Miguel A. Piris

List of Publications by Year in descending order

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421 papers 29,218 citations

81 h-index 7333 152 g-index

431 all docs

431 docs citations

times ranked

431

29177 citing authors

#	Article	IF	CITATIONS
1	PLCÎ ³ 1/PKCÎ, Downstream Signaling Controls Cutaneous T-Cell Lymphoma Development and Progression. Journal of Investigative Dermatology, 2022, 142, 1391-1400.e15.	0.3	5
2	Genetic and phenotypic attributes of splenic marginal zone lymphoma. Blood, 2022, 139, 732-747.	0.6	49
3	Genetic Subtyping and Phenotypic Characterization of the Immune Microenvironment and MYC/BCL2 Double Expression Reveal Heterogeneity in Diffuse Large B-cell Lymphoma. Clinical Cancer Research, 2022, 28, 972-983.	3.2	22
4	Determining clinical course of diffuse large B-cell lymphoma using targeted transcriptome and machine learning algorithms. Blood Cancer Journal, 2022, 12, 25.	2.8	7
5	Localized skinâ€limited blastic plasmacytoid dendritic cell neoplasm. EJHaem, 2022, 3, 560-562.	0.4	2
6	Search for the cause of anaemia and neutropenia in a patient with wellâ€controlled systemic lupus erythematosus. International Journal of Laboratory Hematology, 2022, 44, .	0.7	0
7	CD229 (Ly9) a Novel Biomarker for B-Cell Malignancies and Multiple Myeloma. Cancers, 2022, 14, 2154.	1.7	1
8	An integrated prognostic model for diffuse large Bâ€cell lymphoma treated with immunochemotherapy. EJHaem, 2022, 3, 722-733.	0.4	1
9	The International Consensus Classification of Mature Lymphoid Neoplasms: a report from the Clinical Advisory Committee. Blood, 2022, 140, 1229-1253.	0.6	512
10	Clinical and pathological characteristics of peripheral Tâ€cell lymphomas in a Spanish population: a retrospective study. British Journal of Haematology, 2021, 192, 82-99.	1.2	5
11	Aggressive B-cell Lymphoma with MYC/TP53 Dual Alterations Displays Distinct Clinicopathobiological Features and Response to Novel Targeted Agents. Molecular Cancer Research, 2021, 19, 249-260.	1.5	20
12	Mycosis Fungoides and $S\tilde{A}$ ©zary Syndrome: An Integrative Review of the Pathophysiology, Molecular Drivers, and Targeted Therapy. Cancers, 2021, 13, 1931.	1.7	23
13	Diffuse Large B-Cell Lymphoma: Recognition of Markers for Targeted Therapy. Hemato, 2021, 2, 281-304.	0.2	1
14	Subcutaneous panniculitis-like T-cell lymphoma, lupus erythematosus profundus, and overlapping cases: molecular characterization through the study of 208 genes. Leukemia and Lymphoma, 2021, 62, 2130-2140.	0.6	9
15	Genetic Characterization and Clinical Features of Helicobacter pylori Negative Gastric Mucosa-Associated Lymphoid Tissue Lymphoma. Cancers, 2021, 13, 2993.	1.7	9
16	Peripheral T-cell lymphoma: molecular profiling recognizes subclasses and identifies prognostic markers. Blood Advances, 2021, 5, 5588-5598.	2.5	24
17	Proposal and validation of a method to classify genetic subtypes of diffuse large B cell lymphoma. Scientific Reports, 2021, 11, 1886.	1.6	25
18	Genomic complexity is associated with epigenetic regulator mutations and poor prognosis in diffuse large B-cell lymphoma. Oncolmmunology, 2021, 10, 1928365.	2.1	6

