List of Publications by Year in descending order

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		5558	3021
334	38,906	82	188
papers	citations	h-index	g-index
341	341	341	43325
all docs	docs citations	times ranked	citing authors

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#	Article	IF	CITATIONS
1	Age-Related Clonal Hematopoiesis Associated with Adverse Outcomes. New England Journal of Medicine, 2014, 371, 2488-2498.	13.9	3,474
2	An immunogenic personal neoantigen vaccine for patients with melanoma. Nature, 2017, 547, 217-221.	13.7	2,112
3	Clonal Hematopoiesis and Risk of Atherosclerotic Cardiovascular Disease. New England Journal of Medicine, 2017, 377, 111-121.	13.9	1,738
4	Clinical Effect of Point Mutations in Myelodysplastic Syndromes. New England Journal of Medicine, 2011, 364, 2496-2506.	13.9	1,444
5	Molecular subtypes of diffuse large B cell lymphoma are associated with distinct pathogenic mechanisms and outcomes. Nature Medicine, 2018, 24, 679-690.	15.2	1,224
6	Evolution and Impact of Subclonal Mutations in Chronic Lymphocytic Leukemia. Cell, 2013, 152, 714-726.	13.5	1,202
7	Integrative analysis reveals selective 9p24.1 amplification, increased PD-1 ligand expression, and further induction via JAK2 in nodular sclerosing Hodgkin lymphoma and primary mediastinal large B-cell lymphoma. Blood, 2010, 116, 3268-3277.	0.6	1,122
8	Gene expression signatures define novel oncogenic pathways in T cell acute lymphoblastic leukemia. Cancer Cell, 2002, 1, 75-87.	7.7	1,024
9	<i>SF3B1</i> and Other Novel Cancer Genes in Chronic Lymphocytic Leukemia. New England Journal of Medicine, 2011, 365, 2497-2506.	13.9	1,021
10	Analysis of <i>Fusobacterium</i> persistence and antibiotic response in colorectal cancer. Science, 2017, 358, 1443-1448.	6.0	983
11	Neoantigen vaccine generates intratumoral T cell responses in phase Ib glioblastoma trial. Nature, 2019, 565, 234-239.	13.7	956
12	Mutations driving CLL and their evolution in progression and relapse. Nature, 2015, 526, 525-530.	13.7	868
13	Molecular profiling of diffuse large B-cell lymphoma identifies robust subtypes including one characterized by host inflammatory response. Blood, 2005, 105, 1851-1861.	0.6	778
14	Acute myeloid leukemia ontogeny is defined by distinct somatic mutations. Blood, 2015, 125, 1367-1376.	0.6	747
15	Landscape of genomic alterations in cervical carcinomas. Nature, 2014, 506, 371-375.	13.7	708
16	<i>PD-L1</i> and <i>PD-L2</i> Genetic Alterations Define Classical Hodgkin Lymphoma and Predict Outcome. Journal of Clinical Oncology, 2016, 34, 2690-2697.	0.8	634
17	Immunologic and clinical effects of antibody blockade of cytotoxic T lymphocyte-associated antigen 4 in previously vaccinated cancer patients. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 3005-3010.	3.3	604
18	Prognostic Mutations in Myelodysplastic Syndrome after Stem-Cell Transplantation. New England Journal of Medicine, 2017, 376, 536-547.	13.9	586

