

Randy D Gascoyne

List of Publications by Year in descending order

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711
papers

92,576
citations

317

138
h-index

329

287
g-index

741
all docs

741
docs citations

741
times ranked

60808
citing authors

#	ARTICLE	IF	CITATIONS
1	Circos: An information aesthetic for comparative genomics. <i>Genome Research</i> , 2009, 19, 1639-1645.	5.5	9,003
2	Revised Response Criteria for Malignant Lymphoma. <i>Journal of Clinical Oncology</i> , 2007, 25, 579-586.	1.6	4,061
3	Confirmation of the molecular classification of diffuse large B-cell lymphoma by immunohistochemistry using a tissue microarray. <i>Blood</i> , 2004, 103, 275-282.	1.4	3,574
4	The Use of Molecular Profiling to Predict Survival after Chemotherapy for Diffuse Large-B-Cell Lymphoma. <i>New England Journal of Medicine</i> , 2002, 346, 1937-1947.	27.0	3,474
5	Stromal Gene Signatures in Large-B-Cell Lymphomas. <i>New England Journal of Medicine</i> , 2008, 359, 2313-2323.	27.0	1,564
6	Somatic mutations altering EZH2 (Tyr641) in follicular and diffuse large B-cell lymphomas of germinal-center origin. <i>Nature Genetics</i> , 2010, 42, 181-185.	21.4	1,504
7	Genetics and Pathogenesis of Diffuse Large B-Cell Lymphoma. <i>New England Journal of Medicine</i> , 2018, 378, 1396-1407.	27.0	1,443
8	Frequent mutation of histone-modifying genes in non-Hodgkin lymphoma. <i>Nature</i> , 2011, 476, 298-303.	27.8	1,428
9	Chronic active B-cell-receptor signalling in diffuse large B-cell lymphoma. <i>Nature</i> , 2010, 463, 88-92.	27.8	1,402
10	Prediction of Survival in Follicular Lymphoma Based on Molecular Features of Tumor-Infiltrating Immune Cells. <i>New England Journal of Medicine</i> , 2004, 351, 2159-2169.	27.0	1,293
11	Oncogenically active MYD88 mutations in human lymphoma. <i>Nature</i> , 2011, 470, 115-119.	27.8	1,292
12	Rituximab-CHOP Versus CHOP Alone or With Maintenance Rituximab in Older Patients With Diffuse Large B-Cell Lymphoma. <i>Journal of Clinical Oncology</i> , 2006, 24, 3121-3127.	1.6	1,203
13	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> , 2007, 109, 1857-1861.	1.4	1,193
14	Tumor-Associated Macrophages and Survival in Classic Hodgkin's Lymphoma. <i>New England Journal of Medicine</i> , 2010, 362, 875-885.	27.0	1,141
15	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> , 2003, 198, 851-862.	8.5	1,002
16	Anti-CD47 Antibody Synergizes with Rituximab to Promote Phagocytosis and Eradicate Non-Hodgkin Lymphoma. <i>Cell</i> , 2010, 142, 699-713.	28.9	894
17	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> , 2005, 23, 5027-5033.	1.6	874
18	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> , 2008, 105, 13520-13525.	7.1	868

