Andreas Rosenwald

List of Publications by Year
in descending order

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270 papers	10,627 citations	50276 46 h-index	42399 92 g-index
275 all docs	275 docs citations	275 times ranked	12204 citing authors

#	Article	IF	Citations
1	Molecular subtypes of diffuse large B cell lymphoma are associated with distinct pathogenic mechanisms and outcomes. Nature Medicine, 2018, 24, 679-690.	30.7	1,224
2	The 5th edition of the World Health Organization Classification of Haematolymphoid Tumours: Lymphoid Neoplasms. Leukemia, 2022, 36, 1720-1748.	7.2	1,023
3	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. Lancet Oncology, The, 2015, 16, 1111-1122.	10.7	483
4	Convergent Mutations and Kinase Fusions Lead to Oncogenic STAT3 Activation in Anaplastic Large Cell Lymphoma. Cancer Cell, 2015, 27, 516-532.	16.8	378
5	Safety and activity of ibrutinib plus rituximab for patients with high-risk chronic lymphocytic leukaemia: a single-arm, phase 2 study. Lancet Oncology, The, 2014, 15, 1090-1099.	10.7	315
6	A multiprotein supercomplex controlling oncogenic signalling in lymphoma. Nature, 2018, 560, 387-391.	27.8	276
7	Loss of signalling via GÎ±13 in germinal centre B-cell-derived lymphoma. Nature, 2014, 516, 254-258.	27.8	253
8	IDH2 R172 mutations define a unique subgroup of patients with angioimmunoblastic T-cell lymphoma. Blood, 2015, 126, 1741-1752.	1.4	184
9	Genetic drivers of oncogenic pathways in molecular subgroups of peripheral T-cell lymphoma. Blood, 2019, 133, 1664-1676.	1.4	184
10	Clinicogenetic risk models predict early progression of follicular lymphoma after first-line immunochemotherapy. Blood, 2016, 128, 1112-1120.	1.4	177
11	High-grade B-cell lymphoma with MYC and BCL2 and/or BCL6 rearrangements with diffuse large B-cell lymphoma morphology. Blood, 2018, 131, 2060-2064.	1.4	167
12	Four versus six cycles of CHOP chemotherapy in combination with six applications of rituximab in patients with aggressive B-cell lymphoma with favourable prognosis (FLYER): a randomised, phase 3, non-inferiority trial. Lancet, The, 2019, 394, 2271-2281.	13.7	155
13	Mitotane Inhibits Sterol-O-Acyl Transferase 1 Triggering Lipid-Mediated Endoplasmic Reticulum Stress and Apoptosis in Adrenocortical Carcinoma Cells. Endocrinology, 2015, 156, 3895-3908.	2.8	153
14	Positron Emission Tomographyâ€“Guided Treatment in Early-Stage Favorable Hodgkin Lymphoma: Final Results of the International, Randomized Phase III HD16 Trial by the German Hodgkin Study Group. Journal of Clinical Oncology, 2019, 37, 2835-2845.	1.6	151
15	Survival of human lymphoma cells requires B-cell receptor engagement by self-antigens. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 13447-13454.	7.1	143
16	Homozygous BCMA gene deletion in response to anti-BCMA CAR T cells in a patient with multiple myeloma. Nature Medicine, 2021, 27, 616-619.	30.7	140
17	A MYC-Driven Change in Mitochondrial Dynamics Limits YAP/TAZ Function in Mammary Epithelial Cells and Breast Cancer. Cancer Cell, 2015, 28, 743-757.	16.8	122
18	Genome-wide analysis of pediatric-type follicular lymphoma reveals low genetic complexity and recurrent alterations of TNFRSF14 gene. Blood, 2016, 128, 1101-1111.	1.4	115

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19	Efficacy of Nivolumab and AVD in Early-Stage Unfavorable Classic Hodgkin Lymphoma. JAMA Oncology, 2020, 6, 872.	7.1	112
20	Global microRNA expression profiling uncovers molecular markers for classification and prognosis in aggressive B-cell lymphoma. Blood, 2015, 125, 1137-1145.	1.4	110
21	Expression of TP53 is associated with the outcome of MCL independent of MIPI and Ki-67 in trials of the European MCL Network. Blood, 2018, 131, 417-420.	1.4	108
