

Miguel A. Piris

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403
papers

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74
h-index

139
g-index

431
ext. papers

26,520
ext. citations

5.8
avg. IF

5.89
L-index

#	Paper	IF	Citations
403	International network of cancer genome projects. <i>Nature</i> , 2010 , 464, 993-8	50.4	1613
402	Loss of acetylation at Lys16 and trimethylation at Lys20 of histone H4 is a common hallmark of human cancer. <i>Nature Genetics</i> , 2005 , 37, 391-400	36.3	1492
401	Whole-genome sequencing identifies recurrent mutations in chronic lymphocytic leukaemia. <i>Nature</i> , 2011 , 475, 101-5	50.4	1206
400	A new immunostain algorithm classifies diffuse large B-cell lymphoma into molecular subtypes with high accuracy. <i>Clinical Cancer Research</i> , 2009 , 15, 5494-502	12.9	507
399	Tumours of histiocytes and accessory dendritic cells: an immunohistochemical approach to classification from the International Lymphoma Study Group based on 61 cases. <i>Histopathology</i> , 2002 , 41, 1-29	7.3	488
398	MYC/BCL2 protein coexpression contributes to the inferior survival of activated B-cell subtype of diffuse large B-cell lymphoma and demonstrates high-risk gene expression signatures: a report from The International DLBCL Rituximab-CHOP Consortium Program. <i>Blood</i> , 2013 , 121, 4021-31; quiz 4250	2.2	473
397	Recurrent mutations in epigenetic regulators, RHOA and FYN kinase in peripheral T cell lymphomas. <i>Nature Genetics</i> , 2014 , 46, 166-70	36.3	413
396	Intravascular lymphoma: clinical presentation, natural history, management and prognostic factors in a series of 38 cases, with special emphasis on the 'cutaneous variant'. <i>British Journal of Haematology</i> , 2004 , 127, 173-83	4.5	413
395	Genomic and gene expression profiling defines indolent forms of mantle cell lymphoma. <i>Cancer Research</i> , 2010 , 70, 1408-18	10.1	373
394	Outcome in Hodgkin's lymphoma can be predicted from the presence of accompanying cytotoxic and regulatory T cells. <i>Clinical Cancer Research</i> , 2005 , 11, 1467-73	12.9	360
393	Cell cycle deregulation in B-cell lymphomas. <i>Blood</i> , 2003 , 101, 1220-35	2.2	288
392	Convergent mutations and kinase fusions lead to oncogenic STAT3 activation in anaplastic large cell lymphoma. <i>Cancer Cell</i> , 2015 , 27, 516-32	24.3	283
391	Mantle cell lymphoma: presenting features, response to therapy, and prognostic factors. <i>Cancer</i> , 1998 , 82, 567-75	6.4	254
390	Comprehensive gene expression profiling and immunohistochemical studies support application of immunophenotypic algorithm for molecular subtype classification in diffuse large B-cell lymphoma: a report from the International DLBCL Rituximab-CHOP Consortium Program Study. <i>Leukemia</i> , 2012 , 26, 2103-13	10.7	251
389	A high-throughput study in melanoma identifies epithelial-mesenchymal transition as a major determinant of metastasis. <i>Cancer Research</i> , 2007 , 67, 3450-60	10.1	246
388	The stress-regulated protein p8 mediates cannabinoid-induced apoptosis of tumor cells. <i>Cancer Cell</i> , 2006 , 9, 301-12	24.3	245
387	Mutational profile and prognostic significance of TP53 in diffuse large B-cell lymphoma patients treated with R-CHOP: report from an International DLBCL Rituximab-CHOP Consortium Program Study. <i>Blood</i> , 2012 , 120, 3986-96	2.2	223