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19	The proliferation gene expression signature is a quantitative integrator of oncogenic events that predicts survival in mantle cell lymphoma. <i>Cancer Cell</i> , 2003, 3, 185-197.	16.8	848
20	Molecular Diagnosis of Burkitt's Lymphoma. <i>New England Journal of Medicine</i> , 2006, 354, 2431-2442.	27.0	824
21	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> , 2012, 30, 3452-3459.	1.6	824
22	Oncogenic <i>CARD11</i> Mutations in Human Diffuse Large B Cell Lymphoma. <i>Science</i> , 2008, 319, 1676-1679.	12.6	784
23	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> , 2008, 111, 5496-5504.	1.4	784
24	Burkitt lymphoma pathogenesis and therapeutic targets from structural and functional genomics. <i>Nature</i> , 2012, 490, 116-120.	27.8	759
25	EZH2 Is Required for Germinal Center Formation and Somatic EZH2 Mutations Promote Lymphoid Transformation. <i>Cancer Cell</i> , 2013, 23, 677-692.	16.8	706
26	An enhanced International Prognostic Index (NCCN-IPI) for patients with diffuse large B-cell lymphoma treated in the rituximab era. <i>Blood</i> , 2014, 123, 837-842.	1.4	693
27	Non-Hodgkin lymphoma. <i>Lancet</i> , The, 2017, 390, 298-310.	13.7	615
28	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, 4021-4031.	1.4	596
29	A New Immunostain Algorithm Classifies Diffuse Large B-Cell Lymphoma into Molecular Subtypes with High Accuracy. <i>Clinical Cancer Research</i> , 2009, 15, 5494-5502.	7.0	577
30	MYC gene rearrangements are associated with a poor prognosis in diffuse large B-cell lymphoma patients treated with R-CHOP chemotherapy. <i>Blood</i> , 2009, 114, 3533-3537.	1.4	566
31	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> , 2006, 108, 3295-3301.	1.4	559
32	Somatic mutations at EZH2 Y641 act dominantly through a mechanism of selectively altered PRC2 catalytic activity, to increase H3K27 trimethylation. <i>Blood</i> , 2011, 117, 2451-2459.	1.4	556
33	MHC class II transactivator CIITA is a recurrent gene fusion partner in lymphoid cancers. <i>Nature</i> , 2011, 471, 377-381.	27.8	551
34	Lymphomas with concurrent BCL2 and MYC translocations: the critical factors associated with survival. <i>Blood</i> , 2009, 114, 2273-2279.	1.4	523
35	Determining cell-of-origin subtypes of diffuse large B-cell lymphoma using gene expression in formalin-fixed paraffin-embedded tissue. <i>Blood</i> , 2014, 123, 1214-1217.	1.4	518
36	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</i> , The, 2015, 16, 1111-1122.	10.7	483

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37	Prognostic Significance of Anaplastic Lymphoma Kinase (ALK) Protein Expression in Adults With Anaplastic Large Cell Lymphoma. <i>Blood</i> , 1999, 93, 3913-3921.	1.4	464
38	Prognostic Significance of Bcl-2 Protein Expression and Bcl-2 Gene Rearrangement in Diffuse Aggressive Non-Hodgkin's Lymphoma. <i>Blood</i> , 1997, 90, 244-251.	1.4	451
39	Mantle Cell Lymphoma: A Clinicopathologic Study of 80 Cases. <i>Blood</i> , 1997, 89, 2067-2078.	1.4	448
40	Diffuse large B-cell lymphoma: optimizing outcome in the context of clinical and biologic heterogeneity. <i>Blood</i> , 2015, 125, 22-32.	1.4	445
41	Gene expression signatures delineate biological and prognostic subgroups in peripheral T-cell lymphoma. <i>Blood</i> , 2014, 123, 2915-2923.	1.4	435
42	Analysis of multiple biomarkers shows that lymphoma-associated macrophage (LAM) content is an independent predictor of survival in follicular lymphoma (FL). <i>Blood</i> , 2005, 106, 2169-2174.	1.4	427
43	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> , 2011, 29, 200-207.	1.6	426
44	The tumour microenvironment in B cell lymphomas. <i>Nature Reviews Cancer</i> , 2014, 14, 517-534.	28.4	417
45	Peripheral T-cell lymphoma, not otherwise specified: a report of 340 cases from the International Peripheral T-cell Lymphoma Project. <i>Blood</i> , 2011, 117, 3402-3408.	1.4	376
46	The histone lysine methyltransferase KMT2D sustains a gene expression program that represses B cell lymphoma development. <i>Nature Medicine</i> , 2015, 21, 1199-1208.	30.7	359
47	Molecular Pathogenesis of Hodgkin's Lymphoma: Increasing Evidence of the Importance of the Microenvironment. <i>Journal of Clinical Oncology</i> , 2011, 29, 1812-1826.	1.6	350
48	Mutational and structural analysis of diffuse large B-cell lymphoma using whole-genome sequencing. <i>Blood</i> , 2013, 122, 1256-1265.	1.4	349
49	Diffuse large B-cell lymphoma subgroups have distinct genetic profiles that influence tumor biology and improve gene-expression-based survival prediction. <i>Blood</i> , 2005, 106, 3183-3190.	1.4	348
50	SOX11 expression is highly specific for mantle cell lymphoma and identifies the cyclin D1-negative subtype. <i>Haematologica</i> , 2009, 94, 1555-1562.	3.5	345
51	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> , 2013, 31, 1970-1976.	1.6	335
52	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> , 2015, 33, 2848-2856.	1.6	334
53	Population-Based Analysis of Incidence and Outcome of Transformed Non-Hodgkin's Lymphoma. <i>Journal of Clinical Oncology</i> , 2008, 26, 5165-5169.	1.6	333
54	Cyclin D1-negative mantle cell lymphoma: a clinicopathologic study based on gene expression profiling. <i>Blood</i> , 2005, 106, 4315-4321.	1.4	330