22	Genomic and transcriptomic changes complement each other in the pathogenesis of sporadic Burkitt lymphoma. Nature Communications, 2019, 10, 1459.	12.8	99
23	Odronextamab, a human CD20Ã—CD3 bispecific antibody in patients with CD20-positive B-cell malignancies (ELM-1): results from the relapsed or refractory non-Hodgkin lymphoma cohort in a single-arm, multicentre, phase 1 trial. Lancet Haematology,the, 2022, 9, e327-e339.	4.6	98
24	Targeting Non-proteolytic Protein Ubiquitination for the Treatment of Diffuse Large B Cell Lymphoma. Cancer Cell, 2016, 29, 494-507.	16.8	93
25	Spectrum and functional validation of PSMB5 mutations in multiple myeloma. Leukemia, 2019, 33, 447-456.	7.2	93
26	PET-guided omission of radiotherapy in early-stage unfavourable Hodgkin lymphoma (GHSG HD17): a multicentre, open-label, randomised, phase 3 trial. Lancet Oncology, The, 2021, 22, 223-234.	10.7	93
27	Targeted Molecular Analysis in Adrenocortical Carcinomas: A Strategy Toward Improved Personalized Prognostication. Journal of Clinical Endocrinology and Metabolism, 2018, 103, 4511-4523.	3.6	92
28	MINCR is a MYC-induced lncRNA able to modulate MYCâ€™s transcriptional network in Burkitt lymphoma cells. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, E5261-70.	7.1	91
29	A gene signature that distinguishes conventional and leukemic nonnodal mantle cell lymphoma helps predict outcome. Blood, 2018, 132, 413-422.	1.4	89
30	Frequent NFKBIE deletions are associated with poor outcome in primary mediastinal B-cell lymphoma. Blood, 2016, 128, 2666-2670.	1.4	82
31	Long-term outcomes from the Phase II L-MIND study of tafasitamab (MOR208) plus lenalidomide in patients with relapsed or refractory diffuse large B-cell lymphoma. Haematologica, 2021, 106, 2417-2426.	3.5	81
32	Essential role of IRF4 and MYC signaling for survival of anaplastic large cell lymphoma. Blood, 2015, 125, 124-132.	1.4	79
33	Gemcitabine-Based Chemotherapy in Adrenocortical Carcinoma: A Multicenter Study of Efficacy and Predictive Factors. Journal of Clinical Endocrinology and Metabolism, 2017, 102, 4323-4332.	3.6	79
34	Recurrent intragenic rearrangements of EGFR and BRAF in soft tissue tumors of infants. Nature Communications, 2018, 9, 2378.	12.8	72
35	Mutations of MAP2K1 are frequent in pediatric-type follicular lymphoma and result in ERK pathway activation. Blood, 2017, 130, 323-327.	1.4	69
36	TP53 mutation and survival in aggressive B cell lymphoma. International Journal of Cancer, 2017, 141, 1381-1388.	5.1	69

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37	A 3-cM commonly deleted region in 6q21 in leukemias and lymphomas delineated by fluorescence in situ hybridization. Genes Chromosomes and Cancer, 2000, 27, 52-58.	2.8	67
38	GRP78-directed immunotherapy in relapsed or refractory multiple myeloma - results from a phase 1 trial with the monoclonal immunoglobulin M antibody PAT-SM6. Haematologica, 2015, 100, 377-384.	3.5	64
39	Molecular classification of primary mediastinal large B-cell lymphoma using routinely available tissue specimens. Blood, 2018, 132, 2401-2405.	1.4	64
40	Adult high-grade B-cell lymphoma with Burkitt lymphoma signature: genomic features and potential therapeutic targets. Blood, 2017, 130, 1819-1831.	1.4	62
41	Histiocytic and dendritic cell neoplasms: what have we learnt by studying 67 cases. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2017, 471, 467-489.	2.8	59
42	TRPS1 shapes YAP/TEAD-dependent transcription in breast cancer cells. Nature Communications, 2018, 9, 3115.	12.8	58
43	B-cell receptorâ€“driven MALT1 activity regulates MYC signaling in mantle cell lymphoma. Blood, 2017, 129, 333-346.	1.4	57
44	Active Akt signaling triggers CLL toward Richter transformation via overactivation of Notch1. Blood, 2021, 137, 646-660.	1.4	55
