Joaquim Carreras

List of Publications by Year
in descending order

63 papers	2,217 citations	331259 21 h-index	223531 46 g-index
63 all docs	63 docs citations	63 times ranked	3143 citing authors

#	Article	IF	Citations
1	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
2	High Numbers of Tumor-Infiltrating Programmed Cell Death 1–Positive Regulatory Lymphocytes Are Associated With Improved Overall Survival in Follicular Lymphoma. Journal of Clinical Oncology, 2009, 27, 1470-1476.	0.8	273
3	Double-Stranded RNA of Intestinal Commensal but Not Pathogenic Bacteria Triggers Production of Protective Interferon-β. Immunity, 2013, 38, 1187-1197.	6.6	176
4	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
5	Immunohistochemical analysis of ZAP-70 expression in B-cell lymphoid neoplasms. Journal of Pathology, 2005, 205, 507-513.	2.1	73
6	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
7	Clinical outcome of Epstein–Barr virus–positive diffuse large B–cell lymphoma of the elderly in the rituximab era. Cancer Science, 2014, 105, 1170-1175.	1.7	58
8	Genomic and immunohistochemical profiles of enteropathy-associated T-cell lymphoma in Japan. Modern Pathology, 2015, 28, 1286-1296.	2.9	58
9	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
10	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
11	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
12	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
13	Clinicopathological characteristics and genomic profile of primary sinonasal tract diffuse large B cell lymphoma (DLBCL) reveals gain at 1q31 and RGS1 encoding protein; high RGS1 immunohistochemical expression associates with poor overall survival in DLBCL not otherwise specified (NOS). Histopathology, 2017, 70, 595-621.	1.6	41
14	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
15	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
16	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
17	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
18	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

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19	Secreted phospholipase A2 modifies extracellular vesicles and accelerates B cell lymphoma. Cell Metabolism, 2022, 34, 615-633.e8.	7.2	31
20	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
21	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
22	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
23	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
24	Genomic profiling of pediatric ALK‐positive anaplastic large cell lymphoma: A Children's Cancer and Leukaemia Group Study. Genes Chromosomes and Cancer, 2009, 48, 1018-1026.	1.5	23
25	High <i>PTX3</i> expression is associated with a poor prognosis in diffuse large B‐cell lymphoma. Cancer Science, 2022, 113, 334-348.	1.7	23
26	Monomorphic Epitheliotropic Intestinal T-Cell Lymphoma in Asia Frequently Shows SETD2 Alterations. Cancers, 2020, 12, 3539.	1.7	22
27	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
28	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
29	Intratumoral heterogeneity of <scp>HER</scp>2 protein and amplification of <i><scp>HER</scp>2</i> gene in salivary duct carcinoma. Pathology International, 2014, 64, 453-459.	0.6	18
30	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
31	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
32	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
33	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
34	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
35	Expression of <scp>IL</scp>‐34 correlates with macrophage infiltration and prognosis of diffuse large B‐cell lymphoma. Clinical and Translational Immunology, 2019, 8, e1074.	1.7	13
36	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

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37	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
38	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
39	Mammalian Target of Rapamycin Inhibition Prevents Glomerular Hypertrophy in a Model of Renal Mass Reduction. Transplantation, 2009, 88, 646-652.	0.5	9
40	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
41	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
42	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
43	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
44	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
45	Monomorphic epitheliotropic intestinal T‐cell lymphoma with T‐cell receptor (TCR) of silent phenotype shows rearrangement of TCRÎ² or TCRÎ³ gene. Pathology International, 2019, 69, 117-118.	0.6	7
46	Whole‐genome copy number and immunohistochemical analyses on surgically resected intracholecystic papillary neoplasms. Pathology International, 2021, 71, 823.	0.6	7
47	AID is a poor prognostic marker of high‐grade B‐cell lymphoma with <i>MYC</i> and <i>BCL2</i> and/or <i>BCL6</i> rearrangements. Pathology International, 2022, 72, 35-42.	0.6	7
48	The follicular lymphoma microenvironment: From tumor cell to host immunity. Current Hematologic Malignancy Reports, 2008, 3, 179-186.	1.2	6
49	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
50	Composite Follicular Lymphoma and CD5-Positive Nodal Marginal Zone Lymphoma. Journal of Clinical and Experimental Hematopathology: JCEH, 2016, 56, 55-58.	0.3	5
51	Anti-HBV drug entecavir ameliorates DSS-induced colitis through PD-L1 induction. Pharmacological Research, 2022, 179, 105918.	3.1	5
52	Clinicopathological analysis of follicular lymphoma with BCL2, BCL6, and MYC rearrangements. Pathology International, 2022, 72, 321-331.	0.6	5
53	Over‐expression of <scp>BACH</scp>2 is related to ongoing somatic hypermutation of the immunoglobulin heavy chain gene variable region of <i>de novo</i> diffuse large <scp>B</scp>‐cell lymphoma. Pathology International, 2013, 63, 339-344.	0.6	2
54	A case of diffuse large B-cell lymphoma with<i>MYC</i>gene cluster amplification related to chromothripsis. Leukemia and Lymphoma, 2018, 59, 2460-2464.	0.6	2

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55	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
56	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
57	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
58	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
59	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 ETQq1 1 0.784314 rgBT /O		
60	Genomic and immunohistochemical profiles of enteropathy-associated T-cell lymphoma in Japan. Pathology, 2016, 48, S159-S160.	0.3	0
61	ZAP-70 Expression and Stem Cell Transplantation Results in Patients with CLL.. Blood, 2006, 108, 3670-3670.	0.6	0
62	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
63	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