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19	Interplay of somatic alterations and immune infiltration modulates response to PD-1 blockade in advanced clear cell renal cell carcinoma. Nature Medicine, 2020, 26, 909-918.	15.2	488
20	TET2 mutations predict response to hypomethylating agents in myelodysplastic syndrome patients. Blood, 2014, 124, 2705-2712.	0.6	486
21	High frequency of PTEN, PI3K, and AKT abnormalities in T-cell acute lymphoblastic leukemia. Blood, 2009, 114, 647-650.	0.6	414
22	Assessment of dexrazoxane as a cardioprotectant in doxorubicin-treated children with high-risk acute lymphoblastic leukaemia: long-term follow-up of a prospective, randomised, multicentre trial. Lancet Oncology, The, 2010, 11, 950-961.	5.1	377
23	Implications of TP53 allelic state for genome stability, clinical presentation and outcomes in myelodysplastic syndromes. Nature Medicine, 2020, 26, 1549-1556.	15.2	372
24	Results of the Dana-Farber Cancer Institute ALL Consortium Protocol 95-01 for children with acute lymphoblastic leukemia. Blood, 2007, 109, 896-904.	0.6	362
25	Somatic Mutations Predict Poor Outcome in Patients With Myelodysplastic Syndrome After Hematopoietic Stem-Cell Transplantation. Journal of Clinical Oncology, 2014, 32, 2691-2698.	0.8	359
26	Clonal Hematopoiesis Associated With Adverse Outcomes After Autologous Stem-Cell Transplantation for Lymphoma. Journal of Clinical Oncology, 2017, 35, 1598-1605.	0.8	339
27	Locally Disordered Methylation Forms the Basis of Intratumor Methylome Variation in Chronic Lymphocytic Leukemia. Cancer Cell, 2014, 26, 813-825.	7.7	323
28	Distinct evolution and dynamics of epigenetic and genetic heterogeneity in acute myeloid leukemia. Nature Medicine, 2016, 22, 792-799.	15.2	322
29	Detection by polymerase chain reaction of residual cells with the bcl-2 translocation is associated with increased risk of relapse after autologous bone marrow transplantation for B-cell lymphoma. Blood, 1993, 81, 3449-3457.	0.6	318
30	Toxicity and Efficacy of Defined Doses of CD4+ Donor Lymphocytes for Treatment of Relapse After Allogeneic Bone Marrow Transplant. Blood, 1998, 91, 3671-3680.	0.6	304
31	Increased neutrophil extracellular trap formation promotes thrombosis in myeloproliferative neoplasms. Science Translational Medicine, 2018, 10, .	5.8	299
32	Relative Mitochondrial Priming of Myeloblasts and Normal HSCs Determines Chemotherapeutic Success in AML. Cell, 2012, 151, 344-355.	13.5	294
33	Systematic identification of personal tumor-specific neoantigens in chronic lymphocytic leukemia. Blood, 2014, 124, 453-462.	0.6	286
34	Clonal evolution in patients with chronic lymphocytic leukaemia developing resistance to BTK inhibition. Nature Communications, 2016, 7, 11589.	5.8	285
35	Postinduction Dexamethasone and Individualized Dosing of <i>Escherichia Coli</i> L-Asparaginase Each Improve Outcome of Children and Adolescents With Newly Diagnosed Acute Lymphoblastic Leukemia: Results From a Randomized Study—Dana-Farber Cancer Institute ALL Consortium Protocol 00-01. Journal of Clinical Oncology, 2013, 31, 1202-1210.	0.8	274
36	Major Histocompatibility Complex Class II and Programmed Death Ligand 1 Expression Predict Outcome After Programmed Death 1 Blockade in Classic Hodgkin Lymphoma. Journal of Clinical Oncology, 2018, 36, 942-950.	0.8	273