386	Hodgkin and Reed-Sternberg cells harbor alterations in the major tumor suppressor pathways and cell-cycle checkpoints: analyses using tissue microarrays. <i>Blood</i> , 2003 , 101, 681-9	2.2	208
385	Progression in cutaneous malignant melanoma is associated with distinct expression profiles: a tissue microarray-based study. <i>American Journal of Pathology</i> , 2004 , 164, 193-203	5.8	207
384	Splenic marginal zone lymphoma proposals for a revision of diagnostic, staging and therapeutic criteria. <i>Leukemia</i> , 2008 , 22, 487-95	10.7	205
383	The dynamic DNA methylomes of double-stranded DNA viruses associated with human cancer. <i>Genome Research</i> , 2009 , 19, 438-51	9.7	201
382	Primary cutaneous CD4+ small/medium-sized pleomorphic T-cell lymphoma expresses follicular T-cell markers. <i>American Journal of Surgical Pathology</i> , 2009 , 33, 81-90	6.7	186
381	Mantle-cell lymphoma genotypes identified with CGH to BAC microarrays define a leukemic subgroup of disease and predict patient outcome. <i>Blood</i> , 2005 , 105, 4445-54	2.2	165
380	CD30 expression defines a novel subgroup of diffuse large B-cell lymphoma with favorable prognosis and distinct gene expression signature: a report from the International DLBCL Rituximab-CHOP Consortium Program Study. <i>Blood</i> , 2013 , 121, 2715-24	2.2	157
379	p53 and bcl-2 expression in high-grade B-cell lymphomas: correlation with survival time. <i>British Journal of Cancer</i> , 1994 , 69, 337-41	8.7	157
378	Two main genetic pathways lead to the transformation of chronic lymphocytic leukemia to Richter syndrome. <i>Blood</i> , 2013 , 122, 2673-82	2.2	154
377	PLCG1 mutations in cutaneous T-cell lymphomas. <i>Blood</i> , 2014 , 123, 2034-43	2.2	150
376	Splenic marginal zone lymphoma: clinical characteristics and prognostic factors in a series of 60 patients. <i>Blood</i> , 2002 , 100, 1648-1654	2.2	147
375	Targeted activation of innate immunity for therapeutic induction of autophagy and apoptosis in melanoma cells. <i>Cancer Cell</i> , 2009 , 16, 103-14	24.3	146
374	Genome-wide DNA profiling of marginal zone lymphomas identifies subtype-specific lesions with an impact on the clinical outcome. <i>Blood</i> , 2011 , 117, 1595-604	2.2	145
373	Cytogenetic aberrations and their prognostic value in a series of 330 splenic marginal zone B-cell lymphomas: a multicenter study of the Splenic B-Cell Lymphoma Group. <i>Blood</i> , 2010 , 116, 1479-88	2.2	145
372	EBV-positive diffuse large B-cell lymphoma of the elderly is an aggressive post-germinal center B-cell neoplasm characterized by prominent nuclear factor-kB activation. <i>Modern Pathology</i> , 2012 , 25, 968-82	9.8	140
371	Structural profiles of TP53 gene mutations predict clinical outcome in diffuse large B-cell lymphoma: an international collaborative study. <i>Blood</i> , 2008 , 112, 3088-98	2.2	138
370	Analysis of the IgV(H) somatic mutations in splenic marginal zone lymphoma defines a group of unmutated cases with frequent 7q deletion and adverse clinical course. <i>Blood</i> , 2002 , 99, 1299-304	2.2	138
369	7q31-32 allelic loss is a frequent finding in splenic marginal zone lymphoma. <i>American Journal of Pathology</i> , 1999 , 154, 1583-9	5.8	128

