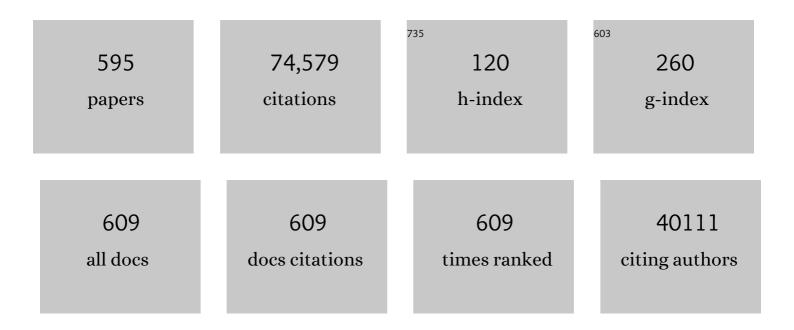
List of Publications by Year in descending order

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HADTMUT DöHNED

#	Article	IF	CITATIONS
1	Diagnosis and management of AML in adults: 2017 ELN recommendations from an international expert panel. Blood, 2017, 129, 424-447.	1.4	4,375
2	Genomic Classification and Prognosis in Acute Myeloid Leukemia. New England Journal of Medicine, 2016, 374, 2209-2221.	27.0	3,067
3	Genomic Aberrations and Survival in Chronic Lymphocytic Leukemia. New England Journal of Medicine, 2000, 343, 1910-1916.	27.0	2,967
4	Diagnosis and management of acute myeloid leukemia in adults: recommendations from an international expert panel, on behalf of the European LeukemiaNet. Blood, 2010, 115, 453-474.	1.4	2,963
5	Guidelines for the diagnosis and treatment of chronic lymphocytic leukemia: a report from the International Workshop on Chronic Lymphocytic Leukemia updating the National Cancer Institute–Working Group 1996 guidelines. Blood, 2008, 111, 5446-5456.	1.4	2,887
6	Acute Myeloid Leukemia. New England Journal of Medicine, 2015, 373, 1136-1152.	27.0	2,466
7	Midostaurin plus Chemotherapy for Acute Myeloid Leukemia with a <i>FLT3</i> Mutation. New England Journal of Medicine, 2017, 377, 454-464.	27.0	1,628
8	Mutations and Treatment Outcome in Cytogenetically Normal Acute Myeloid Leukemia. New England Journal of Medicine, 2008, 358, 1909-1918.	27.0	1,514
9	Azacitidine and Venetoclax in Previously Untreated Acute Myeloid Leukemia. New England Journal of Medicine, 2020, 383, 617-629.	27.0	1,407
10	Retinoic Acid and Arsenic Trioxide for Acute Promyelocytic Leukemia. New England Journal of Medicine, 2013, 369, 111-121.	27.0	1,284
11	Obinutuzumab plus Chlorambucil in Patients with CLL and Coexisting Conditions. New England Journal of Medicine, 2014, 370, 1101-1110.	27.0	1,284
12	Acute myeloid leukaemia. Lancet, The, 2006, 368, 1894-1907.	13.7	1,103
13	iwCLL guidelines for diagnosis, indications for treatment, response assessment, and supportive management of CLL. Blood, 2018, 131, 2745-2760.	1.4	1,069
14	International phase 3 study of azacitidine vs conventional care regimens in older patients with newly diagnosed AML with >30% blasts. Blood, 2015, 126, 291-299.	1.4	982
15	Use of Gene-Expression Profiling to Identify Prognostic Subclasses in Adult Acute Myeloid Leukemia. New England Journal of Medicine, 2004, 350, 1605-1616.	27.0	915
16	Mutations driving CLL and their evolution in progression and relapse. Nature, 2015, 526, 525-530.	27.8	868
17	Management of acute promyelocytic leukemia: recommendations from an expert panel on behalf of the European LeukemiaNet. Blood, 2009, 113, 1875-1891.	1.4	856
18	International Consensus Classification of Myeloid Neoplasms and Acute Leukemias: integrating morphologic, clinical, and genomic data. Blood, 2022, 140, 1200-1228.	1.4	814

#	Article	IF	CITATIONS
19	Diagnosis and management of AML in adults: 2022 recommendations from an international expert panel on behalf of the ELN. Blood, 2022, 140, 1345-1377.	1.4	805
20	Prognostic significance of activating FLT3 mutations in younger adults (16 to 60 years) with acute myeloid leukemia and normal cytogenetics: a study of the AML Study Group Ulm. Blood, 2002, 100, 4372-4380.	1.4	794
21	Mutant nucleophosmin (NPM1) predicts favorable prognosis in younger adults with acute myeloid leukemia and normal cytogenetics: interaction with other gene mutations. Blood, 2005, 106, 3740-3746.	1.4	779
