

Angelo Veronese

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.

65
papers

9,663
citations

36
h-index

71
g-index

71
ext. papers

10,421
ext. citations

9.3
avg, IF

5.16
L-index

#	Paper	IF	Citations
65	MicroRNA gene expression deregulation in human breast cancer. <i>Cancer Research</i> , 2005 , 65, 7065-70	10.1	3315
64	Cyclin G1 is a target of miR-122a, a microRNA frequently down-regulated in human hepatocellular carcinoma. <i>Cancer Research</i> , 2007 , 67, 6092-9	10.1	695
63	MiR-221 controls CDKN1C/p57 and CDKN1B/p27 expression in human hepatocellular carcinoma. <i>Oncogene</i> , 2008 , 27, 5651-61	9.2	545
62	p53 regulates epithelial-mesenchymal transition through microRNAs targeting ZEB1 and ZEB2. <i>Journal of Experimental Medicine</i> , 2011 , 208, 875-83	16.6	423
61	MiR-122/cyclin G1 interaction modulates p53 activity and affects doxorubicin sensitivity of human hepatocarcinoma cells. <i>Cancer Research</i> , 2009 , 69, 5761-7	10.1	346
60	Reprogramming of miRNA networks in cancer and leukemia. <i>Genome Research</i> , 2010 , 20, 589-99	9.7	287
59	Modulation of mismatch repair and genomic stability by miR-155. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010 , 107, 6982-7	11.5	267
58	MicroRNA-221 targets Bmf in hepatocellular carcinoma and correlates with tumor multifocality. <i>Clinical Cancer Research</i> , 2009 , 15, 5073-81	12.9	267
57	Oncogenic role of miR-483-3p at the IGF2/483 locus. <i>Cancer Research</i> , 2010 , 70, 3140-9	10.1	239
56	MicroRNA fingerprints identify miR-150 as a plasma prognostic marker in patients with sepsis. <i>PLoS ONE</i> , 2009 , 4, e7405	3.7	236
55	MicroRNA-135b promotes cancer progression by acting as a downstream effector of oncogenic pathways in colon cancer. <i>Cancer Cell</i> , 2014 , 25, 469-83	24.3	235
54	mRNA/microRNA gene expression profile in microsatellite unstable colorectal cancer. <i>Molecular Cancer</i> , 2007 , 6, 54	42.1	215
53	Micromarkers: miRNAs in cancer diagnosis and prognosis. <i>Expert Review of Molecular Diagnostics</i> , 2010 , 10, 297-308	3.8	207
52	miR-145 participates with TP53 in a death-promoting regulatory loop and targets estrogen receptor-alpha in human breast cancer cells. <i>Cell Death and Differentiation</i> , 2010 , 17, 246-54	12.7	205
51	miR-130a targets MET and induces TRAIL-sensitivity in NSCLC by downregulating miR-221 and 222. <i>Oncogene</i> , 2012 , 31, 634-42	9.2	160
50	Karyotype-specific microRNA signature in chronic lymphocytic leukemia. <i>Blood</i> , 2009 , 114, 3872-9	2.2	159
49	NUP98 is fused to the NSD3 gene in acute myeloid leukemia associated with t(8;11)(p11.2;p15). <i>Blood</i> , 2002 , 99, 3857-60	2.2	148