368	Patients with diffuse large B-cell lymphoma of germinal center origin with BCL2 translocations have poor outcome, irrespective of MYC status: a report from an International DLBCL rituximab-CHOP Consortium Program Study. <i>Haematologica</i> , 2013 , 98, 255-63	6.6	125
367	Expression of two markers of germinal center T cells (SAP and PD-1) in angioimmunoblastic T-cell lymphoma. <i>Haematologica</i> , 2007 , 92, 1059-66	6.6	125
366	Splenic marginal zone lymphoma: proposal of new diagnostic and prognostic markers identified after tissue and cDNA microarray analysis. <i>Blood</i> , 2005 , 106, 1831-8	2.2	123
365	The molecular signature of mantle cell lymphoma reveals multiple signals favoring cell survival. <i>Cancer Research</i> , 2003 , 63, 8226-32	10.1	120
364	Mycosis fungoides shows concurrent deregulation of multiple genes involved in the TNF signaling pathway: an expression profile study. <i>Blood</i> , 2003 , 102, 1042-50	2.2	118
363	FOXP3, a selective marker for a subset of adult T-cell leukaemia/lymphoma. <i>Leukemia</i> , 2005 , 19, 2247-53	10.7	118
362	Expression of the NF-kappaB targets BCL2 and BIRC5/Survivin characterizes small B-cell and aggressive B-cell lymphomas, respectively. <i>Journal of Pathology</i> , 2005 , 206, 123-34	9.4	117
361	Targeting the T cell receptor ζ chain constant region for immunotherapy of T cell malignancies. <i>Nature Medicine</i> , 2017 , 23, 1416-1423	50.5	112
360	Progression to large B-cell lymphoma in splenic marginal zone lymphoma: a description of a series of 12 cases. <i>American Journal of Surgical Pathology</i> , 2001 , 25, 1268-76	6.7	111
359	Exome sequencing reveals novel and recurrent mutations with clinical impact in blastic plasmacytoid dendritic cell neoplasm. <i>Leukemia</i> , 2014 , 28, 823-9	10.7	110
358	Tumor microenvironment and mitotic checkpoint are key factors in the outcome of classic Hodgkin lymphoma. <i>Blood</i> , 2006 , 108, 662-8	2.2	110
357	Inactivation of the lamin A/C gene by CpG island promoter hypermethylation in hematologic malignancies, and its association with poor survival in nodal diffuse large B-cell lymphoma. <i>Journal of Clinical Oncology</i> , 2005 , 23, 3940-7	2.2	110
356	Molecular heterogeneity in MCL defined by the use of specific VH genes and the frequency of somatic mutations. <i>Blood</i> , 2003 , 101, 4042-6	2.2	109
355	Inhibition of poly(ADP-ribose) polymerase modulates tumor-related gene expression, including hypoxia-inducible factor-1 activation, during skin carcinogenesis. <i>Cancer Research</i> , 2006 , 66, 5744-56	10.1	108
354	Aggressive large B-cell lymphoma with plasma cell differentiation: immunohistochemical characterization of plasmablastic lymphoma and diffuse large B-cell lymphoma with partial plasmablastic phenotype. <i>Haematologica</i> , 2010 , 95, 1342-9	6.6	106
353	Loss of p16/INK4A protein expression in non-Hodgkin's lymphomas is a frequent finding associated with tumor progression. <i>American Journal of Pathology</i> , 1998 , 153, 887-97	5.8	103
352	Peripheral T-cell lymphoma with follicular T-cell markers. <i>American Journal of Surgical Pathology</i> , 2008 , 32, 1787-99	6.7	102
351	TCR- β expression in primary cutaneous T-cell lymphomas. <i>American Journal of Surgical Pathology</i> , 2013 , 37, 375-84	6.7	100

350	PD-1, a follicular T-cell marker useful for recognizing nodular lymphocyte-predominant Hodgkin lymphoma. <i>American Journal of Surgical Pathology</i> , 2008 , 32, 1252-7	6.7	100
349	Rearrangements of MYC gene facilitate risk stratification in diffuse large B-cell lymphoma patients treated with rituximab-CHOP. <i>Modern Pathology</i> , 2014 , 27, 958-71	9.8	99
348	Anomalous High p27/KIP1 Expression in a Subset of Aggressive B-Cell Lymphomas Is Associated With Cyclin D3 Overexpression. p27/KIP1/Cyclin D3 Colocalization in Tumor Cells. <i>Blood</i> , 1999 , 94, 765-772	7.2	98
347	Over 30% of patients with splenic marginal zone lymphoma express the same immunoglobulin heavy variable gene: ontogenetic implications. <i>Leukemia</i> , 2012 , 26, 1638-46	10.7	97
346	Prevalence and clinical implications of epstein-barr virus infection in de novo diffuse large B-cell lymphoma in Western countries. <i>Clinical Cancer Research</i> , 2014 , 20, 2338-49	12.9	96
345	Cancer induction by restriction of oncogene expression to the stem cell compartment. <i>EMBO Journal</i> , 2009 , 28, 8-20	13	96
344	miR-33-mediated downregulation of p53 controls hematopoietic stem cell self-renewal. <i>Cell Cycle</i> , 2010 , 9, 3277-85	4.7	92
343	EBV-associated cutaneous NK/T-cell lymphoma: review of a series of 14 cases from peru in children and young adults. <i>American Journal of Surgical Pathology</i> , 2010 , 34, 1773-82	6.7	91
342	Nodal marginal zone lymphoma: a heterogeneous tumor: a comprehensive analysis of a series of 27 cases. <i>American Journal of Surgical Pathology</i> , 2003 , 27, 762-71	6.7	91
341	Identification of genes involved in resistance to interferon-alpha in cutaneous T-cell lymphoma. <i>American Journal of Pathology</i> , 2002 , 161, 1825-37	5.8	91
340	Whole-exome sequencing in splenic marginal zone lymphoma reveals mutations in genes involved in marginal zone differentiation. <i>Leukemia</i> , 2014 , 28, 1334-40	10.7	90
339	P53 protein expression in lymphomas and reactive lymphoid tissue. <i>Journal of Pathology</i> , 1992 , 166, 235-41	4.1	90
338	Identification of genes involved in imatinib resistance in CML: a gene-expression profiling approach. <i>Leukemia</i> , 2006 , 20, 1047-54	10.7	86
337	The inducible T-cell co-stimulator molecule is expressed on subsets of T cells and is a new marker of lymphomas of T follicular helper cell-derivation. <i>Haematologica</i> , 2010 , 95, 432-9	6.6	84
336	Vorinostat interferes with the signaling transduction pathway of T-cell receptor and synergizes with phosphoinositide-3 kinase inhibitors in cutaneous T-cell lymphoma. <i>Haematologica</i> , 2010 , 95, 613-21	6.6	83
335	Hydroa-like cutaneous T-cell lymphoma: a clinicopathologic and molecular genetic study of 16 pediatric cases from Peru. <i>Applied Immunohistochemistry and Molecular Morphology</i> , 2002 , 10, 7-14	1.9	83
334	Large B-cell lymphoma with Hodgkin's features. <i>Histopathology</i> , 2005 , 47, 101-10	7.3	81
333	Evolving concepts in the pathogenesis of hairy-cell leukaemia. <i>Nature Reviews Cancer</i> , 2006 , 6, 437-48	31.3	79