45	Alemtuzumab plus CHOP versus CHOP in elderly patients with peripheral T-cell lymphoma: the DSHNHL2006-1B/ACT-2 trial. Leukemia, 2021, 35, 143-155.	7.2	52
46	Targetable genetic alterations of TCF4 (E2-2) drive immunoglobulin expression in diffuse large B cell lymphoma. Science Translational Medicine, 2019, 11, .	12.4	51
47	USP9X stabilizes XIAP to regulate mitotic cell death and chemoresistance in aggressive Bâ€cell lymphoma. EMBO Molecular Medicine, 2016, 8, 851-862.	6.9	50
48	Selective NFAT targeting in T cells ameliorates GvHD while maintaining antitumor activity. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 1125-1130.	7.1	49
49	Duodenal-type and nodal follicular lymphomas differ by their immune microenvironment rather than their mutation profiles. Blood, 2018, 132, 1695-1702.	1.4	49
50	Inhibition of focal adhesion kinase overcomes resistance of mantle cell lymphoma to ibrutinib in the bone marrow microenvironment. Haematologica, 2018, 103, 116-125.	3.5	48
51	Tumor and microenvironment response but no cytotoxic T-cell activation in classic Hodgkin lymphoma treated with anti-PD1. Blood, 2020, 136, 2851-2863.	1.4	47
52	Halting the vicious cycle within the multiple myeloma ecosystem: blocking JAM-A on bone marrow endothelial cells restores angiogenic homeostasis and suppresses tumor progression. Haematologica, 2021, 106, 1943-1956.	3.5	46
53	The Myb-MuvB Complex Is Required for YAP-Dependent Transcription of Mitotic Genes. Cell Reports, 2019, 27, 3533-3546.e7.	6.4	45
54	MAPK and JAK-STAT pathways dysregulation in plasmablastic lymphoma. Haematologica, 2021, 106, 2682-2693.	3.5	44

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55	Alterations of microRNA and microRNA-regulated messenger RNA expression in germinal center B-cell lymphomas determined by integrative sequencing analysis. Haematologica, 2016, 101, 1380-1389.	3.5	43
56	Clinical impact of recurrently mutated genes on lymphoma diagnostics: state-of-the-art and beyond. Haematologica, 2016, 101, 1002-1009.	3.5	43
57	A GRP78-Directed Monoclonal Antibody Recaptures Response in Refractory Multiple Myeloma with Extramedullary Involvement. Clinical Cancer Research, 2016, 22, 4341-4349.	7.0	43
58	Impact of miR-21, miR-126 and miR-221 as Prognostic Factors of Clear Cell Renal Cell Carcinoma with Tumor Thrombus of the Inferior Vena Cava. PLoS ONE, 2014, 9, e109877.	2.5	42
59	Panel Sequencing Shows Recurrent Genetic FAS Alterations in Primary Cutaneous Marginal Zone Lymphoma. Journal of Investigative Dermatology, 2018, 138, 1573-1581.	0.7	41
60	IKZF1/3 and CRL4^{CRBN} E3 ubiquitin ligase mutations and resistance to immunomodulatory drugs in multiple myeloma. Haematologica, 2020, 105, e237-e241.	3.5	41
61	Chromosomal abnormalities in nodal and extranodal CD30+ anaplastic large cell lymphomas: Infrequent detection of the t(2;5) in extranodal lymphomas. , 1998, 22, 114-121.		40
62	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.	3.5	39
63	A randomized phase 3 trial of auto vs. allo transplantation as part of first-line therapy in poor-risk peripheral T-NHL. Blood, 2021, 137, 2646-2656.	1.4	39
64	Lymphoid Aggregates in the CNS of Progressive Multiple Sclerosis Patients Lack Regulatory T Cells. Frontiers in Immunology, 2019, 10, 3090.	4.8	39
65	ALK-positive anaplastic large-cell lymphoma in adults: an individual patient data pooled analysis of 263 patients. Haematologica, 2019, 104, e562-e565.	3.5	38
66	Interference with ERK-dimerization at the nucleocytosolic interface targets pathological ERK1/2 signaling without cardiotoxic side-effects. Nature Communications, 2020, 11, 1733.	12.8	38
67	A biological role for deletions in chromosomal band 13q14 in mantle cell and peripheral t-cell lymphomas?. , 1999, 26, 210-214.		36
68	BCL2 antibodies targeted at different epitopes detect varying levels of protein expression and correlate with frequent gene amplification in diffuse large B-cell lymphoma. Human Pathology, 2014, 45, 2144-2153.	2.0	34
