

# Angelo Veronese

## List of Publications by Year in descending order

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Version: 2024-02-01

69  
papers

11,041  
citations

87723

38  
h-index

118652

62  
g-index

71  
all docs

71  
docs citations

71  
times ranked

15668  
citing authors

#	ARTICLE	IF	CITATIONS
1	MicroRNA Gene Expression Deregulation in Human Breast Cancer. <i>Cancer Research</i> , 2005, 65, 7065-7070.	0.4	3,719
2	Cyclin G1 Is a Target of miR-122a, a MicroRNA Frequently Down-regulated in Human Hepatocellular Carcinoma. <i>Cancer Research</i> , 2007, 67, 6092-6099.	0.4	782
3	MiR-221 controls CDKN1C/p57 and CDKN1B/p27 expression in human hepatocellular carcinoma. <i>Oncogene</i> , 2008, 27, 5651-5661.	2.6	619
4	p53 regulates epithelial-mesenchymal transition through microRNAs targeting ZEB1 and ZEB2. <i>Journal of Experimental Medicine</i> , 2011, 208, 875-883.	4.2	480
5	MiR-122/Cyclin G1 Interaction Modulates p53 Activity and Affects Doxorubicin Sensitivity of Human Hepatocarcinoma Cells. <i>Cancer Research</i> , 2009, 69, 5761-5767.	0.4	380
6	Reprogramming of miRNA networks in cancer and leukemia. <i>Genome Research</i> , 2010, 20, 589-599.	2.4	331
7	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, 6982-6987.	3.3	306
8	MicroRNA-221 Targets Bmf in Hepatocellular Carcinoma and Correlates with Tumor Multifocality. <i>Clinical Cancer Research</i> , 2009, 15, 5073-5081.	3.2	298
9	MicroRNA Fingerprints Identify miR-150 as a Plasma Prognostic Marker in Patients with Sepsis. <i>PLoS ONE</i> , 2009, 4, e7405.	1.1	273
10	Oncogenic Role of miR-483-3p at the IGF2/483 Locus. <i>Cancer Research</i> , 2010, 70, 3140-3149.	0.4	272
11	MicroRNA-135b Promotes Cancer Progression by Acting as a Downstream Effector of Oncogenic Pathways in Colon Cancer. <i>Cancer Cell</i> , 2014, 25, 469-483.	7.7	267
12	mRNA/microRNA gene expression profile in microsatellite unstable colorectal cancer. <i>Molecular Cancer</i> , 2007, 6, 54.	7.9	240
13	Micromarkers: miRNAs in cancer diagnosis and prognosis. <i>Expert Review of Molecular Diagnostics</i> , 2010, 10, 297-308.	1.5	237
14	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, 246-254.	5.0	231
15	miR-130a targets MET and induces TRAIL-sensitivity in NSCLC by downregulating miR-221 and 222. <i>Oncogene</i> , 2012, 31, 634-642.	2.6	181
16	Karyotype-specific microRNA signature in chronic lymphocytic leukemia. <i>Blood</i> , 2009, 114, 3872-3879.	0.6	179
17	NUP98 is fused to the NSD3 gene in acute myeloid leukemia associated with t(8;11)(p11.2;p15). <i>Blood</i> , 2002, 99, 3857-3860.	0.6	176
18	Epigenetics and MicroRNAs in Cancer. <i>International Journal of Molecular Sciences</i> , 2018, 19, 459.	1.8	135

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19	Identification of NUP98 abnormalities in acute leukemia:JARID1A (12p13) as a new partner gene. <i>Genes Chromosomes and Cancer</i> , 2006, 45, 437-446.	1.5	123
20	MicroRNA profiling for the identification of cancers with unknown primary tissue of origin. <i>Journal of Pathology</i> , 2011, 225, 43-53.	2.1	117
21	miR-181b is a biomarker of disease progression in chronic lymphocytic leukemia. <i>Blood</i> , 2011, 118, 3072-3079.	0.6	115
22	MicroRNAs involvement in fludarabine refractory chronic lymphocytic leukemia. <i>Molecular Cancer</i> , 2010, 9, 123.	7.9	107
23	Chronic lymphocytic leukemia with 6q <sup>+</sup> shows distinct hematological features and intermediate prognosis. <i>Leukemia</i> , 2004, 18, 476-483.	3.3	99
24	Frequent Aberrant Methylation of the CDH4 Gene Promoter in Human Colorectal and Gastric Cancer. <i>Cancer Research</i> , 2004, 64, 8156-8159.	0.4	96
25	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, 5445-5449.	3.3	81
26	MicroRNAs Dysregulation in Human Malignant Pleural Mesothelioma. <i>Journal of Thoracic Oncology</i> , 2011, 6, 844-851.	0.5	77
27	Wnt signalling modulates transcribed-ultraconserved regions in hepatobiliary cancers. <i>Gut</i> , 2017, 66, 1268-1277.	6.1	75
28	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-e1823.	2.7	74
29	Nidogen 1 and 2 gene promoters are aberrantly methylated in human gastrointestinal cancer. <i>Molecular Cancer</i> , 2007, 6, 17.	7.9	64
30	Non-coding RNAs in the reprogramming of glucose metabolism in cancer. <i>Cancer Letters</i> , 2018, 419, 167-174.	3.2	60
31	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, 3873-3878.	3.3	56
32	Loss of methylation at chromosome 11p15.5 is common in human adult tumors. <i>Oncogene</i> , 2002, 21, 2564-2572.	2.6	52
33	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, 214-218.	0.6	50
34	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, 4840-4845.	3.3	48
35	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.	2.1	47
36	MIR-181b: new perspective to evaluate disease progression in chronic lymphocytic leukemia. <i>Oncotarget</i> , 2012, 3, 195-202.	0.8	46