332	p16(INK4a) gene alterations are frequent in lesions of mycosis fungoides. <i>American Journal of Pathology</i> , 2000 , 156, 1565-72	5.8	77
331	PRDM1/BLIMP1 is commonly inactivated in anaplastic large T-cell lymphoma. <i>Blood</i> , 2013 , 122, 2683-93	2.2	75
330	Variability in the expression of polycomb proteins in different normal and tumoral tissues. A pilot study using tissue microarrays. <i>Modern Pathology</i> , 2006 , 19, 684-94	9.8	75
329	Novel genomic imbalances in B-cell splenic marginal zone lymphomas revealed by comparative genomic hybridization and cytogenetics. <i>American Journal of Pathology</i> , 2001 , 158, 1843-50	5.8	74
328	Nodal marginal zone lymphoma: gene expression and miRNA profiling identify diagnostic markers and potential therapeutic targets. <i>Blood</i> , 2012 , 119, e9-e21	2.2	73
327	miRNA expression in diffuse large B-cell lymphoma treated with chemoimmunotherapy. <i>Blood</i> , 2011 , 118, 1034-40	2.2	73
326	Building an outcome predictor model for diffuse large B-cell lymphoma. <i>American Journal of Pathology</i> , 2004 , 164, 613-22	5.8	72
325	Abnormalities on 1q and 7q are associated with poor outcome in sporadic Burkitt's lymphoma. A cytogenetic and comparative genomic hybridization study. <i>Leukemia</i> , 2003 , 17, 2016-24	10.7	70
324	The presence of STAT1-positive tumor-associated macrophages and their relation to outcome in patients with follicular lymphoma. <i>Haematologica</i> , 2006 , 91, 1605-12	6.6	70
323	MicroRNA signatures in B-cell lymphomas. <i>Blood Cancer Journal</i> , 2012 , 2, e57	7	68
322	Comparative genome profiling across subtypes of low-grade B-cell lymphoma identifies type-specific and common aberrations that target genes with a role in B-cell neoplasia. <i>Haematologica</i> , 2008 , 93, 670-9	6.6	68
321	Update on extranodal lymphomas. Conclusions of the Workshop held by the EAHP and the SH in Thessaloniki, Greece. <i>Histopathology</i> , 2006 , 48, 481-504	7.3	68
320	Prognostic impact of concurrent MYC and BCL6 rearrangements and expression in de novo diffuse large B-cell lymphoma. <i>Oncotarget</i> , 2016 , 7, 2401-16	3.3	68
319	Activating mutations and translocations in the guanine exchange factor VAV1 in peripheral T-cell lymphomas. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017 , 114, 764-769	11.5	67
318	Cutaneous follicular B-cell lymphoma: description of a series of 18 cases. <i>American Journal of Surgical Pathology</i> , 2001 , 25, 875-83	6.7	67
317	Identification of a new subclass of ALK-negative ALCL expressing aberrant levels of ERBB4 transcripts. <i>Blood</i> , 2016 , 127, 221-32	2.2	65
316	PRDM1/BLIMP-1 expression in multiple B and T-cell lymphoma. <i>Haematologica</i> , 2006 , 91, 467-74	6.6	65
315	Mantle cell lymphoma: transcriptional regulation by microRNAs. <i>Leukemia</i> , 2010 , 24, 1335-42	10.7	64