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19	Determining Clinical Course of Diffuse Large B-Cell Lymphoma Using Targeted Transcriptome and Machine Learning Algorithms. Blood, 2021, 138, 2395-2395.	0.6	1
20	Advancedâ€stage mycosis fungoides: role of the signal transducer and activator of transcription 3, nuclear factorâ€₽B and nuclear factor of activated T cells pathways. British Journal of Dermatology, 2020, 182, 147-155.	1.4	26
21	Hodgkin lymphoma: a review of pathological features and recent advances in pathogenesis. Pathology, 2020, 52, 154-165.	0.3	58
22	Expansion of PD1-positive T Cells in Nodal Marginal Zone Lymphoma. American Journal of Surgical Pathology, 2020, 44, 657-664.	2.1	21
23	Update on peripheral T-cell lymphomas with T-helper phenotype: ÂAre there too many subtypes?. Seminars in Diagnostic Pathology, 2020, 37, 24-31.	1.0	10
24	Inflammatory Cells in Atypical Eruption of Lymphocyte Recovery Carry the Same Mutations as Neoplastic Myeloid Cells. American Journal of Dermatopathology, 2020, 42, 360-363.	0.3	1
25	Double hit B cell precursor leukemia/lymphoma in a patient with a prior diagnosis of follicular lymphoma: a diagnostic and therapeutic dilemma. Annals of Hematology, 2020, 99, 391-393.	0.8	1
26	The presence of Merkel cell carcinoma polyomavirus is associated with a distinct phenotype in neoplastic Merkel cell carcinoma cells and their tissue microenvironment. PLoS ONE, 2020, 15, e0232517.	1.1	10
27	An analysis of genetic targets for guiding clinical management of follicular lymphoma. Expert Review of Hematology, 2020, 13, 1361-1372.	1.0	0
28	A refined cell-of-origin classifier with targeted NGS and artificial intelligence shows robust predictive value in DLBCL. Blood Advances, 2020, 4, 3391-3404.	2.5	22
29	XPO1 expression worsens the prognosis of unfavorable DLBCL that can be effectively targeted by selinexor in the absence of mutant p53. Journal of Hematology and Oncology, 2020, 13, 148.	6.9	27
30	Prognostic factors, therapeutic approaches, and distinct immunobiologic features in patients with primary mediastinal large B-cell lymphoma on long-term follow-up. Blood Cancer Journal, 2020, 10, 49.	2.8	31
31	Breast implant-associated Epstein-Barr virus-positive large B-cell lymphomas: a report of three cases. Haematologica, 2020, 105, e412-e414.	1.7	17
32	Epstein–Barr virusâ€associated large Bâ€cell lymphoma transformation in marginal zone Bâ€cell lymphoma: a series of four cases. Histopathology, 2020, 77, 112-122.	1.6	7
33	High-mobility group box (TOX) antibody a useful tool for the identification of B and T cell subpopulations. PLoS ONE, 2020, 15, e0229743.	1.1	10
34	Identification of tipifarnib sensitivity biomarkers in T-cell acute lymphoblastic leukemia and T-cell lymphoma. Scientific Reports, 2020, 10, 6721.	1.6	5
35	Three monocytic neoplasms in a single patient. Leukemia and Lymphoma, 2020, 61, 2523-2526.	0.6	3
36	Splenic Marginal Zone Lymphoma. Encyclopedia of Pathology, 2020, , 474-479.	0.0	0

