

# Randy D Gascoyne

## List of Publications by Citations

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695  
papers

73,412  
citations

127  
h-index

261  
g-index

741  
ext. papers

84,265  
ext. citations

5.6  
avg, IF

7.21  
L-index

#	Paper	IF	Citations
695	Circos: an information aesthetic for comparative genomics. <i>Genome Research</i> , <b>2009</b> , 19, 1639-45	9.7	6014
694	Revised response criteria for malignant lymphoma. <i>Journal of Clinical Oncology</i> , <b>2007</b> , 25, 579-86	2.2	3483
693	The use of molecular profiling to predict survival after chemotherapy for diffuse large-B-cell lymphoma. <i>New England Journal of Medicine</i> , <b>2002</b> , 346, 1937-47	59.2	2971
692	Confirmation of the molecular classification of diffuse large B-cell lymphoma by immunohistochemistry using a tissue microarray. <i>Blood</i> , <b>2004</b> , 103, 275-82	2.2	2955
691	Stromal gene signatures in large-B-cell lymphomas. <i>New England Journal of Medicine</i> , <b>2008</b> , 359, 2313-23	59.2	1281
690	Somatic mutations altering EZH2 (Tyr641) in follicular and diffuse large B-cell lymphomas of germinal-center origin. <i>Nature Genetics</i> , <b>2010</b> , 42, 181-5	36.3	1273
689	Frequent mutation of histone-modifying genes in non-Hodgkin lymphoma. <i>Nature</i> , <b>2011</b> , 476, 298-303	50.4	1180
688	Chronic active B-cell-receptor signalling in diffuse large B-cell lymphoma. <i>Nature</i> , <b>2010</b> , 463, 88-92	50.4	1149
687	Prediction of survival in follicular lymphoma based on molecular features of tumor-infiltrating immune cells. <i>New England Journal of Medicine</i> , <b>2004</b> , 351, 2159-69	59.2	1141
686	Oncogenically active MYD88 mutations in human lymphoma. <i>Nature</i> , <b>2011</b> , 470, 115-9	50.4	1068
685	Rituximab-CHOP versus CHOP alone or with maintenance rituximab in older patients with diffuse large B-cell lymphoma. <i>Journal of Clinical Oncology</i> , <b>2006</b> , 24, 3121-7	2.2	1036
684	Tumor-associated macrophages and survival in classic Hodgkin's lymphoma. <i>New England Journal of Medicine</i> , <b>2010</b> , 362, 875-85	59.2	961
683	The revised International Prognostic Index (R-IPI) is a better predictor of outcome than the standard IPI for patients with diffuse large B-cell lymphoma treated with R-CHOP. <i>Blood</i> , <b>2007</b> , 109, 1857-61	2.2	955
682	Molecular diagnosis of primary mediastinal B cell lymphoma identifies a clinically favorable subgroup of diffuse large B cell lymphoma related to Hodgkin lymphoma. <i>Journal of Experimental Medicine</i> , <b>2003</b> , 198, 851-62	16.6	869
681	Genetics and Pathogenesis of Diffuse Large B-Cell Lymphoma. <i>New England Journal of Medicine</i> , <b>2018</b> , 378, 1396-1407	59.2	780
680	Introduction of combined CHOP plus rituximab therapy dramatically improved outcome of diffuse large B-cell lymphoma in British Columbia. <i>Journal of Clinical Oncology</i> , <b>2005</b> , 23, 5027-33	2.2	759
679	The proliferation gene expression signature is a quantitative integrator of oncogenic events that predicts survival in mantle cell lymphoma. <i>Cancer Cell</i> , <b>2003</b> , 3, 185-97	24.3	751