314	Overall survival in aggressive B-cell lymphomas is dependent on the accumulation of alterations in p53, p16, and p27. <i>American Journal of Pathology</i> , 2001 , 159, 205-13	5.8	64
313	PIM2 inhibition as a rational therapeutic approach in B-cell lymphoma. <i>Blood</i> , 2011 , 118, 5517-27	2.2	63
312	Gcet1 (centerin), a highly restricted marker for a subset of germinal center-derived lymphomas. <i>Blood</i> , 2008 , 111, 351-8	2.2	63
311	Risk stratification for Splenic Marginal Zone Lymphoma based on haemoglobin concentration, platelet count, high lactate dehydrogenase level and extrahilar lymphadenopathy: development and validation on 593 cases. <i>British Journal of Haematology</i> , 2012 , 159, 164-71	4.5	62
310	MYD88 (L265P) somatic mutation in marginal zone B-cell lymphoma. <i>American Journal of Surgical Pathology</i> , 2015 , 39, 644-51	6.7	60
309	The role of miRNAs in the pathogenesis and diagnosis of B-cell lymphomas. <i>Blood</i> , 2012 , 120, 1782-90	2.2	60
308	Lymphocyte-rich classical Hodgkin's lymphoma: distinctive tumor and microenvironment markers. <i>Modern Pathology</i> , 2009 , 22, 1006-15	9.8	60
307	Splenic diffuse red pulp small B-cell lymphoma: revision of a series of cases reveals characteristic clinico-pathological features. <i>Haematologica</i> , 2010 , 95, 1122-9	6.6	59
306	Lymphoma microenvironment: culprit or innocent?. <i>Leukemia</i> , 2008 , 22, 49-58	10.7	58
305	Analysis of octamer-binding transcription factors Oct2 and Oct1 and their coactivator BOB.1/OBF.1 in lymphomas. <i>Modern Pathology</i> , 2002 , 15, 211-20	9.8	58
304	Intrafollicular neoplasia/in situ follicular lymphoma: review of a series of 13 cases. <i>Histopathology</i> , 2010 , 56, 658-62	7.3	57
303	A marginal zone pattern may be found in different varieties of non-Hodgkin's lymphoma: the morphology and immunohistology of splenic involvement by B-cell lymphomas simulating splenic marginal zone lymphoma. <i>Histopathology</i> , 1998 , 33, 230-9	7.3	57
302	Combinatorial effects of microRNAs to suppress the Myc oncogenic pathway. <i>Blood</i> , 2011 , 117, 6255-66	2.2	56
301	Genome wide DNA-profiling of HIV-related B-cell lymphomas. <i>British Journal of Haematology</i> , 2010 , 148, 245-55	4.5	56
300	Identification of MNDA as a new marker for nodal marginal zone lymphoma. <i>Leukemia</i> , 2009 , 23, 1847-57	10.7	56
299	Abnormal PcG protein expression in Hodgkin's lymphoma. Relation with E2F6 and NFkappaB transcription factors. <i>Journal of Pathology</i> , 2004 , 204, 528-37	9.4	56
298	Shared Oncogenic Pathways Implicated in Both Virus-Positive and UV-Induced Merkel Cell Carcinomas. <i>Journal of Investigative Dermatology</i> , 2017 , 137, 197-206	4.3	55
297	Silencing of the p18INK4c gene by promoter hypermethylation in Reed-Sternberg cells in Hodgkin lymphomas. <i>Blood</i> , 2004 , 103, 2351-7	2.2	55

