

Angelo Veronese

List of Publications by Year in Descending Order

Source: <https://exaly.com/author-pdf/3090093/angelo-veronese-publications-by-year.pdf>

Version: 2024-04-25

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

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	Pathophysiology roles and translational opportunities of miRNAs in CLL 2022 , 179-186		
64	Tgf- β transcriptionally promotes 90K expression: possible implications for cancer progression. <i>Cell Death Discovery</i> , 2021 , 7, 86	6.9	2
63	Enhanced Expression of in B Cells of CLL Improves the Anti-Tumor Cytotoxic T Cell Response. <i>Cancers</i> , 2021 , 13,	6.6	4
62	A perspective analysis: microRNAs, glucose metabolism, and drug resistance in colon cancer stem cells. <i>Cancer Gene Therapy</i> , 2021 ,	5.4	2
61	Air and surface measurements of SARS-CoV-2 inside a bus during normal operation. <i>PLoS ONE</i> , 2020 , 15, e0235943	3.7	25
60	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
59	HNRNPL Restrains Targeting of BUB1 to Stabilize Aberrant Karyotypes of Transformed Cells in Chronic Lymphocytic Leukemia. <i>Cancers</i> , 2019 , 11,	6.6	7
58	A Developmental Perspective on Paragangliar Tumorigenesis. <i>Cancers</i> , 2019 , 11,	6.6	9
57	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
56	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
55	Non-coding RNAs in the reprogramming of glucose metabolism in cancer. <i>Cancer Letters</i> , 2018 , 419, 167-174	4.4	42
54	Paragangliomas arise through an autonomous vasculo-angio-neurogenic program inhibited by imatinib. <i>Acta Neuropathologica</i> , 2018 , 135, 779-798	14.3	12
53	The Glucose-Regulated Influences Key Signaling Pathways in Cancer. <i>Cancers</i> , 2018 , 10,	6.6	20
52	Epigenetics and MicroRNAs in Cancer. <i>International Journal of Molecular Sciences</i> , 2018 , 19,	6.3	94
51	drives aneuploidy at early stages of cellular transformation. <i>Oncotarget</i> , 2018 , 9, 13036-13047	3.3	9
50	MicroRNAs in Autoimmunity and Hematological Malignancies. <i>International Journal of Molecular Sciences</i> , 2018 , 19,	6.3	19
49	Wnt signalling modulates transcribed-ultraconserved regions in hepatobiliary cancers. <i>Gut</i> , 2017 , 66, 1268-1277	19.2	58

48	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
47	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
46	Impact of BCR Stimulation on Mir-181b in Chronic Lymphocytic Leukemia. <i>Blood</i> , 2016 , 128, 2026-2026	2.2	
45	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
44	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
43	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
42	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	
41	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
40	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
39	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
38	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	
37	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
36	MIR-181b: new perspective to evaluate disease progression in chronic lymphocytic leukemia. <i>Oncotarget</i> , 2012 , 3, 195-202	3.3	41
35	p53 regulates epithelial-mesenchymal transition through microRNAs targeting ZEB1 and ZEB2. <i>Journal of Experimental Medicine</i> , 2011 , 208, 875-83	16.6	423
34	miR-181b is a biomarker of disease progression in chronic lymphocytic leukemia. <i>Blood</i> , 2011 , 118, 3072-2	2.2	103
33	MicroRNAs dysregulation in human malignant pleural mesothelioma. <i>Journal of Thoracic Oncology</i> , 2011 , 6, 844-51	8.9	72
32	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
31	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

30	p53 regulates epithelial-mesenchymal transition through microRNAs targeting ZEB1 and ZEB2. <i>Journal of Cell Biology</i> , 2011 , 193, i8-i8	7.3	
29	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
28	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
27	Oncogenic role of miR-483-3p at the IGF2/483 locus. <i>Cancer Research</i> , 2010 , 70, 3140-9	10.1	239
26	MicroRNAs involvement in fludarabine refractory chronic lymphocytic leukemia. <i>Molecular Cancer</i> , 2010 , 9, 123	42.1	87
25	Micromarkers: miRNAs in cancer diagnosis and prognosis. <i>Expert Review of Molecular Diagnostics</i> , 2010 , 10, 297-308	3.8	207
24	Reprogramming of miRNA networks in cancer and leukemia. <i>Genome Research</i> , 2010 , 20, 589-99	9.7	287
23	MicroRNA fingerprints identify miR-150 as a plasma prognostic marker in patients with sepsis. <i>PLoS ONE</i> , 2009 , 4, e7405	3.7	236
22	MicroRNA-221 targets Bmf in hepatocellular carcinoma and correlates with tumor multifocality. <i>Clinical Cancer Research</i> , 2009 , 15, 5073-81	12.9	267
21	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
20	Karyotype-specific microRNA signature in chronic lymphocytic leukemia. <i>Blood</i> , 2009 , 114, 3872-9	2.2	159
19	MiR-221 controls CDKN1C/p57 and CDKN1B/p27 expression in human hepatocellular carcinoma. <i>Oncogene</i> , 2008 , 27, 5651-61	9.2	545
18	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
17	Genome wide identification of recessive cancer genes by combinatorial mutation analysis. <i>PLoS ONE</i> , 2008 , 3, e3380	3.7	10
16	Nidogen 1 and 2 gene promoters are aberrantly methylated in human gastrointestinal cancer. <i>Molecular Cancer</i> , 2007 , 6, 17	42.1	52
15	mRNA/microRNA gene expression profile in microsatellite unstable colorectal cancer. <i>Molecular Cancer</i> , 2007 , 6, 54	42.1	215
14	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
13	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

12	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
11	MicroRNA gene expression deregulation in human breast cancer. <i>Cancer Research</i> , 2005 , 65, 7065-70	10.1	3315
10	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
9	Chronic lymphocytic leukemia with 6q- shows distinct hematological features and intermediate prognosis. <i>Leukemia</i> , 2004 , 18, 476-83	10.7	93
8	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
7	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
6	Multigene Methylation Analysis of Gastrointestinal Tumors. <i>Molecular Diagnosis and Therapy</i> , 2003 , 7, 201-207		32
5	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
4	Loss of methylation at chromosome 11p15.5 is common in human adult tumors. <i>Oncogene</i> , 2002 , 21, 2564-72	9.2	49
3	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
2	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
1	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