678	Molecular subtypes of diffuse large B-cell lymphoma arise by distinct genetic pathways. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2008</b> , 105, 13520-5	11.5	746
677	Molecular diagnosis of Burkitt's lymphoma. <i>New England Journal of Medicine</i> , <b>2006</b> , 354, 2431-42	59.2	700
676	Anti-CD47 antibody synergizes with rituximab to promote phagocytosis and eradicate non-Hodgkin lymphoma. <i>Cell</i> , <b>2010</b> , 142, 699-713	56.2	672
675	Concurrent expression of MYC and BCL2 in diffuse large B-cell lymphoma treated with rituximab plus cyclophosphamide, doxorubicin, vincristine, and prednisone. <i>Journal of Clinical Oncology</i> , <b>2012</b> , 30, 3452-9	2.2	669
674	ALK- anaplastic large-cell lymphoma is clinically and immunophenotypically different from both ALK+ ALCL and peripheral T-cell lymphoma, not otherwise specified: report from the International Peripheral T-Cell Lymphoma Project. <i>Blood</i> , <b>2008</b> , 111, 5496-504	2.2	664
673	Oncogenic CARD11 mutations in human diffuse large B cell lymphoma. <i>Science</i> , <b>2008</b> , 319, 1676-9	33.3	660
672	Burkitt lymphoma pathogenesis and therapeutic targets from structural and functional genomics. <i>Nature</i> , <b>2012</b> , 490, 116-20	50.4	600
671	EZH2 is required for germinal center formation and somatic EZH2 mutations promote lymphoid transformation. <i>Cancer Cell</i> , <b>2013</b> , 23, 677-92	24.3	547
670	An enhanced International Prognostic Index (NCCN-IPI) for patients with diffuse large B-cell lymphoma treated in the rituximab era. <i>Blood</i> , <b>2014</b> , 123, 837-42	2.2	517
669	A new immunostain algorithm classifies diffuse large B-cell lymphoma into molecular subtypes with high accuracy. <i>Clinical Cancer Research</i> , <b>2009</b> , 15, 5494-502	12.9	507
668	Rituximab maintenance improves clinical outcome of relapsed/resistant follicular non-Hodgkin lymphoma in patients both with and without rituximab during induction: results of a prospective randomized phase 3 intergroup trial. <i>Blood</i> , <b>2006</b> , 108, 3295-301	2.2	499
667	MYC gene rearrangements are associated with a poor prognosis in diffuse large B-cell lymphoma patients treated with R-CHOP chemotherapy. <i>Blood</i> , <b>2009</b> , 114, 3533-7	2.2	484
666	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> , <b>2013</b> , 121, 4021-31; quiz 4250	2.2	473
665	MHC class II transactivator CIITA is a recurrent gene fusion partner in lymphoid cancers. <i>Nature</i> , <b>2011</b> , 471, 377-81	50.4	467
664	Somatic mutations at EZH2 Y641 act dominantly through a mechanism of selectively altered PRC2 catalytic activity, to increase H3K27 trimethylation. <i>Blood</i> , <b>2011</b> , 117, 2451-9	2.2	458
663	Lymphomas with concurrent BCL2 and MYC translocations: the critical factors associated with survival. <i>Blood</i> , <b>2009</b> , 114, 2273-9	2.2	439
662	Prognostic Significance of Anaplastic Lymphoma Kinase (ALK) Protein Expression in Adults With Anaplastic Large Cell Lymphoma. <i>Blood</i> , <b>1999</b> , 93, 3913-3921	2.2	434
661	Prognostic Significance of Bcl-2 Protein Expression and Bcl-2 Gene Rearrangement in Diffuse Aggressive Non-Hodgkin's Lymphoma. <i>Blood</i> , <b>1997</b> , 90, 244-251	2.2	422