69	CCL3 chemokine expression by chronic lymphocytic leukemia cells orchestrates the composition of the microenvironment in lymph node infiltrates. Leukemia and Lymphoma, 2016, 57, 563-571.	1.3	34
70	Gene expression profiling reveals a close relationship between follicular lymphoma grade 3A and 3B, but distinct profiles of follicular lymphoma grade 1 and 2. Haematologica, 2018, 103, 1182-1190.	3.5	34
71	Mutational mechanisms shaping the coding and noncoding genome of germinal center derived B-cell lymphomas. Leukemia, 2021, 35, 2002-2016.	7.2	34
72	SYK expression in monomorphic epitheliotropic intestinal T-cell lymphoma. Modern Pathology, 2018, 31, 505-516.	5.5	31

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73	FOXP1 expression is a prognostic biomarker in follicular lymphoma treated with rituximab and chemotherapy. Blood, 2018, 131, 226-235.	1.4	31
74	Cognate Nonlytic Interactions between CD8+ T Cells and Breast Cancer Cells Induce Cancer Stem Cellâ€“like Properties. Cancer Research, 2019, 79, 1507-1519.	0.9	31
75	Subgroup-Independent Mapping of Renal Cell Carcinomaâ€”Machine Learning Reveals Prognostic Mitochondrial Gene Signature Beyond Histopathologic Boundaries. Frontiers in Oncology, 2021, 11, 621278.	2.8	31
76	Repression of <scp>SRF</scp> target genes is critical for <scp>M</scp> ycâ€dependent apoptosis of epithelial cells. EMBO Journal, 2015, 34, 1554-1571.	7.8	30
77	Single- and double-hit events in genes encoding immune targets before and after T cellâ€“engaging antibody therapy in MM. Blood Advances, 2021, 5, 3794-3798.	5.2	30
78	Blocking TWEAK-Fn14 interaction inhibits hematopoietic stem cell transplantation-induced intestinal cell death and reduces GVHD. Blood, 2015, 126, 437-444.	1.4	29
79	RSPO2 gene rearrangement: a powerful driver of Î²-catenin activation in liver tumours. Gut, 2019, 68, 1287-1296.	12.1	29
80	Organ manifestations of COVID-19: what have we learned so far (not only) from autopsies?. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2022, 481, 139-159.	2.8	28
81	The clinicopathologic spectrum of mature aggressive B cell lymphomas. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2017, 471, 453-466.	2.8	27
82	Hyper-N-glycosylated SAMD14 and neurabin-I as driver autoantigens of primary central nervous system lymphoma. Blood, 2018, 132, 2744-2753.	1.4	27
83	Livin/BIRC7 expression as malignancy marker in adrenocortical tumors. Oncotarget, 2017, 8, 9323-9338.	1.8	27
84	Molecular and functional profiling identifies therapeutically targetable vulnerabilities in plasmablastic lymphoma. Nature Communications, 2021, 12, 5183.	12.8	26
85	Validation of the <scp>MCL</scp>35 gene expression proliferation assay in randomized trials of the European Mantle Cell Lymphoma Network. British Journal of Haematology, 2019, 184, 616-624.	2.5	25
86	The time to relapse correlates with the histopathological growth pattern in nodular lymphocyte predominant Hodgkin lymphoma. American Journal of Hematology, 2019, 94, 1208-1213.	4.1	25
87	Potential influence of concomitant chemotherapy on <scp>CXCR</scp>4 expression in receptor directed endoradiotherapy. British Journal of Haematology, 2019, 184, 440-443.	2.5	25
88	The exomic landscape of t(14;18)â€negative diffuse follicular lymphoma with 1p36 deletion. British Journal of Haematology, 2018, 180, 391-394.	2.5	24
89	The <scp>MCL</scp>35 gene expression proliferation assay predicts highâ€risk <scp>MCL</scp> patients in a Norwegian cohort of younger patients given intensive first line therapy. British Journal of Haematology, 2018, 183, 225-234.	2.5	24
90	In-depth cell-free DNA sequencing reveals genomic landscape of Hodgkinâ€™s lymphoma and facilitates ultrasensitive residual disease detection. Med, 2021, 2, 1171-1193.e11.	4.4	24

