Joaquim Carreras

List of Publications by Year
in descending order

63	2,217	331259 21	223531 46
papers	citations	h-index	g-index

63	63	63	3143
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	High PTX3 expression is associated with a poor prognosis in diffuse large Bâ€cell lymphoma. Cancer Science, 2022, 113, 334-348.	1.7	23
2	AID is a poor prognostic marker of highâ€grade Bâ€cell lymphoma with MYC and BCL2 and/or BCL6 rearrangements. Pathology International, 2022, 72, 35-42.	0.6	7
3	Anti-HBV drug entecavir ameliorates DSS-induced colitis through PD-L1 induction. Pharmacological Research, 2022, 179, 105918.	3.1	5
4	Artificial Intelligence Analysis of Gene Expression Predicted the Overall Survival of Mantle Cell Lymphoma and a Large Pan-Cancer Series. Healthcare (Switzerland), 2022, 10, 155.	1.0	19
5	Clinicopathological analysis of follicular lymphoma with BCL2, BCL6, and MYC rearrangements. Pathology International, 2022, 72, 321-331.	0.6	5
6	Secreted phospholipase A2 modifies extracellular vesicles and accelerates B cell lymphoma. Cell Metabolism, 2022, 34, 615-633.e8.	7.2	31
7	The Use of the Random Number Generator and Artificial Intelligence Analysis for Dimensionality Reduction of Follicular Lymphoma Transcriptomic Data. BioMedInformatics, 2022, 2, 268-280.	1.0	8
8	9p24.1 Genetic Alteration and PD-L1 Expression Are Characteristic of De Novo and Methotrexate-associated Epstein-Barr Virusâ€“positive Hodgkin Lymphoma, But Not Methotrexate-associated Hodgkin-like Lesions. American Journal of Surgical Pathology, 2022, 46, 1017-1024.	2.1	2
9	The significance of tyrosine kinase receptor B and brain-derived neurotrophic factor expression in salivary duct carcinoma. Annals of Diagnostic Pathology, 2021, 50, 151673.	0.6	2
10	The receptor of the colony-stimulating factor-1 (CSF-1R) is a novel prognostic factor and therapeutic target in follicular lymphoma. Leukemia, 2021, 35, 2635-2649.	3.3	32
11	A Combination of Multilayer Perceptron, Radial Basis Function Artificial Neural Networks and Machine Learning Image Segmentation for the Dimension Reduction and the Prognosis Assessment of Diffuse Large B-Cell Lymphoma. AI, 2021, 2, 106-134.	2.1	24
12	Lack of expression of LMO2 clone SP51 identifies MYC rearrangements in aggressive large B-cell lymphomas. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2021, , 1.	1.4	1
13	Integrative Statistics, Machine Learning and Artificial Intelligence Neural Network Analysis Correlated CSF1R with the Prognosis of Diffuse Large B-Cell Lymphoma. Hemato, 2021, 2, 182-206.	0.2	13
14	High Expression of Caspase-8 Associated with Improved Survival in Diffuse Large B-Cell Lymphoma: Machine Learning and Artificial Neural Networks Analyses. BioMedInformatics, 2021, 1, 18-46.	1.0	14
15	Artificial Neural Network Analysis of Gene Expression Data Predicted Non-Hodgkin Lymphoma Subtypes with High Accuracy. Machine Learning and Knowledge Extraction, 2021, 3, 720-739.	3.2	12
16	Wholeâ€genome copy number and immunohistochemical analyses on surgically resected intracholecystic papillary neoplasms. Pathology International, 2021, 71, 823.	0.6	7
17	Artificial Neural Networks Predicted the Overall Survival and Molecular Subtypes of Diffuse Large B-Cell Lymphoma Using a Pancancer Immune-Oncology Panel. Cancers, 2021, 13, 6384.	1.7	24
18	Monomorphic Epitheliotropic Intestinal T-Cell Lymphoma in Asia Frequently Shows SETD2 Alterations. Cancers, 2020, 12, 3539.	1.7	22