296	Clinical and biological significance of de novo CD5+ diffuse large B-cell lymphoma in Western countries. <i>Oncotarget</i> , 2015 , 6, 5615-33	3.3	54
295	Dysregulated CXCR4 expression promotes lymphoma cell survival and independently predicts disease progression in germinal center B-cell-like diffuse large B-cell lymphoma. <i>Oncotarget</i> , 2015 , 6, 5597-614	3.3	53
294	Influence of biologic markers on the outcome of Hodgkin's lymphoma: a study by the Spanish Hodgkin's Lymphoma Study Group. <i>Journal of Clinical Oncology</i> , 2004 , 22, 1664-73	2.2	53
293	Immune Profiling and Quantitative Analysis Decipher the Clinical Role of Immune-Checkpoint Expression in the Tumor Immune Microenvironment of DLBCL. <i>Cancer Immunology Research</i> , 2019 , 7, 644-657	12.5	52
292	Clinical implications of phosphorylated STAT3 expression in De Novo diffuse large B-cell lymphoma. <i>Clinical Cancer Research</i> , 2014 , 20, 5113-23	12.9	52
291	Frequent involvement of chromosomes 1, 3, 7 and 8 in splenic marginal zone B-cell lymphoma. <i>British Journal of Haematology</i> , 1997 , 98, 446-9	4.5	52
290	Splenic small B-cell lymphoma with predominant red pulp involvement: a diffuse variant of splenic marginal zone lymphoma?. <i>Histopathology</i> , 2002 , 40, 22-30	7.3	52
289	Angioimmunoblastic T-cell lymphoma with hyperplastic germinal centres: a neoplasia with origin in the outer zone of the germinal centre? Clinicopathological and immunohistochemical study of 10 cases with follicular T-cell markers. <i>Modern Pathology</i> , 2009 , 22, 753-61	9.8	51
288	Polycomb proteins in hematologic malignancies. <i>Blood</i> , 2010 , 116, 5465-75	2.2	51
287	Immunohistochemical markers for tumor associated macrophages and survival in advanced classical Hodgkin's lymphoma. <i>Haematologica</i> , 2012 , 97, 1080-4	6.6	50
286	miR-217 is an oncogene that enhances the germinal center reaction. <i>Blood</i> , 2014 , 124, 229-39	2.2	49
285	Transcriptional signature of Ecteinascidin 743 (Yondelis, Trabectedin) in human sarcoma cells explanted from chemo-naive patients. <i>Molecular Cancer Therapeutics</i> , 2005 , 4, 814-23	6.1	49
284	Increasing genomic and epigenomic complexity in the clonal evolution from in situ to manifest t(14;18)-positive follicular lymphoma. <i>Leukemia</i> , 2014 , 28, 1103-12	10.7	48
283	T-cell/histiocyte-rich large B-cell lymphoma is a disseminated aggressive neoplasm: differential diagnosis from Hodgkin's lymphoma. <i>Histopathology</i> , 2002 , 41, 216-29	7.3	48
282	Molecular heterogeneity in chronic lymphocytic leukemia is dependent on BCR signaling: clinical correlation. <i>Leukemia</i> , 2007 , 21, 1984-91	10.7	47
281	Aberrant Bcl6 protein expression in mantle cell lymphoma. <i>American Journal of Surgical Pathology</i> , 2004 , 28, 1051-6	6.7	47
280	p14(ARF) nuclear overexpression in aggressive B-cell lymphomas is a sensor of malfunction of the common tumor suppressor pathways. <i>Blood</i> , 2002 , 99, 1411-8	2.2	47
279	NIK controls classical and alternative NF- κ B activation and is necessary for the survival of human T-cell lymphoma cells. <i>Clinical Cancer Research</i> , 2013 , 19, 2319-30	12.9	46

