

Ibai Goicoechea

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.

22
papers

490
citations

11
h-index

22
g-index

25
ext. papers

669
ext. citations

7.3
avg, IF

3
L-index

#	Paper	IF	Citations
22	Preneoplastic somatic mutations including in lymphoplasmacytic lymphoma.. <i>Science Advances</i> , 2022 , 8, eabl4644	14.3	4
21	Tumor cells in light-chain amyloidosis and myeloma show distinct transcriptional rewiring of normal plasma cell development. <i>Blood</i> , 2021 , 138, 1583-1589	2.2	4
20	Deep MRD profiling defines outcome and unveils different modes of treatment resistance in standard- and high-risk myeloma. <i>Blood</i> , 2021 , 137, 49-60	2.2	28
19	Circulating tumor cells for comprehensive and multiregional non-invasive genetic characterization of multiple myeloma. <i>Leukemia</i> , 2020 , 34, 3007-3018	10.7	7
18	Immunogenomic identification and characterization of granulocytic myeloid-derived suppressor cells in multiple myeloma. <i>Blood</i> , 2020 , 136, 199-209	2.2	31
17	The Urinary Transcriptome as a Source of Biomarkers for Prostate Cancer. <i>Cancers</i> , 2020 , 12,	6.6	8
16	Biological and clinical significance of dysplastic hematopoiesis in patients with newly diagnosed multiple myeloma. <i>Blood</i> , 2020 , 135, 2375-2387	2.2	11
15	Transcriptional profiling of circulating tumor cells in multiple myeloma: a new model to understand disease dissemination. <i>Leukemia</i> , 2020 , 34, 589-603	10.7	19
14	Single-Cell Characterization of the Multiple Myeloma (MM) Immune Microenvironment Identifies CD27-Negative T Cells As Potential Source of Tumor-Reactive Lymphocytes. <i>Blood</i> , 2019 , 134, 506-506	2.2	4
13	The Circulating Transcriptome as a Source of Biomarkers for Melanoma. <i>Cancers</i> , 2019 , 11,	6.6	28
12	Mutational profiling can identify laryngeal dysplasia at risk of progression to invasive carcinoma. <i>Scientific Reports</i> , 2018 , 8, 6613	4.9	13
11	Noncoding RNA Expression and Targeted Next-Generation Sequencing Distinguish Tubulocystic Renal Cell Carcinoma (TC-RCC) from Other Renal Neoplasms. <i>Journal of Molecular Diagnostics</i> , 2018 , 20, 34-45	5.1	16
10	Spatial intratumoural heterogeneity in the expression of GIT1 is associated with poor prognostic outcome in oestrogen receptor positive breast cancer patients with synchronous lymph node metastases. <i>F1000Research</i> , 2017 , 6, 1606	3.6	4
9	Spatial intratumoural heterogeneity in the expression of GIT1 is associated with poor prognostic outcome in oestrogen receptor positive breast cancer patients with synchronous lymph node metastases. <i>F1000Research</i> , 2017 , 6, 1606	3.6	4
8	Aberrant Expression of MicroRNAs in B-cell Lymphomas. <i>MicroRNA (Shariqah, United Arab Emirates)</i> , 2016 , 5, 87-105	2.9	4
7	New Concepts in Cancer Biomarkers: Circulating miRNAs in Liquid Biopsies. <i>International Journal of Molecular Sciences</i> , 2016 , 17,	6.3	162
6	The circulating transcriptome as a source of non-invasive cancer biomarkers: concepts and controversies of non-coding and coding RNA in body fluids. <i>Journal of Cellular and Molecular Medicine</i> , 2015 , 19, 2307-23	5.6	64

5	MicroRNAs as B-cell lymphoma biomarkers. <i>Blood and Lymphatic Cancer: Targets and Therapy</i> , 2015 , 25	2.6	1
4	MGMT Expression Predicts PARP-Mediated Resistance to Temozolomide. <i>Molecular Cancer Therapeutics</i> , 2015 , 14, 1236-46	6.1	26
3	Targeted next-generation sequencing and non-coding RNA expression analysis of clear cell papillary renal cell carcinoma suggests distinct pathological mechanisms from other renal tumour subtypes. <i>Journal of Pathology</i> , 2014 , 232, 32-42	9.4	39
2	Differential chemosensitivity to antifolate drugs between RAS and BRAF melanoma cells. <i>Molecular Cancer</i> , 2014 , 13, 154	42.1	2
1	Characterization of herpes simplex virus 1 strains as platforms for the development of oncolytic viruses against liver cancer. <i>Liver International</i> , 2011 , 31, 1542-53	7.9	10