22	Allogeneic Stem Cell Transplantation for Acute Myeloid Leukemia in First Complete Remission. JAMA - Journal of the American Medical Association, 2009, 301, 2349.	7.4	758
23	High-Dose Daunorubicin in Older Patients with Acute Myeloid Leukemia. New England Journal of Medicine, 2009, 361, 1235-1248.	27.0	745
24	Genome Sequencing of Pediatric Medulloblastoma Links Catastrophic DNA Rearrangements with TP53 Mutations. Cell, 2012, 148, 59-71.	28.9	743
25	<i>IDH1</i> and <i>IDH2</i> Mutations Are Frequent Genetic Alterations in Acute Myeloid Leukemia and Confer Adverse Prognosis in Cytogenetically Normal Acute Myeloid Leukemia With <i>NPM1</i> Mutation Without <i>FLT3</i> Internal Tandem Duplication. Journal of Clinical Oncology, 2010, 28, 3636-3643.	1.6	728
26	A Single Oncogenic Enhancer Rearrangement Causes Concomitant EVI1 and GATA2 Deregulation in Leukemia. Cell, 2014, 157, 369-381.	28.9	571
27	Long-term remissions after FCR chemoimmunotherapy in previously untreated patients with CLL: updated results of the CLL8 trial. Blood, 2016, 127, 208-215.	1.4	571
28	Reduced-intensity chemotherapy and PET-guided radiotherapy in patients with advanced stage Hodgkin's lymphoma (HD15 trial): a randomised, open-label, phase 3 non-inferiority trial. Lancet, The, 2012, 379, 1791-1799.	13.7	564
29	Selective BCL-2 Inhibition by ABT-199 Causes On-Target Cell Death in Acute Myeloid Leukemia. Cancer Discovery, 2014, 4, 362-375.	9.4	561
30	TP53 alterations in acute myeloid leukemia with complex karyotype correlate with specific copy number alterations, monosomal karyotype, and dismal outcome. Blood, 2012, 119, 2114-2121.	1.4	553
31	Detection of complete and partial chromosome gains and losses by comparative genomic in situ hybridization. Human Genetics, 1993, 90, 590-610.	3.8	544
32	Fludarabine plus cyclophosphamide versus fludarabine alone in first-line therapy of younger patients with chronic lymphocytic leukemia. Blood, 2005, 107, 885-891.	1.4	524
33	<i>TP53</i> Mutation and Survival in Chronic Lymphocytic Leukemia. Journal of Clinical Oncology, 2010, 28, 4473-4479.	1.6	523
34	Molecular Genetics of Adult Acute Myeloid Leukemia: Prognostic and Therapeutic Implications. Journal of Clinical Oncology, 2011, 29, 475-486.	1.6	510
35	From pathogenesis to treatment of chronic lymphocytic leukaemia. Nature Reviews Cancer, 2010, 10, 37-50.	28.4	503
36	A phase 1 study of SU11248 in the treatment of patients with refractory or resistant acute myeloid leukemia (AML) or not amenable to conventional therapy for the disease. Blood, 2005, 105, 986-993.	1.4	481

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37	Gene mutations and treatment outcome in chronic lymphocytic leukemia: results from the CLL8 trial. Blood, 2014, 123, 3247-3254.	1.4	428
38	<i>CEBPA</i> Mutations in Younger Adults With Acute Myeloid Leukemia and Normal Cytogenetics: Prognostic Relevance and Analysis of Cooperating Mutations. Journal of Clinical Oncology, 2004, 22, 624-633.	1.6	427
39	Intensified Chemotherapy and Dose-Reduced Involved-Field Radiotherapy in Patients With Early Unfavorable Hodgkin's Lymphoma: Final Analysis of the German Hodgkin Study Group HD11 Trial. Journal of Clinical Oncology, 2010, 28, 4199-4206.	1.6	397
40	Minimal Residual Disease Quantification Is an Independent Predictor of Progression-Free and Overall Survival in Chronic Lymphocytic Leukemia: A Multivariate Analysis From the Randomized GCLLSG CLL8 Trial. Journal of Clinical Oncology, 2012, 30, 980-988.	1.6	397
