

Luciane R Cavalli

List of Publications by Year in descending order

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Version: 2024-02-01

31
papers

1,031
citations

516710

16
h-index

477307

29
g-index

32
all docs

32
docs citations

32
times ranked

1446
citing authors

#	ARTICLE	IF	CITATIONS
1	Increased Steroidogenic Factor-1 Dosage Triggers Adrenocortical Cell Proliferation and Cancer. <i>Molecular Endocrinology</i> , 2007, 21, 2968-2987.	3.7	194
2	Amplification of the Steroidogenic Factor 1 Gene in Childhood Adrenocortical Tumors. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2005, 90, 615-619.	3.6	120
3	ERR β Mediates Tamoxifen Resistance in Novel Models of Invasive Lobular Breast Cancer. <i>Cancer Research</i> , 2008, 68, 8908-8917.	0.9	97
4	SF-1 overexpression in childhood adrenocortical tumours. <i>European Journal of Cancer</i> , 2006, 42, 1040-1043.	2.8	90
5	Deregulated miRNA expression is associated with endothelial dysfunction in post-mortem lung biopsies of COVID-19 patients. <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 2021, 320, L405-L412.	2.9	59
6	Detection of LOH and Mitochondrial DNA Alterations in Ductal Lavage and Nipple Aspirate Fluids from High-risk Patients. <i>Breast Cancer Research and Treatment</i> , 2004, 84, 99-105.	2.5	49
7	Differentially expressed miRNAs in triple negative breast cancer between African-American and non-Hispanic white women. <i>Oncotarget</i> , 2016, 7, 79274-79291.	1.8	43
8	Loss of heterozygosity in normal breast epithelial tissue and benign breast lesions in BRCA1/2 carriers with breast cancer. <i>Cancer Genetics and Cytogenetics</i> , 2004, 149, 38-43.	1.0	40
9	Identification of miRNAs Enriched in Extracellular Vesicles Derived from Serum Samples of Breast Cancer Patients. <i>Biomolecules</i> , 2020, 10, 150.	4.0	38
10	The role of microRNAs in modulating SARS-CoV-2 infection in human cells: a systematic review. <i>Infection, Genetics and Evolution</i> , 2021, 91, 104832.	2.3	35
11	Genetic and epigenetic alterations in sentinel lymph nodes metastatic lesions compared to their corresponding primary breast tumors. <i>Cancer Genetics and Cytogenetics</i> , 2003, 146, 33-40.	1.0	33
12	Conditionally Reprogrammed Normal and Transformed Mouse Mammary Epithelial Cells Display a Progenitor-Cell-Like Phenotype. <i>PLoS ONE</i> , 2014, 9, e97666.	2.5	27
13	Genomic comparison of early-passage conditionally reprogrammed breast cancer cells to their corresponding primary tumors. <i>PLoS ONE</i> , 2017, 12, e0186190.	2.5	24
14	Integrated molecular analysis of Tamoxifen-resistant invasive lobular breast cancer cells identifies MAPK and GRM/mGluR signaling as therapeutic vulnerabilities. <i>Molecular and Cellular Endocrinology</i> , 2018, 471, 105-117.	3.2	22
15	Increased copy number of the DLX4 homeobox gene in breast axillary lymph node metastasis. <i>Cancer Genetics</i> , 2014, 207, 177-187.	0.4	19
16	Copy number and expression analysis of FOSL1, GSTP1, NTSR1, FADD and CCND1 genes in primary breast tumors with axillary lymph node metastasis. <i>Cancer Genetics</i> , 2016, 209, 331-339.	0.4	19
17	Molecular markers of breast axillary lymph node metastasis. <i>Expert Review of Molecular Diagnostics</i> , 2009, 9, 441-454.	3.1	16
18	Amplification of the BP1 homeobox gene in breast cancer. <i>Cancer Genetics and Cytogenetics</i> , 2008, 187, 19-24.	1.0	15

#	ARTICLE	IF	CITATIONS
19	Integrated copy number and miRNA expression analysis in triple negative breast cancer of Latin American patients. <i>Oncotarget</i> , 2019, 10, 6184-6203.	1.8	15
20	MiR-150-5p Overexpression in Triple-Negative Breast Cancer Contributes to the In Vitro Aggressiveness of This Breast Cancer Subtype. <i>Cancers</i> , 2022, 14, 2156.	3.7	12
21	Genome interaction of the virus and the host genes and non-coding RNAs in SARS-CoV-2 infection. <i>Immunobiology</i> , 2021, 226, 152130.	1.9	10
22	QNBC Is Associated with High Genomic Instability Characterized by Copy Number Alterations and miRNA Dereglulation. <i>International Journal of Molecular Sciences</i> , 2021, 22, 11548.	4.1	10
23	The orphan nuclear receptor estrogen-related receptor beta (ERR β) in triple-negative breast cancer. <i>Breast Cancer Research and Treatment</i> , 2020, 179, 585-604.	2.5	8
24	Frequency of the TP53 R337H variant in sporadic breast cancer and its impact on genomic instability. <i>Scientific Reports</i> , 2020, 10, 16614.	3.3	8
25	A panel of miRNAs as prognostic markers for African-American patients with triple negative breast cancer. <i>BMC Cancer</i> , 2021, 21, 861.	2.6	8
26	Correlates of Triple Negative Breast Cancer and Chemotherapy Patterns in Black and White Women With Breast Cancer. <i>Clinical Breast Cancer</i> , 2017, 17, 232-238.	2.4	6
27	Copy Number Analysis of the <i>DLX4</i> and <i>ERBB2</i> Genes in South African Breast Cancer Patients. <i>Cytogenetic and Genome Research</i> , 2015, 146, 195-203.	1.1	5
28	Patterns of copy number alterations in primary breast tumors of South African patients and their impact on functional cellular pathways. <i>International Journal of Oncology</i> , 2018, 53, 2745-2757.	3.3	5
29	Abstract 327: Genomic profiling of sentinel lymph node breast cancer metastasis. , 2010, , .		3
30	Breast Axillary Lymph Node Metastasis. <i>International Journal of Breast Cancer</i> , 2011, 2011, 1-2.	1.2	1
31	Molecular Classification and Prognostic Signatures of Breast Tumors. , 2019, , 129-138.		0