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37	Lymphoplasmacytic lymphoma associated with diffuse large B-cell lymphoma: Progression or divergent evolution?. PLoS ONE, 2020, 15, e0241634.	1.1	5
38	Molecular Genetics in the Diagnosis and Biology of Lymphoid Neoplasms. American Journal of Clinical Pathology, 2019, 152, 277-301.	0.4	6
39	Double hit B cell precursor leukemia/lymphoma in a patient with a prior diagnosis of follicular lymphoma: a diagnostic and therapeutic dilemma. Annals of Hematology, 2019, 98, 2837-2839.	0.8	0
40	Large cells with CD30 expression and Hodgkin-like features in primary cutaneous marginal zone B-cell lymphoma: a study of 13 cases. European Journal of Cancer, 2019, 119, S23.	1.3	0
41	Immunoglobulin somatic hypermutation has clinical impact in DLBCL and potential implications for immune checkpoint blockade and neoantigen-based immunotherapies., 2019, 7, 272.		22
42	PD-1/PD-L1 expression and interaction by automated quantitative immunofluorescent analysis show adverse prognostic impact in patients with diffuse large B-cell lymphoma having T-cell infiltration: a study from the International DLBCL Consortium Program. Modern Pathology, 2019, 32, 741-754.	2.9	39
43	Clonal dynamics monitoring during clinical evolution in chronic lymphocytic leukaemia. Scientific Reports, 2019, 9, 975.	1.6	8
44	Genomic analyses of microdissected Hodgkin and Reed-Sternberg cells: mutations in epigenetic regulators and p53 are frequent in refractory classic Hodgkin lymphoma. Blood Cancer Journal, 2019, 9, 34.	2.8	23
45	The Spectrum of EBV-Positive Mucocutaneous Ulcer. American Journal of Surgical Pathology, 2019, 43, 201-210.	2.1	41
46	Immune Profiling and Quantitative Analysis Decipher the Clinical Role of Immune-Checkpoint Expression in the Tumor Immune Microenvironment of DLBCL. Cancer Immunology Research, 2019, 7, 644-657.	1.6	106
47	Unraveling transformation of follicular lymphoma to diffuse large B-cell lymphoma. PLoS ONE, 2019, 14, e0212813.	1.1	31
48	Large Cells With CD30 Expression and Hodgkin-like Features in Primary Cutaneous Marginal Zone B-Cell Lymphoma. American Journal of Surgical Pathology, 2019, 43, 1191-1202.	2.1	11
49	Atypical Histiocytic Lesion Preceding a Peripheral T-Cell Lymphoma Involving the Skin Exhibiting the Same Molecular Alterations. American Journal of Dermatopathology, 2019, 41, 148-154.	0.3	2
50	Mycosis Fungoides Associated With Lesions in the Spectrum of Primary Cutaneous CD30+ Lymphoproliferative Disorders: The Same Process or 3 Coexisting Lymphomas?. American Journal of Dermatopathology, 2019, 41, 846-850.	0.3	7
51	CD30-positive primary cutaneous lymphoproliferative disorders: molecular alterations and targeted therapies. Haematologica, 2019, 104, 226-235.	1.7	38
52	DUSP22-rearranged anaplastic lymphomas are characterized by specific morphological features and a lack of cytotoxic and JAK/STAT surrogate markers. Haematologica, 2019, 104, e158-e162.	1.7	28
53	Diseaseâ€biased and shared characteristics of the immunoglobulin gene repertoires in marginal zone B cell lymphoproliferations. Journal of Pathology, 2019, 247, 416-421.	2.1	25
54	MYC and BCL2 mRNA Expression As Determined By NGS Predicts Survival in DLBCL in GCB but Not in ABC Subgroup. Blood, 2019, 134, 5092-5092.	0.6	1