41	Management of acute promyelocytic leukemia: updated recommendations from an expert panel of the European LeukemiaNet. Blood, 2019, 133, 1630-1643.	1.4	393
42	The impact of therapy-related acute myeloid leukemia (AML) on outcome in 2853 adult patients with newly diagnosed AML. Blood, 2011, 117, 2137-2145.	1.4	392
43	Bendamustine in Combination With Rituximab for Previously Untreated Patients With Chronic Lymphocytic Leukemia: A Multicenter Phase II Trial of the German Chronic Lymphocytic Leukemia Study Group. Journal of Clinical Oncology, 2012, 30, 3209-3216.	1.6	388
44	Genomics of Acute Myeloid Leukemia Diagnosis and Pathways. Journal of Clinical Oncology, 2017, 35, 934-946.	1.6	372
45	Monitoring of Minimal Residual Disease in <i>NPM1</i> -Mutated Acute Myeloid Leukemia: A Study From the German-Austrian Acute Myeloid Leukemia Study Group. Journal of Clinical Oncology, 2011, 29, 2709-2716.	1.6	355
46	Differential impact of allelic ratio and insertion site in FLT3-ITD–positive AML with respect to allogeneic transplantation. Blood, 2014, 124, 3441-3449.	1.4	350
47	<i>RUNX1</i> Mutations in Acute Myeloid Leukemia: Results From a Comprehensive Genetic and Clinical Analysis From the AML Study Group. Journal of Clinical Oncology, 2011, 29, 1364-1372.	1.6	349
48	Monoallelic TP53 inactivation is associated with poor prognosis in chronic lymphocytic leukemia: results from a detailed genetic characterization with long-term follow-up. Blood, 2008, 112, 3322-3329.	1.4	348
49	Prognostic impact, concurrent genetic mutations, and gene expression features of AML with CEBPA mutations in a cohort of 1182 cytogenetically normal AML patients: further evidence for CEBPA double mutant AML as a distinctive disease entity. Blood, 2011, 117, 2469-2475.	1.4	341
50	Distinct evolution and dynamics of epigenetic and genetic heterogeneity in acute myeloid leukemia. Nature Medicine, 2016, 22, 792-799.	30.7	322
51	Prognostic Significance of Partial Tandem Duplications of the MLL Gene in Adult Patients 16 to 60 Years Old With Acute Myeloid Leukemia and Normal Cytogenetics: A Study of the Acute Myeloid Leukemia Study Group Ulm. Journal of Clinical Oncology, 2002, 20, 3254-3261.	1.6	291
52	Allogeneic stem cell transplantation provides durable disease control in poor-risk chronic lymphocytic leukemia: long-term clinical and MRD results of the German CLL Study Group CLL3X trial. Blood, 2010, 116, 2438-2447.	1.4	273
53	Cytogenetics and age are major determinants of outcome in intensively treated acute myeloid leukemia patients older than 60 years: results from AMLSG trial AML HD98-B. Blood, 2006, 108, 3280-3288.	1.4	269
54	A dominant-negative effect drives selection of <i>TP53</i> missense mutations in myeloid malignancies. Science, 2019, 365, 599-604.	12.6	265

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55	Oral Azacitidine Maintenance Therapy for Acute Myeloid Leukemia in First Remission. New England Journal of Medicine, 2020, 383, 2526-2537.	27.0	265
56	miR-34a as part of the resistance network in chronic lymphocytic leukemia. Blood, 2009, 113, 3801-3808.	1.4	258
57	Subcutaneous Alemtuzumab in Fludarabine-Refractory Chronic Lymphocytic Leukemia: Clinical Results and Prognostic Marker Analyses From the CLL2H Study of the German Chronic Lymphocytic Leukemia Study Group. Journal of Clinical Oncology, 2009, 27, 3994-4001.	1.6	257
58	DNA methylation dynamics during B cell maturation underlie a continuum of disease phenotypes in chronic lymphocytic leukemia. Nature Genetics, 2016, 48, 253-264.	21.4	254