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19	A Single Gene Expression Set Derived from Artificial Intelligence Predicted the Prognosis of Several Lymphoma Subtypes; and High Immunohistochemical Expression of TNFAIP8 Associated with Poor Prognosis in Diffuse Large B-Cell Lymphoma. AI, 2020, 1, 342-360.	2.1	15
20	Artificial Intelligence Analysis of the Gene Expression of Follicular Lymphoma Predicted the Overall Survival and Correlated with the Immune Microenvironment Response Signatures. Machine Learning and Knowledge Extraction, 2020, 2, 647-671.	3.2	14
21	PD-L1/L2 protein levels rapidly increase on monocytes via trogocytosis from tumor cells in classical Hodgkin lymphoma. Leukemia, 2020, 34, 2405-2417.	3.3	31
22	PD-L1 is induced on the hepatocyte surface via CKLF-like MARVEL transmembrane domain-containing protein 6 up-regulation by the anti-HBV drug Entecavir. International Immunology, 2020, 32, 519-531.	1.8	8
23	Methotrexate-associated lymphoproliferative disorder demonstrating composite lymphoma of EBV-negative diffuse large B-cell lymphoma and EBV-positive mucocutaneous ulcer. Journal of Clinical and Experimental Hematopathology: JCEH, 2020, 60, 11-16.	0.3	9
24	Clinicopathological evaluation of methotrexate-associated lymphoproliferative disorders with special focus on Epstein-Barr virus-positive mucocutaneous lesions. Journal of Clinical and Experimental Hematopathology: JCEH, 2020, 60, 159-168.	0.3	8
25	Artificial Intelligence Analysis of Gene Expression Data Predicted the Prognosis of Patients with Diffuse Large B-Cell Lymphoma. Tokai Journal of Experimental and Clinical Medicine, 2020, 45, 37-48.	0.4	14
26	Expression of <scp>IL</scp>â€34 correlates with macrophage infiltration and prognosis of diffuse large Bâ€cell lymphoma. Clinical and Translational Immunology, 2019, 8, e1074.	1.7	13
27	Oncogenic Rag GTPase signalling enhances B cell activation and drives follicular lymphoma sensitive to pharmacological inhibition of mTOR. Nature Metabolism, 2019, 1, 775-789.	5.1	40
28	Prediction of steroid demand in the treatment of patients with ulcerative colitis by immunohistochemical analysis of the mucosal microenvironment and immune checkpoint: role of macrophages and regulatory markers in disease severity. Pathology International, 2019, 69, 260-271.	0.6	10
29	High TNFRSF14 and low BTLA are associated with poor prognosis in Follicular Lymphoma and in Diffuse Large B-cell Lymphoma transformation. Journal of Clinical and Experimental Hematopathology: JCEH, 2019, 59, 1-16.	0.3	36
30	Monomorphic epitheliotropic intestinal Tâ€cell lymphoma with Tâ€cell receptor (TCR) of silent phenotype shows rearrangement of TCRÎ² or TCRÎ³ gene. Pathology International, 2019, 69, 117-118.	0.6	7
31	A case of diffuse large B-cell lymphoma withMYCgene cluster amplification related to chromothripsis. Leukemia and Lymphoma, 2018, 59, 2460-2464.	0.6	2
32	Clinicopathological and genomic analysis of double-hit follicular lymphoma: comparison with high-grade B-cell lymphoma with MYC and BCL2 and/or BCL6 rearrangements. Modern Pathology, 2018, 31, 313-326.	2.9	42
33	Classical Hodgkin lymphoma-type and monomorphic-type post-transplant lymphoproliferative disorder following liver transplantation: a case report. Surgical Case Reports, 2018, 4, 72.	0.2	1
34	Genomic Profile and Pathologic Features of Diffuse Large B-Cell Lymphoma Subtype of Methotrexate-associated Lymphoproliferative Disorder in Rheumatoid Arthritis Patients. American Journal of Surgical Pathology, 2018, 42, 936-950.	2.1	26
35	Clinicopathological characteristics and genomic profile of primary sinonasal tract diffuse large B cell lymphoma (<scp>DLBCL</scp>) reveals gain at 1q31 and <scp>RGS</scp>1 encoding protein; high <scp>RGS</scp>1 immunohistochemical expression associates with poor overall survival in <scp>DLBCL</scp> not otherwise specified (<scp>NOS</scp>). Histopathology, 2017, 70, 595-621.	1.6	41
36	T-cell subsets in lymph nodes identify a subgroup of follicular lymphoma patients with favorable outcome. Leukemia and Lymphoma, 2017, 58, 842-850.	0.6	6