278	Retinoblastoma (Rb) gene product expression in lymphomas. Correlation with Ki67 growth fraction. <i>Journal of Pathology</i> , 1993 , 169, 405-12	9.4	46
277	The RHOA G17V gene mutation occurs frequently in peripheral T-cell lymphoma and is associated with a characteristic molecular signature. <i>Blood</i> , 2014 , 123, 2893-4	2.2	45
276	Plasmablastic lymphoma phenotype is determined by genetic alterations in MYC and PRDM1. <i>Modern Pathology</i> , 2017 , 30, 85-94	9.8	45
275	Single nucleotide polymorphism-arrays provide new insights in the pathogenesis of post-transplant diffuse large B-cell lymphoma. <i>British Journal of Haematology</i> , 2010 , 149, 569-77	4.5	45
274	Genomic lesions associated with a different clinical outcome in diffuse large B-Cell lymphoma treated with R-CHOP-21. <i>British Journal of Haematology</i> , 2010 , 151, 221-31	4.5	45
273	Nodal and splenic marginal zone B cell lymphomas. <i>Hematological Oncology</i> , 2005 , 23, 108-18	1.3	45
272	p16(INK4a) is selectively silenced in the tumoral progression of mycosis fungoides. <i>Laboratory Investigation</i> , 2002 , 82, 123-32	5.9	44
271	B-cell lymphoma mutations: improving diagnostics and enabling targeted therapies. <i>Haematologica</i> , 2014 , 99, 222-31	6.6	43
270	A cyclin-D1 interaction with BAX underlies its oncogenic role and potential as a therapeutic target in mantle cell lymphoma. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 12461-6	11.5	43
269	A short mutational hot spot in the first intron of BCL-6 is associated with increased BCL-6 expression and with longer overall survival in large B-cell lymphomas. <i>American Journal of Pathology</i> , 2002 , 160, 1371-80	5.8	43
268	Long-term follow-up of dose-adjusted EPOCH plus rituximab (DA-EPOCH-R) in untreated patients with poor prognosis large B-cell lymphoma. A phase II study conducted by the Spanish PETHEMA Group. <i>British Journal of Haematology</i> , 2015 , 169, 188-98	4.5	42
267	The Epstein Barr-encoded BART-6-3p microRNA affects regulation of cell growth and immune response in Burkitt lymphoma. <i>Infectious Agents and Cancer</i> , 2014 , 9, 12	3.5	42
266	Crystal-storing histiocytosis and immunocytoma associated with multifocal fibrosclerosis. <i>Histopathology</i> , 1998 , 33, 459-64	7.3	42
265	Composite Hodgkin lymphoma and mantle cell lymphoma: two clonally unrelated tumors. <i>American Journal of Surgical Pathology</i> , 2003 , 27, 1577-80	6.7	42
264	. <i>Applied Immunohistochemistry & Molecular Morphology</i> , 2002 , 10, 7-14		42
263	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. <i>Leukemia</i> , 2016 , 30, 361-72	10.7	41
262	MDM2 phenotypic and genotypic profiling, respective to 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. <i>Blood</i> , 2013 , 122, 2630-40	2.2	41
261	Large B-cell lymphomas with plasmablastic differentiation: a biological and therapeutic challenge. <i>Leukemia and Lymphoma</i> , 2012 , 53, 185-94	1.9	41

260	Splenic marginal zone lymphoma with increased number of blasts: an aggressive variant?. <i>Human Pathology</i> , 1999 , 30, 1153-60	3.7	41
259	Detection of the bcl-1 rearrangement at the major translocation cluster in frozen and paraffin-embedded tissues of mantle cell lymphomas by polymerase chain reaction. <i>American Journal of Clinical Pathology</i> , 1996 , 105, 532-7	1.9	41
258	Lymph node involvement by splenic marginal zone lymphoma: morphological and immunohistochemical features. <i>American Journal of Surgical Pathology</i> , 1997 , 21, 772-80	6.7	41
257	Assessment of CD37 B-cell antigen and cell of origin significantly improves risk prediction in diffuse large B-cell lymphoma. <i>Blood</i> , 2016 , 128, 3083-3100	2.2	41
256	Clinical features, tumor biology, and prognosis associated with MYC rearrangement and Myc overexpression in diffuse large B-cell lymphoma patients treated with rituximab-CHOP. <i>Modern Pathology</i> , 2015 , 28, 1555-73	9.8	40
255	DNA methylation profiling identifies two splenic marginal zone lymphoma subgroups with different clinical and genetic features. <i>Blood</i> , 2015 , 125, 1922-31	2.2	40
254	Persistent and generalized lymphadenopathy: a lesion of follicular dendritic cells? An immunohistologic and ultrastructural study. <i>American Journal of Clinical Pathology</i> , 1987 , 87, 716-24	1.9	40
253	SPIB, a novel immunohistochemical marker for human blastic plasmacytoid dendritic cell neoplasms: characterization of its expression in major hematolymphoid neoplasms. <i>Blood</i> , 2013 , 121, 643-7	2.2	39
252	A molecular risk score based on 4 functional pathways for advanced classical Hodgkin lymphoma. <i>Blood</i> , 2010 , 116, e12-7	2.2	39
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