59	Development of a comprehensive prognostic index for patients with chronic lymphocytic leukemia. Blood, 2014, 124, 49-62.	1.4	244
60	Biallelic mutations in the ATM gene in T-prolymphocytic leukemia. Nature Medicine, 1997, 3, 1155-1159.	30.7	243
61	Insertion of FLT3 internal tandem duplication in the tyrosine kinase domain-1 is associated with resistance to chemotherapy and inferior outcome. Blood, 2009, 114, 2386-2392.	1.4	242
62	Clonal evolution in relapsed NPM1-mutated acute myeloid leukemia. Blood, 2013, 122, 100-108.	1.4	242
63	Measurable residual disease monitoring by NGS before allogeneic hematopoietic cell transplantation in AML. Blood, 2018, 132, 1703-1713.	1.4	237
64	Precision oncology for acute myeloid leukemia using a knowledge bank approach. Nature Genetics, 2017, 49, 332-340.	21.4	229
65	Midostaurin added to chemotherapy and continued single-agent maintenance therapy in acute myeloid leukemia with FLT3-ITD. Blood, 2019, 133, 840-851.	1.4	228
66	High <i>EVI1</i> Expression Predicts Outcome in Younger Adult Patients With Acute Myeloid Leukemia and Is Associated With Distinct Cytogenetic Abnormalities. Journal of Clinical Oncology, 2010, 28, 2101-2107.	1.6	222
67	Automated array-based genomic profiling in chronic lymphocytic leukemia: Development of a clinical tool and discovery of recurrent genomic alterations. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 1039-1044.	7.1	221
68	A multicenter phase II trial of decitabine as first-line treatment for older patients with acute myeloid leukemia judged unfit for induction chemotherapy. Haematologica, 2012, 97, 393-401.	3.5	219
69	Clinical, Molecular, and Prognostic Significance of WHO Type inv(3)(q21q26.2)/t(3;3)(q21;q26.2) and Various Other 3q Abnormalities in Acute Myeloid Leukemia. Journal of Clinical Oncology, 2010, 28, 3890-3898.	1.6	217
70	The genomic landscape of core-binding factor acute myeloid leukemias. Nature Genetics, 2016, 48, 1551-1556.	21.4	215
71	An Inv(16)(p13.3q24.3)-Encoded CBFA2T3-GLIS2 Fusion Protein Defines an Aggressive Subtype of Pediatric Acute Megakaryoblastic Leukemia. Cancer Cell, 2012, 22, 683-697.	16.8	213
72	V(H) mutation status, CD38 expression level, genomic aberrations, and survival in chronic lymphocytic leukemia. Blood, 2002, 100, 1410-6.	1.4	206

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73	Prospective Evaluation of Allogeneic Hematopoietic Stem-Cell Transplantation From Matched Related and Matched Unrelated Donors in Younger Adults With High-Risk Acute Myeloid Leukemia: German-Austrian Trial AMLHD98A. Journal of Clinical Oncology, 2010, 28, 4642-4648.	1.6	205
74	Quizartinib, an FLT3 inhibitor, as monotherapy in patients with relapsed or refractory acute myeloid leukaemia: an open-label, multicentre, single-arm, phase 2 trial. Lancet Oncology, The, 2018, 19, 889-903.	10.7	205
75	Randomized, phase 2 trial of low-dose cytarabine with or without volasertib in AML patients not suitable for induction therapy. Blood, 2014, 124, 1426-1433.	1.4	204
76	Microarray Gene Expression Profiling of B-Cell Chronic Lymphocytic Leukemia Subgroups Defined by Genomic Aberrations and <i>VH</i> Mutation Status. Journal of Clinical Oncology, 2004, 22, 3937-3949.	1.6	200
77	<i>TET2</i> Mutations in Acute Myeloid Leukemia (AML): Results From a Comprehensive Genetic and Clinical Analysis of the AML Study Group. Journal of Clinical Oncology, 2012, 30, 1350-1357.	1.6	198