660	Determining cell-of-origin subtypes of diffuse large B-cell lymphoma using gene expression in formalin-fixed paraffin-embedded tissue. <i>Blood</i> , <b>2014</b> , 123, 1214-7	2.2	404
659	Mantle Cell Lymphoma: A Clinicopathologic Study of 80 Cases. <i>Blood</i> , <b>1997</b> , 89, 2067-2078	2.2	402
658	Analysis of multiple biomarkers shows that lymphoma-associated macrophage (LAM) content is an independent predictor of survival in follicular lymphoma (FL). <i>Blood</i> , <b>2005</b> , 106, 2169-74	2.2	386
657	Immunohistochemical methods for predicting cell of origin and survival in patients with diffuse large B-cell lymphoma treated with rituximab. <i>Journal of Clinical Oncology</i> , <b>2011</b> , 29, 200-7	2.2	360
656	Integration of gene mutations in risk prognostication for patients receiving first-line immunochemotherapy for follicular lymphoma: a retrospective analysis of a prospective clinical trial and validation in a population-based registry. <i>Lancet Oncology, The</i> , <b>2015</b> , 16, 1111-1122	21.7	347
655	Diffuse large B-cell lymphoma: optimizing outcome in the context of clinical and biologic heterogeneity. <i>Blood</i> , <b>2015</b> , 125, 22-32	2.2	340
654	The tumour microenvironment in B cell lymphomas. <i>Nature Reviews Cancer</i> , <b>2014</b> , 14, 517-34	31.3	308
653	Diffuse large B-cell lymphoma subgroups have distinct genetic profiles that influence tumor biology and improve gene-expression-based survival prediction. <i>Blood</i> , <b>2005</b> , 106, 3183-90	2.2	304
652	Gene expression signatures delineate biological and prognostic subgroups in peripheral T-cell lymphoma. <i>Blood</i> , <b>2014</b> , 123, 2915-23	2.2	299
651	SOX11 expression is highly specific for mantle cell lymphoma and identifies the cyclin D1-negative subtype. <i>Haematologica</i> , <b>2009</b> , 94, 1555-62	6.6	299
650	Population-based analysis of incidence and outcome of transformed non-Hodgkin's lymphoma. <i>Journal of Clinical Oncology</i> , <b>2008</b> , 26, 5165-9	2.2	291
649	Peripheral T-cell lymphoma, not otherwise specified: a report of 340 cases from the International Peripheral T-cell Lymphoma Project. <i>Blood</i> , <b>2011</b> , 117, 3402-8	2.2	290
648	Mutational and structural analysis of diffuse large B-cell lymphoma using whole-genome sequencing. <i>Blood</i> , <b>2013</b> , 122, 1256-65	2.2	289
647	Cooperative signaling through the signal transducer and activator of transcription 3 and nuclear factor- $\kappa$ B pathways in subtypes of diffuse large B-cell lymphoma. <i>Blood</i> , <b>2008</b> , 111, 3701-13	2.2	287
646	Molecular pathogenesis of Hodgkin's lymphoma: increasing evidence of the importance of the microenvironment. <i>Journal of Clinical Oncology</i> , <b>2011</b> , 29, 1812-26	2.2	283
645	Cyclin D1-negative mantle cell lymphoma: a clinicopathologic study based on gene expression profiling. <i>Blood</i> , <b>2005</b> , 106, 4315-21	2.2	279
644	Lenalidomide combined with R-CHOP overcomes negative prognostic impact of non-germinal center B-cell phenotype in newly diagnosed diffuse large B-Cell lymphoma: a phase II study. <i>Journal of Clinical Oncology</i> , <b>2015</b> , 33, 251-7	2.2	268
643	Whole transcriptome sequencing reveals recurrent NOTCH1 mutations in mantle cell lymphoma. <i>Blood</i> , <b>2012</b> , 119, 1963-71	2.2	264

