

Anna Enjuanes

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.

48
papers

4,338
citations

27
h-index

56
g-index

56
ext. papers

5,028
ext. citations

9.9
avg. IF

3.84
L-index

#	Paper	IF	Citations
48	Revised International Prognostic Index and genetic alterations are associated with early failure to R-CHOP in patients with diffuse large B-cell lymphoma. <i>British Journal of Haematology</i> , 2021 , 196, 589	4.5	0
47	MAPK and JAK-STAT pathways dysregulation in plasmablastic lymphoma. <i>Haematologica</i> , 2021 , 106, 2682-2693	6.6	12
46	Comparative analysis of targeted next-generation sequencing panels for the detection of gene mutations in chronic lymphocytic leukemia: an ERIC multi-center study. <i>Haematologica</i> , 2021 , 106, 682-691	6.6	3
45	A Cyclin D1-Dependent Transcriptional Program Predicts Clinical Outcome in Mantle Cell Lymphoma. <i>Clinical Cancer Research</i> , 2021 , 27, 213-225	12.9	2
44	Mutational Landscape and Tumor Burden Assessed by Cell-free DNA in Diffuse Large B-Cell Lymphoma in a Population-Based Study. <i>Clinical Cancer Research</i> , 2021 , 27, 513-521	12.9	11
43	IgCaller for reconstructing immunoglobulin gene rearrangements and oncogenic translocations from whole-genome sequencing in lymphoid neoplasms. <i>Nature Communications</i> , 2020 , 11, 3390	17.4	8
42	PI3K inhibition reshapes follicular lymphoma-immune microenvironment cross talk and unleashes the activity of venetoclax. <i>Blood Advances</i> , 2020 , 4, 4217-4231	7.8	12
41	Distinct molecular profile of IRF4-rearranged large B-cell lymphoma. <i>Blood</i> , 2020 , 135, 274-286	2.2	24
40	Clinical impact of the subclonal architecture and mutational complexity in chronic lymphocytic leukemia. <i>Leukemia</i> , 2018 , 32, 645-653	10.7	64
39	Cyclin D1 overexpression induces global transcriptional downregulation in lymphoid neoplasms. <i>Journal of Clinical Investigation</i> , 2018 , 128, 4132-4147	15.9	14
38	Integrating genomic alterations in diffuse large B-cell lymphoma identifies new relevant pathways and potential therapeutic targets. <i>Leukemia</i> , 2018 , 32, 675-684	10.7	95
37	Dissection of DLBCL microenvironment provides a gene expression-based predictor of survival applicable to formalin-fixed paraffin-embedded tissue. <i>Annals of Oncology</i> , 2018 , 29, 2363-2370	10.3	47
36	The reference epigenome and regulatory chromatin landscape of chronic lymphocytic leukemia. <i>Nature Medicine</i> , 2018 , 24, 868-880	50.5	103
35	Decoding the DNA Methylome of Mantle Cell Lymphoma in the Light of the Entire B Cell Lineage. <i>Cancer Cell</i> , 2016 , 30, 806-821	24.3	73
34	NOTCH1, TP53, and MAP2K1 Mutations in Splenic Diffuse Red Pulp Small B-cell Lymphoma Are Associated With Progressive Disease. <i>American Journal of Surgical Pathology</i> , 2016 , 40, 192-201	6.7	24
33	MYD88 L265P Mutations, But No Other Variants, Identify a Subpopulation of DLBCL Patients of Activated B-cell Origin, Extranodal Involvement, and Poor Outcome. <i>Clinical Cancer Research</i> , 2016 , 22, 2755-64	12.9	42
32	Blocking interferon γ reduces expression of chemokines CXCL9, CXCL10 and CXCL11 and decreases macrophage infiltration in ex vivo cultured arteries from patients with giant cell arteritis. <i>Annals of the Rheumatic Diseases</i> , 2016 , 75, 1177-86	2.4	57

