranked

1349
citing authors

#	ARTICLE	IF	CITATIONS
1	Neuroprotective effects of human amniotic fluid stem cells-derived secretome in an ischemia/reperfusion model. <i>Stem Cells Translational Medicine</i> , 2021, 10, 251-266.	3.3	31
2	Mechanisms involved in selecting and maintaining neuroblastoma cancer stem cell populations, and perspectives for therapeutic targeting. <i>World Journal of Stem Cells</i> , 2021, 13, 685-736.	2.8	3
3	Emerging Role of isomiRs in Cancer: State of the Art and Recent Advances. <i>Genes</i> , 2021, 12, 1447.	2.4	11
4	Transcriptome of Male Breast Cancer Matched with Germline Profiling Reveals Novel Molecular Subtypes with Possible Clinical Relevance. <i>Cancers</i> , 2021, 13, 4515.	3.7	6
5	Circulating MicroRNAs as Prognostic and Therapeutic Biomarkers in Breast Cancer Molecular Subtypes. <i>Journal of Personalized Medicine</i> , 2020, 10, 98.	2.5	16
6	Multidisciplinary Treatment, Including Locoregional Chemotherapy, for Merkel-Polyomavirus-Positive Merkel Cell Carcinomas: Perspectives for Patients Exhibiting Oncogenic Alternative 1 st exon 6â€“7 TrkAIII Splicing of Neurotrophin Receptor Tropomyosin-Related Kinase A. <i>International Journal of Molecular Sciences</i> , 2020, 21, 8222.	4.1	4
7	Hypoxia-induced alternative splicing: the 11th Hallmark of Cancer. <i>Journal of Experimental and Clinical Cancer Research</i> , 2020, 39, 110.	8.6	79
8	Applications of Next Generation Sequencing to the Analysis of Familial Breast/Ovarian Cancer. <i>High-Throughput</i> , 2020, 9, 1.	4.4	22
9	MiRNAs as Potential Prognostic Biomarkers for Metastasis in Thin and Thick Primary Cutaneous Melanomas. <i>Anticancer Research</i> , 2019, 39, 4085-4093.	1.1	11
10	The antiquity of hydrocephalus: the first full palaeo-neuropathological description. <i>Neurological Sciences</i> , 2019, 40, 1315-1322.	1.9	3
11	Insight into genetic susceptibility to male breast cancer by multigene panel testing: Results from a multicenter study in Italy. <i>International Journal of Cancer</i> , 2019, 145, 390-400.	5.1	40
12	Evaluation of CYP17A1 and CYP1B1 polymorphisms in male breast cancer risk. <i>Endocrine Connections</i> , 2019, 8, 1224-1229.	1.9	6
13	A possible role of FANCM mutations in male breast cancer susceptibility: Results from a multicenter study in Italy. <i>Breast</i> , 2018, 38, 92-97.	2.2	23
14	Smoking and FGFR2 rs2981582 variant independently modulate male breast cancer survival: A population-based study in Tuscany, Italy. <i>Breast</i> , 2018, 40, 85-91.	2.2	7
15	Contribution of MUTYH Variants to Male Breast Cancer Risk: Results From a Multicenter Study in Italy. <i>Frontiers in Oncology</i> , 2018, 8, 583.	2.8	25
16	<i>c.3140A>G mutation in a patient with suspected Proteus Syndrome: a case report. Clinical Case Reports (discontinued)</i> , 2018, 6, 1358-1363.	0.5	4
17	Gene-specific methylation profiles in BRCA-mutation positive and BRCA-mutation negative male breast cancers. <i>Oncotarget</i> , 2018, 9, 19783-19792.	1.8	8
18	Metastases risk in thin cutaneous melanoma: prognostic value of clinical-pathologic characteristics and mutation profile. <i>Oncotarget</i> , 2018, 9, 32173-32181.	1.8	10

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19	Whole-exome sequencing and targeted gene sequencing provide insights into the role of <i>PALB2</i> as a male breast cancer susceptibility gene. <i>Cancer</i> , 2017, 123, 210-218.	4.1	31
20	Prediction of Breast and Prostate Cancer Risks in Male <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers Using Polygenic Risk Scores. <i>Journal of Clinical Oncology</i> , 2017, 35, 2240-2250.	1.6	152
21	EMSY copy number variation in male breast cancers characterized for <i>BRCA1</i> and <i>BRCA2</i> mutations. <i>Breast Cancer Research and Treatment</i> , 2016, 160, 181-186.	2.5	6
22	Somatic alterations of targetable oncogenes are frequently observed in <i>BRCA1/2</i> mutation negative male breast cancers. <i>Oncotarget</i> , 2016, 7, 74097-74106.	1.8	8
23	Novel and known genetic variants for male breast cancer risk at 8q24.21, 9p21.3, 11q13.3 and 14q24.1: Results from a multicenter study in Italy. <i>European Journal of Cancer</i> , 2015, 51, 2289-2295.	2.8	25