

Adriane F Evangelista

List of Publications by Year in descending order

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Version: 2024-02-01

74
papers

1,358
citations

304368

22
h-index

377514

34
g-index

81
all docs

81
docs citations

81
times ranked

2815
citing authors

#	ARTICLE	IF	CITATIONS
1	Identifying common and specific microRNAs expressed in peripheral blood mononuclear cell of type 1, type 2, and gestational diabetes mellitus patients. <i>BMC Research Notes</i> , 2013, 6, 491.	0.6	132
2	Role of glioblastoma stem cells in cancer therapeutic resistance: a perspective on antineoplastic agents from natural sources and chemical derivatives. <i>Stem Cell Research and Therapy</i> , 2021, 12, 206.	2.4	91
3	Vemurafenib resistance increases melanoma invasiveness and modulates the tumor microenvironment by MMP-2 upregulation. <i>Pharmacological Research</i> , 2016, 111, 523-533.	3.1	70
4	MicroRNA expression profiling and functional annotation analysis of their targets in patients with type 1 diabetes mellitus. <i>Gene</i> , 2014, 539, 213-223.	1.0	65
5	Autoimmune regulator (Aire) controls the expression of microRNAs in medullary thymic epithelial cells. <i>Immunobiology</i> , 2013, 218, 554-560.	0.8	57
6	Gene expression profiles displayed by peripheral blood mononuclear cells from patients with type 2 diabetes mellitus focusing on biological processes implicated on the pathogenesis of the disease. <i>Gene</i> , 2012, 511, 151-160.	1.0	54
7	MiR-21 as prognostic biomarker in head and neck squamous cell carcinoma patients undergoing an organ preservation protocol. <i>Oncotarget</i> , 2017, 8, 9911-9921.	0.8	48
8	MicroRNA-27a-5p regulation by promoter methylation and MYC signaling in prostate carcinogenesis. <i>Cell Death and Disease</i> , 2018, 9, 167.	2.7	48
9	MicroRNA expression as risk biomarker of breast cancer metastasis: a pilot retrospective case-cohort study. <i>BMC Cancer</i> , 2014, 14, 739.	1.1	45
10	Molecular characterization of breast cancer cell lines by clinical immunohistochemical markers. <i>Oncology Letters</i> , 2017, 13, 4708-4712.	0.8	40
11	Overexpression of mir-183 and mir-494 promotes proliferation and migration in human breast cancer cell lines. <i>Oncology Letters</i> , 2017, 14, 1054-1060.	0.8	40
12	Accuracy of microRNAs as markers for the detection of neck lymph node metastases in patients with head and neck squamous cell carcinoma. <i>BMC Medicine</i> , 2015, 13, 108.	2.3	33
13	Euphol, a tetracyclic triterpene, from <i>Euphorbia tirucalli</i> induces autophagy and sensitizes temozolomide cytotoxicity on glioblastoma cells. <i>Investigational New Drugs</i> , 2019, 37, 223-237.	1.2	33
14	A microRNA signature profile in EBV+ diffuse large B-cell lymphoma of the elderly. <i>Oncotarget</i> , 2014, 5, 11813-11826.	0.8	32
15	HER Family Receptors are Important Theranostic Biomarkers for Cervical Cancer: Blocking Glucose Metabolism Enhances the Therapeutic Effect of HER Inhibitors. <i>Theranostics</i> , 2017, 7, 717-732.	4.6	31
16	Mutation profiling of cancer drivers in Brazilian colorectal cancer. <i>Scientific Reports</i> , 2019, 9, 13687.	1.6	31
17	Detection of ALK fusion transcripts in FFPE lung cancer samples by NanoString technology. <i>BMC Pulmonary Medicine</i> , 2017, 17, 86.	0.8	30
18	Identification of Cell-Free Circulating MicroRNAs for the Detection of Early Breast Cancer and Molecular Subtyping. <i>Journal of Oncology</i> , 2019, 2019, 1-11.	0.6	30

