

# Randy D Gascoyne

## List of Publications by Year in descending order

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711  
papers

92,576  
citations

317

138  
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326

287  
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741  
all docs

741  
docs citations

741  
times ranked

60808  
citing authors

#	ARTICLE	IF	CITATIONS
1	Tumor-associated antigen PRAME exhibits dualistic functions that are targetable in diffuse large B cell lymphoma. <i>Journal of Clinical Investigation</i> , 2022, 132, .	3.9	12
2	Impact of MYC and BCL2 structural variants in tumors of DLBCL morphology and mechanisms of false-negative MYC IHC. <i>Blood</i> , 2021, 137, 2196-2208.	0.6	18
3	Genomic predictors of central nervous system relapse in primary testicular diffuse large B-cell lymphoma. <i>Blood</i> , 2021, 137, 1256-1259.	0.6	13
4	BCL2 Expression in First-Line Diffuse Large B-Cell Lymphoma Identifies a Patient Population With Poor Prognosis. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , 2021, 21, 267-278.e10.	0.2	8
5	Molecular attributes underlying central nervous system and systemic relapse in diffuse large B-cell lymphoma. <i>Haematologica</i> , 2021, 106, 1466-1471.	1.7	9
6	Long-term outcomes of R-CEOP show curative potential in patients with DLBCL and a contraindication to anthracyclines. <i>Blood Advances</i> , 2021, 5, 1483-1489.	2.5	17
7	Characterization of DLBCL with a PMBL gene expression signature. <i>Blood</i> , 2021, 138, 136-148.	0.6	19
8	ROBUST: A Phase III Study of Lenalidomide Plus R-CHOP Versus Placebo Plus R-CHOP in Previously Untreated Patients With ABC-Type Diffuse Large B-Cell Lymphoma. <i>Journal of Clinical Oncology</i> , 2021, 39, 1317-1328.	0.8	132
9	MAPK and JAK-STAT pathways dysregulation in plasmablastic lymphoma. <i>Haematologica</i> , 2021, 106, 2682-2693.	1.7	44
10	Outcome of limited-stage nodular lymphocyte-predominant Hodgkin lymphoma and the impact of a PET-adapted approach. <i>Blood Advances</i> , 2021, 5, 3647-3655.	2.5	4
11	Gene expression-based model predicts outcome in children with intermediate-risk classical Hodgkin lymphoma. <i>Blood</i> , 2021, , .	0.6	9
12	Variable global distribution of cell-of-origin from the ROBUST phase III study in diffuse large B-cell lymphoma. <i>Haematologica</i> , 2020, 105, e72-e75.	1.7	11
13	Single Cell Phenotypic Profiling of 27 DLBCL Cases Reveals Marked Intertumoral and Intratumoral Heterogeneity. <i>Cytometry Part A: the Journal of the International Society for Analytical Cytology</i> , 2020, 97, 620-629.	1.1	12
14	Expansion of PD1-positive T Cells in Nodal Marginal Zone Lymphoma. <i>American Journal of Surgical Pathology</i> , 2020, 44, 657-664.	2.1	21
15	A Three-Arm Randomized Phase II Study of Bendamustine/Rituximab with Bortezomib Induction or Lenalidomide Continuation in Untreated Follicular Lymphoma: ECOG-ACRIN E2408. <i>Clinical Cancer Research</i> , 2020, 26, 4468-4477.	3.2	16
16	TMEM30A loss-of-function mutations drive lymphomagenesis and confer therapeutically exploitable vulnerability in B-cell lymphoma. <i>Nature Medicine</i> , 2020, 26, 577-588.	15.2	46
17	Coding and noncoding drivers of mantle cell lymphoma identified through exome and genome sequencing. <i>Blood</i> , 2020, 136, 572-584.	0.6	44
18	The Tumor Associated Antigen PRAME Exhibits Dualistic Functions That Are Targetable in Diffuse Large B-Cell Lymphoma. <i>Blood</i> , 2020, 136, 34-34.	0.6	1