642	Survival of patients with peripheral T-cell lymphoma after first relapse or progression: spectrum of disease and rare long-term survivors. <i>Journal of Clinical Oncology</i> , <b>2013</b> , 31, 1970-6	2.2	260
641	Loss of MHC class II gene and protein expression in diffuse large B-cell lymphoma is related to decreased tumor immunosurveillance and poor patient survival regardless of other prognostic factors: a follow-up study from the Leukemia and Lymphoma Molecular Profiling Project. <i>Blood</i> , <b>2004</b> , 103, 4251-8	2.2	256
640	Enteropathy-associated T-cell lymphoma: clinical and histological findings from the international peripheral T-cell lymphoma project. <i>Blood</i> , <b>2011</b> , 118, 148-55	2.2	254
639	The histone lysine methyltransferase KMT2D sustains a gene expression program that represses B cell lymphoma development. <i>Nature Medicine</i> , <b>2015</b> , 21, 1199-208	50.5	247
638	ALK-positive diffuse large B-cell lymphoma is associated with Clathrin-ALK rearrangements: report of 6 cases. <i>Blood</i> , <b>2003</b> , 102, 2568-73	2.2	243
637	Immunohistochemical prognostic markers in diffuse large B-cell lymphoma: validation of tissue microarray as a prerequisite for broad clinical applications--a study from the Lunenburg Lymphoma Biomarker Consortium. <i>Journal of Clinical Oncology</i> , <b>2007</b> , 25, 805-12	2.2	241
636	Maintenance rituximab after cyclophosphamide, vincristine, and prednisone prolongs progression-free survival in advanced indolent lymphoma: results of the randomized phase III ECOG1496 Study. <i>Journal of Clinical Oncology</i> , <b>2009</b> , 27, 1607-14	2.2	238
635	BCL2 expression is a prognostic marker for the activated B-cell-like type of diffuse large B-cell lymphoma. <i>Journal of Clinical Oncology</i> , <b>2006</b> , 24, 961-8	2.2	238
634	Pharmacological and genomic profiling identifies NF- $\kappa$ B-targeted treatment strategies for mantle cell lymphoma. <i>Nature Medicine</i> , <b>2014</b> , 20, 87-92	50.5	237
633	Prognostic Significance of Diffuse Large B-Cell Lymphoma Cell of Origin Determined by Digital Gene Expression in Formalin-Fixed Paraffin-Embedded Tissue Biopsies. <i>Journal of Clinical Oncology</i> , <b>2015</b> , 33, 2848-56	2.2	236
632	Phase II study of oral vinorelbine in first-line advanced breast cancer chemotherapy. <i>Journal of Clinical Oncology</i> , <b>2003</b> , 21, 35-40	2.2	235
631	BCL2 translocation defines a unique tumor subset within the germinal center B-cell-like diffuse large B-cell lymphoma. <i>American Journal of Pathology</i> , <b>2004</b> , 165, 159-66	5.8	232
630	Cooperative epigenetic modulation by cancer amplicon genes. <i>Cancer Cell</i> , <b>2010</b> , 18, 590-605	24.3	230
629	Peripheral T-cell lymphoma. <i>Blood</i> , <b>2011</b> , 117, 6756-67	2.2	228
628	EZH2-mediated epigenetic silencing in germinal center B cells contributes to proliferation and lymphomagenesis. <i>Blood</i> , <b>2010</b> , 116, 5247-55	2.2	227
627	Prognostic significance of Bcl-6 protein expression in DLBCL treated with CHOP or R-CHOP: a prospective correlative study. <i>Blood</i> , <b>2006</b> , 107, 4207-13	2.2	227
626	Non-Hodgkin lymphoma. <i>Lancet, The</i> , <b>2017</b> , 390, 298-310	40	225
625	Signal transducer and activator of transcription 6 is frequently activated in Hodgkin and Reed-Sternberg cells of Hodgkin lymphoma. <i>Blood</i> , <b>2002</b> , 99, 618-26	2.2	223