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91	Identification of <scp> Candida albicans </scp> regulatory genes governing mucosal infection. Cellular Microbiology, 2018, 20, e12841.	2.1	23
92	CD40L mediated alternative NFÎºB-signaling induces resistance to BCR-inhibitors in patients with mantle cell lymphoma. Cell Death and Disease, 2018, 9, 86.	6.3	23
93	Targeted Gene Expression Profile Reveals CDK4 as Therapeutic Target for Selected Patients With Adrenocortical Carcinoma. Frontiers in Endocrinology, 2020, 11, 219.	3.5	23
94	Prognostic value of tumour-infiltrating CD8+â€‰lymphocytes in rectal cancer after neoadjuvant chemoradiation: is indoleamine-2,3-dioxygenase (IDO1) a friend or foe?. Cancer Immunology, Immunotherapy, 2019, 68, 563-575.	4.2	22
95	Aggressive genomic features in clinically indolent primary HHV8-negative effusion-based lymphoma. Blood, 2019, 133, 377-380.	1.4	22
96	T-cell repertoires in refractory coeliac disease. Gut, 2018, 67, gutjnl-2016-311816.	12.1	21
97	Establishing Pure Cancer Organoid Cultures: Identification, Selection and Verification of Cancer Phenotypes and Genotypes. Journal of Molecular Biology, 2019, 431, 2884-2893.	4.2	21
98	Hexokinase-2 Expression in ¹¹C-Methionineâ€“Positive, ¹⁸F-FDGâ€“Negative Multiple Myeloma. Journal of Nuclear Medicine, 2019, 60, 348-352.	5.0	21
99	HSP90 promotes Burkitt lymphoma cell survival by maintaining tonic B-cell receptor signaling. Blood, 2017, 129, 598-608.	1.4	20
100	A 70% cut-off for MYC protein expression in diffuse large B cell lymphoma identifies a high-risk group of patients. Haematologica, 2020, 105, 2667-2670.	3.5	20
101	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. Blood, 2018, 132, 998-998.	1.4	19
102	Ten-year follow-up of a prospective trial for the targeted therapy of gastric cancer with the human monoclonal antibody PAT-SC1. Oncology Reports, 2014, 31, 1059-1066.	2.6	18
103	Rituximab plus high-dose chemotherapy (MegaCHOEP) or conventional chemotherapy (CHOEP-14) in young, high-risk patients with aggressive B-cell lymphoma: 10-year follow-up of a randomised, open-label, phase 3 trial. Lancet Haematology,the, 2021, 8, e267-e277.	4.6	18
104	MB3W1 is an orthotopic xenograft model for anaplastic medulloblastoma displaying cancer stem cell- and Group 3-properties. BMC Cancer, 2016, 16, 115.	2.6	17
105	Genome-Wide miRNA Expression Profiling of Molecular Subgroups of Peripheral T-cell Lymphoma. Clinical Cancer Research, 2021, 27, 6039-6053.	7.0	17
106	Targeting CXCR4 with [68Ga]Pentixafor: a suitable theranostic approach in pleural mesothelioma?. Oncotarget, 2017, 8, 96732-96737.	1.8	17
107	Gene Expression Signatures for the Accurate Diagnosis of Peripheral T-Cell Lymphoma Entities in the Routine Clinical Practice. Journal of Clinical Oncology, 2022, 40, 4261-4275.	1.6	17
108	Numerical and Structural Genomic Aberrations Are Reliably Detectable in Tissue Microarrays of Formalin-Fixed Paraffin-Embedded Tumor Samples by Fluorescence In-Situ Hybridization. PLoS ONE, 2014, 9, e95047.	2.5	16

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109	Cutaneous CD8+ Cytotoxic T-Cell Lymphoma Infiltrates: Clinicopathological Correlation and Outcome of 35 Cases. Oncology and Therapy, 2016, 4, 199-210.	2.6	16
110	The identification of patientâ€specific mutations reveals dual pathway activation in most patients with melanoma and activated receptor tyrosine kinases in BRAF/NRAS wildâ€type melanomas. Cancer, 2019, 125, 586-600.	4.1	16
111	Abemaciclib, a CDK4/6 inhibitor, exerts preclinical activity against aggressive germinal centerâ€derived Bâ€cell lymphomas. Cancer Science, 2020, 111, 749-759.	3.9	16
