

Arianna Nicolussi

List of Publications by Year in descending order

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Version: 2024-02-01

12
papers

305
citations

1163117

8
h-index

1125743

13
g-index

13
all docs

13
docs citations

13
times ranked

696
citing authors

#	ARTICLE	IF	CITATIONS
1	The role of peroxiredoxins in cancer. <i>Molecular and Clinical Oncology</i> , 2017, 6, 139-153.	1.0	145
2	Validation of the Ion Torrent PGM sequencing for the prospective routine molecular diagnostic of colorectal cancer. <i>Clinical Biochemistry</i> , 2015, 48, 908-910.	1.9	30
3	Optimizing the identification of risk-relevant mutations by multigene panel testing in selected hereditary breast/ovarian cancer families. <i>Cancer Medicine</i> , 2018, 7, 46-55.	2.8	28
4	PRDX1 and PRDX6 are repressed in papillary thyroid carcinomas via BRAF V600E-dependent and -independent mechanisms. <i>International Journal of Oncology</i> , 2014, 44, 548-556.	3.3	27
5	Next-generation sequencing of <i>BRCA1</i> and <i>BRCA2</i> genes for rapid detection of germline mutations in hereditary breast/ovarian cancer. <i>PeerJ</i> , 2019, 7, e6661.	2.0	21
6	A Simplified Genomic Profiling Approach Predicts Outcome in Metastatic Colorectal Cancer. <i>Cancers</i> , 2019, 11, 147.	3.7	15
7	Novel and recurrent BRCA2 mutations in Italian breast/ovarian cancer families widen the ovarian cancer cluster region boundaries to exons 13 and 14. <i>Breast Cancer Research and Treatment</i> , 2014, 148, 629-635.	2.5	12
8	Clinical Multigene Panel Sequencing Identifies Distinct Mutational Association Patterns in Metastatic Colorectal Cancer. <i>Frontiers in Oncology</i> , 2020, 10, 560.	2.8	12
9	Detection of ATM germline variants by the p53 mitotic centrosomal localization test in BRCA1/2-negative patients with early-onset breast cancer. <i>Journal of Experimental and Clinical Cancer Research</i> , 2016, 35, 135.	8.6	9
10	CTNNB1 p.L31P mutation in an ovarian endometrioid carcinoma with synchronous uterine endometrioid carcinoma. <i>Pathology Research and Practice</i> , 2020, 216, 153260.	2.3	2
11	Identification of novel <i>BRCA1</i> large genomic rearrangements by a computational algorithm of amplicon-based Next-Generation Sequencing data. <i>PeerJ</i> , 2019, 7, e7972.	2.0	2
12	A novel <i>BRCA2</i> splice variant identified in a young woman. <i>Molecular Genetics & Genomic Medicine</i> , 2020, 8, e1513.	1.2	1