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19	Integrative genomic analysis identifies key pathogenic mechanisms in primary mediastinal large B-cell lymphoma. <i>Blood</i> , 2019, 134, 802-813.	0.6	96
20	Targetable genetic alterations of <i>TCF4</i> ( <i>E2-2</i> ) drive immunoglobulin expression in diffuse large B cell lymphoma. <i>Science Translational Medicine</i> , 2019, 11, .	5.8	51
21	Prognostic Significance of <i>MYC</i> 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.	0.8	161
22	The whole-genome landscape of Burkitt lymphoma subtypes. <i>Blood</i> , 2019, 134, 1598-1607.	0.6	113
23	Molecular and Genetic Characterization of MHC Deficiency Identifies EZH2 as Therapeutic Target for Enhancing Immune Recognition. <i>Cancer Discovery</i> , 2019, 9, 546-563.	7.7	213
24	Cell of origin in diffuse large B-cell lymphoma in systemic lupus erythematosus: molecular and clinical factors associated with survival. <i>Lupus Science and Medicine</i> , 2019, 6, e000324.	1.1	16
25	Novel insights into the genetics and epigenetics of MALT lymphoma unveiled by next generation sequencing analyses. <i>Haematologica</i> , 2019, 104, e558-e561.	1.7	55
26	Identification of high-risk <i>DUSP22</i> rearranged <i>ALK</i> -negative anaplastic large cell lymphoma. <i>British Journal of Haematology</i> , 2019, 186, e28-e31.	1.2	56
27	Follicular lymphoma patients with <i>KIR2DL2</i> and <i>KIR3DL1</i> and their ligands ( <i>HLA-C1</i> and <i>HLA-Bw4</i> ) show improved outcome when receiving rituximab. , 2019, 7, 70.		19
28	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.	0.8	257
29	Genetic drivers of oncogenic pathways in molecular subgroups of peripheral T-cell lymphoma. <i>Blood</i> , 2019, 133, 1664-1676.	0.6	184
30	Molecular features of a large cohort of primary central nervous system lymphoma using tissue microarray. <i>Blood Advances</i> , 2019, 3, 3953-3961.	2.5	22
31	<i>JUNB</i> , <i>DUSP2</i> , <i>SGK1</i> , <i>SOCS1</i> and <i>CREBBP</i> are frequently mutated in T-cell/histiocyte-rich large B-cell lymphoma. <i>Haematologica</i> , 2019, 104, 330-337.	1.7	45
32	Convergence of risk prediction models in follicular lymphoma. <i>Haematologica</i> , 2019, 104, e252-e255.	1.7	9
33	TP53 Expression Correlates with TP53 Mutations and Is an Independent Predictor of Clinical Outcome in Patients with DLBCL Treated with R-CHOP. <i>Blood</i> , 2019, 134, 3964-3964.	0.6	2
34	Potential Factors That Impact Lenalidomide/R-CHOP Efficacy in Previously Untreated Diffuse Large B-Cell Lymphoma in the ROBUST and ECOG-ACRIN 1412 Studies. <i>Blood</i> , 2019, 134, 4092-4092.	0.6	4
35	Longitudinal toxicity analysis with novel summary metrics of lenalidomide maintenance in follicular lymphoma in ECOG-ACRIN 2408.. <i>Journal of Clinical Oncology</i> , 2019, 37, 6511-6511.	0.8	0
36	Abstract 3765: Somatic JAK-STAT mutations in subtypes of aggressive B-cell lymphomas with DLBCL morphology. , 2019, , .		0