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55	Proof of Concept for Tipifarnib in Relapsed or Refractory Angioimmunoblastic T-Cell Lymphoma (AITL) and CXCL12+ Peripheral T-Cell Lymphoma (PTCL): Preliminary Results from an Open-Label, Phase 2 Study. Blood, 2019, 134, 468-468.	0.6	8
56	Abstract 4725: Efficacy of a new small-molecule inhibitor of histone deacetylase 6 (HDAC6) in preclinical models of B-cell lymphoma and acute myeloid leukemia., 2019,,.		0
57	Richter transformation driven by Epstein–Barr virus reactivation during therapyâ€related immunosuppression in chronic lymphocytic leukaemia. Journal of Pathology, 2018, 245, 61-73.	2.1	24
58	Whole-exome sequencing reveals acquisition of mutations leading to the onset of donor cell leukemia after hematopoietic transplantation: a model of leukemogenesis. Leukemia, 2018, 32, 1822-1826.	3.3	10
59	Castleman Disease and Rosai-Dorfman Disease. Seminars in Diagnostic Pathology, 2018, 35, 44-53.	1.0	15
60	Re-Defining â€~Reactive' lymphadenopathies: How molecular lessons have changed our minds. Seminars in Diagnostic Pathology, 2018, 35, 1-3.	1.0	0
61	Concordant bone marrow involvement of diffuse large B-cell lymphoma represents a distinct clinical and biological entity in the era of immunotherapy. Leukemia, 2018, 32, 353-363.	3.3	36
62	Mutual regulation between <scp>BCL</scp> 6 and a specific set of mi <scp>RNA</scp> s controls <scp>T_{FH}</scp> phenotype in peripheral Tâ€cell lymphoma. British Journal of Haematology, 2018, 182, 587-590.	1.2	6
63	Mutations in the <scp>JAK</scp> / <scp>STAT</scp> pathway genes and activation of the pathway, a relevant finding in nodal Peripheral Tâ€cell lymphoma. British Journal of Haematology, 2018, 183, 497-501.	1.2	17
64	Applied diagnostics in liver cancer. Efficient combinations of sorafenib with targeted inhibitors blocking AKT/mTOR. Oncotarget, 2018, 9, 30869-30882.	0.8	9
65	Overlap at the molecular and immunohistochemical levels between angioimmunoblastic T-cell lymphoma and a subgroup of peripheral T-cell lymphomas without specific morphological features. Oncotarget, 2018, 9, 16124-16133.	0.8	30
66	Updates from the central pathology review in patients with advanced stage mycosis fungoides (MF) and Sezary syndrome (SS) for the Global PROCLIPI study. European Journal of Cancer, 2018, 101, S16-S17.	1.3	0
67	Spontaneously Ruptured Spleen Samples in Patients With Infectious Mononucleosis. American Journal of Clinical Pathology, 2018, 150, 310-317.	0.4	10
68	Clinical Significance of PTEN Deletion, Mutation, and Loss of PTEN Expression in De Novo Diffuse Large B-Cell Lymphoma. Neoplasia, 2018, 20, 574-593.	2.3	64
69	Mycosis fungoides progression could be regulated by microRNAs. PLoS ONE, 2018, 13, e0198477.	1.1	14
70	<i>In vitro</i> and <i>in vivo</i> activity of a new small-molecule inhibitor of HDAC6 in mantle cell lymphoma. Haematologica, 2018, 103, e537-e540.	1.7	15
71	Alteraciones moleculares en leucemia mieloide aguda y sus implicaciones clÃnicas y terapéuticas. Medicina ClÃnica, 2018, 151, 362-367.	0.3	21
72	Molecular Subtypes of Splenic Marginal Zone Lymphoma (SMZL) Are Associated with Distinct Pathogenic Mechanisms and Outcomes - Interim Analysis of the IELSG46 Study. Blood, 2018, 132, 922-922.	0.6	2

