

Claudia A Rainho

List of Publications by Year in descending order

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49
papers

1,119
citations

471509

17
h-index

395702

33
g-index

49
all docs

49
docs citations

49
times ranked

1836
citing authors

#	ARTICLE	IF	CITATIONS
1	Brazilian green propolis: A novel tool to improve the cytotoxic and immunomodulatory action of docetaxel on MCF-7 breast cancer cells and on women monocyte. <i>Phytotherapy Research</i> , 2022, 36, 448-461.	5.8	12
2	The Long Non-Coding RNA SNHG12 as a Mediator of Carboplatin Resistance in Ovarian Cancer via Epigenetic Mechanisms. <i>Cancers</i> , 2022, 14, 1664.	3.7	4
3	Relevance of PD-L1 Non-Coding Polymorphisms on the Prognosis of a Genetically Admixed NSCLC Cohort. <i>Pharmacogenomics and Personalized Medicine</i> , 2021, Volume 14, 239-252.	0.7	3
4	Master Regulators of Epithelial-Mesenchymal Transition and WNT Signaling Pathways in Juvenile Nasopharyngeal Angiofibromas. <i>Biomedicines</i> , 2021, 9, 1258.	3.2	3
5	The Interplay between Long Noncoding RNAs and Proteins of the Epigenetic Machinery in Ovarian Cancer. <i>Cancers</i> , 2020, 12, 2701.	3.7	6
6	Comprehensive Analysis of DNA Methylation and Prediction of Response to Neoadjuvant Therapy in Locally Advanced Rectal Cancer. <i>Cancers</i> , 2020, 12, 3079.	3.7	13
7	Epigenetics in Inflammatory Breast Cancer: Biological Features and Therapeutic Perspectives. <i>Cells</i> , 2020, 9, 1164.	4.1	18
8	Effects of Propolis and Phenolic Acids on Triple-Negative Breast Cancer Cell Lines: Potential Involvement of Epigenetic Mechanisms. <i>Molecules</i> , 2020, 25, 1289.	3.8	27
9	The long non-coding RNA ANRASSF1 in the regulation of alternative protein-coding transcripts RASSF1A and RASSF1C in human breast cancer cells: implications to epigenetic therapy. <i>Epigenetics</i> , 2019, 14, 741-750.	2.7	12
10	Expression and promoter methylation status of two DNA repair genes in leukocytes from patients undergoing propofol or isoflurane anaesthesia. <i>Mutagenesis</i> , 2018, 33, 147-152.	2.6	10
11	Abstract A12: Brazilian propolis as a source of novel DNA methyltransferase inhibitors: A computer-aided discovery and in vitro approaches. , 2018, , .		1
12	Triploidy in a sexually dimorphic passerine provides new evidence for the effect of the W chromosome on secondary sexual traits in birds. <i>Journal of Avian Biology</i> , 2017, 48, 1475-1480.	1.2	4
13	PHF21B as a candidate tumor suppressor gene in head and neck squamous cell carcinomas. <i>Molecular Oncology</i> , 2015, 9, 450-462.	4.6	18
14	Adhesion study of cultured human lens capsule cells on hydrophilic intraocular lenses coated with polyethylene glycol. <i>Journal of Cataract and Refractive Surgery</i> , 2015, 41, 1478-1483.	1.5	4
15	The Th17 pathway in the peripheral lung microenvironment interacts with expression of collagen V in the late state of experimental pulmonary fibrosis. <i>Immunobiology</i> , 2015, 220, 124-135.	1.9	15
16	Defining suitable reference genes for RT-qPCR analysis on human sertoli cells after 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD) exposure. <i>Molecular Biology Reports</i> , 2014, 41, 7063-7066.	2.3	5
17	DNA methylation patterns of steroid receptor genes ESR1, ESR2 and PGR in deep endometriosis compromising the rectum. <i>International Journal of Molecular Medicine</i> , 2014, 33, 897-904.	4.0	36
18	Dicephalus dipus dibrachius twins: report of an autopsy case. <i>Autopsy and Case Reports</i> , 2014, 4, 21-26.	0.6	2

