Daniela F Gradia

List of Publications by Year
in descending order

35	645	623574	610775
papers	citations	14	24
		h-index	g-index

37	37	37	1004
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Long nonâ€coding RNAs in cancer: Another layer of complexity. Journal of Gene Medicine, 2019, 21, e3065.	1.4	92
2	Predicting the Proteins of Angomonas deanei, Strigomonas culicis and Their Respective Endosymbionts Reveals New Aspects of the Trypanosomatidae Family. PLoS ONE, 2013, 8, e60209.	1.1	55
3	Long Non-Coding RNAs in Multifactorial Diseases: Another Layer of Complexity. Non-coding RNA, 2018, 4, 13.	1.3	55
4	Characterization of a novel Obg-like ATPase in the protozoan Trypanosoma cruzi. International Journal for Parasitology, 2009, 39, 49-58.	1.3	45
5	Identification of miRNAs Enriched in Extracellular Vesicles Derived from Serum Samples of Breast Cancer Patients. Biomolecules, 2020, 10, 150.	1.8	38
6	Long nonâ€coding RNAs differential expression in breast cancer subtypes: What do we know?. Clinical Genetics, 2019, 95, 558-568.	1.0	37
7	Unraveling Immune-Related lncRNAs in Breast Cancer Molecular Subtypes. Frontiers in Oncology, 2021, 11, 692170.	1.3	34
8	Analysis of Proteasomal Proteolysis during the In Vitro Metacyclogenesis of Trypanosoma cruzi. PLoS ONE, 2011, 6, e21027.	1.1	26
9	NEAT1 and MALAT1 are highly expressed in saliva and nasopharyngeal swab samples of COVIDâ€19 patients. Molecular Oral Microbiology, 2021, 36, 291-294.	1.3	25
10	The zinc finger protein TcZFP2 binds target mRNAs enriched during Trypanosoma cruzi metacyclogenesis. Memorias Do Instituto Oswaldo Cruz, 2012, 107, 790-799.	0.8	22
11	Stage-Regulated GFP Expression in Trypanosoma cruzi: Applications from Host-Parasite Interactions to Drug Screening. PLoS ONE, 2013, 8, e67441.	1.1	22
12	Long Non-Coding RNA TUG1 Expression Is Associated with Different Subtypes in Human Breast Cancer. Non-coding RNA, 2017, 3, 26.	1.3	17
13	Highlighting transcribed ultraconserved regions in human diseases. Wiley Interdisciplinary Reviews RNA, 2020, 11, e1567.	3.2	17
14	SARS-CoV-2 Delta and Omicron Variants Surge in Curitiba, Southern Brazil, and Its Impact on Overall COVID-19 Lethality. Viruses, 2022, 14, 809.	1.5	17
15	PBX1: a key character of the hallmarks of cancer. Journal of Molecular Medicine, 2021, 99, 1667-1680.	1.7	16
16	Large-Scale Screening of Asymptomatic Persons for SARS-CoV-2 Variants of Concern and Gamma Takeover, Brazil. Emerging Infectious Diseases, 2021, 27, 3124-3127.	2.0	14
17	Genome interaction of the virus and the host genes and non-coding RNAs in SARS-CoV-2 infection. Immunobiology, 2021, 226, 152130.	0.8	10
18	MicroRNAs miR-142-5p, miR-150-5p, miR-320a-3p, and miR-4433b-5p in Serum and Tissue: Potential Biomarkers in Sporadic Breast Cancer. Frontiers in Genetics, 0, 13, .	1.1	10

#	Article	IF	Citations
19	Association between SNP rs527616 in lncRNA AQP4-AS1 and susceptibility to breast cancer in a southern Brazilian population. Genetics and Molecular Biology, 2021, 44, e20200216.	0.6	9
20	A novel lncRNA derived from an ultraconserved region: lnc-uc.147, a potential biomarker in luminal A breast cancer. RNA Biology, 2021, , 1-14.	1.5	9
21	Comprehensive analysis of ceRNA networks in HPV16- and HPV18-mediated cervical cancers reveals XIST as a pivotal competing endogenous RNA. Biochimica Et Biophysica Acta - Molecular Basis of Disease, 2021, 1867, 166172.	1.8	9
22	Comparison of SARS-CoV-2 molecular detection in nasopharyngeal swab, saliva, and gargle samples. Diagnostic Microbiology and Infectious Disease, 2022, 103, 115678.	0.8	9
23	Frequency of the TP53 R337H variant in sporadic breast cancer and its impact on genomic instability. Scientific Reports, 2020, 10, 16614.	1.6	8
24	From Micro to Long: Non-Coding RNAs in Tamoxifen Resistance of Breast Cancer Cells. Cancers, 2021, 13, 3688.	1.7	8
25	Polymorphism of lncRNAs in breast cancer: Metaâ€analysis shows no association with susceptibility. Journal of Gene Medicine, 2020, 22, e3271.	1.4	7
26	So alike yet so different. Differential expression of the long non-coding RNAs NORAD and HCG11 in breast cancer subtypes. Genetics and Molecular Biology, 2021, 44, e20200153.	0.6	7
27	A genetic variant in microRNA-146a is associated with sporadic breast cancer in a Southern Brazilian Population. Genetics and Molecular Biology, 2019, 42, e20190278.	0.6	7
28	PUMILIO competes with AUF1 to control DICER1 RNA levels and miRNA processing. Nucleic Acids Research, 2022, 50, 7048-7066.	6.5	5
29	Novel lncRNAs Co-Expression Networks Identifies LINC00504 with Oncogenic Role in Luminal A Breast Cancer Cells. International Journal of Molecular Sciences, 2021, 22, 2420.	1.8	4
30	Transcribed Ultraconserved Regions Are Associated with Clinicopathological Features in Breast Cancer. Biomolecules, 2022, 12, 214.	1.8	3
31	lncRNAs in Hallmarks of Cancer and Clinical Applications. , 0, , .		2
32	Effects of PUMILIO1 and PUMILIO2 knockdown on cardiomyogenic differentiation of human embryonic stem cells culture. PLoS ONE, 2020, 15, e0222373.	1.1	2
33	Severe acute respiratory syndrome coronavirus 2 infection among healthcare workers in a tertiary public hospital in Curitiba, Brazil. Revista Da Sociedade Brasileira De Medicina Tropical, 2022, 55, e0265.	0.4	1
34	COVID-19: The question of genetic diversity and therapeutic intervention approaches. Genetics and Molecular Biology, 2021, 44, e20200452.	0.6	1
35	Sirtuins and the hallmarks of cancer. , 2021, , 129-152.		0