

Rebeca Sanz-Pamplona

List of Publications by Year in Descending Order

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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 | 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 |
| 50 | Future Prospects of Colorectal Cancer Screening: Characterizing Interval Cancers. <i>Cancers</i> , 2021 , 13, | 6.6 | 3 |
| 49 | Additive Role of Immune System Infiltration and Angiogenesis in Uveal Melanoma Progression. <i>International Journal of Molecular Sciences</i> , 2021 , 22, | 6.3 | 5 |
| 48 | 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 |
| 47 | Positive impact of a faecal-based screening programme on colorectal cancer mortality risk. <i>PLoS ONE</i> , 2021 , 16, e0253369 | 3.7 | 0 |
| 46 | 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 |
| 45 | Tumor immune infiltration estimated from gene expression profiles predicts colorectal cancer relapse. <i>Onc Immunology</i> , 2021 , 10, 1862529 | 7.2 | 2 |
| 44 | 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 |
| 43 | 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 |
| 42 | 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 |
| 41 | Lung metastases share common immune features regardless of primary tumor origin 2020 , 8, | | 22 |
| 40 | Analysis of Killer Immunoglobulin-Like Receptor Genes in Colorectal Cancer. <i>Cells</i> , 2020 , 9, | 7.9 | 4 |
| 39 | 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 |
| 38 | Lymphocytic infiltration in stage II microsatellite stable colorectal tumors: A retrospective prognosis biomarker analysis. <i>PLoS Medicine</i> , 2020 , 17, e1003292 | 11.6 | 8 |
| 37 | DNA methylation events in transcription factors and gene expression changes in colon cancer. <i>Epigenomics</i> , 2020 , 12, 1593-1610 | 4.4 | 6 |
| 36 | Extracellular Granzyme A Promotes Colorectal Cancer Development by Enhancing Gut Inflammation. <i>Cell Reports</i> , 2020 , 32, 107847 | 10.6 | 18 |
| 35 | Role of POLE and POLD1 in familial cancer. <i>Genetics in Medicine</i> , 2020 , 22, 2089-2100 | 8.1 | 23 |

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| 34 | GRP94 promotes brain metastasis by engaging pro-survival autophagy. <i>Neuro-Oncology</i> , 2020 , 22, 652-664 | 11 |
| 33 | Uveal Melanoma, Angiogenesis and Immunotherapy, Is There Any Hope?. <i>Cancers</i> , 2019 , 11, | 6.6 25 |
| 32 | Colorectal cancer: A paradigmatic model for cancer immunology and immunotherapy. <i>Molecular Aspects of Medicine</i> , 2019 , 69, 123-129 | 16.7 14 |
| 31 | Noncanonical TGF β Pathway Relieves the Blockade of IL1 β /TGF β Mediated 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 | 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 |
| 29 | Colon-specific eQTL analysis to inform on functional SNPs. <i>British Journal of Cancer</i> , 2018 , 119, 971-977 | 8.7 13 |
| 28 | Molecular approaches for classifying endometrial carcinoma. <i>Gynecologic Oncology</i> , 2017 , 145, 200-207 | 4.9 88 |
| 27 | 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 |
| 26 | Glyceraldehyde-3-phosphate dehydrogenase is overexpressed in colorectal cancer onset. <i>Translational Medicine Communications</i> , 2017 , 2, | 4 7 |
| 25 | 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 |
| 24 | 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 |
| 23 | Mutanome and expression of immune response genes in microsatellite stable colon cancer. <i>Oncotarget</i> , 2016 , 7, 17711-25 | 3.3 5 |
| 22 | Exome Sequencing Reveals AMER1 as a Frequently Mutated Gene in Colorectal Cancer. <i>Clinical Cancer Research</i> , 2015 , 21, 4709-18 | 12.9 35 |
| 21 | Germline Mutations in FAN1 Cause Hereditary Colorectal Cancer by Impairing DNA Repair. <i>Gastroenterology</i> , 2015 , 149, 563-6 | 13.3 75 |
| 20 | Altered pathways and colorectal cancer prognosis. <i>BMC Medicine</i> , 2015 , 13, 76 | 11.4 4 |
| 19 | AMER1 Is a Frequently Mutated Gene in Colorectal Cancer--Letter. <i>Clinical Cancer Research</i> , 2015 , 21, 4985 | 12.9 4 |
| 18 | Intrinsic cancer subtypes--next steps into personalized medicine. <i>Cellular Oncology (Dordrecht)</i> , 2015 , 38, 3-16 | 7.2 20 |
| 17 | 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 |

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| 16 | Discovery and validation of new potential biomarkers for early detection of colon cancer. <i>PLoS ONE</i> , 2014 , 9, e106748 | 3.7 | 61 |
| 15 | 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 |
| 14 | Large differences in global transcriptional regulatory programs of normal and tumor colon cells. <i>BMC Cancer</i> , 2014 , 14, 708 | 4.8 | 25 |
| 13 | Identification of candidate susceptibility genes for colorectal cancer through eQTL analysis. <i>Carcinogenesis</i> , 2014 , 35, 2039-46 | 4.6 | 45 |
| 12 | 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 |
| 11 | 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 |
| 10 | 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 |
| 9 | 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 |
| 8 | Tools for protein-protein interaction network analysis in cancer research. <i>Clinical and Translational Oncology</i> , 2012 , 14, 3-14 | 3.6 | 31 |
| 7 | A taxonomy of organ-specific breast cancer metastases based on a protein-protein interaction network. <i>Molecular BioSystems</i> , 2012 , 8, 2085-96 | | 9 |
| 6 | Understanding Cancer Progression Using Protein Interaction Networks 2012 , 167-195 | | 1 |
| 5 | Clinical value of prognosis gene expression signatures in colorectal cancer: a systematic review. <i>PLoS ONE</i> , 2012 , 7, e48877 | 3.7 | 69 |
| 4 | 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 |
| 3 | Gene expression differences between colon and rectum tumors. <i>Clinical Cancer Research</i> , 2011 , 17, 7303-12 | | 59 |
| 2 | Algorithmic Methods to Infer the Evolutionary Trajectories in Cancer Progression | | 2 |
| 1 | Detection of Merkel cell polyomavirus using whole exome sequencing data | | 1 |