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37	Diffuse Large B-Cell Lymphomas with a Molecular PMBCL Expression Signature Represent a Distinct Molecular Subtype Associated with Poor Clinical Outcome. <i>Blood</i> , 2019, 134, 922-922.	0.6	1
38	Quality of Life Was Not Negatively Impacted By the Addition of Lenalidomide to R-CHOP Chemotherapy (R2-CHOP) Compared with Placebo Plus R-CHOP Chemotherapy in Patients with Previously Untreated Activated B-Cell (ABC)-Type Diffuse Large B-Cell Lymphoma (DLBCL): Health-Related Quality of Life (HRQoL) Analysis of the International Robust Study. <i>Blood</i> , 2019, 134, 3475-3475.	0.6	0
39	Molecular Correlates of Central Nervous System Relapse in Diffuse Large B-Cell Lymphoma. <i>Blood</i> , 2019, 134, 2763-2763.	0.6	0
40	Mutations Affecting RNA Binding Proteins Are a Novel Feature of Mantle Cell Lymphoma. <i>Blood</i> , 2019, 134, 1478-1478.	0.6	0
41	Abstract 3480: <i>TMEM30A</i> loss-of-function mutations drive lymphomagenesis and confer therapeutically exploitable vulnerability in B-cell lymphoma. , 2019, , .		0
42	Somatic IL4R mutations in primary mediastinal large B-cell lymphoma lead to constitutive JAK-STAT signaling activation. <i>Blood</i> , 2018, 131, 2036-2046.	0.6	39
43	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.	0.6	167
44	Assessment of Capture and Amplicon-Based Approaches for the Development of a Targeted Next-Generation Sequencing Pipeline to Personalize Lymphoma Management. <i>Journal of Molecular Diagnostics</i> , 2018, 20, 203-214.	1.2	58
45	Genetics and Pathogenesis of Diffuse Large B-Cell Lymphoma. <i>New England Journal of Medicine</i> , 2018, 378, 1396-1407.	13.9	1,443
46	AICDA drives epigenetic heterogeneity and accelerates germinal center-derived lymphomagenesis. <i>Nature Communications</i> , 2018, 9, 222.	5.8	51
47	Impact of age on genetics and treatment efficacy in follicular lymphoma. <i>Haematologica</i> , 2018, 103, e364-e367.	1.7	10
48	Rapid, real time pathology review for ECOG/ACRIN 1412: a novel and successful paradigm for future lymphoma clinical trials in the precision medicine era. <i>Blood Cancer Journal</i> , 2018, 8, 27.	2.8	10
49	Outcomes in adolescents and young adults with Hodgkin lymphoma treated on US cooperative group protocols: An adult intergroup (E2496) and Children's Oncology Group (COG AHOD0031) comparative analysis. <i>Cancer</i> , 2018, 124, 136-144.	2.0	47
50	FOXP1 expression is a prognostic biomarker in follicular lymphoma treated with rituximab and chemotherapy. <i>Blood</i> , 2018, 131, 226-235.	0.6	31
51	High-resolution architecture and partner genes of MYC rearrangements in lymphoma with DLBCL morphology. <i>Blood Advances</i> , 2018, 2, 2755-2765.	2.5	74
52	Molecular classification of primary mediastinal large B-cell lymphoma using routinely available tissue specimens. <i>Blood</i> , 2018, 132, 2401-2405.	0.6	64
53	Genome-wide discovery of somatic regulatory variants in diffuse large B-cell lymphoma. <i>Nature Communications</i> , 2018, 9, 4001.	5.8	102
54	A gene signature that distinguishes conventional and leukemic nonnodal mantle cell lymphoma helps predict outcome. <i>Blood</i> , 2018, 132, 413-422.	0.6	89