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55	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> , 2015, 33, 251-257.	1.6	319
56	Cooperative signaling through the signal transducer and activator of transcription 3 and nuclear factor- κ B pathways in subtypes of diffuse large B-cell lymphoma. <i>Blood</i> , 2008, 111, 3701-3713.	1.4	315
57	Whole transcriptome sequencing reveals recurrent NOTCH1 mutations in mantle cell lymphoma. <i>Blood</i> , 2012, 119, 1963-1971.	1.4	313
58	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> , 2016, 34, 3150-3156.	1.6	313
59	Enteropathy-associated T-cell lymphoma: clinical and histological findings from the International Peripheral T-Cell Lymphoma Project. <i>Blood</i> , 2011, 118, 148-155.	1.4	308
60	Pharmacological and genomic profiling identifies NF- κ B-targeted treatment strategies for mantle cell lymphoma. <i>Nature Medicine</i> , 2014, 20, 87-92.	30.7	303
61	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> , 2004, 103, 4251-4258.	1.4	296
62	ALK-positive diffuse large B-cell lymphoma is associated with Clathrin-ALK rearrangements: report of 6 cases. <i>Blood</i> , 2003, 102, 2568-2573.	1.4	281
63	Peripheral T-cell lymphoma. <i>Blood</i> , 2011, 117, 6756-6767.	1.4	278
64	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> , 2006, 24, 961-968.	1.6	277
65	A multiprotein supercomplex controlling oncogenic signalling in lymphoma. <i>Nature</i> , 2018, 560, 387-391.	27.8	276
66	EZH2 mutations are frequent and represent an early event in follicular lymphoma. <i>Blood</i> , 2013, 122, 3165-3168.	1.4	274
67	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> , 2007, 25, 805-812.	1.6	271
68	Etiologic Heterogeneity Among Non-Hodgkin Lymphoma Subtypes: The InterLymph Non-Hodgkin Lymphoma Subtypes Project. <i>Journal of the National Cancer Institute Monographs</i> , 2014, 2014, 130-144.	2.1	265
69	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> , 2009, 27, 1607-1614.	1.6	264
70	Pathogenesis of follicular lymphoma. <i>Journal of Clinical Investigation</i> , 2012, 122, 3424-3431.	8.2	264
71	Cooperative Epigenetic Modulation by Cancer Amplicon Genes. <i>Cancer Cell</i> , 2010, 18, 590-605.	16.8	263
72	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> , 2004, 165, 159-166.	3.8	262