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37	Clinicopathological Analysis of 320 Cases of Diffuse Large B-cell Lymphoma Using the Hans Classifier. Journal of Clinical and Experimental Hematopathology: JCEH, 2017, 57, 54-63.	0.3	20
38	A Case of Pedunculated Esophageal Leiomyoma Successfully Treated by Endoscopic Mucosal Resection. Tokai Journal of Experimental and Clinical Medicine, 2017, 42, 121-125.	0.4	0
39	Composite Follicular Lymphoma and CD5-Positive Nodal Marginal Zone Lymphoma. Journal of Clinical and Experimental Hematopathology: JCEH, 2016, 56, 55-58.	0.3	5
40	Genomic and immunohistochemical profiles of enteropathy-associated T-cell lymphoma in Japan. Pathology, 2016, 48, S159-S160.	0.3	0
41	Clinicopathologic Analysis of Angioimmunoblastic T-cell Lymphoma With or Without RHOA G17V Mutation Using Formalin-fixed Paraffin-embedded Sections. American Journal of Surgical Pathology, 2016, 40, 1041-1050.	2.1	38
42	Genomic and immunohistochemical profiles of enteropathy-associated T-cell lymphoma in Japan. Modern Pathology, 2015, 28, 1286-1296.	2.9	58
43	Intratumoral heterogeneity of <scp>HER</scp>2 protein and amplification of <scp>HER</scp>2 gene in salivary duct carcinoma. Pathology International, 2014, 64, 453-459.	0.6	18
44	Clinical outcome of Epsteinâ€“Barr virusâ€positive diffuse large Bâ€cell lymphoma of the elderly in the rituximab era. Cancer Science, 2014, 105, 1170-1175.	1.7	58
45	The reliability of immunohistochemical analysis of the tumor microenvironment in follicular lymphoma: a validation study from the Lunenburg Lymphoma Biomarker Consortium. Haematologica, 2014, 99, 715-725.	1.7	52
46	Clinicopathological analysis of 502 patients with oral squamous cell carcinoma with special interest to distant metastasis. Tokai Journal of Experimental and Clinical Medicine, 2014, 39, 178-85.	0.4	14
47	Double-Stranded RNA of Intestinal Commensal but Not Pathogenic Bacteria Triggers Production of Protective Interferon-Î². Immunity, 2013, 38, 1187-1197.	6.6	176
48	Overâ€expression of <scp>BACH</scp>2 is related to ongoing somatic hypermutation of the immunoglobulin heavy chain gene variable region of de novo diffuse large <scp>B</scp>â€cell lymphoma. Pathology International, 2013, 63, 339-344.	0.6	2
49	MYD88 (L265P) Mutation in Malignant Lymphoma Using Formalin-Fixed Paraffin-Embedded Section. Journal of Clinical and Experimental Hematopathology: JCEH, 2013, 53, 175-177.	0.3	8
50	P1-008â€ƒâ€ƒDefective immune homeostasis mechanisms in Celiac Disease (CD), in its progression to Refractory Celiac Disease (RCD) and transformation to Enteropathy-Associated T-Cell Lymphoma (EATL) Tj ETQq0 0 0 rgBT /Overlock 10		
51	Incidence and prognostic impact of secondary cytogenetic aberrations in a series of 145 patients with mantle cell lymphoma. Genes Chromosomes and Cancer, 2010, 49, 439-451.	1.5	68
52	Continual monitoring of intraepithelial lymphocyte immunophenotype and clonality is more important than snapshot analysis in the surveillance of refractory coeliac disease. Gut, 2010, 59, 452-460.	6.1	51
53	Genomic profiling of pediatric ALKâ€positive anaplastic large cell lymphoma: A Children's Cancer and Leukaemia Group Study. Genes Chromosomes and Cancer, 2009, 48, 1018-1026.	1.5	23
54	High Numbers of Tumor-Infiltrating Programmed Cell Death 1â€“Positive Regulatory Lymphocytes Are Associated With Improved Overall Survival in Follicular Lymphoma. Journal of Clinical Oncology, 2009, 27, 1470-1476.	0.8	273

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55	Mammalian Target of Rapamycin Inhibition Prevents Glomerular Hypertrophy in a Model of Renal Mass Reduction. Transplantation, 2009, 88, 646-652.	0.5	9
56	T-Cell Subpopulations Quantified by Flow Cytometry in Lymph Node Cell Suspensions Identify a Group of Patients with Follicular Lymphoma with Good Prognosis.. Blood, 2009, 114, 1945-1945.	0.6	0
57	The follicular lymphoma microenvironment: From tumor cell to host immunity. Current Hematologic Malignancy Reports, 2008, 3, 179-186.	1.2	6
58	Primary Cutaneous Small/Medium CD4⁺ T-Cell Lymphomas: A Heterogeneous Group of Tumors With Different Clinicopathologic Features and Outcome. Journal of Clinical Oncology, 2008, 26, 3364-3371.	0.8	163
59	Redistribution of FOXP3-Positive Regulatory T Cells From Lymphoid Tissues to Peripheral Blood in HIV-Infected Patients. Journal of Acquired Immune Deficiency Syndromes (1999), 2007, 46, 529-537.	0.9	28
60	Mammalian Target of Rapamycin Inhibition Halts the Progression of Proteinuria in a Rat Model of Reduced Renal Mass. Journal of the American Society of Nephrology: JASN, 2007, 18, 2653-2660.	3.0	52
61	High numbers of tumor-infiltrating FOXP3-positive regulatory T cells are associated with improved overall survival in follicular lymphoma. Blood, 2006, 108, 2957-2964.	0.6	448
62	ZAP-70 Expression and Stem Cell Transplantation Results in Patients with CLL.. Blood, 2006, 108, 3670-3670.	0.6	0
63	Immunohistochemical analysis of ZAP-70 expression in B-cell lymphoid neoplasms. Journal of Pathology, 2005, 205, 507-513.	2.1	73