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37	Over-expression of the <i>miR-483-3p</i> overcomes the miR-145/TP53 pro-apoptotic loop in hepatocellular carcinoma. <i>Oncotarget</i> , 2016, 7, 31361-31371.	0.8	45
38	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, 203-216.	1.5	43
39	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-e328.	2.1	36
40	Air and surface measurements of SARS-CoV-2 inside a bus during normal operation. <i>PLoS ONE</i> , 2020, 15, e0235943.	1.1	36
41	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-e1948.	2.7	35
42	The Glucose-Regulated MiR-483-3p Influences Key Signaling Pathways in Cancer. <i>Cancers</i> , 2018, 10, 181.	1.7	35
43	Cryptic insertion producing twoNUP98/NSD1 chimeric transcripts in adult refractory anemia with an excess of blasts. <i>Genes Chromosomes and Cancer</i> , 2004, 41, 395-399.	1.5	34
44	Multigene Methylation Analysis of Gastrointestinal Tumors. <i>Molecular Diagnosis and Therapy</i> , 2003, 7, 201-207.	1.3	33
45	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.	1.0	29
46	Integrative genetic, epigenetic and pathological analysis of paraganglioma reveals complex dysregulation of NOTCH signaling. <i>Acta Neuropathologica</i> , 2013, 126, 575-594.	3.9	27
47	Allele-specific loss and transcription of the miR-15a/16-1 cluster in chronic lymphocytic leukemia. <i>Leukemia</i> , 2015, 29, 86-95.	3.3	27
48	MicroRNAs in Autoimmunity and Hematological Malignancies. <i>International Journal of Molecular Sciences</i> , 2018, 19, 3139.	1.8	26
49	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.	1.2	24
50	Paragangliomas arise through an autonomous vasculo-angio-neurogenic program inhibited by imatinib. <i>Acta Neuropathologica</i> , 2018, 135, 779-798.	3.9	20
51	Multigene Methylation Analysis of Gastrointestinal Tumors. <i>Molecular Diagnosis and Therapy</i> , 2003, 7, 201-207.	1.3	18
52	Genome Wide Identification of Recessive Cancer Genes by Combinatorial Mutation Analysis. <i>PLoS ONE</i> , 2008, 3, e3380.	1.1	12
53	<i>Hsa-miR-155-5p</i> drives aneuploidy at early stages of cellular transformation. <i>Oncotarget</i> , 2018, 9, 13036-13047.	0.8	12
54	HNRNPL Restrains miR-155 Targeting of BUB1 to Stabilize Aberrant Karyotypes of Transformed Cells in Chronic Lymphocytic Leukemia. <i>Cancers</i> , 2019, 11, 575.	1.7	11

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55	A Developmental Perspective on Paragangliar Tumorigenesis. <i>Cancers</i> , 2019, 11, 273.	1.7	11
56	Enhanced Expression of miR-181b in B Cells of CLL Improves the Anti-Tumor Cytotoxic T Cell Response. <i>Cancers</i> , 2021, 13, 257.	1.7	10
57	Anticancer activity of an adenoviral vector expressing short hairpin RNA against BK virus T-ag. <i>Cancer Gene Therapy</i> , 2007, 14, 297-305.	2.2	7
58	A perspective analysis: microRNAs, glucose metabolism, and drug resistance in colon cancer stem cells. <i>Cancer Gene Therapy</i> , 2021, , .	2.2	6
59	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.	6.9	5
60	Genome DNA methylation, aneuploidy and immunity in cancer. <i>Epigenomics</i> , 2020, 12, .	1.0	3
61	Tgf- $\beta$ 1 transcriptionally promotes 90K expression: possible implications for cancer progression. <i>Cell Death Discovery</i> , 2021, 7, 86.	2.0	2
62	Abstract 3313: Epigenetic biomarkers of prognosis in stage IIA colon cancer. <i>Cancer Research</i> , 2018, 78, 3313-3313.	0.4	1
63	Abstract 4051: MiR-181b expression levels decreases during the progression of the Chronic Lymphocytic Leukemia: a new potential prognostic tool. , 2010, , .		0
64	Abstract 2087: miR-483-3p is an oncogene involved in nephroblastoma and in adult tumors with activated $\beta$ -catenin. , 2010, , .		0
65	p53 regulates epithelial- $\rightarrow$ mesenchymal transition through microRNAs targeting ZEB1 and ZEB2. <i>Journal of Cell Biology</i> , 2011, 193, i8-i8.	2.3	0
66	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.	0.6	0
67	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.	0.6	0
68	Impact of BCR Stimulation on Mir-181b in Chronic Lymphocytic Leukemia. <i>Blood</i> , 2016, 128, 2026-2026.	0.6	0
69	Pathophysiology roles and translational opportunities of miRNAs in CLL. , 2022, , 179-186.		0