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73	EZH2-mediated epigenetic silencing in germinal center B cells contributes to proliferation and lymphomagenesis. <i>Blood</i> , 2010, 116, 5247-5255.	1.4	262
74	Flavopiridol in Untreated or Relapsed Mantle-Cell Lymphoma: Results of a Phase II Study of the National Cancer Institute of Canada Clinical Trials Group. <i>Journal of Clinical Oncology</i> , 2003, 21, 1740-1745.	1.6	261
75	Genomic rearrangements involving programmed death ligands are recurrent in primary mediastinal large B-cell lymphoma. <i>Blood</i> , 2014, 123, 2062-2065.	1.4	259
76	Signal transducer and activator of transcription 6 is frequently activated in Hodgkin and Reed-Sternberg cells of Hodgkin lymphoma. <i>Blood</i> , 2002, 99, 618-626.	1.4	257
77	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> , 2019, 37, 190-201.	1.6	257
78	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> , 2013, 31, 684-691.	1.6	256
79	Activating mutations in genes related to TCR signaling in angioimmunoblastic and other follicular helper T-cell-derived lymphomas. <i>Blood</i> , 2016, 128, 1490-1502.	1.4	255
80	Loss of signalling via $\text{C}13$ in germinal centre B-cell-derived lymphoma. <i>Nature</i> , 2014, 516, 254-258.	27.8	253
81	Prognostic significance of Bcl-6 protein expression in DLBCL treated with CHOP or R-CHOP: a prospective correlative study. <i>Blood</i> , 2006, 107, 4207-4213.	1.4	248
82	Whole-Genome Analysis and HLA Genotyping of Enteropathy-Type T-Cell Lymphoma Reveals 2 Distinct Lymphoma Subtypes. <i>Gastroenterology</i> , 2007, 132, 1902-1911.	1.3	240
83	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> , 2016, 34, 2020-2027.	1.6	239
84	MALT1 Small Molecule Inhibitors Specifically Suppress ABC-DLBCL In Vitro and In Vivo. <i>Cancer Cell</i> , 2012, 22, 812-824.	16.8	229
85	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> , 2007, 109, 4599-4606.	1.4	226
86	Differentiation stage-specific expression of microRNAs in B lymphocytes and diffuse large B-cell lymphomas. <i>Blood</i> , 2009, 113, 3754-3764.	1.4	226
87	<i>CREBBP</i> Inactivation Promotes the Development of HDAC3-Dependent Lymphomas. <i>Cancer Discovery</i> , 2017, 7, 38-53.	9.4	218
88	Molecular and Genetic Characterization of MHC Deficiency Identifies EZH2 as Therapeutic Target for Enhancing Immune Recognition. <i>Cancer Discovery</i> , 2019, 9, 546-563.	9.4	213
89	Favorable outcome of primary mediastinal large B-cell lymphoma in a single institution: the British Columbia experience. <i>Annals of Oncology</i> , 2006, 17, 123-130.	1.2	212
90	Loss of BAF250a (<i>ARID1A</i>) is frequent in high-grade endometrial carcinomas. <i>Journal of Pathology</i> , 2011, 224, 328-333.	4.5	210