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73	Peripheral T-Cell Lymphomas in Spain: Profiling Clinical, Phenotypic and Genetic Characteristics in Spanish Population. Blood, 2018, 132, 2938-2938.	0.6	0
74	Identification of Tipifarnib Sensitivity Biomarkers in T-Cell Tumor Cell Lines. Blood, 2018, 132, 2851-2851.	0.6	0
75	Efficacy of a New Small-Molecule Inhibitor of Histone Deacetylase 6 (HDAC6) in Preclinical Models of B-Cell Lymphoma and Acute Myeloid Leukemia. Blood, 2018, 132, 5383-5383.	0.6	0
76	pâ€< scp>MAPK1 expression associated with poor prognosis in angioimmunoblastic Tâ€eell lymphoma patients. British Journal of Haematology, 2017, 176, 661-664.	1.2	2
77	B-cell lymphoblastic lymphoma presenting as solitary temporal mass with amplification of AML1/RUNX1: case report. Hematological Oncology, 2017, 35, 380-384.	0.8	1
78	Activating mutations and translocations in the guanine exchange factor VAV1 in peripheral T-cell lymphomas. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 764-769.	3.3	100
79	Splenic diffuse red pulp small B-cell lymphoma displays increased expression of cyclin D3 and recurrent CCND3 mutations. Blood, 2017, 129, 1042-1045.	0.6	52
80	Splenic marginal zone lymphoma. Best Practice and Research in Clinical Haematology, 2017, 30, 56-64.	0.7	33
81	AKT Hyperactivation and the Potential of AKT-Targeted Therapy in Diffuse Large B-Cell Lymphoma. American Journal of Pathology, 2017, 187, 1700-1716.	1.9	39
82	Clinical and diagnostic relevance of <i>NOTCH2</i> -and <i>KLF2</i> -mutations in splenic marginal zone lymphoma. Haematologica, 2017, 102, e310-e312.	1.7	31
83	Hepatitis C virus positive diffuse large B-cell lymphomas have distinct molecular features and lack BCL2 translocations. British Journal of Cancer, 2017, 117, 1685-1688.	2.9	13
84	Targeting the T cell receptor \hat{I}^2 -chain constant region for immunotherapy of T cell malignancies. Nature Medicine, 2017, 23, 1416-1423.	15.2	196
85	Loss of PRDM1/BLIMP-1 function contributes to poor prognosis of activated B-cell-like diffuse large B-cell lymphoma. Leukemia, 2017, 31, 625-636.	3.3	47
86	Plasmablastic lymphoma phenotype is determined by genetic alterations in MYC and PRDM1. Modern Pathology, 2017, 30, 85-94.	2.9	63
87	Shared Oncogenic Pathways Implicated in Both Virus-Positive and UV-Induced Merkel Cell Carcinomas. Journal of Investigative Dermatology, 2017, 137, 197-206.	0.3	78
88	Mutational profile of primary breast diffuse large B-cell lymphoma. Oncotarget, 2017, 8, 102888-102897.	0.8	22
89	Molecular basis of targeted therapy in T/NK-cell lymphoma/leukemia: A comprehensive genomic and immunohistochemical analysis of a panel of 33 cell lines. PLoS ONE, 2017, 12, e0177524.	1.1	4
90	Analysis of the mutational landscape of classic Hodgkin lymphoma identifies disease heterogeneity and potential therapeutic targets. Oncotarget, 2017, 8, 111386-111395.	0.8	33

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91	Prognostic impact of concurrent <i>MYC</i> and <i>BCL6</i> rearrangements and expression in <i>de novo</i> diffuse large B-cell lymphoma. Oncotarget, 2016, 7, 2401-2416.	0.8	93
92	Stratifying diffuse large B-cell lymphoma patients treated with chemoimmunotherapy: GCB/non-GCB by immunohistochemistry is still a robust and feasible marker. Oncotarget, 2016, 7, 18036-18049.	0.8	26
93	CD30 Expression by B and T Cells. American Journal of Surgical Pathology, 2016, 40, 378-385.	2.1	37
94	Incidental and Isolated Follicular Lymphoma In Situ and Mantle Cell Lymphoma In Situ Lack Clinical Significance. American Journal of Surgical Pathology, 2016, 40, 943-949.	2.1	30
95	Primary cutaneous follicular helper Tâ€cell lymphoma. Journal of Cutaneous Pathology, 2016, 43, 164-170.	0.7	9
96	Assessment of CD37 B-cell antigen and cell of origin significantly improves risk prediction in diffuse large B-cell lymphoma. Blood, 2016, 128, 3083-3100.	0.6	59
97	Clinical and Biologic Significance of <i>MYC</i> Genetic Mutations in <i>De Novo</i> Diffuse Large B-cell Lymphoma. Clinical Cancer Research, 2016, 22, 3593-3605.	3.2	48
98	Nodal marginal zone mutational signature. Blood, 2016, 128, 1315-1316.	0.6	0
99	C-MYC is related to GATA3 expression and associated with poor prognosis in nodal peripheral T-cell lymphomas. Haematologica, 2016, 101, e336-e338.	1.7	25
100	Angioimmunoblastic Tâ€cell lymphoma with a clonal plasma cell proliferation that underwent immunoglobulin isotype switch in the skin, coinciding with cutaneous disease progression. Journal of Cutaneous Pathology, 2016, 43, 1203-1210.	0.7	7
101	Identification of a new subclass of ALK-negative ALCL expressing aberrant levels of ERBB4 transcripts. Blood, 2016, 127, 221-232.	0.6	97
102	An Immunogenetic Signature of Ongoing Antigen Interactions in Splenic Marginal Zone Lymphoma Expressing IGHV1-2*04 Receptors. Clinical Cancer Research, 2016, 22, 2032-2040.	3.2	17
103	Epstein-Barr virus-associated diffuse large B-cell lymphoma: diagnosis, difficulties and therapeutic options. Expert Review of Anticancer Therapy, 2016, 16, 411-421.	1.1	14
104	Primary testicular diffuse large B-cell lymphoma displays distinct clinical and biological features for treatment failure in rituximab era: a report from the International PTL Consortium. Leukemia, 2016, 30, 361-372.	3.3	55
105	Contribution of JAK2 mutations to T-cell lymphoblastic lymphoma development. Leukemia, 2016, 30, 94-103.	3.3	27
106	Two distinct molecular subtypes of chronic lymphocytic leukemia give new insights on the pathogenesis of the disease and identify novel therapeutic targets. Leukemia and Lymphoma, 2016, 57, 134-142.	0.6	3
107	p63 expression confers significantly better survival outcomes in high-risk diffuse large B-cell lymphoma and demonstrates p53-like and p53-independent tumor suppressor function. Aging, 2016, 8, 345-365.	1.4	19
108	RelA NF-ÎB subunit activation as a therapeutic target in diffuse large B-cell lymphoma. Aging, 2016, 8, 3321-3340.	1.4	29