78	Campath-1H–Induced Complete Remission of Chronic Lymphocytic Leukemia despitep53Gene Mutation and Resistance to Chemotherapy. New England Journal of Medicine, 2002, 347, 452-453.	27.0	195
79	Clonal evolution in chronic lymphocytic leukemia: acquisition of high-risk genomic aberrations associated with unmutated VH, resistance to therapy, and short survival. Haematologica, 2007, 92, 1242-1245.	3.5	195
80	Gene mutations and response to treatment with all-trans retinoic acid in elderly patients with acute myeloid leukemia. Results from the AMLSG Trial AML HD98B. Haematologica, 2009, 94, 54-60.	3.5	195
81	V617F mutation in JAK2 is associated with poorer survival in idiopathic myelofibrosis. Blood, 2006, 107, 2098-2100.	1.4	194
82	Chromosomal Abnormalities in Cancer. New England Journal of Medicine, 2008, 359, 722-734.	27.0	188
83	lvosidenib and Azacitidine in <i>IDH1</i> -Mutated Acute Myeloid Leukemia. New England Journal of Medicine, 2022, 386, 1519-1531.	27.0	186
84	Human Chromosome 7: DNA Sequence and Biology. Science, 2003, 300, 767-772.	12.6	185
85	miRNA-130a Targets <i>ATG2B</i> and <i>DICER1</i> to Inhibit Autophagy and Trigger Killing of Chronic Lymphocytic Leukemia Cells. Cancer Research, 2012, 72, 1763-1772.	0.9	185
86	APO-1 mediated apoptosis or proliferation in human chronic B lymphocytic leukemia: Correlation with bcl-2 oncogene expression. European Journal of Immunology, 1993, 23, 702-708.	2.9	178
87	Additional Genetic High-Risk Features Such As 11q Deletion, 17p Deletion, and <i>V3-21</i> Usage Characterize Discordance of ZAP-70 and <i>VH</i> Mutation Status in Chronic Lymphocytic Leukemia. Journal of Clinical Oncology, 2006, 24, 969-975.	1.6	177
88	Identification of Driver and Passenger Mutations of FLT3 by High-Throughput DNA Sequence Analysis and Functional Assessment of Candidate Alleles. Cancer Cell, 2007, 12, 501-513.	16.8	174
89	Molecular Imaging of Proliferation in Malignant Lymphoma. Cancer Research, 2006, 66, 11055-11061.	0.9	173
90	Strikingly homologous immunoglobulin gene rearrangements and poor outcome in VH3-21-using chronic lymphocytic leukemia patients independent of geographic origin and mutational status. Blood, 2006, 107, 2889-2894.	1.4	167

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91	Mutations in the cohesin complex in acute myeloid leukemia: clinical and prognostic implications. Blood, 2014, 123, 914-920.	1.4	167
92	Prognostic impact of WT1 mutations in cytogenetically normal acute myeloid leukemia: a study of the German-Austrian AML Study Group. Blood, 2009, 113, 4505-4511.	1.4	164
93	Secondary genetic lesions in acute myeloid leukemia with inv(16) or t(16;16): a study of the German-Austrian AML Study Group (AMLSG). Blood, 2013, 121, 170-177.	1.4	164
94	Clinical impact of DNMT3A mutations in younger adult patients with acute myeloid leukemia: results of the AML Study Group (AMLSG). Blood, 2013, 121, 4769-4777.	1.4	162
95	Venetoclax resistance and acquired <i>BCL2</i> mutations in chronic lymphocytic leukemia. Haematologica, 2019, 104, e434-e437.	3.5	144
96	Molecular characterization of acute myeloid leukemia. Haematologica, 2008, 93, 976-982.	3.5	143
97	Minimal Residual Disease Assessment Improves Prediction of Outcome in Patients With Chronic Lymphocytic Leukemia (CLL) Who Achieve Partial Response: Comprehensive Analysis of Two Phase III Studies of the German CLL Study Group. Journal of Clinical Oncology, 2016, 34, 3758-3765.	1.6	142