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19	Integrative analysis of the transcriptome profiles observed in type 1, type 2 and gestational diabetes mellitus reveals the role of inflammation. <i>BMC Medical Genomics</i> , 2014, 7, 28.	0.7	28
20	Changes in Expression Profiles Revealed by Transcriptomic Analysis in Peripheral Blood Mononuclear Cells of Alzheimer's Disease Patients. <i>Journal of Alzheimer's Disease</i> , 2018, 66, 1483-1495.	1.2	28
21	Evidence for a network transcriptional control of promiscuous gene expression in medullary thymic epithelial cells. <i>Molecular Immunology</i> , 2009, 46, 3240-3244.	1.0	26
22	Reproducibility of the NanoString 22-gene molecular subgroup assay for improved prognostic prediction of medulloblastoma. <i>Neuropathology</i> , 2018, 38, 475-483.	0.7	26
23	Transcriptome meta-analysis of peripheral lymphomononuclear cells indicates that gestational diabetes is closer to type 1 diabetes than to type 2 diabetes mellitus. <i>Molecular Biology Reports</i> , 2013, 40, 5351-5358.	1.0	24
24	Expression of tumor suppressors miR-195 and let-7a as potential biomarkers of invasive breast cancer. <i>Clinics</i> , 2018, 73, e184.	0.6	24
25	Assessment of DNA damage and mRNA/miRNA transcriptional expression profiles in hyperglycemic versus non-hyperglycemic patients with type 2 diabetes mellitus. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2015, 776, 98-110.	0.4	22
26	Expression of genes related to apoptosis, cell cycle and signaling pathways are independent of TP53 status in urinary bladder cancer cells. <i>Molecular Biology Reports</i> , 2011, 38, 4159-4170.	1.0	21
27	Multiple myeloma cell lines and primary tumors proteome: protein biosynthesis and Immune system as potential therapeutic targets. <i>Genes and Cancer</i> , 2015, 6, 462-471.	0.6	16
28	Establishment, molecular and biological characterization of HCB-514: a novel human cervical cancer cell line. <i>Scientific Reports</i> , 2019, 9, 1913.	1.6	14
29	Gene Expression Profiles Stratified according to Type 1 Diabetes Mellitus Susceptibility Regions. <i>Annals of the New York Academy of Sciences</i> , 2008, 1150, 282-289.	1.8	13
30	The combination of coffee compounds attenuates early fibrosis-associated hepatocarcinogenesis in mice: involvement of miRNA profile modulation. <i>Journal of Nutritional Biochemistry</i> , 2020, 85, 108479.	1.9	13
31	Differential Transcript Profiles of MHC Class Ib (Qa-1, Qa-2, and Qa-10) and <i>Aire</i> Genes during the Ontogeny of Thymus and Other Tissues. <i>Journal of Immunology Research</i> , 2014, 2014, 1-12.	0.9	12
32	Somatic Copy Number Alterations and Associated Genes in Clear-Cell Renal-Cell Carcinoma in Brazilian Patients. <i>International Journal of Molecular Sciences</i> , 2021, 22, 2265.	1.8	12
33	Integrated analysis of mRNA and miRNA profiles revealed the role of miR-193 and miR-210 as potential regulatory biomarkers in different molecular subtypes of breast cancer. <i>BMC Cancer</i> , 2021, 21, 76.	1.1	12
34	Expression of GNAS, TP53, and PTEN Improves the Patient Prognostication in Sonic Hedgehog (SHH) Medulloblastoma Subgroup. <i>Journal of Molecular Diagnostics</i> , 2020, 22, 957-966.	1.2	11
35	miRNA expression profiling of hereditary breast tumors from BRCA1- and BRCA2-germline mutation carriers in Brazil. <i>BMC Cancer</i> , 2020, 20, 143.	1.1	10
36	Bioprospecting of Natural Compounds from Brazilian Cerrado Biome Plants in Human Cervical Cancer Cell Lines. <i>International Journal of Molecular Sciences</i> , 2021, 22, 3383.	1.8	9

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37	MicroRNA Biomarkers of High-Grade Cervical Intraepithelial Neoplasia in Liquid Biopsy. <i>BioMed Research International</i> , 2021, 2021, 1-9.	0.9	9
38	Comprehensive Molecular Landscape of Cetuximab Resistance in Head and Neck Cancer Cell Lines. <i>Cells</i> , 2022, 11, 154.	1.8	9
39	The role of single-nucleotide polymorphism (SNPs) in toxicity of induction chemotherapy based on cisplatin and paclitaxel in patients with advanced head and neck cancer. <i>Oral Oncology</i> , 2019, 98, 48-52.	0.8	8
40	Semi-Synthetic Ingenol Derivative from <i>Euphorbia tirucalli</i> Inhibits Protein Kinase C Isotypes and Promotes Autophagy and S-phase Arrest on Glioma Cell Lines. <i>Molecules</i> , 2019, 24, 4265.	1.7	8
41	Phospholipase gene expression during <i>Paracoccidioides brasiliensis</i> morphological transition and infection. <i>Memorias Do Instituto Oswaldo Cruz</i> , 2013, 108, 808-811.	0.8	7
42	MicroRNA profiling in human breast cancer cell lines exposed to the anti-neoplastic drug cediranib. <i>Oncology Reports</i> , 2016, 36, 3197-3206.	1.2	7
43	A 4-Gene Signature Associated With Recurrence in Low- and Intermediate-Risk Endometrial Cancer. <i>Frontiers in Oncology</i> , 2021, 11, 729219.	1.3	7
44	Cutaneous and acral melanoma cross-OMICs reveals prognostic cancer drivers associated with pathobiology and ultraviolet exposure. <i>Nature Communications</i> , 2022, 13, .	5.8	7
45	Identification and performance evaluation of housekeeping genes for microRNA expression normalization by reverse transcription-quantitative PCR using liquid-based cervical cytology samples. <i>Oncology Letters</i> , 2019, 18, 4753-4761.	0.8	6
46	Transcript Expression Profiles and MicroRNA Regulation Indicate an Upregulation of Processes Linked to Oxidative Stress, DNA Repair, Cell Death, and Inflammation in Type 1 Diabetes Mellitus Patients. <i>Journal of Diabetes Research</i> , 2022, 2022, 1-15.	1.0	6
47	Using Co-segregation and Loss of Heterozygosity Analysis to Define the Pathogenicity of Unclassified Variants in Hereditary Breast Cancer Patients. <i>Frontiers in Oncology</i> , 2020, 10, 571330.	1.3	5
48	Deregulated microRNAs Are Associated with Patient Survival and Predicted to Target Genes That Modulate Lung Cancer Signaling Pathways. <i>Cancers</i> , 2020, 12, 2711.	1.7	5
49	Analysis of RPL37A, MTSS1, and HTRA1 expression as potential markers for pathologic complete response and survival. <i>Breast Cancer</i> , 2021, 28, 307-320.	1.3	5
50	Approaches for the identification of driver mutations in cancer: A tutorial from a computational perspective. <i>Journal of Bioinformatics and Computational Biology</i> , 2020, 18, 2050016.	0.3	4
51	Simultaneous analysis of ALK, RET, and ROS1 gene fusions by NanoString in Brazilian lung adenocarcinoma patients. <i>Translational Lung Cancer Research</i> , 2021, 10, 292-303.	1.3	4
52	Genetic alterations detected by comparative genomic hybridization in BRCA1 breast and ovarian cancers of Brazilian population. <i>Oncotarget</i> , 2018, 9, 27525-27534.	0.8	4
53	Behavior of α -tomatine and tomatidine against several genera of trypanosomatids from insects and plants and <i>Trypanosoma cruzi</i> . <i>Acta Scientiarum - Biological Sciences</i> , 2018, 40, 41853.	0.3	3
54	Pyknon-Containing Transcripts Are Downregulated in Colorectal Cancer Tumors, and Loss of PYK4 Is Associated With Worse Patient Outcome. <i>Frontiers in Genetics</i> , 2020, 11, 581454.	1.1	3