112	Ephrin receptor A2, the epithelial receptor for Epstein-Barr virus entry, is not available for efficient infection in human gastric organoids. PLoS Pathogens, 2021, 17, e1009210.	4.7	16
113	Actin cytoskeleton deregulation confers midostaurin resistance in FLT3-mutant acute myeloid leukemia. Communications Biology, 2021, 4, 799.	4.4	16
114	Lack of NFATc1 SUMOylation prevents autoimmunity and alloreactivity. Journal of Experimental Medicine, 2021, 218, .	8.5	15
115	Alleleâ€specific <scp>PCR</scp> is a powerful tool for the detection of the <scp>MYD</scp>88 L265P mutation in diffuse large B cell lymphoma and decalcified bone marrow samples. British Journal of Haematology, 2015, 171, 145-148.	2.5	14
116	Reverted exhaustion phenotype of circulating lymphocytes as immune correlate of anti-PD1 first-line treatment in Hodgkin lymphoma. Leukemia, 2022, 36, 760-771.	7.2	14
117	Alemtuzumab Added to CHOP for Treatment of Peripheral T-Cell Lymphoma (PTCL) in Previously Untreated Young and Elderly Patients: Pooled Analysis of the International ACT-1/2 Phase III Trials. Blood, 2018, 132, 1622-1622.	1.4	14
118	Altered Cellular Protein Levels of Tumor Suppressor Genes and Heat Shock Elements (TRAP1) Indicate Sensitivity to the Proteasome Inhibitor Bortezomib (VelcadeÂ®) in Mantle Cell Lymphoma.. Blood, 2005, 106, 2424-2424.	1.4	14
119	The G protein-coupled estrogen receptor 1 (GPER-1) contributes to the proliferation and survival of mantle cell lymphoma cells. Haematologica, 2015, 100, e458-e461.	3.5	13
120	Diffuse large B-cell lymphoma cell-of-origin classification using the Lymph2Cx assay in the context of BCL2 and MYC expression status. Leukemia and Lymphoma, 2016, 57, 717-720.	1.3	13
121	[¹¹C]Methionine emerges as a new biomarker for tracking active myeloma lesions. British Journal of Haematology, 2018, 181, 701-703.	2.5	13
122	ATM activity in T cells is critical for immune surveillance of lymphoma in vivo. Leukemia, 2020, 34, 771-786.	7.2	13
123	⁶⁸Ga-Pentixafor PET/CT for Detection of Chemokine Receptor CXCR4 Expression in Myeloproliferative Neoplasms. Journal of Nuclear Medicine, 2022, 63, 96-99.	5.0	13
124	Prognostic Significance of MYC Single, Double, Triple Hit and MYC-Translocation Partner Status in Diffuse Large B-Cell Lymphoma - a Study By the Lunenburg Lymphoma Biomarker Consortium (LLBC). Blood, 2018, 132, 344-344.	1.4	13
125	Alemtuzumab added to CHOP for treatment of peripheral T-cell lymphoma (pTNHL) of the elderly: Final results of 116 patients treated in the international ACT-2 phase III trial.. Journal of Clinical Oncology, 2016, 34, 7500-7500.	1.6	13
126	An analysis of the role of follicular lymphoma-associated fibroblasts to promote tumor cell viability following drug-induced apoptosis. Leukemia and Lymphoma, 2017, 58, 1922-1930.	1.3	12

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127	Differential expression of long nonâ€coding <scp>RNA</scp>s are related to proliferation and histological diversity in follicular lymphomas. British Journal of Haematology, 2019, 184, 373-383.	2.5	12
128	RAL GTPases mediate multiple myeloma cell survival and are activated independently of oncogenic RAS. Haematologica, 2020, 105, 2316-2326.	3.5	12
129	Mantle cell lymphomas with concomitant MYC and CCND1 breakpoints are recurrently TdT positive and frequently show high-grade pathological and genetic features. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2021, 479, 133-145.	2.8	12
130	PET-Guided Treatment of Early-Stage Favorable Hodgkin Lymphoma: Final Results of the International, Randomized Phase 3 Trial HD16 By the German Hodgkin Study Group. Blood, 2018, 132, 925-925.	1.4	12
131	Diverse PSMA expression in primary prostate cancer: reason for negative [68Ga]Ga-PSMA PET/CT scans? Immunohistochemical validation in 40 surgical specimens. European Journal of Nuclear Medicine and Molecular Imaging, 2022, , 1.	6.4	12