48	Identification of NUP98 abnormalities in acute leukemia: JARID1A (12p13) as a new partner gene. <i>Genes Chromosomes and Cancer</i> , 2006 , 45, 437-46	5	112
47	miR-181b is a biomarker of disease progression in chronic lymphocytic leukemia. <i>Blood</i> , 2011 , 118, 3072-9.2		103
46	MicroRNA profiling for the identification of cancers with unknown primary tissue-of-origin. <i>Journal of Pathology</i> , 2011 , 225, 43-53	9.4	101
45	Epigenetics and MicroRNAs in Cancer. <i>International Journal of Molecular Sciences</i> , 2018 , 19,	6.3	94
44	Chronic lymphocytic leukemia with 6q- shows distinct hematological features and intermediate prognosis. <i>Leukemia</i> , 2004 , 18, 476-83	10.7	93
43	Frequent aberrant methylation of the CDH4 gene promoter in human colorectal and gastric cancer. <i>Cancer Research</i> , 2004 , 64, 8156-9	10.1	89
42	MicroRNAs involvement in fludarabine refractory chronic lymphocytic leukemia. <i>Molecular Cancer</i> , 2010 , 9, 123	42.1	87
41	Gain of imprinting at chromosome 11p15: A pathogenetic mechanism identified in human hepatocarcinomas. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000 , 97, 5445-9	11.5	74
40	MicroRNAs dysregulation in human malignant pleural mesothelioma. <i>Journal of Thoracic Oncology</i> , 2011 , 6, 844-51	8.9	72
39	Wnt signalling modulates transcribed-ultraconserved regions in hepatobiliary cancers. <i>Gut</i> , 2017 , 66, 1268-1277	19.2	58
38	miR-205-5p-mediated downregulation of ErbB/HER receptors in breast cancer stem cells results in targeted therapy resistance. <i>Cell Death and Disease</i> , 2015 , 6, e1823	9.8	55
37	Transcriptional map of 170-kb region at chromosome 11p15.5: identification and mutational analysis of the BWR1A gene reveals the presence of mutations in tumor samples. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1998 , 95, 3873-8	11.5	54
36	Nidogen 1 and 2 gene promoters are aberrantly methylated in human gastrointestinal cancer. <i>Molecular Cancer</i> , 2007 , 6, 17	42.1	52
35	Loss of methylation at chromosome 11p15.5 is common in human adult tumors. <i>Oncogene</i> , 2002 , 21, 2564-72	9.2	49
34	The methylator phenotype in microsatellite stable colorectal cancers is characterized by a distinct gene expression profile. <i>Journal of Pathology</i> , 2008 , 214, 594-602	9.4	45
33	Mutated beta-catenin evades a microRNA-dependent regulatory loop. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 4840-5	11.5	43
32	Exon structure and promoter identification of STIM1 (alias GOK), a human gene causing growth arrest of the human tumor cell lines G401 and RD. <i>Cytogenetic and Genome Research</i> , 1999 , 86, 214-8	1.9	43
31	Non-coding RNAs in the reprogramming of glucose metabolism in cancer. <i>Cancer Letters</i> , 2018 , 419, 167-174	17.4	42