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55	Profile of esophageal squamous cell carcinoma mutations in Brazilian patients. Scientific Reports, 2021, 11, 20596.	1.6	2
56	Abstract 3386: Brazilian melanoma genome project: mutational landscape based on whole-genome sequencing. , 2017, , .		2
57	Combining Mutation and Gene Network Data in a Machine Learning Approach for False-Positive Cancer Driver Gene Discovery. Lecture Notes in Computer Science, 2020, , 81-92.	1.0	2
58	Differential expression profile of microRNAs associated with human breast cancer progression. BMC Proceedings, 2013, 7, .	1.8	1
59	MicroRNAs from peripheral blood mononuclear cells as biomarkers for detection of preclinical fibrosarcoma. BMC Proceedings, 2013, 7, P2.	1.8	1
60	Micro-RNA Expression Profile Reveals MiR-222 As a Potential Biomarker For EBV-Positive Diffuse Large B-Cell Lymphoma. Blood, 2013, 122, 4269-4269.	0.6	1
61	Transcriptome Analysis During Normal Human Mesenchymal Stem Cell Differentiation. , 2014, , 109-119.		1
62	A proposal of a graph-based computational method for ranking significant set of related genes in cancer. , 0, , .		1
63	A computational approach for the discovery of significant cancer genes by weighted mutation and asymmetric spreading strength in networks. Scientific Reports, 2021, 11, 23551.	1.6	1
64	Correction, article no. 3671: Importance of the patient's clinical questionnaire for the diagnosis of skin cancer through teledermatology in remote areas of Brazil. Rural and Remote Health, 2017, 17, 4496.	0.4	1
65	P102. Human Immunology, 2014, 75, 122.	1.2	0
66	P041 Serum and tissue expression of tumor suppressors miR-195 and let-7A in breast cancer. Breast, 2015, 24, S40.	0.9	0
67	Transcriptome Metanalysis of Peripheral Blood Mononuclear Cells of Type 1, Type 2 and Gestational Diabetes Mellitus Patients Features Modulated Immune System and Metabolism Genes. , 2011, , P3-463-P3-463.		0
68	MicroRNAs in Cancer. , 2014, , 181-193.		0
69	Abstract A33: Molecular classification of Brazilian medulloblastoma by a NanoString gene panel. , 2018, , .		0
70	Abstract A34: SNPs as predictors in the tailoring of treatment for HNSCC patients undergoing organ preservation protocols. , 2018, , .		0
71	Abstract A29: Omics profile of two immortalized Brazilian glioblastoma cell cultures. , 2018, , .		0
72	Abstract A35: Somatic mutation profile of a panel of oncogenes and tumor suppressor genes in colorectal cancer in a Brazilian population. , 2018, , .		0

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73	Abstract 5380: Whole-exome sequencing of Brazilian non-small cell lung cancer. , 2018, , .		0
74	Abstract 3998: Integrative molecular analysis uncovers key molecules and signaling pathways regulated by RKIP in gastrointestinal stromal tumors (GISTs). , 2018, , .		0