## Marco Galasso

## List of Publications by Year in descending order

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304743 395702 2,125 33 22 33 h-index citations g-index papers 34 34 34 4369 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Papillary Thyroid Carcinoma: Molecular Distinction by MicroRNA Profiling. Frontiers in Endocrinology, 2022, 13, 834075.	3.5	5
2	UC.183, UC.110, and UC.84 Ultra-Conserved RNAs Are Mutually Exclusive with miR-221 and Are Engaged in the Cell Cycle Circuitry in Breast Cancer Cell Lines. Genes, 2021, 12, 1978.	2.4	5
3	The network of non-coding RNAs and their molecular targets in breast cancer. Molecular Cancer, 2020, 19, 61.	19.2	36
4	miR-129-5p: A key factor and therapeutic target in amyotrophic lateral sclerosis. Progress in Neurobiology, 2020, 190, 101803.	5.7	31
5	MiRâ€16â€1â€3p and miRâ€16â€2â€3p possess strong tumor suppressive and antimetastatic properties in osteosarcoma. International Journal of Cancer, 2019, 145, 3052-3063.	5.1	27
6	An Ultraconserved Element Containing IncRNA Preserves Transcriptional Dynamics and Maintains ESC Self-Renewal. Stem Cell Reports, 2018, 10, 1102-1114.	4.8	17
7	A long non-coding RNA inside the type 2 transglutaminase gene tightly correlates with the expression of its transcriptional variants. Amino Acids, 2018, 50, 421-438.	2.7	7
8	Loss of miR-204 expression is a key event in melanoma. Molecular Cancer, 2018, 17, 71.	19.2	25
9	Heterogeneous expression of EPCAM in human circulating tumour cells from patient-derived xenografts. Biomarker Research, 2018, 6, 31.	6.8	17
10	Aptamer-miR-34c Conjugate Affects Cell Proliferation of Non-Small-Cell Lung Cancer Cells. Molecular Therapy - Nucleic Acids, 2018, 13, 334-346.	5.1	43
11	Screen for MicroRNA and Drug Interactions in Breast Cancer Cell Lines Points to miR-126 as a Modulator of CDK4/6 and PIK3CA Inhibitors. Frontiers in Genetics, 2018, 9, 174.	2.3	46
12	Levels of miR-126 and miR-218 are elevated in ductal carcinoma <i>in situ</i> (DCIS) and inhibit malignant potential of DCIS derived cells. Oncotarget, 2018, 9, 23543-23553.	1.8	12
13	Blood to skin recirculation of CD4 + memory T cells associates with cutaneous and systemic manifestations of psoriatic disease. Clinical Immunology, 2017, 180, 84-94.	3.2	26
14	miRâ€130A as a diagnostic marker to differentiate malignant mesothelioma from lung adenocarcinoma in pleural effusion cytology. Cancer Cytopathology, 2017, 125, 635-643.	2.4	18
15	Risk factors associated with relapse of eyelid basal cell carcinoma: results from a retrospective study of 142 patients. European Journal of Dermatology, 2017, 27, 363-368.	0.6	5
16	Profiling of the Predicted Circular RNAs in Ductal In Situ and Invasive Breast Cancer: A Pilot Study. International Journal of Genomics, 2016, 2016, 1-7.	1.6	30
17	miR-27a and miR-27a* contribute to metastatic properties of osteosarcoma cells. Oncotarget, 2015, 6, 4920-4935.	1.8	58
18	A MiRNA Signature for Defining Aggressive Phenotype and Prognosis in Gliomas. PLoS ONE, 2014, 9, e108950.	2.5	60

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19	Transcribed ultraconserved noncoding RNAs (T-UCR) are involved in Barrett's esophagus carcinogenesis. Oncotarget, 2014, 5, 7162-7171.	1.8	35
20	A large scale expression study associates uc.283-plus lncRNA with pluripotent stem cells and human glioma. Genome Medicine, 2014, 6, 76.	8.2	32
21	Pluripotent Stem Cell miRNAs and Metastasis in Invasive Breast Cancer. Journal of the National Cancer Institute, 2014, 106, .	6.3	37
22	Next generation analysis of breast cancer genomes for precision medicine. Cancer Letters, 2013, 339, 1-7.	7.2	19
23	Estrogen Mediated-Activation of miR-191/425 Cluster Modulates Tumorigenicity of Breast Cancer Cells Depending on Estrogen Receptor Status. PLoS Genetics, 2013, 9, e1003311.	3.5	139
24	Association between idiopathic hearing loss and mitochondrial DNA mutations: A study on 169 hearing-impaired subjects. International Journal of Molecular Medicine, 2013, 32, 785-794.	4.0	16
25	MicroRNA Expression Signatures in Solid Malignancies. Cancer Journal (Sudbury, Mass ), 2012, 18, 238-243.	2.0	72
26	Breast cancer signatures for invasiveness and prognosis defined by deep sequencing of microRNA. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 3024-3029.	7.1	334
27	miR-155 targets histone deacetylase 4 (HDAC4) and impairs transcriptional activity of B-cell lymphoma 6 (BCL6) in the Eµ-miR-155 transgenic mouse model. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 20047-20052.	7.1	121
28	miRNA Signatures Associate with Pathogenesis and Progression of Osteosarcoma. Cancer Research, 2012, 72, 1865-1877.	0.9	341
29	Prion proteins (PRNP and PRND) are overâ€expressed in osteosarcoma. Journal of Orthopaedic Research, 2012, 30, 1004-1012.	2.3	15
30	GAMES identifies and annotates mutations in next-generation sequencing projects. Bioinformatics, 2011, 27, 9-13.	4.1	28
31	Reprogramming of miRNA networks in cancer and leukemia. Genome Research, 2010, 20, 589-599.	5.5	331
32	Identification of microRNA activity by Targets' Reverse EXpression. Bioinformatics, 2010, 26, 91-97.	4.1	39
33	Non-coding RNAs: a key to future personalized molecular therapy?. Genome Medicine, 2010, 2, 12.	8.2	97