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55	Interim PET-directed therapy in limited-stage Hodgkin lymphoma initially treated with ABVD. <i>Haematologica</i> , 2018, 103, e590-e593.	1.7	16
56	Duodenal-type and nodal follicular lymphomas differ by their immune microenvironment rather than their mutation profiles. <i>Blood</i> , 2018, 132, 1695-1702.	0.6	49
57	A multiprotein supercomplex controlling oncogenic signalling in lymphoma. <i>Nature</i> , 2018, 560, 387-391.	13.7	276
58	Final Analysis of the Front-Line Phase III Randomized ACT-1 Trial in Younger Patients with Systemic Peripheral T-Cell Lymphoma Treated with CHOP Chemotherapy with or without Alemtuzumab and Consolidated By Autologous Hematopoietic Stem Cell Transplant. <i>Blood</i> , 2018, 132, 998-998.	0.6	19
59	Molecular and Genetic Characterization of MHC Deficiency Identifies EZH2 As a Therapeutic Target for Restoring MHC Expression in Diffuse Large B-Cell Lymphoma. <i>Blood</i> , 2018, 132, 1560-1560.	0.6	2
60	The Prognostic Impact of Baseline Positron Emission Tomography (PET) Imaging in Untreated High Risk (HR) Follicular Lymphoma (FL): Analysis from E2408, the Bortezomib Induction or Novel Imidazole Continuation (BIONIC) Study. <i>Blood</i> , 2018, 132, 1615-1615.	0.6	1
61	Molecular Features of Primary Central Nervous System Lymphoma in a Large Tissue Microarray. <i>Blood</i> , 2018, 132, 348-348.	0.6	1
62	The Double-Hit Gene Expression Signature Defines a Clinically and Biologically Distinct Subgroup within GCB-DLBCL. <i>Blood</i> , 2018, 132, 921-921.	0.6	1
63	Results of real-time cell-of-origin subtype identification by gene expression profiling in patients with ABC-type diffuse large B-cell lymphoma in the phase III trial of lenalidomide plus R-CHOP vs placebo plus R-CHOP (ROBUST).. <i>Journal of Clinical Oncology</i> , 2018, 36, 7548-7548.	0.8	1
64	Single-Cell Profiling Reveals Distinct Tumor Subtypes and Their Associated T-Cell Environments in Follicular Lymphoma. <i>Blood</i> , 2018, 132, 1577-1577.	0.6	0
65	UBR5 Mutations in Mantle Cell Lymphoma Lead to Increased Proliferation through a Cyclin D1-Dependent Mechanism. <i>Blood</i> , 2018, 132, 2849-2849.	0.6	0
66	A Longitudinal Toxicity over Time (ToxT) Analysis of Bortezomib When Added to Bendamustine-Rituximab (BR) in Previously Untreated High Risk (HR) Follicular Lymphoma (FL) from in E2408. <i>Blood</i> , 2018, 132, 4157-4157.	0.6	0
67	Recurrent IL4R Somatic Mutations in Diffuse Large B-Cell Lymphoma Lead to an Altered Gene Expression Profile and Changes in Tumor Microenvironment Composition. <i>Blood</i> , 2018, 132, 669-669.	0.6	1
68	Gemcitabine, dexamethasone, and cisplatin (GDP) is an effective and well-tolerated salvage therapy for relapsed/refractory diffuse large B-cell lymphoma and Hodgkin lymphoma. <i>Leukemia and Lymphoma</i> , 2017, 58, 324-332.	0.6	32
69	Combined copy number and mutation analysis identifies oncogenic pathways associated with transformation of follicular lymphoma. <i>Leukemia</i> , 2017, 31, 83-91.	3.3	87
70	Non-Hodgkin lymphoma. <i>Lancet, The</i> , 2017, 390, 298-310.	6.3	615
71	The Genetic Basis of Hepatosplenic T-cell Lymphoma. <i>Cancer Discovery</i> , 2017, 7, 369-379.	7.7	163
72	Genetic polymorphism at BCL2 as a predictor for rituximab, cyclophosphamide, doxorubicin, vincristine and prednisone efficacy in patients with diffuse large B-cell lymphoma. <i>Haematologica</i> , 2017, 102, e199-e202.	1.7	4