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91	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> , 1995, 103, 472-478.	0.7	207
92	The BCL6 transcriptional program features repression of multiple oncogenes in primary B cells and is deregulated in DLBCL. <i>Blood</i> , 2009, 113, 5536-5548.	1.4	205
93	Loss of the HVEM Tumor Suppressor in Lymphoma and Restoration by Modified CAR-T Cells. <i>Cell</i> , 2016, 167, 405-418.e13.	28.9	204
94	EZH2 and BCL6 Cooperate to Assemble CBX8-BCOR Complex to Repress Bivalent Promoters, Mediate Germinal Center Formation and Lymphomagenesis. <i>Cancer Cell</i> , 2016, 30, 197-213.	16.8	200
95	Distinctive patterns of BCL6 molecular alterations and their functional consequences in different subgroups of diffuse large B-cell lymphoma. <i>Leukemia</i> , 2007, 21, 2332-2343.	7.2	198
96	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> , 2011, 29, 1452-1457.	1.6	197
97	HVCN1 modulates BCR signal strength via regulation of BCR-dependent generation of reactive oxygen species. <i>Nature Immunology</i> , 2010, 11, 265-272.	14.5	196
98	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> , 2007, 18, 129-135.	1.2	190
99	The molecular pathogenesis of primary mediastinal large B-cell lymphoma. <i>Blood</i> , 2011, 118, 2659-2669.	1.4	189
100	MALT1 is deregulated by both chromosomal translocation and amplification in B-cell non-Hodgkin lymphoma. <i>Blood</i> , 2003, 101, 4539-4546.	1.4	188
101	Tumor-associated macrophages predict inferior outcomes in classic Hodgkin lymphoma: a correlative study from the E2496 Intergroup trial. <i>Blood</i> , 2012, 120, 3280-3287.	1.4	188
102	Histological Transformation and Progression in Follicular Lymphoma: A Clonal Evolution Study. <i>PLoS Medicine</i> , 2016, 13, e1002197.	8.4	185
103	<i>Helicobacter pylori</i> and MALT Lymphoma. <i>Gastroenterology</i> , 2005, 128, 1579-1605.	1.3	184
104	IDH2 R172 mutations define a unique subgroup of patients with angioimmunoblastic T-cell lymphoma. <i>Blood</i> , 2015, 126, 1741-1752.	1.4	184
105	Genetic drivers of oncogenic pathways in molecular subgroups of peripheral T-cell lymphoma. <i>Blood</i> , 2019, 133, 1664-1676.	1.4	184
106	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> , 2010, 21, 1046-1052.	1.2	182
107	Mechanism-Based Epigenetic Chemosensitization Therapy of Diffuse Large B-Cell Lymphoma. <i>Cancer Discovery</i> , 2013, 3, 1002-1019.	9.4	180
108	Recurrent somatic mutations of PTPN1 in primary mediastinal B cell lymphoma and Hodgkin lymphoma. <i>Nature Genetics</i> , 2014, 46, 329-335.	21.4	180

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109	Transformation to Aggressive Lymphoma in Nodular Lymphocyte-Predominant Hodgkin's Lymphoma. <i>Journal of Clinical Oncology</i> , 2010, 28, 793-799.	1.6	178
110	Follicular lymphomas with and without translocation t(14;18) differ in gene expression profiles and genetic alterations. <i>Blood</i> , 2009, 114, 826-834.	1.4	177
111	Clinicogenetic risk models predict early progression of follicular lymphoma after first-line immunochemotherapy. <i>Blood</i> , 2016, 128, 1112-1120.	1.4	177
112	Aberrant immunoglobulin class switch recombination and switch translocations in activated B cell-like diffuse large B cell lymphoma. <i>Journal of Experimental Medicine</i> , 2007, 204, 633-643.	8.5	176
113	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> , 2013, 31, 692-700.	1.6	176
114	Prediction of survival in diffuse large B-cell lymphoma based on the expression of 2 genes reflecting tumor and microenvironment. <i>Blood</i> , 2011, 118, 1350-1358.	1.4	175
115	Structural profiles of TP53 gene mutations predict clinical outcome in diffuse large B-cell lymphoma: an international collaborative study. <i>Blood</i> , 2008, 112, 3088-3098.	1.4	173
116	The architectural pattern of FOXP3-positive T cells in follicular lymphoma is an independent predictor of survival and histologic transformation. <i>Blood</i> , 2010, 115, 289-295.	1.4	173
117	Genome-wide DNA profiling of marginal zone lymphomas identifies subtype-specific lesions with an impact on the clinical outcome. <i>Blood</i> , 2011, 117, 1595-1604.	1.4	173
118	Molecular Pathogenesis of Mucosa-Associated Lymphoid Tissue Lymphoma. <i>Journal of Clinical Oncology</i> , 2005, 23, 6370-6378.	1.6	172
119	International Prognostic Score in Advanced-Stage Hodgkin's Lymphoma: Altered Utility in the Modern Era. <i>Journal of Clinical Oncology</i> , 2012, 30, 3383-3388.	1.6	171
120	Small Noncleaved, Non-Burkitt's (Burkitt-Like) Lymphoma: Cytogenetics Predict Outcome and Reflect Clinical Presentation. <i>Journal of Clinical Oncology</i> , 1999, 17, 1558-1558.	1.6	169
121	Prognostic significance of immunohistochemical biomarkers in diffuse large B-cell lymphoma: a study from the Lunenburg Lymphoma Biomarker Consortium. <i>Blood</i> , 2011, 117, 7070-7078.	1.4	168
122	High-grade B-cell lymphoma with MYC and BCL2 and/or BCL6 rearrangements with diffuse large B-cell lymphoma morphology. <i>Blood</i> , 2018, 131, 2060-2064.	1.4	167
123	MCL1 transgenic mice exhibit a high incidence of B-cell lymphoma manifested as a spectrum of histologic subtypes. <i>Blood</i> , 2001, 97, 3902-3909.	1.4	166
124	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> , 2007, 25, 1216-1222.	1.6	166
125	The Genetic Basis of Hepatosplenic T-cell Lymphoma. <i>Cancer Discovery</i> , 2017, 7, 369-379.	9.4	163
126	Prognostic Significance of MYC Rearrangement and Translocation Partner in Diffuse Large B-Cell Lymphoma: A Study by the Lunenburg Lymphoma Biomarker Consortium. <i>Journal of Clinical Oncology</i> , 2019, 37, 3359-3368.	1.6	161