132	Identification of the atypically modified autoantigen Ars2 as the target of B-cell receptors from activated B-cell-type diffuse large B-cell lymphoma. Haematologica, 2021, 106, 2224-2232.	3.5	11
133	Localized- and advanced-stage follicular lymphomas differ in their gene expression profiles. Blood, 2020, 135, 181-190.	1.4	11
134	Whole-slide image analysis of the tumor microenvironment identifies low B-cell content as a predictor of adverse outcome in patients with advanced-stage classical Hodgkin lymphoma treated with BEACOPP. Haematologica, 2021, 106, 1684-1692.	3.5	11
135	Long-term outcomes from the phase II L-MIND study of tafasitamab (MOR208) plus lenalidomide in patients with relapsed or refractory diffuse large B-cell lymphoma. Haematologica, 2021, , .	3.5	11
136	Fluorescence in situ analysis of soft tissue tumor associated genetic alterations in formalin-fixed paraffin-embedded tissue. Pathology Research and Practice, 2014, 210, 804-811.	2.3	10
137	Detection of an activated JAK3 variant and a Xq26.3 microdeletion causing loss of PHF6 and miR-424 expression in myelodysplastic syndromes by combined targeted next generation sequencing and SNP array analysis. Pathology Research and Practice, 2014, 210, 369-376.	2.3	10
138	Targeting protein kinase C in mantle cell lymphoma. British Journal of Haematology, 2016, 173, 394-403.	2.5	10
139	Novel IGH and MYC Translocation Partners in Diffuse Large Bâ€Cell Lymphomas. Genes Chromosomes and Cancer, 2016, 55, 932-943.	2.8	10
140	Recurrent Oncogenic JAK and STAT AlterationsÂin Cutaneous CD30-Positive Lymphoproliferative Disorders. Journal of Investigative Dermatology, 2020, 140, 2023-2031.e1.	0.7	10
141	The impact of <scp>SAMHD1</scp> expression and mutation status in mantle cell lymphoma: An analysis of the <scp>MCL</scp> Younger and Elderly trial. International Journal of Cancer, 2021, 148, 150-160.	5.1	10
142	A Cyclin D1â€“Dependent Transcriptional Program Predicts Clinical Outcome in Mantle Cell Lymphoma. Clinical Cancer Research, 2021, 27, 213-225.	7.0	10
143	Evolutionary clonal trajectories in nodular lymphocyte-predominant Hodgkin lymphoma with high risk of transformation. Haematologica, 2021, 106, 2654-2666.	3.5	10
144	Primary mediastinal germ cell tumours: an immunohistochemical and molecular diagnostic approach. Histopathology, 2022, 80, 381-396.	2.9	10

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145	First-line therapy of T-cell lymphoma: Allogeneic or autologous transplantation for consolidationâ€"Final results of the AATT study.. Journal of Clinical Oncology, 2019, 37, 7503-7503.	1.6	10
146	CD19 expression is maintained in DLBCL patients after treatment with tafasitamab plus lenalidomide in the L-MIND study. Leukemia and Lymphoma, 2022, 63, 468-472.	1.3	10
147	Memory CD4+ T cells lacking expression of CCR7 promote pro-inflammatory cytokine production in patients with diffuse cutaneous systemic sclerosis. European Journal of Dermatology, 2019, 29, 468-476.	0.6	9
148	Doubling rituximab in highâ€risk patients with aggressive Bâ€cell lymphoma â€results of the <scp>DENSE</scp>â€Râ€Mega<scp>CHOEP</scp> trial. British Journal of Haematology, 2019, 184, 760-768.	2.5	9
149	9p24.1 alterations and programmed cell death 1 ligand 1 expression in early stage unfavourable classical Hodgkin lymphoma: an analysis from the German Hodgkin Study Group NIVAHL trial. British Journal of Haematology, 2022, 196, 116-126.	2.5	9
150	First case of human peritoneal cysticercosis mimicking peritoneal carcinosis: necessity of laparoscopy and histologic assessment for the correct diagnosis. JMM Case Reports, 2017, 4, e005097.	1.3	9
151	Nivolumab and AVD for Early-Stage Unfavorable Hodgkin Lymphoma (NIVAHL). Blood, 2019, 134, 236-236.	1.4	9
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