624	Genomic rearrangements involving programmed death ligands are recurrent in primary mediastinal large B-cell lymphoma. <i>Blood</i> , <b>2014</b> , 123, 2062-5	2.2	215
623	Pathogenesis of follicular lymphoma. <i>Journal of Clinical Investigation</i> , <b>2012</b> , 122, 3424-31	15.9	215
622	Differentiation stage-specific expression of microRNAs in B lymphocytes and diffuse large B-cell lymphomas. <i>Blood</i> , <b>2009</b> , 113, 3754-64	2.2	213
621	EZH2 mutations are frequent and represent an early event in follicular lymphoma. <i>Blood</i> , <b>2013</b> , 122, 3165-8		208
620	Randomized phase III trial of ABVD versus Stanford V with or without radiation therapy in locally extensive and advanced-stage Hodgkin lymphoma: an intergroup study coordinated by the Eastern Cooperative Oncology Group (E2496). <i>Journal of Clinical Oncology</i> , <b>2013</b> , 31, 684-91	2.2	204
619	CNS International Prognostic Index: A Risk Model for CNS Relapse in Patients With Diffuse Large B-Cell Lymphoma Treated With R-CHOP. <i>Journal of Clinical Oncology</i> , <b>2016</b> , 34, 3150-6	2.2	201
618	Whole-genome analysis and HLA genotyping of enteropathy-type T-cell lymphoma reveals 2 distinct lymphoma subtypes. <i>Gastroenterology</i> , <b>2007</b> , 132, 1902-11	13.3	200
617	Etiologic heterogeneity among non-Hodgkin lymphoma subtypes: the InterLymph Non-Hodgkin Lymphoma Subtypes Project. <i>Journal of the National Cancer Institute Monographs</i> , <b>2014</b> , 2014, 130-44	4.8	199
616	Loss of signalling via G $\beta$ 3 in germinal centre B-cell-derived lymphoma. <i>Nature</i> , <b>2014</b> , 516, 254-8	50.4	192
615	Comparison of cytogenetic analysis, southern analysis, and polymerase chain reaction for the detection of t(14; 18) in follicular lymphoma. <i>American Journal of Clinical Pathology</i> , <b>1995</b> , 103, 472-8	1.9	190
614	US Intergroup Trial of Response-Adapted Therapy for Stage III to IV Hodgkin Lymphoma Using Early Interim Fluorodeoxyglucose-Positron Emission Tomography Imaging: Southwest Oncology Group S0816. <i>Journal of Clinical Oncology</i> , <b>2016</b> , 34, 2020-7	2.2	184
613	Point mutations and genomic deletions in CCND1 create stable truncated cyclin D1 mRNAs that are associated with increased proliferation rate and shorter survival. <i>Blood</i> , <b>2007</b> , 109, 4599-606	2.2	183
612	MALT1 small molecule inhibitors specifically suppress ABC-DLBCL in vitro and in vivo. <i>Cancer Cell</i> , <b>2012</b> , 22, 812-24	24.3	182
611	The BCL6 transcriptional program features repression of multiple oncogenes in primary B cells and is deregulated in DLBCL. <i>Blood</i> , <b>2009</b> , 113, 5536-48	2.2	179
610	Loss of BAF250a (ARID1A) is frequent in high-grade endometrial carcinomas. <i>Journal of Pathology</i> , <b>2011</b> , 224, 328-33	9.4	174
609	A multiprotein supercomplex controlling oncogenic signalling in lymphoma. <i>Nature</i> , <b>2018</b> , 560, 387-391	50.4	172
608	Favorable outcome of primary mediastinal large B-cell lymphoma in a single institution: the British Columbia experience. <i>Annals of Oncology</i> , <b>2006</b> , 17, 123-30	10.3	171
607	Activating mutations in genes related to TCR signaling in angioimmunoblastic and other follicular helper T-cell-derived lymphomas. <i>Blood</i> , <b>2016</b> , 128, 1490-502	2.2	170



