

Assunta Sellitto

List of Publications by Year in descending order

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Version: 2024-02-01

13
papers

312
citations

1039880

9
h-index

1125617

13
g-index

14
all docs

14
docs citations

14
times ranked

542
citing authors

#	ARTICLE	IF	CITATIONS
1	Inhibition of histone methyltransferase DOT1L silences ER ⁺ gene and blocks proliferation of antiestrogen-resistant breast cancer cells. <i>Science Advances</i> , 2019, 5, eaav5590.	4.7	70
2	Specific patterns of PIWI-interacting small noncoding RNA expression in dysplastic liver nodules and hepatocellular carcinoma. <i>Oncotarget</i> , 2016, 7, 54650-54661.	0.8	63
3	Insights into the Role of Estrogen Receptor β in Triple-Negative Breast Cancer. <i>Cancers</i> , 2020, 12, 1477.	1.7	33
4	WIND (Workflow for piRNAs aNd beyond): a strategy for in-depth analysis of small RNA-seq data. <i>F1000Research</i> , 2021, 10, 1.	0.8	22
5	Histone Methyltransferase DOT1L as a Promising Epigenetic Target for Treatment of Solid Tumors. <i>Frontiers in Genetics</i> , 2022, 13, 864612.	1.1	22
6	Metabolic Regulation of Epigenetic Modifications and Cell Differentiation in Cancer. <i>Cancers</i> , 2020, 12, 3788.	1.7	21
7	An Overview of Candidate Therapeutic Target Genes in Ovarian Cancer. <i>Cancers</i> , 2020, 12, 1470.	1.7	20
8	Molecular and Functional Characterization of the Somatic PIWI1/piRNA Pathway in Colorectal Cancer Cells. <i>Cells</i> , 2019, 8, 1390.	1.8	16
9	Small-bowel carcinomas associated with celiac disease: transcriptomic profiling shows predominance of microsatellite instability-immune and mesenchymal subtypes. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2020, 476, 711-723.	1.4	13
10	Regulation of Metabolic Reprogramming by Long Non-Coding RNAs in Cancer. <i>Cancers</i> , 2021, 13, 3485.	1.7	12
11	NGS analysis of nasopharyngeal microbiota in SARS-CoV-2 positive patients during the first year of the pandemic in the Campania Region of Italy. <i>Microbial Pathogenesis</i> , 2022, 165, 105506.	1.3	12
12	WIND (Workflow for piRNAs aNd beyond): a strategy for in-depth analysis of small RNA-seq data. <i>F1000Research</i> , 2021, 10, 1.	0.8	5
13	Whole-genome sequencing of <i>Pseudomonas</i> sp. TAE6080, a strain capable of inhibiting <i>Staphylococcus epidermidis</i> biofilm. <i>Marine Genomics</i> , 2021, 60, 100887.	0.4	3