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127	Acquired <i>TNFRSF14</i> Mutations in Follicular Lymphoma Are Associated with Worse Prognosis. <i>Cancer Research</i> , 2010, 70, 9166-9174.	0.9	160
128	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> , 2008, 26, 447-454.	1.6	159
129	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> , 2014, 32, 3096-3102.	1.6	159
130	Homozygous Deletions at Chromosome 9p21 Involving p16 and p15 Are Associated With Histologic Progression in Follicle Center Lymphoma. <i>Blood</i> , 1998, 91, 4677-4685.	1.4	158
131	<i>MYC</i> and <i>BCL2</i> protein expression predicts survival in patients with diffuse large B-cell lymphoma treated with rituximab. <i>British Journal of Haematology</i> , 2014, 165, 382-391.	2.5	157
132	Anaplastic Lymphoma Kinase-Positive Diffuse Large B-Cell Lymphoma: A Rare Clinicopathologic Entity With Poor Prognosis. <i>Journal of Clinical Oncology</i> , 2009, 27, 4211-4216.	1.6	154
133	Genome-wide copy number analysis of Hodgkin Reed-Sternberg cells identifies recurrent imbalances with correlations to treatment outcome. <i>Blood</i> , 2010, 116, 418-427.	1.4	152
134	<i>BCL2</i> 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> , 2011, 17, 7785-7795.	7.0	152
135	Analysis of Heritability and Shared Heritability Based on Genome-Wide Association Studies for Thirteen Cancer Types. <i>Journal of the National Cancer Institute</i> , 2015, 107, djv279.	6.3	152
136	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> , 2007, 18, 116-121.	1.2	151
137	<i>FOXO1</i> is a tumor suppressor in classical Hodgkin lymphoma. <i>Blood</i> , 2012, 119, 3503-3511.	1.4	149
138	Genomic profiling reveals different genetic aberrations in systemic ALK-positive and ALK-negative anaplastic large cell lymphomas. <i>British Journal of Haematology</i> , 2008, 140, 516-526.	2.5	145
139	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> , 2012, 119, 5118-5125.	1.4	145
140	Impact of dual expression of <i>MYC</i> and <i>BCL2</i> by immunohistochemistry on the risk of CNS relapse in DLBCL. <i>Blood</i> , 2016, 127, 2182-2188.	1.4	145
141	Enteropathy-associated T cell lymphoma subtypes are characterized by loss of function of <i>SETD2</i> . <i>Journal of Experimental Medicine</i> , 2017, 214, 1371-1386.	8.5	144
142	Survival of human lymphoma cells requires B-cell receptor engagement by self-antigens. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 13447-13454.	7.1	143
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