

Rebeca Sanz-Pamplona

List of Publications by Citations

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Version: 2024-04-27

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

51
papers

1,131
citations

20
h-index

32
g-index

65
ext. papers

1,529
ext. citations

7.5
avg, IF

3.96
L-index

#	Paper	IF	Citations
51	Molecular approaches for classifying endometrial carcinoma. <i>Gynecologic Oncology</i> , 2017 , 145, 200-207	4.9	88
50	Aberrant gene expression in mucosa adjacent to tumor reveals a molecular crosstalk in colon cancer. <i>Molecular Cancer</i> , 2014 , 13, 46	42.1	87
49	Comprehensive analysis of copy number aberrations in microsatellite stable colon cancer in view of stromal component. <i>British Journal of Cancer</i> , 2017 , 117, 421-431	8.7	83
48	Germline Mutations in FAN1 Cause Hereditary Colorectal Cancer by Impairing DNA Repair. <i>Gastroenterology</i> , 2015 , 149, 563-6	13.3	75
47	Clinical value of prognosis gene expression signatures in colorectal cancer: a systematic review. <i>PLoS ONE</i> , 2012 , 7, e48877	3.7	69
46	Differences between CAFs and their paired NCF from adjacent colonic mucosa reveal functional heterogeneity of CAFs, providing prognostic information. <i>Molecular Oncology</i> , 2014 , 8, 1290-305	7.9	68
45	Discovery and validation of new potential biomarkers for early detection of colon cancer. <i>PLoS ONE</i> , 2014 , 9, e106748	3.7	61
44	Gene expression differences between colon and rectum tumors. <i>Clinical Cancer Research</i> , 2011 , 17, 7303-12	11.2	59
43	Algorithmic methods to infer the evolutionary trajectories in cancer progression. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, E4025-34	11.5	49
42	Identification of candidate susceptibility genes for colorectal cancer through eQTL analysis. <i>Carcinogenesis</i> , 2014 , 35, 2039-46	4.6	45
41	Exome Sequencing Reveals AMER1 as a Frequently Mutated Gene in Colorectal Cancer. <i>Clinical Cancer Research</i> , 2015 , 21, 4709-18	12.9	35
40	Expression of endoplasmic reticulum stress proteins is a candidate marker of brain metastasis in both ErbB-2+ and ErbB-2- primary breast tumors. <i>American Journal of Pathology</i> , 2011 , 179, 564-79	5.8	34
39	Tools for protein-protein interaction network analysis in cancer research. <i>Clinical and Translational Oncology</i> , 2012 , 14, 3-14	3.6	31
38	A 5-gene classifier from the carcinoma-associated fibroblast transcriptomic profile and clinical outcome in colorectal cancer. <i>Oncotarget</i> , 2014 , 5, 6437-52	3.3	28
37	Uveal Melanoma, Angiogenesis and Immunotherapy, Is There Any Hope?. <i>Cancers</i> , 2019 , 11,	6.6	25
36	Large differences in global transcriptional regulatory programs of normal and tumor colon cells. <i>BMC Cancer</i> , 2014 , 14, 708	4.8	25
35	A transcriptome-proteome integrated network identifies endoplasmic reticulum thiol oxidoreductase (ERp57) as a hub that mediates bone metastasis. <i>Molecular and Cellular Proteomics</i> , 2013 , 12, 2111-25	7.6	25