98	Disclosure of Candidate Genes in Acute Myeloid Leukemia With Complex Karyotypes Using Microarray-Based Molecular Characterization. Journal of Clinical Oncology, 2006, 24, 3887-3894.	1.6	141
99	Genomic DNA-Chip Hybridization Reveals a Higher Incidence of Genomic Amplifications in Pancreatic Cancer than Conventional Comparative Genomic Hybridization and Leads to the Identification of Novel Candidate Genes. Cancer Research, 2004, 64, 4428-4433.	0.9	140
100	Perspectives on the use of new diagnostic tools in the treatment of chronic lymphocytic leukemia. Blood, 2005, 107, 859-861.	1.4	140
101	Monosomal karyotype in adult acute myeloid leukemia: prognostic impact and outcome after different treatment strategies. Blood, 2012, 119, 551-558.	1.4	140
102	Expressed sequences as candidates for a novel tumor suppressor gene at band 13q14 in B-cell chronic lymphocytic leukemia and mantle cell lymphoma. Oncogene, 1998, 16, 1891-1897.	5.9	139
103	Gain of chromosome arm 9p is characteristic of primary mediastinal b-cell lymphoma (MBL): Comprehensive molecular cytogenetic analysis and presentation of a novel MBL cell line. Genes Chromosomes and Cancer, 2001, 30, 393-401.	2.8	138
104	The value of allogeneic and autologous hematopoietic stem cell transplantation in prognostically favorable acute myeloid leukemia with double mutant CEBPA. Blood, 2013, 122, 1576-1582.	1.4	138
105	Towards precision medicine for AML. Nature Reviews Clinical Oncology, 2021, 18, 577-590.	27.6	138
106	Quantitative DNA methylation predicts survival in adult acute myeloid leukemia. Blood, 2010, 115, 636-642.	1.4	137
107	VH mutation status and VDJ rearrangement structure in mantle cell lymphoma: correlation with genomic aberrations, clinical characteristics, and outcome. Blood, 2003, 102, 3003-3009.	1.4	136
108	Evolution of DNA Methylation Is Linked to Genetic Aberrations in Chronic Lymphocytic Leukemia. Cancer Discovery, 2014, 4, 348-361.	9.4	135

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109	Epigenetic Upregulation of IncRNAs at 13q14.3 in Leukemia Is Linked to the In Cis Downregulation of a Gene Cluster That Targets NF-kB. PLoS Genetics, 2013, 9, e1003373.	3.5	134
110	Genomic Classification in Acute Myeloid Leukemia. New England Journal of Medicine, 2016, 375, 900-901.	27.0	134
111	Gemtuzumab ozogamicin as postremission treatment in AML at 60 years of age or more: results of a multicenter phase 3 study. Blood, 2010, 115, 2586-2591.	1.4	131
112	Commonly altered genomic regions in acute myeloid leukemia are enriched for somatic mutations involved in chromatin remodeling and splicing. Blood, 2012, 120, e83-e92.	1.4	131
113	Serum microRNAs as a novel class of biomarkers: a comprehensive review of the literature. Experimental Hematology, 2010, 38, 1126-1130.	0.4	129
114	Mutated regions of nucleophosmin 1 elicit both CD4+ and CD8+ T-cell responses in patients with acute myeloid leukemia. Blood, 2012, 120, 1282-1289.	1.4	129
115	Impact of NPM1/FLT3-ITD genotypes defined by the 2017 European LeukemiaNet in patients with acute myeloid leukemia. Blood, 2020, 135, 371-380.	1.4	127
116	The homeobox gene CDX2 is aberrantly expressed in most cases of acute myeloid leukemia and promotes leukemogenesis. Journal of Clinical Investigation, 2007, 117, 1037-1048.	8.2	127
117	Receptor for hyaluronan acid–mediated motility (RHAMM) is a new immunogenic leukemia-associated antigen in acute and chronic myeloid leukemia. Experimental Hematology, 2002, 30, 1029-1035.	0.4	126
118	Prognostic Impact of Minimal Residual Disease in <i>CBFB-MYH11</i> –Positive Acute Myeloid Leukemia. Journal of Clinical Oncology, 2010, 28, 3724-3729.	1.6	126