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109	Akt activation confers an inferior survival in patients with activated B-cell subtype of diffuse large B-cell lymphoma: a report from The International DLBCL Rituximab-CHOP Consortium Program. Clinical Lymphoma, Myeloma and Leukemia, 2015, 15, S220-S221.	0.2	o
110	Clinical and Biological significance of MYC/BCL6 dual gene rearrangements and protein co-expression in de novo diffuse large B-cell lymphoma: a report from the International DLBCL Rituximab-CHOP Consortium Program. Clinical Lymphoma, Myeloma and Leukemia, 2015, 15, S228.	0.2	0
111	MYC Signatures and Characterization of MYC-Driven Aggressive B-Cell Lymphoma. Clinical Lymphoma, Myeloma and Leukemia, 2015, 15, S223.	0.2	O
112	DNA methylation profiling identifies two splenic marginal zone lymphoma subgroups with different clinical and genetic features. Blood, 2015, 125, 1922-1931.	0.6	53
113	Chronic lymphocytic leukemia cells in lymph nodes show frequent NOTCH1 activation. Haematologica, 2015, 100, e200-e203.	1.7	21
114	Nuclear coexpression of NF-κB subunit c-Rel and p53 mutants confers significantly poor survival in diffuse large B-cell lymphoma patients treated with rituximab-CHOP immunochemotherapy: A Report from the International DLBCL Rituximab-CHOP Consortium. Clinical Lymphoma, Myeloma and Leukemia, 2015, 15, S224.	0.2	O
115	Primary cutaneous anaplastic large cell lymphomas with 6p25.3 rearrangement exhibit particular histological features. Histopathology, 2015, 66, 846-855.	1.6	50
116	CD 30-positive transformed follicular lymphoma: two case reports and literature review. Histopathology, 2015, 67, 918-922.	1.6	3
117	Localized lymphomatoid papulosis. International Journal of Dermatology, 2015, 54, e98-100.	0.5	3
118	MYD88 (L265P) Somatic Mutation in Marginal Zone B-cell Lymphoma. American Journal of Surgical Pathology, 2015, 39, 644-651.	2.1	76
119	Dysregulated CXCR4 expression promotes lymphoma cell survival and independently predicts disease progression in germinal center B-cell-like diffuse large B-cell lymphoma. Oncotarget, 2015, 6, 5597-5614.	0.8	61
120	Colorectal Adenomas Contain Multiple Somatic Mutations That Do Not Coincide with Synchronous Adenocarcinoma Specimens. PLoS ONE, 2015, 10, e0119946.	1.1	11
121	Toll-like receptor stimulation in splenic marginal zone lymphoma can modulate cell signaling, activation and proliferation. Haematologica, 2015, 100, 1460-1468.	1.7	19
122	Convergent Mutations and Kinase Fusions Lead to Oncogenic STAT3 Activation in Anaplastic Large Cell Lymphoma. Cancer Cell, 2015, 27, 744.	7.7	2
123	Advances in the diagnosis and treatment of Hodgkin lymphoma and systemic anaplastic large cell lymphoma. Cancer Treatment Communications, 2015, 4, S1-S11.	0.4	1
124	A novel patient-derived tumorgraft model with TRAF1-ALK anaplastic large-cell lymphoma translocation. Leukemia, 2015, 29, 1390-1401.	3.3	42
125	Convergent Mutations and Kinase Fusions Lead to Oncogenic STAT3 Activation in Anaplastic Large Cell Lymphoma. Cancer Cell, 2015, 27, 516-532.	7.7	378
126	Clinical features, tumor biology, and prognosis associated with MYC rearrangement and Myc overexpression in diffuse large B-cell lymphoma patients treated with rituximab-CHOP. Modern Pathology, 2015, 28, 1555-1573.	2.9	48

