

Joan Climent

List of Publications by Year in descending order

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Version: 2024-02-01

32
papers

2,376
citations

279487

23
h-index

476904

29
g-index

34
all docs

34
docs citations

34
times ranked

4528
citing authors

#	ARTICLE	IF	CITATIONS
1	FBXW7 Targets mTOR for Degradation and Cooperates with PTEN in Tumor Suppression. <i>Science</i> , 2008, 321, 1499-1502.	6.0	375
2	Homozygous deletions localize novel tumor suppressor genes in B-cell lymphomas. <i>Blood</i> , 2007, 109, 271-280.	0.6	227
3	Sequence and analysis of chromosome 3 of the plant <i>Arabidopsis thaliana</i> . <i>Nature</i> , 2000, 408, 820-823.	13.7	188
4	MALT1 is deregulated by both chromosomal translocation and amplification in B-cell non-Hodgkin lymphoma. <i>Blood</i> , 2003, 101, 4539-4546.	0.6	188
5	Mantle-cell lymphoma genotypes identified with CGH to BAC microarrays define a leukemic subgroup of disease and predict patient outcome. <i>Blood</i> , 2005, 105, 4445-4454.	0.6	180
6	Inverse and Direct Cancer Comorbidity in People with Central Nervous System Disorders: A Meta-Analysis of Cancer Incidence in 577,013 Participants of 50 Observational Studies. <i>Psychotherapy and Psychosomatics</i> , 2014, 83, 89-105.	4.0	164
7	Characterization of 8p21.3 chromosomal deletions in B-cell lymphoma: TRAIL-R1 and TRAIL-R2 as candidate dosage-dependent tumor suppressor genes. <i>Blood</i> , 2005, 106, 3214-3222.	0.6	137
8	No paradox, no progress: inverse cancer comorbidity in people with other complex diseases. <i>Lancet Oncology</i> , The, 2011, 12, 604-608.	5.1	122
9	Co-amplified genes at 8p12 and 11q13 in breast tumors cooperate with two major pathways in oncogenesis. <i>Oncogene</i> , 2009, 28, 1892-1903.	2.6	94
10	Loss of a novel tumor suppressor gene locus at chromosome 8p is associated with leukemic mantle cell lymphoma. <i>Blood</i> , 2001, 98, 3479-3482.	0.6	86
11	Deletion of Chromosome 11q Predicts Response to Anthracycline-Based Chemotherapy in Early Breast Cancer. <i>Cancer Research</i> , 2007, 67, 818-826.	0.4	75
12	Deletion of the <i>PER3</i> Gene on Chromosome 1p36 in Recurrent ER-Positive Breast Cancer. <i>Journal of Clinical Oncology</i> , 2010, 28, 3770-3778.	0.8	57
13	MicroRNA profile in very young women with breast cancer. <i>BMC Cancer</i> , 2014, 14, 529.	1.1	56
14	Homeobox NKX2-3 promotes marginal-zone lymphomagenesis by activating B-cell receptor signalling and shaping lymphocyte dynamics. <i>Nature Communications</i> , 2016, 7, 11889.	5.8	42
15	Pleckstrin homology domain-interacting protein (PHIP) as a marker and mediator of melanoma metastasis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 7067-7072.	3.3	39
16	Characterization of breast cancer by array comparative genomic hybridization This paper is one of a selection of papers published in this Special Issue, entitled 28th International West Coast Chromatin and Chromosome Conference, and has undergone the Journal's usual peer review process.. <i>Biochemistry and Cell Biology</i> , 2007, 85, 497-508.	0.9	38
17	Hipk2 cooperates with p53 to suppress $\hat{\gamma}$ -ray radiation-induced mouse thymic lymphoma. <i>Oncogene</i> , 2012, 31, 1176-1180.	2.6	36
18	Homozygous deletion of SOCS1 in primary mediastinal B-cell lymphoma detected by CGH to BAC microarrays. <i>Leukemia</i> , 2005, 19, 1082-1084.	3.3	35

#	ARTICLE	IF	CITATIONS
19	MicroRNA Profile in Response to Doxorubicin Treatment in Breast Cancer. <i>Journal of Cellular Biochemistry</i> , 2015, 116, 2061-2073.	1.2	31
20	Transcriptomic metaanalyses of autistic brains reveals shared gene expression and biological pathway abnormalities with cancer. <i>Molecular Autism</i> , 2019, 10, 17.	2.6	30
21	Genomic loss of 18p predicts an adverse clinical outcome in patients with high-risk breast cancer. <i>Clinical Cancer Research</i> , 2002, 8, 3863-9.	3.2	29
22	Selective concomitant inhibition of mTORC1 and mTORC2 activity in estrogen receptor negative breast cancer cells by BN107 and oleanolic acid. <i>International Journal of Cancer</i> , 2010, 127, 1209-1219.	2.3	28
23	Transcriptomic and Genetic Associations between Alzheimer's Disease, Parkinson's Disease, and Cancer. <i>Cancers</i> , 2021, 13, 2990.	1.7	26
24	Genomic Abnormalities Acquired in the Blastic Transformation of Splenic Marginal Zone B-cell Lymphoma. <i>Leukemia and Lymphoma</i> , 2003, 44, 459-464.	0.6	24
25	Applications of computational homology to the analysis of treatment response in breast cancer patients. <i>Topology and Its Applications</i> , 2010, 157, 157-164.	0.2	24
26	A genomic approach to study down syndrome and cancer inverse comorbidity: untangling the chromosome 21. <i>Frontiers in Physiology</i> , 2015, 6, 10.	1.3	17
27	A COVID-19 Drug Repurposing Strategy through Quantitative Homological Similarities Using a Topological Data Analysis-Based Framework. <i>Pharmaceutics</i> , 2021, 13, 488.	2.0	13
28	Analysis of the CDKN2A and CDK4 Genes and HLA-DR and HLA-DQ Alleles in Two Spanish Familial Melanoma Kindreds. <i>Acta Dermato-Venereologica</i> , 2000, 80, 440-442.	0.6	10
29	Circadian Performance in breast cancer: a germline and somatic genetic study of PER3VNTR polymorphisms and gene co-expression. <i>Npj Breast Cancer</i> , 2021, 7, 118.	2.3	3
30	DNA Methylation in Breast Cancer. , 2016, , 297-312.		1
31	miRNA Expression Analysis: Cell Lines HCC1500 and HCC1937 as Models for Breast Cancer in Young Women and the miR-23a as a Poor Prognostic Biomarker. <i>Breast Cancer: Basic and Clinical Research</i> , 2020, 14, 117822342097784.	0.6	0
32	Mantle Cell Lymphoma Genotypes Identified with CGH to BAC Microarrays Define Clinical Subgroups of Disease and Strongly Predict Patient Outcome.. <i>Blood</i> , 2004, 104, 695-695.	0.6	0