119	Evidence for distinct pathomechanisms in B-cell chronic lymphocytic leukemia and mantle cell lymphoma by quantitative expression analysis of cell cycle and apoptosis-associated genes. Blood, 2002, 99, 4554-4561.	1.4	125
120	Exclusive Detection of the t(11;18)(q21;q21) in Extranodal Marginal Zone B Cell Lymphomas (MZBL) of MALT Type in Contrast to other MZBL and Extranodal Large B Cell Lymphomas. American Journal of Pathology, 1999, 155, 1817-1821.	3.8	124
121	High-dose RHAMM-R3 peptide vaccination for patients with acute myeloid leukemia, myelodysplastic syndrome and multiple myeloma. Haematologica, 2010, 95, 1191-1197.	3.5	124
122	lvosidenib or enasidenib combined with intensive chemotherapy in patients with newly diagnosed AML: a phase 1 study. Blood, 2021, 137, 1792-1803.	1.4	123
123	Short telomeres are associated with genetic complexity, high-risk genomic aberrations, and short survival in chronic lymphocytic leukemia. Blood, 2008, 111, 2246-2252.	1.4	122
124	Acute Myeloid Leukemia (AML): Different Treatment Strategies Versus a Common Standard Arm—Combined Prospective Analysis by the German AML Intergroup. Journal of Clinical Oncology, 2012, 30, 3604-3610.	1.6	121
125	Prognostic Value of Minimal Residual Disease Quantification by Real-Time Reverse Transcriptase Polymerase Chain Reaction in Patients With Core Binding Factor Leukemias. Journal of Clinical Oncology, 2003, 21, 4413-4422.	1.6	120
126	Quantitative DNA Methylation Analysis Identifies a Single CpG Dinucleotide Important for ZAP-70 Expression and Predictive of Prognosis in Chronic Lymphocytic Leukemia. Journal of Clinical Oncology, 2012, 30, 2483-2491.	1.6	120

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127	Deletions below 10 megabasepairs are detected in comparative genomic hybridization by standard reference intervals. Genes Chromosomes and Cancer, 1999, 25, 410-413.	2.8	119
128	mRNA expression of leukemiaâ€associated antigens in patients with acute myeloid leukemia for the development of specific immunotherapies. International Journal of Cancer, 2004, 108, 704-711.	5.1	118
129	Circulating microRNAs in hematological diseases: principles, challenges, and perspectives. Blood, 2013, 121, 4977-4984.	1.4	118
130	Mutant Isocitrate Dehydrogenase 1 Inhibitor Ivosidenib in Combination With Azacitidine for Newly Diagnosed Acute Myeloid Leukemia. Journal of Clinical Oncology, 2021, 39, 57-65.	1.6	118
131	Unmutated immunoglobulin variable heavy-chain gene status remains an adverse prognostic factor after autologous stem cell transplantation for chronic lymphocytic leukemia. Blood, 2003, 101, 2049-2053.	1.4	116
132	Impact of Fluoroquinolone Prophylaxis on Reduced Infectionâ€Related Mortality among Patients with Neutropenia and Hematologic Malignancies. Clinical Infectious Diseases, 2005, 40, 1087-1093.	5.8	116
133	Inactivating CUX1 mutations promote tumorigenesis. Nature Genetics, 2014, 46, 33-38.	21.4	111
134	Fludarabine plus cyclophosphamide is an efficient treatment for advanced chronic lymphocytic leukaemia (CLL): results of a phase II study of the German CLL Study Group. British Journal of Haematology, 2001, 114, 342-348.	2.5	108
135	Epidemiological, genetic, and clinical characterization by age of newly diagnosed acute myeloid leukemia based on an academic population-based registry study (AMLSG BiO). Annals of Hematology, 2017, 96, 1993-2003.	1.8	108