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127	Prognostic and biological significance of survivin expression in patients with diffuse large B-cell lymphoma treated with rituximab-CHOP therapy. Modern Pathology, 2015, 28, 1297-1314.	2.9	21
128	Evaluation of NF-ÎB subunit expression and signaling pathway activation demonstrates that p52 expression confers better outcome in germinal center B-cell-like diffuse large B-cell lymphoma in association with CD30 and BCL2 functions. Modern Pathology, 2015, 28, 1202-1213.	2.9	17
129	Prognostic Significance of Survivin Expression in Patients with Diffuse Large B-Cell Lymphoma Treated with R-CHOP Therapy: A Report from the International DLBCL Rituximab-CHOP Consortium Program. Clinical Lymphoma, Myeloma and Leukemia, 2015, 15, S216.	0.2	1
130	Longâ€ŧerm followâ€up of doseâ€adjusted EPOCH plus rituximab (DAâ€EPOCHâ€R) in untreated patients with poor prognosis large Bâ€cell lymphoma. A phase II study conducted by the Spanish PETHEMA Group. British Journal of Haematology, 2015, 169, 188-198.	1.2	49
131	Recurrent presence of the PLCG1 S345F mutation in nodal peripheral T-cell lymphomas. Haematologica, 2015, 100, e25-e27.	1.7	37
132	Unique Versus Common: Disease-Biased Immunoglobulin Gene Repertoires Along with Public Antigen Receptor Stereotypes in Marginal Zone B-Cell Lymphoproliferations. Blood, 2015, 126, 1479-1479.	0.6	2
133	CSF1R Protein Expression in Reactive Lymphoid Tissues and Lymphoma: Its Relevance in Classical Hodgkin Lymphoma. PLoS ONE, 2015, 10, e0125203.	1.1	30
134	Clinical and biological significance of <i>de novo</i> CD5+ diffuse large B-cell lymphoma in Western countries. Oncotarget, 2015, 6, 5615-5633.	0.8	72
135	Prognostic impact of c-Rel nuclear expression and <i>REL</i> amplification and crosstalk between c-Rel and the p53 pathway in diffuse large B-cell lymphoma. Oncotarget, 2015, 6, 23157-23180.	0.8	35
136	Age cutoff for Epstein-Barr virus-positive diffuse large B-cell lymphoma-is it necessary?. Oncotarget, 2015, 6, 13933-13945.	0.8	33
137	Mutated JAK kinases and deregulated STAT activity are potential therapeutic targets in cutaneous T-cell lymphoma. Haematologica, 2015, 100, e450-e453.	1.7	59
138	Individualized strategies to target specific mechanisms of disease in malignant melanoma patients displaying unique mutational signatures. Oncotarget, 2015, 6, 25452-25465.	0.8	3
139	Analysis of the Genomic Heterogeneity in Hodgkin Lymphoma Using Next Generation Sequencing. Blood, 2015, 126, 178-178.	0.6	0
140	PIM Kinases as Potential Therapeutic Targets in a Subset of Peripheral T Cell Lymphoma Cases. PLoS ONE, 2014, 9, e112148.	1,1	18
141	FAS system deregulation in T-cell lymphoblastic lymphoma. Cell Death and Disease, 2014, 5, e1110-e1110.	2.7	15
142	Increasing genomic and epigenomic complexity in the clonal evolution from in situ to manifest t(14;18)-positive follicular lymphoma. Leukemia, 2014, 28, 1103-1112.	3.3	60
143	NFκB expression is a feature of both activated B-cell-like and germinal center B-cell-like subtypes of diffuse large B-cell lymphoma. Modern Pathology, 2014, 27, 1331-1337.	2.9	27
144	Adult pityriasis lichenoidesâ€like mycosis fungoides: a clinical variant of mycosis fungoides. International Journal of Dermatology, 2014, 53, 1331-1338.	0.5	12

