

# 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	CDH1 promoter hypermethylation and E-cadherin protein expression in infiltrating breast cancer. BMC Cancer, 2006, 6, 48.	2.6	137
2	The contribution of 700,000 ORF sequence tags to the definition of the human transcriptome. Proceedings of the National Academy of Sciences of the United States of America, 2001, 98, 12103-12108.	7.1	123
3	The generation and utilization of a cancer-oriented representation of the human transcriptome by using expressed sequence tags. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 13418-13423.	7.1	105
4	Identification of human chromosome 22 transcribed sequences with ORF expressed sequence tags. Proceedings of the National Academy of Sciences of the United States of America, 2000, 97, 12690-12693.	7.1	70
5	Large-scale Transcriptome Analyses Reveal New Genetic Marker Candidates of Head, Neck, and Thyroid Cancer. Cancer Research, 2005, 65, 1693-1699.	0.9	55
6	CYP1A2*1C, CYP2E1*5B, and GSTM1 polymorphisms are predictors of risk and poor outcome in head and neck squamous cell carcinoma patients. Oral Oncology, 2009, 45, e73-e79.	1.5	48
7	Independent clonal origin of multiple uterine leiomyomas that was determined by X chromosome inactivation and microsatellite analysis. American Journal of Obstetrics and Gynecology, 2005, 193, 1395-1403.	1.3	47
8	Chromosomes in the genesis and progression of ependymomas. Cancer Genetics and Cytogenetics, 1993, 69, 146-152.	1.0	46
9	Evidence of epigenetic regulation of the tumor suppressor gene cluster flanking <i>RASSF1</i> in breast cancer cell lines. Epigenetics, 2011, 6, 1413-1424.	2.7	41
10	DNA methylation patterns in bladder cancer and washing cell sediments: a perspective for tumor recurrence detection. BMC Cancer, 2008, 8, 238.	2.6	37
11	DNA methylation patterns of steroid receptor genes ESR1, ESR2 and PGR in deep endometriosis compromising the rectum. International Journal of Molecular Medicine, 2014, 33, 897-904.	4.0	36
12	Cytogenetic study of a pineocytoma. Cancer Genetics and Cytogenetics, 1992, 64, 127-132.	1.0	28
13	Effects of Propolis and Phenolic Acids on Triple-Negative Breast Cancer Cell Lines: Potential Involvement of Epigenetic Mechanisms. Molecules, 2020, 25, 1289.	3.8	27
14	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?. Molecular Carcinogenesis, 2005, 44, 102-110.	2.7	24
15	A Transcript Finishing Initiative for Closing Gaps in the Human Transcriptome. Genome Research, 2004, 14, 1413-1423.	5.5	22
16	<i>PHF21B</i> as a candidate tumor suppressor gene in head and neck squamous cell carcinomas. Molecular Oncology, 2015, 9, 450-462.	4.6	18
17	Epigenetics in Inflammatory Breast Cancer: Biological Features and Therapeutic Perspectives. Cells, 2020, 9, 1164.	4.1	18
18	Genetic polymorphisms associated with steroids metabolism and insulin action in polycystic ovary syndrome. Gynecological Endocrinology, 2012, 28, 190-194.	1.7	17

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19	Hemangioendothelioma of Bone in a Patient with a Constitutional Supernumerary Marker. <i>Cancer Genetics and Cytogenetics</i> , 1999, 110, 23-27.	1.0	16
20	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
21	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
22	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
23	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
24	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
25	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
26	Cytogenetic report of a male breast cancer. <i>Cancer Genetics and Cytogenetics</i> , 1995, 81, 66-71.	1.0	12
27	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
28	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
29	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
30	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
31	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
32	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
33	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
34	The Interplay between Long Noncoding RNAs and Proteins of the Epigenetic Machinery in Ovarian Cancer. <i>Cancers</i> , 2020, 12, 2701.	3.7	6
35	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
36	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

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37	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
38	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
39	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
40	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
41	Master Regulators of Epithelial-Mesenchymal Transition and WNT Signaling Pathways in Juvenile Nasopharyngeal Angiofibromas. <i>Biomedicines</i> , 2021, 9, 1258.	3.2	3
42	Dicephalus dipus dibrachius twins: report of an autopsy case. <i>Autopsy and Case Reports</i> , 2014, 4, 21-26.	0.6	2
43	Abstract A12: Brazilian propolis as a source of novel DNA methyltransferase inhibitors: A computer-aided discovery and in vitro approaches. , 2018, , .		1
44	Chromosomes in the genesis and progression of ependymomas. <i>Cancer Genetics and Cytogenetics</i> , 1992, 63, 177.	1.0	0
45	Velocardiofacial syndrome with a rare t(2;22). <i>Clinical Dysmorphology</i> , 2007, 16, 181-183.	0.3	0
46	Analysis of the DNA methylation of the H19 gene in human bladder cancer. <i>BMC Proceedings</i> , 2013, 7, .	1.6	0
47	ACVR1 (activin A receptor, type I). <i>Atlas of Genetics and Cytogenetics in Oncology and Haematology</i> , 2013, , .	0.1	0
48	Clonal chromosome abnormalities found in three non-neoplastic proliferative brain lesions. <i>Genetics and Molecular Biology</i> , 1999, 22, 25-28.	1.3	0
49	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