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73	Can histologic transformation of follicular lymphoma be predicted and prevented?. Blood, 2017, 130, 258-266.	0.6	52
74	Prognostic relevance of CD163 and CD8 combined with EZH2 and gain of chromosome 18 in follicular lymphoma: a study by the Lunenburg Lymphoma Biomarker Consortium. Haematologica, 2017, 102, 1413-1423.	1.7	39
75	Enteropathy-associated T cell lymphoma subtypes are characterized by loss of function of SETD2. Journal of Experimental Medicine, 2017, 214, 1371-1386.	4.2	144
76	Mapping the human T cell repertoire to recurrent driver mutations in MYD88 and EZH2 in lymphoma. Oncoimmunology, 2017, 6, e1321184.	2.1	23
77	Physical activity, obesity and survival in diffuse large B-cell and follicular lymphoma cases. British Journal of Haematology, 2017, 178, 442-447.	1.2	21
78	Follicular lymphoma: State-of-the-art ICML workshop in Lugano 2015. Hematological Oncology, 2017, 35, 397-407.	0.8	11
79	A Phase 2/3 Multicenter, Randomized, Open-Label Study to Compare the Efficacy and Safety of Lenalidomide Versus Investigator's Choice in Patients with Relapsed or Refractory Diffuse Large B-Cell Lymphoma. Clinical Cancer Research, 2017, 23, 4127-4137.	3.2	135
80	Genetic profiling of MYC and BCL2 in diffuse large B-cell lymphoma determines cell-of-origin-specific clinical impact. Blood, 2017, 129, 2760-2770.	0.6	112
81	Adult high-grade B-cell lymphoma with Burkitt lymphoma signature: genomic features and potential therapeutic targets. Blood, 2017, 130, 1819-1831.	0.6	62
82	Reliable subtype classification of diffuse large B-cell lymphoma samples from GELA LNH2003 trials using the Lymph2Cx gene expression assay. Haematologica, 2017, 102, e404-e406.	1.7	16
83	Observation as the initial management strategy in patients with mantle cell lymphoma. Annals of Oncology, 2017, 28, 2489-2495.	0.6	67
84	Diffuse large B-cell lymphoma with testicular involvement: outcome and risk of CNS relapse in the rituximab era. British Journal of Haematology, 2017, 176, 210-221.	1.2	78
85	CREBBP Inactivation Promotes the Development of HDAC3-Dependent Lymphomas. Cancer Discovery, 2017, 7, 38-53.	7.7	218
86	Outcome of primary cutaneous anaplastic large cell lymphoma: a 20-year British Columbia Cancer Agency experience. British Journal of Haematology, 2017, 176, 234-240.	1.2	20
87	Mantle cell lymphoma initial therapy with abbreviated R-CHOP followed by 90Y-ibritumomab tiuxetan: 10-year follow-up of the phase 2 ECOG-ACRIN study E1499. Leukemia, 2017, 31, 517-519.	3.3	14
88	Site of central nervous system (CNS) relapse in patients with diffuse large B-cell lymphoma (DLBCL) by the CNS-IPI risk model. British Journal of Haematology, 2017, 179, 508-510.	1.2	26
89	Aberrant cytoplasmic expression of MHCII confers worse progression free survival in diffuse large B-cell lymphoma. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2017, 470, 113-117.	1.4	5
90	Clinicopathologic consensus study of gray zone lymphoma with features intermediate between DLBCL and classical HL. Blood Advances, 2017, 1, 2600-2609.	2.5	62

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91	New Molecular Assay for the Proliferation Signature in Mantle Cell Lymphoma Applicable to Formalin-Fixed Paraffin-Embedded Biopsies. <i>Journal of Clinical Oncology</i> , 2017, 35, 1668-1677.	0.8	102
92	Prognostic Model to Predict Post-Autologous Stem-Cell Transplantation Outcomes in Classical Hodgkin Lymphoma. <i>Journal of Clinical Oncology</i> , 2017, 35, 3722-3733.	0.8	48
93	Rapid, real-time central pathology review for E1412: A novel and successful paradigm for future National Clinical Trials Network diffuse large B cell lymphoma studies.. <i>Journal of Clinical Oncology</i> , 2017, 35, 7547-7547.	0.8	1
94	Abstract 2445: Integrative genetic analysis identifies therapeutic relevance of cell of origin-specific genetic alterations in diffuse large B-cell lymphoma. , 2017, , .		0
95	9. The biology of primary mediastinal large B-cell lymphoma. , 2016, , 171-192.		0
96	ROBUST: Lenalidomide-R-CHOP versus placebo-R-CHOP in previously untreated ABC-type diffuse large B-cell lymphoma. <i>Future Oncology</i> , 2016, 12, 1553-1563.	1.1	85
97	Personalized risk prediction for event-free survival at 24 months in patients with diffuse large B-cell lymphoma. <i>American Journal of Hematology</i> , 2016, 91, 179-184.	2.0	41
98	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.	0.8	313
99	Evaluation of the Risk of Relapse in Classical Hodgkin Lymphoma at Event-Free Survival Time Points and Survival Comparison With the General Population in British Columbia. <i>Journal of Clinical Oncology</i> , 2016, 34, 2493-2500.	0.8	56
100	Rituximab extended schedule or retreatment trial for low tumour burden non-follicular indolent B-cell non-Hodgkin lymphomas: Eastern Cooperative Oncology Group Protocol E4402. <i>British Journal of Haematology</i> , 2016, 173, 867-875.	1.2	36
101	General Biomarker Recommendations for Lymphoma. <i>Journal of the National Cancer Institute</i> , 2016, 108, djw250.	3.0	2
102	Molecular etiology of an indolent lymphoproliferative disorder determined by whole-genome sequencing. <i>Journal of Physical Education and Sports Management</i> , 2016, 2, a000679.	0.5	3
103	Targeting Non-proteolytic Protein Ubiquitination for the Treatment of Diffuse Large B Cell Lymphoma. <i>Cancer Cell</i> , 2016, 29, 494-507.	7.7	93
104	Role of the tumor microenvironment in mature B-cell lymphoid malignancies. <i>Haematologica</i> , 2016, 101, 531-540.	1.7	75
105	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.	0.8	239
106	Impact of dual expression of MYC and BCL2 by immunohistochemistry on the risk of CNS relapse in DLBCL. <i>Blood</i> , 2016, 127, 2182-2188.	0.6	145
107	Comprehensive characterization of programmed death ligand structural rearrangements in B-cell non-Hodgkin lymphomas. <i>Blood</i> , 2016, 128, 1206-1213.	0.6	47
108	Clinicogenetic risk models predict early progression of follicular lymphoma after first-line immunochemotherapy. <i>Blood</i> , 2016, 128, 1112-1120.	0.6	177