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145	Phenotypic and Genotypic Profiling of MDM2, Respective to the TP53 Genetic Status, in Diffuse Large B-cell Lymphoma Patients Treated With Rituximab-CHOP Immunochemotherapy: A Report from the International DLBCL Rituximab-CHOP Consortium Program. Clinical Lymphoma, Myeloma and Leukemia, 2014. 14. S146-S147.	0.2	o
146	Prevalence and Clinical Implications of Epstein-Barr Virus Infection in de novo Diffuse Large B-Cell Lymphoma in Western Countries: A report from The International DLBCL Rituximab-CHOP Consortium Program. Clinical Lymphoma, Myeloma and Leukemia, 2014, 14, S144-S145.	0.2	0
147	NF-κB directly mediates epigenetic deregulation of common microRNAs in Epstein-Barr virus-mediated transformation of B-cells and in lymphomas. Nucleic Acids Research, 2014, 42, 11025-11039.	6.5	27
148	Childhood florid follicular hyperplasia with immunoglobulin lightâ€chain restriction in the gastrointestinal tract. Histopathology, 2014, 65, 805-813.	1.6	4
149	Simplification of risk stratification for splenic marginal zone lymphoma: a point-based score for practical use. Leukemia and Lymphoma, 2014, 55, 929-931.	0.6	40
150	Exome sequencing reveals novel and recurrent mutations with clinical impact in blastic plasmacytoid dendritic cell neoplasm. Leukemia, 2014, 28, 823-829.	3.3	148
151	Recurrent mutations in epigenetic regulators, RHOA and FYN kinase in peripheral T cell lymphomas. Nature Genetics, 2014, 46, 166-170.	9.4	534
152	HLA-partially matched cellular therapy (stem-cell microtransplantation) for acute myeloid leukaemia: description of four cases. British Journal of Haematology, 2014, 165, 580-581.	1.2	10
153	Prevalence and Clinical Implications of Epstein–Barr Virus Infection in <i>De Novo</i> Diffuse Large B-Cell Lymphoma in Western Countries. Clinical Cancer Research, 2014, 20, 2338-2349.	3.2	117
154	B-cell lymphoma mutations: improving diagnostics and enabling targeted therapies. Haematologica, 2014, 99, 222-231.	1.7	52
155	Hepatitis C virus-related lymphoproliferative disorders encompass a broader clinical and morphological spectrum than previously recognized: a clinicopathological study. Modern Pathology, 2014, 27, 281-293.	2.9	22
156	The RHOA G17V gene mutation occurs frequently in peripheral T-cell lymphoma and is associated with a characteristic molecular signature. Blood, 2014, 123, 2893-2894.	0.6	53
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