34	Role of POLE and POLD1 in familial cancer. <i>Genetics in Medicine</i> , 2020 , 22, 2089-2100	8.1	23
33	Lung metastases share common immune features regardless of primary tumor origin 2020 , 8,		22
32	Intrinsic cancer subtypes--next steps into personalized medicine. <i>Cellular Oncology (Dordrecht)</i> , 2015 , 38, 3-16	7.2	20
31	Noncanonical TGFβPathway Relieves the Blockade of IL1β/TGFβMediated Crosstalk between Tumor and Stroma: TGFBR1 and TAK1 Inhibition in Colorectal Cancer. <i>Clinical Cancer Research</i> , 2019 , 25, 4466-4479	12.9	18
30	Extracellular Granzyme A Promotes Colorectal Cancer Development by Enhancing Gut Inflammation. <i>Cell Reports</i> , 2020 , 32, 107847	10.6	18
29	Colorectal cancer: A paradigmatic model for cancer immunology and immunotherapy. <i>Molecular Aspects of Medicine</i> , 2019 , 69, 123-129	16.7	14
28	Colon-specific eQTL analysis to inform on functional SNPs. <i>British Journal of Cancer</i> , 2018 , 119, 971-977	8.7	13
27	T-Type Calcium Channels as Potential Therapeutic Targets in Vemurafenib-Resistant BRAF Melanoma. <i>Journal of Investigative Dermatology</i> , 2020 , 140, 1253-1265	4.3	11
26	GRP94 promotes brain metastasis by engaging pro-survival autophagy. <i>Neuro-Oncology</i> , 2020 , 22, 652-664		11
25	A taxonomy of organ-specific breast cancer metastases based on a protein-protein interaction network. <i>Molecular BioSystems</i> , 2012 , 8, 2085-96		9
24	Lymphocytic infiltration in stage II microsatellite stable colorectal tumors: A retrospective prognosis biomarker analysis. <i>PLoS Medicine</i> , 2020 , 17, e1003292	11.6	8
23	High Cysteinyl Leukotriene Receptor 1 Expression Correlates with Poor Survival of Uveal Melanoma Patients and Cognate Antagonist Drugs Modulate the Growth, Cancer Secretome, and Metabolism of Uveal Melanoma Cells. <i>Cancers</i> , 2020 , 12,	6.6	7
22	Genetic and Immune Changes Associated with Disease Progression under the Pressure of Oncolytic Therapy in A Neuroblastoma Outlier Patient. <i>Cancers</i> , 2020 , 12,	6.6	7
21	Comprehensive molecular characterisation of hereditary non-polyposis colorectal tumours with mismatch repair proficiency. <i>European Journal of Cancer</i> , 2014 , 50, 1964-72	7.5	7
20	Glyceraldehyde-3-phosphate dehydrogenase is overexpressed in colorectal cancer onset. <i>Translational Medicine Communications</i> , 2017 , 2,	4	7
19	DNA methylation events in transcription factors and gene expression changes in colon cancer. <i>Epigenomics</i> , 2020 , 12, 1593-1610	4.4	6
18	Mutanome and expression of immune response genes in microsatellite stable colon cancer. <i>Oncotarget</i> , 2016 , 7, 17711-25	3.3	5
17	Additive Role of Immune System Infiltration and Angiogenesis in Uveal Melanoma Progression. <i>International Journal of Molecular Sciences</i> , 2021 , 22,	6.3	5

16	Altered pathways and colorectal cancer prognosis. <i>BMC Medicine</i> , 2015 , 13, 76	11.4	4
15	AMER1 Is a Frequently Mutated Gene in Colorectal Cancer--Letter. <i>Clinical Cancer Research</i> , 2015 , 21, 4985	12.9	4
14	Analysis of Killer Immunoglobulin-Like Receptor Genes in Colorectal Cancer. <i>Cells</i> , 2020 , 9,	7.9	4
13	Telomere length alterations in microsatellite stable colorectal cancer and association with the immune response. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2018 , 1864, 2992-3000	6.9	3
12	Future Prospects of Colorectal Cancer Screening: Characterizing Interval Cancers. <i>Cancers</i> , 2021 , 13,	6.6	3
11	Unsupervised analyses reveal molecular subtypes associated to prognosis and response to therapy in colorectal cancer. <i>Colorectal Cancer</i> , 2014 , 3, 277-288	0.8	2
10	Algorithmic Methods to Infer the Evolutionary Trajectories in Cancer Progression		2
9	Tumor immune infiltration estimated from gene expression profiles predicts colorectal cancer relapse. <i>Onc Immunology</i> , 2021 , 10, 1862529	7.2	2
8	Identifying causal models between genetically regulated methylation patterns and gene expression in healthy colon tissue. <i>Clinical Epigenetics</i> , 2021 , 13, 162	7.7	2
7	Are Gene Signatures Ready for Use in the Selection of Patients for Adjuvant Treatment?. <i>Current Colorectal Cancer Reports</i> , 2016 , 12, 18-26	1	1
6	Understanding Cancer Progression Using Protein Interaction Networks 2012 , 167-195		1
5	Detection of Merkel cell polyomavirus using whole exome sequencing data		1
4	Predicting MHC I restricted T cell epitopes in mice with NAP-CNB, a novel online tool. <i>Scientific Reports</i> , 2021 , 11, 10780	4.9	1
3	New advances in the clinical management of RAS and BRAF mutant colorectal cancer patients. <i>Expert Review of Gastroenterology and Hepatology</i> , 2021 , 15, 65-79	4.2	1
2	Integrated analysis of circulating immune cellular and soluble mediators reveals specific COVID19 signatures at hospital admission with utility for prediction of clinical outcomes.. <i>Theranostics</i> , 2022 , 12, 290-306	12.1	0
1	Positive impact of a faecal-based screening programme on colorectal cancer mortality risk. <i>PLoS ONE</i> , 2021 , 16, e0253369	3.7	0