31	Clinical impact of clonal and subclonal TP53, SF3B1, BIRC3, NOTCH1, and ATM mutations in chronic lymphocytic leukemia. <i>Blood</i> , 2016 , 127, 2122-30	2.2	188
30	Non-coding recurrent mutations in chronic lymphocytic leukaemia. <i>Nature</i> , 2015 , 526, 519-24	50.4	565
29	Risk of Central Nervous System (CNS) Involvement in Patients with Mantle Cell Lymphoma (MCL): Analysis of Clinico-Biological Factors in a Series of 283 Cases. <i>Blood</i> , 2014 , 124, 1677-1677	2.2	1
28	Genome-wide methylation analyses identify a subset of mantle cell lymphoma with a high number of methylated CpGs and aggressive clinicopathological features. <i>International Journal of Cancer</i> , 2013 , 133, 2852-63	7.5	11
27	Landscape of somatic mutations and clonal evolution in mantle cell lymphoma. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 18250-5	11.5	377
26	SOX11 regulates PAX5 expression and blocks terminal B-cell differentiation in aggressive mantle cell lymphoma. <i>Blood</i> , 2013 , 121, 2175-85	2.2	109
25	Association of NOS2 and potential effect of VEGF, IL6, CCL2 and IL1RN polymorphisms and haplotypes on susceptibility to GCA—a simultaneous study of 130 potentially functional SNPs in 14 candidate genes. <i>Rheumatology</i> , 2012 , 51, 841-51	3.9	31
24	Identification of methylated genes associated with aggressive clinicopathological features in mantle cell lymphoma. <i>PLoS ONE</i> , 2011 , 6, e19736	3.7	32
23	Epigenetic activation of SOX11 in lymphoid neoplasms by histone modifications. <i>PLoS ONE</i> , 2011 , 6, e21382	3.2	32
22	Effects of bilirubin and sera from jaundiced patients on osteoblasts: contribution to the development of osteoporosis in liver diseases. <i>Hepatology</i> , 2011 , 54, 2104-13	11.2	47
21	A putative "hepitype" in the ATM gene associated with chronic lymphocytic leukemia risk. <i>Genes Chromosomes and Cancer</i> , 2011 , 50, 887-95	5	5
20	Whole-genome sequencing identifies recurrent mutations in chronic lymphocytic leukaemia. <i>Nature</i> , 2011 , 475, 101-5	50.4	1206
19	Verification that common variation at 2q37.1, 6p25.3, 11q24.1, 15q23, and 19q13.32 influences chronic lymphocytic leukaemia risk. <i>British Journal of Haematology</i> , 2010 , 150, 473-9	4.5	33
18	Lithocholic acid downregulates vitamin D effects in human osteoblasts. <i>European Journal of Clinical Investigation</i> , 2010 , 40, 25-34	4.6	40
17	Common variants at 2q37.3, 8q24.21, 15q21.3 and 16q24.1 influence chronic lymphocytic leukemia risk. <i>Nature Genetics</i> , 2010 , 42, 132-6	36.3	196
16	535 HIGH BILIRUBIN REDUCES CELL SURVIVAL AND DIFFERENTIATION OF PRIMARY HUMAN OSTEOBLASTS. IS THIS EFFECT APPLICABLE TO THE SERUM OF CHRONIC CHOLESTATIC PATIENTS?. <i>Journal of Hepatology</i> , 2010 , 52, S214-S215	13.4	2
15	The effect of the alendronate on OPG/RANKL system in differentiated primary human osteoblasts. <i>Endocrine</i> , 2010 , 37, 180-6	4	12
14	The effect of the alendronate on OPG/RANKL system in differentiated primary human osteoblasts. <i>Endocrine</i> , 2010 , 37, 322-8	4	14

13	High osteoprotegerin serum levels in primary biliary cirrhosis are associated with disease severity but not with the mRNA gene expression in liver tissue. <i>Journal of Bone and Mineral Metabolism</i> , 2009 , 27, 347-54	2.9	18
12	Efectos del tabaquismo sobre los niveles plasmáticos de osteoprotegerina en adultos jóvenes sanos. <i>Revista Española De Enfermedades Metabólicas</i> , 2008 , 17, 55-58		
11	Genetic variants in apoptosis and immunoregulation-related genes are associated with risk of chronic lymphocytic leukemia. <i>Cancer Research</i> , 2008 , 68, 10178-86	10.1	64
10	Simvastatin and atorvastatin enhance gene expression of collagen type 1 and osteocalcin in primary human osteoblasts and MG-63 cultures. <i>Journal of Cellular Biochemistry</i> , 2007 , 101, 1430-8	4.7	73
9	COL1A1, ESR1, VDR and TGF β 1 polymorphisms and haplotypes in relation to BMD in Spanish postmenopausal women. <i>Osteoporosis International</i> , 2007 , 18, 235-43	5.3	45
8	A new SNP in a negative regulatory region of the CYP19A1 gene is associated with lumbar spine BMD in postmenopausal women. <i>Bone</i> , 2006 , 38, 738-43	4.7	24
7	The association between common vitamin D receptor gene variations and osteoporosis: a participant-level meta-analysis. <i>Annals of Internal Medicine</i> , 2006 , 145, 255-64	8	185
6	In vitro functional assay of alleles and haplotypes of two COL1A1-promoter SNPs. <i>Bone</i> , 2005 , 36, 902-8	4.7	32
5	Functional analysis of the I.3, I.6, pII and I.4 promoters of CYP19 (aromatase) gene in human osteoblasts and their role in vitamin D and dexamethasone stimulation. <i>European Journal of Endocrinology</i> , 2005 , 153, 981-8	6.5	26
4	Differential genetic effects of ESR1 gene polymorphisms on osteoporosis outcomes. <i>JAMA - Journal of the American Medical Association</i> , 2004 , 292, 2105-14	27.4	238
3	Regulation of CYP19 gene expression in primary human osteoblasts: effects of vitamin D and other treatments. <i>European Journal of Endocrinology</i> , 2003 , 148, 519-26	6.5	45
2	Two new single-nucleotide polymorphisms in the COL1A1 upstream regulatory region and their relationship to bone mineral density. <i>Journal of Bone and Mineral Research</i> , 2002 , 17, 384-93	6.3	77
1	Leptin receptor (OB-R) gene expression in human primary osteoblasts: confirmation. <i>Journal of Bone and Mineral Research</i> , 2002 , 17, 1135; author reply 1136	6.3	16