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109	Loss of the HVEM Tumor Suppressor in Lymphoma and Restoration by Modified CAR-T Cells. <i>Cell</i> , 2016, 167, 405-418.e13.	13.5	204
110	Maintenance rituximab following induction R-CHOP chemotherapy in patients with composite or discordant, indolent and aggressive, B-cell non-Hodgkin lymphomas. <i>Haematologica</i> , 2016, 101, e411-e414.	1.7	11
111	The combined role of biomarkers and interim PET scan in prediction of treatment outcome in classical Hodgkin's lymphoma: a retrospective, European, multicentre cohort study. <i>Lancet Haematology</i> , 2016, 3, e467-e479.	2.2	63
112	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.	7.7	200
113	Impaired functional responses in follicular lymphoma CD8 <sup>+</sup> TIM-3 <sup>+</sup> T lymphocytes following TCR engagement. <i>OncImmunology</i> , 2016, 5, e1224044.	2.1	32
114	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.	0.6	255
115	Diffuse large B-cell lymphoma patient-derived xenograft models capture the molecular and biological heterogeneity of the disease. <i>Blood</i> , 2016, 127, 2203-2213.	0.6	68
116	Randomized phase 3 study in low-grade lymphoma comparing maintenance anti-CD20 antibody with observation after induction therapy: A trial of the ECOG-ACRIN Cancer Research Group (E1496). <i>Cancer</i> , 2016, 122, 2996-3004.	2.0	31
117	Toward Personalized Lymphoma Immunotherapy: Identification of Common Driver Mutations Recognized by Patient CD8+ T Cells. <i>Clinical Cancer Research</i> , 2016, 22, 2226-2236.	3.2	26
118	Impact of time from diagnosis to initiation of curative-intent chemotherapy on clinical outcomes in patients with classical Hodgkin lymphoma. <i>Leukemia and Lymphoma</i> , 2016, 57, 872-879.	0.6	10
119	The value of routine bone marrow biopsy in patients with diffuse large B-cell lymphoma staged with PET/CT: a Danish-Canadian study. <i>Annals of Oncology</i> , 2016, 27, 1095-1099.	0.6	43
120	Fc Gamma Receptor 3A and 2A Polymorphisms Do Not Predict Response to Rituximab in Follicular Lymphoma. <i>Clinical Cancer Research</i> , 2016, 22, 821-826.	3.2	26
121	Diffuse large B-cell lymphoma cell-of-origin classification using the Lymph2Cx assay in the context of BCL2 and MYC expression status. <i>Leukemia and Lymphoma</i> , 2016, 57, 717-720.	0.6	13
122	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> , 2016, 128, 1093-1093.	0.6	12
123	DNA Copy Number Gains of TCF4 (E2-2) Are Associated with Poor Outcome in Diffuse Large B-Cell Lymphoma. <i>Blood</i> , 2016, 128, 2686-2686.	0.6	1
124	Serum Biomarkers Predict Outcomes in Advanced Hodgkin Lymphoma Independent of International Prognostic Score (IPS) and Treatment: Correlative Analysis from a Large North American Cooperative Group Trial. <i>Blood</i> , 2016, 128, 2992-2992.	0.6	5
125	Molecular Subgroups of Peripheral T-Cell Lymphoma Evolve By Distinct Genetic Pathways. <i>Blood</i> , 2016, 128, 4096-4096.	0.6	1
126	EFS24 as a predictor of outcome in a population-based cohort of patients with DLBCL in British Columbia (BC).. <i>Journal of Clinical Oncology</i> , 2016, 34, 7569-7569.	0.8	2