606	Distinctive patterns of BCL6 molecular alterations and their functional consequences in different subgroups of diffuse large B-cell lymphoma. <i>Leukemia</i> , <b>2007</b> , 21, 2332-43	10.7	169
605	MALT1 is deregulated by both chromosomal translocation and amplification in B-cell non-Hodgkin lymphoma. <i>Blood</i> , <b>2003</b> , 101, 4539-46	2.2	167
604	HVCN1 modulates BCR signal strength via regulation of BCR-dependent generation of reactive oxygen species. <i>Nature Immunology</i> , <b>2010</b> , 11, 265-72	19.1	164
603	Impact of concordant and discordant bone marrow involvement on outcome in diffuse large B-cell lymphoma treated with R-CHOP. <i>Journal of Clinical Oncology</i> , <b>2011</b> , 29, 1452-7	2.2	163
602	Aberrant immunoglobulin class switch recombination and switch translocations in activated B cell-like diffuse large B cell lymphoma. <i>Journal of Experimental Medicine</i> , <b>2007</b> , 204, 633-43	16.6	160
601	CREBBP Inactivation Promotes the Development of HDAC3-Dependent Lymphomas. <i>Cancer Discovery</i> , <b>2017</b> , 7, 38-53	24.4	159
600	Tumor-associated macrophages predict inferior outcomes in classic Hodgkin lymphoma: a correlative study from the E2496 Intergroup trial. <i>Blood</i> , <b>2012</b> , 120, 3280-7	2.2	159
599	Small noncleaved, non-Burkitt's (Burkit-Like) lymphoma: cytogenetics predict outcome and reflect clinical presentation. <i>Journal of Clinical Oncology</i> , <b>1999</b> , 17, 1558-67	2.2	158
598	The molecular pathogenesis of primary mediastinal large B-cell lymphoma. <i>Blood</i> , <b>2011</b> , 118, 2659-69	2.2	155
597	MCL1 transgenic mice exhibit a high incidence of B-cell lymphoma manifested as a spectrum of histologic subtypes. <i>Blood</i> , <b>2001</b> , 97, 3902-9	2.2	154
596	Loss of the HVEM Tumor Suppressor in Lymphoma and Restoration by Modified CAR-T Cells. <i>Cell</i> , <b>2016</b> , 167, 405-418.e13	56.2	154
595	Prediction of survival in diffuse large B-cell lymphoma based on the expression of 2 genes reflecting tumor and microenvironment. <i>Blood</i> , <b>2011</b> , 118, 1350-8	2.2	152
594	Homozygous Deletions at Chromosome 9p21 Involving p16 and p15 Are Associated With Histologic Progression in Follicle Center Lymphoma. <i>Blood</i> , <b>1998</b> , 91, 4677-4685	2.2	151
593	EZH2 and BCL6 Cooperate to Assemble CBX8-BCOR Complex to Repress Bivalent Promoters, Mediate Germinal Center Formation and Lymphomagenesis. <i>Cancer Cell</i> , <b>2016</b> , 30, 197-213	24.3	150
592	Specific secondary genetic alterations in mantle cell lymphoma provide prognostic information independent of the gene expression-based proliferation signature. <i>Journal of Clinical Oncology</i> , <b>2007</b> , 25, 1216-22	2.2	148
591	Gene expression-based model using formalin-fixed paraffin-embedded biopsies predicts overall survival in advanced-stage classical Hodgkin lymphoma. <i>Journal of Clinical Oncology</i> , <b>2013</b> , 31, 692-700	2.2	145
590	Genome-wide DNA profiling of marginal zone lymphomas identifies subtype-specific lesions with an impact on the clinical outcome. <i>Blood</i> , <b>2011</b> , 117, 1595-604	2.2	145
589	The architectural pattern of FOXP3-positive T cells in follicular lymphoma is an independent predictor of survival and histologic transformation. <i>Blood</i> , <b>2010</b> , 115, 289-95	2.2	145

