## Barbara Pasculli

List of Publications by Year in descending order

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516215 794141 1,282 32 16 19 citations h-index g-index papers 33 33 33 2671 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	MicroRNAome genome: A treasure for cancer diagnosis and therapy. Ca-A Cancer Journal for Clinicians, 2014, 64, 311-336.	157.7	428
2	Allele-Specific Reprogramming of Cancer Metabolism by the Long Non-coding RNA CCAT2. Molecular Cell, 2016, 61, 520-534.	4.5	142
3	Epigenetics of breast cancer: Biology and clinical implication in the era of precision medicine. Seminars in Cancer Biology, 2018, 51, 22-35.	4.3	115
4	N-BLR, a primate-specific non-coding transcript leads to colorectal cancer invasion and migration. Genome Biology, 2017, 18, 98.	3.8	97
5	Aberrant $\langle i \rangle$ Keap $1 \langle i \rangle$ methylation in breast cancer and association with clinicopathological features. Epigenetics, 2013, 8, 105-112.	1.3	77
6	A MiRNA Signature for Defining Aggressive Phenotype and Prognosis in Gliomas. PLoS ONE, 2014, 9, e108950.	1.1	60
7	MiR-1287-5p inhibits triple negative breast cancer growth by interaction with phosphoinositide 3-kinase CB, thereby sensitizing cells for PI3Kinase inhibitors. Breast Cancer Research, 2019, 21, 20.	2.2	52
8	Stepwise analysis of MIR9 loci identifies miR-9-5p to be involved in Oestrogen regulated pathways in breast cancer patients. Scientific Reports, 2017, 7, 45283.	1.6	45
9	Evaluation of microRNA-10b prognostic significance in a prospective cohort of breast cancer patients. Molecular Cancer, 2014, 13, 142.	7.9	40
10	Targeting the microRNA-regulating DNA damage/repair pathways in cancer. Expert Opinion on Biological Therapy, 2014, 14, 1667-1683.	1.4	36
11	Hsa-miR-155-5p Up-Regulation in Breast Cancer and Its Relevance for Treatment With Poly[ADP-Ribose] Polymerase 1 (PARP-1) Inhibitors. Frontiers in Oncology, 2020, 10, 1415.	1.3	31
12	Competitive allele-specific TaqMan PCR (Cast-PCR) is a sensitive, specific and fast method for BRAF V600 mutation detection in Melanoma patients. Scientific Reports, 2015, 5, 18592.	1.6	27
13	Can Epigenetics of Endothelial Dysfunction Represent the Key to Precision Medicine in Type 2 Diabetes Mellitus?. International Journal of Molecular Sciences, 2019, 20, 2949.	1.8	27
14	Combined analysis of miR-200 family and its significance for breast cancer. Scientific Reports, 2021, 11, 2980.	1.6	22
15	Hsa-miR-210-3p expression in breast cancer and its putative association with worse outcome in patients treated with Docetaxel. Scientific Reports, 2019, 9, 14913.	1.6	19
16	ALYREF, a novel factor involved in breast carcinogenesis, acts through transcriptional and post-transcriptional mechanisms selectively regulating the short NEAT1 isoform. Cellular and Molecular Life Sciences, 2022, 79, .	2.4	17
17	Carbonic Anhydrase XII Expression Is Modulated during Epithelial Mesenchymal Transition and Regulated through Protein Kinase C Signaling. International Journal of Molecular Sciences, 2020, 21, 715.	1.8	12
18	Evaluation of pre-analytical procedures for the detection of BRAF V600 mutations in melanoma patients: comparison between Sanger sequencing and Competitive allele-specific TaqMan PCR (Cast-PCR). European Journal of Cancer, 2016, 61, S127-S128.	1.3	1

#	Article	IF	CITATIONS
19	Abstract 1479: A miRNA signature distinguishing low-grade and high-grade gliomas shows miR-21 and 210 as promising biomarkers of aggressive phenotype and prognosis. , 2014, , .		1
20	Abstract 4734: miR-9-5p expression in breast cancer correlates with hormone receptor status and affects patients survival. , $2017$ , , .		1
21	602 Frequent Epigenetic Inactivation of KEAP1 Gene in Breast Cancer. European Journal of Cancer, 2012, 48, S143.	1.3	O
22	Predictive Value of Epigenetic Signatures. , 2018, , 275-311.		0
23	Abstract 664: Aberrant KEAP1 promoter methylation is associated with disease progression in breast cancer patients treated with epirubicin/cyclophosfamide and docetaxel chemotherapy, 2013, , .		O
24	Evaluation of microRNA-10b expression as a novel predictive marker of metastases development and patients' survival in breast cancer Journal of Clinical Oncology, 2013, 31, 576-576.	0.8	0
25	Hypermethylation of the KEAP1 gene in colorectal cancer and association with disease progression Journal of Clinical Oncology, 2013, 31, e14655-e14655.	0.8	O
26	Abstract 1477: Evaluation of microRNA-10b prognostic significance in a prospective cohort of breast cancer patients., 2014,,.		0
27	Abstract 2251: nrf2-keap1 axis molecular profile in small cell lung cancer cell lines. , 2014, , .		O
28	Abstract 3977: Evaluation of miR10b and miR9 expression in breast cancer and correlations with distant metastases development., 2015,,.		0
29	Abstract 5394: Initial results from TRANSCAN ERA-NET BREMIR project: MicroRNAs expression profiling for identification of breast cancer patients at high risk to develop distant metastases. , 2018, , .		O
30	Abstract 4904: High levels of microRNA-210-3p are associated with increased risk of disease progression in breast cancer patients treated with docetaxel., 2019,,.		0
31	Abstract 1422: Clinical association of miR-155-5p with breast cancer and its relevance for treatment with PARP inhibitors. , 2020, , .		0
32	Abstract 4904: High levels of microRNA-210-3p are associated with increased risk of disease progression in breast cancer patients treated with docetaxel., 2019,,.		O