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19	Analysis of the DNA methylation of the H19 gene in human bladder cancer. <i>BMC Proceedings</i> , 2013, 7, .	1.6	0
20	ACVR1 (activin A receptor, type I). <i>Atlas of Genetics and Cytogenetics in Oncology and Haematology</i> , 2013, , .	0.1	0
21	Genetic polymorphisms associated with steroids metabolism and insulin action in polycystic ovary syndrome. <i>Gynecological Endocrinology</i> , 2012, 28, 190-194.	1.7	17
22	Evidence of epigenetic regulation of the tumor suppressor gene cluster flanking <i>RASSF1</i> in breast cancer cell lines. <i>Epigenetics</i> , 2011, 6, 1413-1424.	2.7	41
23	CYP1A2*1C, CYP2E1*5B, and GSTM1 polymorphisms are predictors of risk and poor outcome in head and neck squamous cell carcinoma patients. <i>Oral Oncology</i> , 2009, 45, e73-e79.	1.5	48
24	DNA methylation patterns in bladder cancer and washing cell sediments: a perspective for tumor recurrence detection. <i>BMC Cancer</i> , 2008, 8, 238.	2.6	37
25	PSA and Androgen-Related Gene (AR, CYP17, and CYP19) Polymorphisms and the Risk of Adenocarcinoma at Prostate Biopsy. <i>DNA and Cell Biology</i> , 2008, 27, 497-503.	1.9	15
26	Polymorphisms of CYP17A1, CYP19, and androgen in Brazilian women with uterine leiomyomas. <i>Clinical Chemistry and Laboratory Medicine</i> , 2008, 46, 814-23.	2.3	9
27	Velocardiofacial syndrome with a rare t(2;22). <i>Clinical Dysmorphology</i> , 2007, 16, 181-183.	0.3	0
28	DNA methylation patterns of the CDH1, RARB, and SFN genes in choroid plexus tumors. <i>Cancer Genetics and Cytogenetics</i> , 2007, 179, 140-145.	1.0	14
29	Shorter CAG repeat length in the AR gene is associated with poor outcome in head and neck cancer. <i>Archives of Oral Biology</i> , 2007, 52, 732-739.	1.8	8
30	Shorter CAG repeat in the AR gene is associated with atypical hyperplasia and breast carcinoma. <i>Anticancer Research</i> , 2007, 27, 1199-205.	1.1	12
31	CDH1 promoter hypermethylation and E-cadherin protein expression in infiltrating breast cancer. <i>BMC Cancer</i> , 2006, 6, 48.	2.6	137
32	H19-DMR allele-specific methylation analysis reveals epigenetic heterogeneity of CTCF binding site 6 but not of site 5 in head-and-neck carcinomas: a pilot case-control analysis. <i>International Journal of Molecular Medicine</i> , 2006, 17, 397-404.	4.0	8
33	Independent clonal origin of multiple uterine leiomyomas that was determined by X chromosome inactivation and microsatellite analysis. <i>American Journal of Obstetrics and Gynecology</i> , 2005, 193, 1395-1403.	1.3	47
34	DNA methylation in the CTCF-binding site I and the expression pattern of the H19 gene: Does positive expression predict poor prognosis in early stage head and neck carcinomas?. <i>Molecular Carcinogenesis</i> , 2005, 44, 102-110.	2.7	24
35	Large-scale Transcriptome Analyses Reveal New Genetic Marker Candidates of Head, Neck, and Thyroid Cancer. <i>Cancer Research</i> , 2005, 65, 1693-1699.	0.9	55
36	A Transcript Finishing Initiative for Closing Gaps in the Human Transcriptome. <i>Genome Research</i> , 2004, 14, 1413-1423.	5.5	22

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37	Genotyping of AR and PSA polymorphisms in a patient with Klinefelter syndrome, non-Hodgkin lymphoma, and adenocarcinoma of the prostate. <i>Cancer Genetics and Cytogenetics</i> , 2004, 153, 165-169.	1.0	4
38	Identification and complete sequencing of novel human transcripts through the use of mouse orthologs and testis cDNA sequences. <i>Genetics and Molecular Research</i> , 2004, 3, 493-511.	0.2	0
39	The generation and utilization of a cancer-oriented representation of the human transcriptome by using expressed sequence tags. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003, 100, 13418-13423.	7.1	105
40	Loss of imprinting and loss of heterozygosity on 11p15.5 in head and neck squamous cell carcinomas. <i>Head and Neck</i> , 2001, 23, 851-859.	2.0	15
41	The contribution of 700,000 ORF sequence tags to the definition of the human transcriptome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001, 98, 12103-12108.	7.1	123
42	Identification of human chromosome 22 transcribed sequences with ORF expressed sequence tags. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000, 97, 12690-12693.	7.1	70
43	Expression and Imprinting of Insulin-like Growth Factor II (IGF2) and H19 Genes in Uterine Leiomyomas. <i>Gynecologic Oncology</i> , 1999, 74, 375-380.	1.4	13
44	Hemangioendothelioma of Bone in a Patient with a Constitutional Supernumerary Marker. <i>Cancer Genetics and Cytogenetics</i> , 1999, 110, 23-27.	1.0	16
45	Clonal chromosome abnormalities found in three non-neoplastic proliferative brain lesions. <i>Genetics and Molecular Biology</i> , 1999, 22, 25-28.	1.3	0
46	Cytogenetic report of a male breast cancer. <i>Cancer Genetics and Cytogenetics</i> , 1995, 81, 66-71.	1.0	12
47	Chromosomes in the genesis and progression of ependymomas. <i>Cancer Genetics and Cytogenetics</i> , 1993, 69, 146-152.	1.0	46
48	Cytogenetic study of a pineocytoma. <i>Cancer Genetics and Cytogenetics</i> , 1992, 64, 127-132.	1.0	28
49	Chromosomes in the genesis and progression of ependymomas. <i>Cancer Genetics and Cytogenetics</i> , 1992, 63, 177.	1.0	0