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127	Histological Transformation and Progression in Follicular Lymphoma: A Clonal Evolution Study. PLoS Medicine, 2016, 13, e1002197.	3.9	185
128	Abstract IA18: Functional characterization of the tumor suppressor lysine-specific methyltransferase KMT2D in lymphoma. , 2016, , .		0
129	Classification of diffuse large b-cell lymphoma (DLBCL) FFPE samples of the GELA LNH2003 program, using Lymph2Cx assay on the nCounter analysis system.. Journal of Clinical Oncology, 2016, 34, 7547-7547.	0.8	0
130	Gray Zone Lymphoma (GZL) with Features Intermediate Between Diffuse Large B-Cell Lymphoma (DLBCL) and Classical Hodgkin Lymphoma (cHL): Pathologic Classification and Clinical Outcomes from a Multicenter Consensus Study. Blood, 2016, 128, 4145-4145.	0.6	0
131	Targeted Sequencing Reveals Novel Gene Mutations Associated with Transformation and Early Progression in Follicular Lymphoma. Blood, 2016, 128, 2919-2919.	0.6	5
132	Crebbp Mutations Disrupt Dynamic Enhancer Acetylation in B-Cells, Enabling HDAC3 to Drive Lymphomagenesis. Blood, 2016, 128, 735-735.	0.6	0
133	Frequent Genetic Alterations of PI3K-AKT Pathway and Their Clinical Significance in Germinal Center B-Cell-like Diffuse Large B-Cell Lymphoma. Blood, 2016, 128, 607-607.	0.6	1
134	Revealing the Tumor Ecosystem in Follicular Lymphoma By Mass Cytometry. Blood, 2016, 128, 2939-2939.	0.6	0
135	The Role of UBR5 Mutations in the Pathogenesis of Mantle CELL Lymphoma. Blood, 2016, 128, 4124-4124.	0.6	0
136	Divergent Modes of Tumor Evolution Underlie Histological Transformation and Early Progression of Follicular Lymphoma. Blood, 2016, 128, 1091-1091.	0.6	0
137	Characterization of Genomic Rearrangements Involving CIITA and SOCS1 Using Targeted Capture Sequencing of Archival Tissue Specimens. Blood, 2016, 128, 2925-2925.	0.6	0
138	Comprehensive Genomic Analysis of Adult Burkitt Lymphoma Identifies the B-Cell Receptor Signaling Pathway As a Potential Therapeutic Target. Blood, 2016, 128, 4095-4095.	0.6	0
139	Genomic Alterations in CIITA Are Frequent in Primary Mediastinal Large B Cell Lymphoma and Are Associated with Diminished MHC Class II Expression. Cell Reports, 2015, 13, 1418-1431.	2.9	112
140	An RCOR1 loss-associated gene expression signature identifies a prognostically significant DLBCL subgroup. Blood, 2015, 125, 959-966.	0.6	24
141	IDH2 R172 mutations define a unique subgroup of patients with angioimmunoblastic T-cell lymphoma. Blood, 2015, 126, 1741-1752.	0.6	184
142	Global microRNA expression profiling uncovers molecular markers for classification and prognosis in aggressive B-cell lymphoma. Blood, 2015, 125, 1137-1145.	0.6	110
143	Genetic inactivation of TRAF3 in canine and human B-cell lymphoma. Blood, 2015, 125, 999-1005.	0.6	67
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413	Prognostic Factors in Follicular Lymphoma. <i>Journal of Clinical Oncology</i> , 2010, 28, 2902-2913.	0.8	136
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419	A Randomized Phase III Trial of ABVD Vs. Stanford V +/â” Radiation Therapy In Locally Extensive and Advanced Stage Hodgkin's Lymphoma: An Intergroup Study Coordinated by the Eastern Cooperative Oncology Group (E2496). <i>Blood</i> , 2010, 116, 415-415.	0.6	18
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