136	Risk categories and refractory CLL in the era of chemoimmunotherapy. Blood, 2012, 119, 4101-4107.	1.4	107
137	CDNA microarray gene expression analysis of B-cell chronic lymphocytic leukemia proposes potential new prognostic markers involved in lymphocyte trafficking. International Journal of Cancer, 2001, 91, 474-480.	5.1	106
138	The prognostic impact of autologous stem cell transplantation in patients with chronic lymphocytic leukemia: a risk-matched analysis based on the VH gene mutational status. Blood, 2004, 103, 2850-2858.	1.4	101
139	Interactions between comorbidity and treatment of chronic lymphocytic leukemia: results of German Chronic Lymphocytic Leukemia Study Group trials. Haematologica, 2014, 99, 1095-1100.	3.5	101
140	Cytogenetics and gene mutations influence survival in older patients with acute myeloid leukemia treated with azacitidine or conventional care. Leukemia, 2018, 32, 2546-2557.	7.2	101
141	Hidden gene amplifications in aggressive B-cell non-Hodgkin lymphomas detected by microarray-based comparative genomic hybridization. Oncogene, 2003, 22, 1425-1429.	5.9	99
142	Comparison of Cytogenetic and Molecular Cytogenetic Detection of Chromosome Abnormalities in 240 Consecutive Adult Patients With Acute Myeloid Leukemia. Journal of Clinical Oncology, 2002, 20, 2480-2485.	1.6	98
143	Automated Screening for Genomic Imbalances using Matrix-Based Comparative Genomic Hybridization. Laboratory Investigation, 2002, 82, 47-60.	3.7	96
144	TP53, SF3B1, and NOTCH1 mutations and outcome of allotransplantation for chronic lymphocytic leukemia: six-year follow-up of the GCLLSG CLL3X trial. Blood, 2013, 121, 3284-3288.	1.4	96

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145	Deregulated Expression of <i>EVI1</i> Defines a Poor Prognostic Subset of <i>MLL</i> -Rearranged Acute Myeloid Leukemias: A Study of the German-Austrian Acute Myeloid Leukemia Study Group and the Dutch-Belgian-Swiss HOVON/SAKK Cooperative Group. Journal of Clinical Oncology, 2013, 31, 95-103.	1.6	95
146	An FLT3 gene-expression signature predicts clinical outcome in normal karyotype AML. Blood, 2008, 111, 4490-4495.	1.4	94
147	188Re or 90Y-labelled anti-CD66 antibody as part of a dose-reduced conditioning regimen for patients with acute leukaemia or myelodysplastic syndrome over the age of 55: results of a phase I-II study. British Journal of Haematology, 2005, 130, 604-613.	2.5	92
148	Down-regulation of candidate tumor suppressor genes within chromosome band 13q14.3 is independent of the DNA methylation pattern in B-cell chronic lymphocytic leukemia. Blood, 2002, 99, 4116-4121.	1.4	91
149	Contrasting requirements during disease evolution identify EZH2 as a therapeutic target in AML. Journal of Experimental Medicine, 2019, 216, 966-981.	8.5	91
150	Clinical practice recommendation on hematopoietic stem cell transplantation for acute myeloid leukemia patients with <i>FLT3</i> -internal tandem duplication: a position statement from the Acute Leukemia Working Party of the European Society for Blood and Marrow Transplantation. Haematologica, 2020, 105, 1507-1516.	3.5	91
151	Clonal evolution of acute myeloid leukemia with <i>FLT3</i> -ITD mutation under treatment with midostaurin. Blood, 2021, 137, 3093-3104.	1.4	91
152	A phase I/II study of sunitinib and intensive chemotherapy in patients over 60 years of age with acute myeloid leukaemia and activating <i>FLT3</i> mutations. British Journal of Haematology, 2015, 169, 694-700.	2.5	90
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