## Francesco Nicassio

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5245763/publications.pdf

Version: 2024-02-01

38 papers

2,386 citations

279798 23 h-index 39 g-index

43 all docs 43 docs citations

43 times ranked 4512 citing authors

#	Article	IF	CITATIONS
1	A serum circulating miRNA diagnostic test to identify asymptomatic highâ€risk individuals with early stage lung cancer. EMBO Molecular Medicine, 2011, 3, 495-503.	6.9	322
2	Human USP3 Is a Chromatin Modifier Required for S Phase Progression and Genome Stability. Current Biology, 2007, 17, 1972-1977.	3.9	251
3	miR-Test: A Blood Test for Lung Cancer Early Detection. Journal of the National Cancer Institute, 2015, 107, djv063.	6.3	221
4	Np95 Is a Histone-Binding Protein Endowed with Ubiquitin Ligase Activity. Molecular and Cellular Biology, 2004, 24, 2526-2535.	2.3	174
5	Degradation dynamics of microRNAs revealed by a novel pulse-chase approach. Genome Research, 2016, 26, 554-565.	5.5	155
6	Endogenous transcripts control miRNA levels and activity in mammalian cells by target-directed miRNA degradation. Nature Communications, 2018, 9, 3119.	12.8	121
7	Survival prediction of stage I lung adenocarcinomas by expression of 10 genes. Journal of Clinical Investigation, 2007, 117, 3436-3444.	8.2	103
8	Np95 is regulated by E1A during mitotic reactivation of terminally differentiated cells and is essential for S phase entry. Journal of Cell Biology, 2002, 157, 909-914.	5,2	86
9	DEPDC1B Coordinates De-adhesion Events and Cell-Cycle Progression at Mitosis. Developmental Cell, 2014, 31, 420-433.	7.0	76
10	Microglia-specific overexpression of $\hat{l}_{\pm}$ -synuclein leads to severe dopaminergic neurodegeneration by phagocytic exhaustion and oxidative toxicity. Nature Communications, 2021, 12, 6237.	12.8	74
11	The many faces of ubiquitinated histone H2A: insights from the DUBs. Cell Division, 2008, 3, 8.	2.4	68
12	LncRNA EPR controls epithelial proliferation by coordinating Cdkn1a transcription and mRNA decay response to TGF- $\hat{l}^2$ . Nature Communications, 2019, 10, 1969.	12.8	68
13	IsomiRage: From Functional Classification to Differential Expression of miRNA Isoforms. Frontiers in Bioengineering and Biotechnology, 2014, 2, 38.	4.1	61
14	Optimization and Standardization of Circulating MicroRNA Detection for Clinical Application: The miR-Test Case. Clinical Chemistry, 2016, 62, 743-754.	3.2	53
15	Synergic Functions of miRNAs Determine Neuronal Fate of Adult Neural Stem Cells. Stem Cell Reports, 2017, 8, 1046-1061.	4.8	49
16	MicroRNAâ€independent functions of DGCR8 are essential for neocortical development and TBR1 expression. EMBO Reports, 2017, 18, 603-618.	<b>4.</b> 5	47
17	p53 Loss in Breast Cancer Leads to Myc Activation, Increased Cell Plasticity, and Expression of a Mitotic Signature with Prognostic Value. Cell Reports, 2019, 26, 624-638.e8.	6.4	47
18	The role of non-coding RNAs in the regulation of stem cells and progenitors in the normal mammary gland and in breast tumors. Frontiers in Genetics, 2015, 6, 72.	2.3	44

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19	Frataxin gene editing rescues Friedreich's ataxia pathology in dorsal root ganglia organoid-derived sensory neurons. Nature Communications, 2020, 11, 4178.	12.8	42
20	Differentiation-associated microRNAs antagonize the Rb–E2F pathway to restrict proliferation. Journal of Cell Biology, 2012, 199, 77-95.	5.2	39
21	Dual role for miR-34a in the control of early progenitor proliferation and commitment in the mammary gland and in breast cancer. Oncogene, 2019, 38, 360-374.	5.9	39
22	MiR-135a-5p Is Critical for Exercise-Induced Adult Neurogenesis. Stem Cell Reports, 2019, 12, 1298-1312.	4.8	37
23	Insc:LGN tetramers promote asymmetric divisions of mammary stem cells. Nature Communications, 2018, 9, 1025.	12.8	27
24	Parent-of-origin genetic background affects the transcriptional levels of circadian and neuronal plasticity genes following sleep loss. Philosophical Transactions of the Royal Society B: Biological Sciences, 2014, 369, 20120471.	4.0	21
25	Recessive Cancer Genes Engage in Negative Genetic Interactions with Their Functional Paralogs. Cell Reports, 2013, 5, 1519-1526.	6.4	19
26	Delivery of biologically active miR-34a in normal and cancer mammary epithelial cells by synthetic nanoparticles. Nanomedicine: Nanotechnology, Biology, and Medicine, 2019, 19, 95-105.	3.3	19
27	MicroRNAs in Cancer Management: Big Challenges for Small Molecules. BioMed Research International, 2015, 2015, 1-2.	1.9	17
28	Prediction and pan-cancer analysis of mammalian transcripts involved in target directed miRNA degradation. Nucleic Acids Research, 2022, 50, 2019-2035.	14.5	16
29	A cancer-specific transcriptional signature in human neoplasia. Journal of Clinical Investigation, 2005, 115, 3015-3025.	8.2	14
30	Unbiased vs. biased approaches to the identification of cancer signatures: the case of lung cancer. Cell Cycle, 2008, 7, 729-734.	2.6	13
31	Nanopore ReCappable sequencing maps SARS-CoV-2 $5\hat{a}\in^2$ capping sites and provides new insights into the structure of sgRNAs. Nucleic Acids Research, 2022, 50, 3475-3489.	14.5	12
32	miR-146 connects stem cell identity with metabolism and pharmacological resistance in breast cancer. Journal of Cell Biology, 2021, 220, .	5.2	8
33	A multifunctional locus controls motor neuron differentiation through short and long noncoding RNAs. EMBO Journal, 2022, 41, .	7.8	8
34	Mining cancer gene expression databases for latent information on intronic microRNAs. Molecular Oncology, 2015, 9, 473-487.	4.6	6
35	Microenvironment Stimuli HGF and Hypoxia Differently Affected miR-125b and Ets-1 Function with Opposite Effects on the Invasiveness of Bone Metastatic Cells: A Comparison with Breast Carcinoma Cells. International Journal of Molecular Sciences, 2018, 19, 258.	4.1	5
36	In silico prediction and experimental validation of natural antisense transcripts in two cancer-associated regions of human chromosome 6. International Journal of Oncology, 2009, 34, 1099-108.	3.3	4

#	Article	lF	CITATIONS
37	Uncovering the Stability of Mature miRNAs by 4-Thio-Uridine Metabolic Labeling. Methods in Molecular Biology, 2018, 1823, 141-152.	0.9	4
38	microRNAs transcriptional profiling of mammary stem cells isolated by PKH26 staining and FACS sorting. Methods in Cell Biology, 2022, , 59-79.	1.1	0