588	LMO2 protein expression predicts survival in patients with diffuse large B-cell lymphoma treated with anthracycline-based chemotherapy with and without rituximab. <i>Journal of Clinical Oncology</i> , <b>2008</b> , 26, 447-54	2.2	143
587	Recurrent somatic mutations of PTPN1 in primary mediastinal B cell lymphoma and Hodgkin lymphoma. <i>Nature Genetics</i> , <b>2014</b> , 46, 329-35	36.3	141
586	Mechanism-based epigenetic chemosensitization therapy of diffuse large B-cell lymphoma. <i>Cancer Discovery</i> , <b>2013</b> , 3, 1002-19	24.4	140
585	Incidence and risk factors for central nervous system relapse in patients with diffuse large B-cell lymphoma: the impact of the addition of rituximab to CHOP chemotherapy. <i>Annals of Oncology</i> , <b>2010</b> , 21, 1046-52	10.3	139
584	Prognostic significance of immunohistochemical biomarkers in diffuse large B-cell lymphoma: a study from the Lunenburg Lymphoma Biomarker Consortium. <i>Blood</i> , <b>2011</b> , 117, 7070-8	2.2	138
583	Structural profiles of TP53 gene mutations predict clinical outcome in diffuse large B-cell lymphoma: an international collaborative study. <i>Blood</i> , <b>2008</b> , 112, 3088-98	2.2	138
582	Double-Hit Gene Expression Signature Defines a Distinct Subgroup of Germinal Center B-Cell-Like Diffuse Large B-Cell Lymphoma. <i>Journal of Clinical Oncology</i> , <b>2019</b> , 37, 190-201	2.2	137
581	Molecular pathogenesis of mucosa-associated lymphoid tissue lymphoma. <i>Journal of Clinical Oncology</i> , <b>2005</b> , 23, 6370-8	2.2	137
580	Follicular lymphomas with and without translocation t(14;18) differ in gene expression profiles and genetic alterations. <i>Blood</i> , <b>2009</b> , 114, 826-34	2.2	136
579	A phase II study of bortezomib in mantle cell lymphoma: the National Cancer Institute of Canada Clinical Trials Group trial IND.150. <i>Annals of Oncology</i> , <b>2007</b> , 18, 116-121	10.3	136
578	Transformation to aggressive lymphoma in nodular lymphocyte-predominant Hodgkin's lymphoma. <i>Journal of Clinical Oncology</i> , <b>2010</b> , 28, 793-9	2.2	135
577	IDH2R172 mutations define a unique subgroup of patients with angioimmunoblastic T-cell lymphoma. <i>Blood</i> , <b>2015</b> , 126, 1741-52	2.2	133
576	Acquired TNFRSF14 mutations in follicular lymphoma are associated with worse prognosis. <i>Cancer Research</i> , <b>2010</b> , 70, 9166-74	10.1	133
575	A clinicopathological retrospective study of 131 patients with primary bone lymphoma: a population-based study of successively treated cohorts from the British Columbia Cancer Agency. <i>Annals of Oncology</i> , <b>2007</b> , 18, 129-135	10.3	133
574	A phase 1 study of obinutuzumab induction followed by 2 years of maintenance in patients with relapsed CD20-positive B-cell malignancies. <i>Blood</i> , <b>2012</b> , 119, 5118-25	2.2	132
573	Anaplastic lymphoma kinase-positive diffuse large B-cell lymphoma: a rare clinicopathologic entity with poor prognosis. <i>Journal of Clinical Oncology</i> , <b>2009</b> , 27, 4211-6	2.2	132
572	BCL2 predicts survival in germinal center B-cell-like diffuse large B-cell lymphoma treated with CHOP-like therapy and rituximab. <i>Clinical Cancer Research</i> , <b>2011</b> , 17, 7785-95	12.9	132
571	Rituximab extended schedule or re-treatment trial for low-tumor burden follicular lymphoma: eastern cooperative oncology group protocol e4402. <i>Journal of Clinical Oncology</i> , <b>2014</b> , 32, 3096-102	2.2	131



570	Helicobacter pylori and MALT lymphoma. <i>Gastroenterology</i> , <b>2005</b> , 128, 1579-605	13.3	131
569	TNFR-associated factor family protein expression in normal tissues and lymphoid malignancies. <i>Journal of Immunology</i> , <b>2000</b> , 165, 5084-96	5.3	129
568	International Prognostic Score in advanced-stage Hodgkin's lymphoma: altered utility in the modern era. <i>Journal of Clinical Oncology</i> , <b>2012</b> , 30, 3383-8	2.2	127
567	FOXO1 is a tumor suppressor in classical Hodgkin lymphoma. <i>Blood</i> , <b>2012</b> , 119, 3503-11	2.2	127
566	Analysis of secondary chromosomal alterations in 165 cases of follicular lymphoma with t(14;18). <i>Genes Chromosomes and Cancer</i> , <b>2001</b> , 30, 375-82	5	127
565	Expression of the FOXP1 transcription factor is strongly associated with inferior survival in patients with diffuse large B-cell lymphoma. <i>Clinical Cancer Research</i> , <b>2005</b> , 11, 1065-72	12.9	127
564	MYC and BCL2 protein expression predicts survival in patients with diffuse large B-cell lymphoma treated with rituximab. <i>British Journal of Haematology</i> , <b>2014</b> , 165, 382-91	4.5	126
563	Genomic profiling reveals different genetic aberrations in systemic ALK-positive and ALK-negative anaplastic large cell lymphomas. <i>British Journal of Haematology</i> , <b>2008</b> , 140, 516-26	4.5	125
562	DNA methylation signatures define molecular subtypes of diffuse large B-cell lymphoma. <i>Blood</i> , <b>2010</b> , 116, e81-9	2.2	124
561	Transformation of follicular lymphoma to diffuse large B-cell lymphoma proceeds by distinct oncogenic mechanisms. <i>British Journal of Haematology</i> , <b>2007</b> , 136, 286-93	4.5	124
560	Molecular and Genetic Characterization of MHC Deficiency Identifies EZH2 as Therapeutic Target for Enhancing Immune Recognition. <i>Cancer Discovery</i> , <b>2019</b> , 9, 546-563	24.4	123
559	Genome-wide copy number analysis of Hodgkin Reed-Sternberg cells identifies recurrent imbalances with correlations to treatment outcome. <i>Blood</i> , <b>2010</b> , 116, 418-27	2.2	123
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157	Array-CGH Identifies Both Common and Subtype-Specific Genomic Aberrations in Marginal Zone Lymphomas. <i>Blood</i> , <b>2008</b> , 112, 622-622	2.2	1