30	MiR-181b: new perspective to evaluate disease progression in chronic lymphocytic leukemia. <i>Oncotarget</i> , 2012 , 3, 195-202	3.3	41
29	p53/mdm2 feedback loop sustains miR-221 expression and dictates the response to anticancer treatments in hepatocellular carcinoma. <i>Molecular Cancer Research</i> , 2014 , 12, 203-16	6.6	36
28	Over-expression of the miR-483-3p overcomes the miR-145/TP53 pro-apoptotic loop in hepatocellular carcinoma. <i>Oncotarget</i> , 2016 , 7, 31361-71	3.3	33
27	Multigene Methylation Analysis of Gastrointestinal Tumors. <i>Molecular Diagnosis and Therapy</i> , 2003 , 7, 201-207		32
26	Cryptic insertion producing two NUP98/NSD1 chimeric transcripts in adult refractory anemia with an excess of blasts. <i>Genes Chromosomes and Cancer</i> , 2004 , 41, 395-9	5	30
25	A novel miR-371a-5p-mediated pathway, leading to BAG3 upregulation in cardiomyocytes in response to epinephrine, is lost in Takotsubo cardiomyopathy. <i>Cell Death and Disease</i> , 2015 , 6, e1948	9.8	29
24	Regulation of miR-483-3p by the O-linked N-acetylglucosamine transferase links chemosensitivity to glucose metabolism in liver cancer cells. <i>Oncogenesis</i> , 2017 , 6, e328	6.6	27
23	Air and surface measurements of SARS-CoV-2 inside a bus during normal operation. <i>PLoS ONE</i> , 2020 , 15, e0235943	3.7	25
22	Integrative genetic, epigenetic and pathological analysis of paraganglioma reveals complex dysregulation of NOTCH signaling. <i>Acta Neuropathologica</i> , 2013 , 126, 575-94	14.3	22
21	Allele-specific loss and transcription of the miR-15a/16-1 cluster in chronic lymphocytic leukemia. <i>Leukemia</i> , 2015 , 29, 86-95	10.7	22
20	The Glucose-Regulated Influences Key Signaling Pathways in Cancer. <i>Cancers</i> , 2018 , 10,	6.6	20
19	MicroRNAs in Autoimmunity and Hematological Malignancies. <i>International Journal of Molecular Sciences</i> , 2018 , 19,	6.3	19
18	Multigene methylation analysis of gastrointestinal tumors: TPEF emerges as a frequent tumor-specific aberrantly methylated marker that can be detected in peripheral blood. <i>Molecular Diagnosis and Therapy</i> , 2003 , 7, 201-7		17
17	Impact of primary tumor location in patients with RAS wild-type metastatic colon cancer treated with first-line chemotherapy plus anti-EGFR or anti-VEGF monoclonal antibodies: a retrospective multicenter study. <i>Journal of Cancer</i> , 2019 , 10, 5926-5934	4.5	16
16	DNA methylation of shelf, shore and open sea CpG positions distinguish high microsatellite instability from low or stable microsatellite status colon cancer stem cells. <i>Epigenomics</i> , 2019 , 11, 587-604	4.4	12
15	Paragangliomas arise through an autonomous vasculo-angio-neurogenic program inhibited by imatinib. <i>Acta Neuropathologica</i> , 2018 , 135, 779-798	14.3	12
14	Genome wide identification of recessive cancer genes by combinatorial mutation analysis. <i>PLoS ONE</i> , 2008 , 3, e3380	3.7	10
13	A Developmental Perspective on Paragangliar Tumorigenesis. <i>Cancers</i> , 2019 , 11,	6.6	9

12	drives aneuploidy at early stages of cellular transformation. <i>Oncotarget</i> , 2018 , 9, 13036-13047	3.3	9
11	HNRNPL Restrains Targeting of BUB1 to Stabilize Aberrant Karyotypes of Transformed Cells in Chronic Lymphocytic Leukemia. <i>Cancers</i> , 2019 , 11,	6.6	7
10	Anticancer activity of an adenoviral vector expressing short hairpin RNA against BK virus T-ag. <i>Cancer Gene Therapy</i> , 2007 , 14, 297-305	5.4	7
9	Enhanced Expression of in B Cells of CLL Improves the Anti-Tumor Cytotoxic T Cell Response. <i>Cancers</i> , 2021 , 13,	6.6	4
8	Genetic dynamics in untreated CLL patients with either stable or progressive disease: a longitudinal study. <i>Journal of Hematology and Oncology</i> , 2019 , 12, 114	22.4	3
7	Tgf- β transcriptionally promotes 90K expression: possible implications for cancer progression. <i>Cell Death Discovery</i> , 2021 , 7, 86	6.9	2
6	A perspective analysis: microRNAs, glucose metabolism, and drug resistance in colon cancer stem cells. <i>Cancer Gene Therapy</i> , 2021 ,	5.4	2
5	MiR-181b in Chronic Lymphocytic Leukemia B Cells Is Regulated By Cellular Interaction with CD4+ T Cells and Increases the CTL Toxicity Versus the Leukemic Clone. <i>Blood</i> , 2015 , 126, 4134-4134	2.2	
4	Impact of BCR Stimulation on Mir-181b in Chronic Lymphocytic Leukemia. <i>Blood</i> , 2016 , 128, 2026-2026	2.2	
3	p53 regulates epithelial-mesenchymal transition through microRNAs targeting ZEB1 and ZEB2. <i>Journal of Cell Biology</i> , 2011 , 193, i8-i8	7.3	
2	Allele-Specific Loss Of The Mir-15a/16-1 Cluster Correlates With ZAP70 Expression In CLL Patients With 13q Deletion. <i>Blood</i> , 2013 , 122, 3753-3753	2.2	
1	Pathophysiology roles and translational opportunities of miRNAs in CLL 2022 , 179-186		