156	A Comparison of Gene Expression Pattern in Major Histocompatibility Class II-Low Diffuse Large B-Cell Lymphoma with Plasmablastic Lymphoma.. <i>Blood</i> , <b>2009</b> , 114, 1941-1941	2.2	1
155	The Percentage of Cytotoxic T-Cells in Mantle Cell Lymphoma (MCL) Biopsies Predicts Response to Rituximab.. <i>Blood</i> , <b>2009</b> , 114, 2923-2923	2.2	1
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147	Single Cell Mass Cytometry for Phenotypic Analysis of Diffuse Large B-Cell Lymphoma. <i>Blood</i> , <b>2014</b> , 124, 2976-2976	2.2	1
146	Accurate Diagnosis of Aggressive B Cell Non-Hodgkin Lymphomas Using Gene Expression Profiling of Formalin-Fixed, Paraffin-Embedded Tissues. <i>Blood</i> , <b>2014</b> , 124, 3016-3016	2.2	1
145	FAS Mutations Accelerate Lymphoma Growth and Induce Therapeutic Resistance. <i>Blood</i> , <b>2014</b> , 124, 3018-3018	2.2	1
144	Comprehensive MYC and BCL2 Genetic Profiling in De Novo Diffuse Large B-Cell Lymphoma Demonstrates Clinically Relevant Genetic Alterations According to Cell of Origin Subtype. <i>Blood</i> , <b>2015</b> , 126, 109-109	2.2	1
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142	A Novel Prognostic Model Based on Tumor Microenvironment Biology in Relapse Biopsies Predicts Post-Autologous Stem Cell Transplantation Outcomes in Classical Hodgkin Lymphoma. <i>Blood</i> , <b>2016</b> , 128, 1093-1093	2.2	1
141	DNA Copy Number Gains of TCF4 (E2-2) Are Associated with Poor Outcome in Diffuse Large B-Cell Lymphoma. <i>Blood</i> , <b>2016</b> , 128, 2686-2686	2.2	1
140	Molecular Subgroups of Peripheral T-Cell Lymphoma Evolve By Distinct Genetic Pathways. <i>Blood</i> , <b>2016</b> , 128, 4096-4096	2.2	1
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131	Coding and non-coding drivers of mantle cell lymphoma identified through exome and genome sequencing		1
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128	Targetable genetic alterations of TCF4 (E2-2) drive immunoglobulin expression in the activated B-cell subtype of diffuse large B-cell lymphoma		1
127	Gene Expression Profiling of Microdissected Hodgkin Reed Sternberg Cells: Molecular Subtypes and Treatment Outcome Correlations.. <i>Blood</i> , <b>2009</b> , 114, 268-268	2.2	1
126	TNFRSF14 Is Mutated in a Subset of Follicular Lymphoma and Correlated with Inferior Prognosis.. <i>Blood</i> , <b>2009</b> , 114, 1919-1919	2.2	1
125	Prediction of Survival in Diffuse Large B-Cell Lymphoma Based On the Expression of Two Genes: Integration of Tumor and Microenvironment Contributions.. <i>Blood</i> , <b>2009</b> , 114, 622-622	2.2	1
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