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37	A dominant-negative effect drives selection of <i>TP53</i> missense mutations in myeloid malignancies. Science, 2019, 365, 599-604.	6.0	265
38	Topological analysis reveals a PD-L1-associated microenvironmental niche for Reed-Sternberg cells in Hodgkin lymphoma. Blood, 2017, 130, 2420-2430.	0.6	262
39	Chronic lymphocytic leukemia cells induce changes in gene expression of CD4 and CD8 T cells. Journal of Clinical Investigation, 2005, 115, 1797-1805.	3.9	259
40	Personal neoantigen vaccines induce persistent memory T cell responses and epitope spreading in patients with melanoma. Nature Medicine, 2021, 27, 515-525.	15.2	248
41	Phenotype, specificity and avidity of antitumour CD8+ T cells in melanoma. Nature, 2021, 596, 119-125.	13.7	239
42	MHC class I chain-related protein A antibodies and shedding are associated with the progression of multiple myeloma. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 1285-1290.	3.3	235
43	The Public Repository of Xenografts Enables Discovery and Randomized Phase II-like Trials in Mice. Cancer Cell, 2016, 29, 574-586.	7.7	227
44	Mitochondrial Reprogramming Underlies Resistance to BCL-2 Inhibition in Lymphoid Malignancies. Cancer Cell, 2019, 36, 369-384.e13.	7.7	224
45	Phase I Trial of Autologous CAR T Cells Targeting NKG2D Ligands in Patients with AML/MDS and Multiple Myeloma. Cancer Immunology Research, 2019, 7, 100-112.	1.6	220
46	The frequency and management of asparaginaseâ€related thrombosis in paediatric and adult patients with acute lymphoblastic leukaemia treated on Danaâ€Farber Cancer Institute consortium protocols. British Journal of Haematology, 2011, 152, 452-459.	1.2	216
47	TP53 mutation status divides myelodysplastic syndromes with complex karyotypes into distinct prognostic subgroups. Leukemia, 2019, 33, 1747-1758.	3.3	195
48	<i>SF3B1</i> -mutant MDS as a distinct disease subtype: a proposal from the International Working Group for the Prognosis of MDS. Blood, 2020, 136, 157-170.	0.6	195
49	Intravenous pegylated asparaginase versus intramuscular native Escherichia coli l-asparaginase in newly diagnosed childhood acute lymphoblastic leukaemia (DFCI 05-001): a randomised, open-label phase 3 trial. Lancet Oncology, The, 2015, 16, 1677-1690.	5.1	193
50	Outcome in Patients With Myelodysplastic Syndrome After Autologous Bone Marrow Transplantation for Non-Hodgkin's Lymphoma. Journal of Clinical Oncology, 1999, 17, 3128-3135.	0.8	180
51	Integrative Analysis Reveals an Outcome-Associated and Targetable Pattern of p53 and Cell Cycle Deregulation in Diffuse Large B Cell Lymphoma. Cancer Cell, 2012, 22, 359-372.	7.7	179
52	Biophysical determinants for cellular uptake of hydrocarbon-stapled peptide helices. Nature Chemical Biology, 2016, 12, 845-852.	3.9	178
53	Transcriptomic Characterization of SF3B1 Mutation Reveals Its Pleiotropic Effects in Chronic Lymphocytic Leukemia. Cancer Cell, 2016, 30, 750-763.	7.7	173
54	Mutations in epigenetic regulators including SETD2 are gained during relapse in paediatric acute lymphoblastic leukaemia. Nature Communications, 2014, 5, 3469.	5.8	171

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55	Marked hyperferritinemia does not predict for HLH in the adult population. Blood, 2015, 125, 1548-1552.	0.6	170
56	SYK Inhibition Modulates Distinct PI3K/AKT- Dependent Survival Pathways and Cholesterol Biosynthesis in Diffuse Large B Cell Lymphomas. Cancer Cell, 2013, 23, 826-838.	7.7	152
57	The evolutionary landscape of chronic lymphocytic leukemia treated with ibrutinib targeted therapy. Nature Communications, 2017, 8, 2185.	5.8	148
58	Classical Hodgkin Lymphoma with Reduced β2M/MHC Class I Expression Is Associated with Inferior Outcome Independent of 9p24.1 Status. Cancer Immunology Research, 2016, 4, 910-916.	1.6	146
59	T-Lymphoblastic Lymphoma Cells Express High Levels of BCL2, S1P1, and ICAM1, Leading to a Blockade of Tumor Cell Intravasation. Cancer Cell, 2010, 18, 353-366.	7.7	141
60	Sequence-Intrinsic Mechanisms that Target AID Mutational Outcomes on Antibody Genes. Cell, 2015, 163, 1124-1137.	13.5	136
61	Intracranial hemorrhage in patients with brain metastases treated with therapeutic enoxaparin: a matched cohort study. Blood, 2015, 126, 494-499.	0.6	128
62	Quantitative analysis of minimal residual disease predicts relapse in children with B-lineage acute lymphoblastic leukemia in DFCI ALL Consortium Protocol 95-01. Blood, 2007, 110, 1607-1611.	0.6	126
63	Pediatric-type nodal follicular lymphoma: a biologically distinct lymphoma with frequent MAPK pathway mutations. Blood, 2016, 128, 1093-1100.	0.6	126
64	Genetic Basis for PD-L1 Expression in Squamous Cell Carcinomas of the Cervix and Vulva. JAMA Oncology, 2016, 2, 518.	3.4	121
65	Mass cytometry of Hodgkin lymphoma reveals a CD4+ regulatory T-cell–rich and exhausted T-effector microenvironment. Blood, 2018, 132, 825-836.	0.6	121
66	Triplication of a 21q22 region contributes to B cell transformation through HMGN1 overexpression and loss of histone H3 Lys27 trimethylation. Nature Genetics, 2014, 46, 618-623.	9.4	117
67	Prediction and prevention of thromboembolic events with enoxaparin in cancer patients with elevated tissue factorâ€bearing microparticles: a randomizedâ€controlled phase II trial (the Microtec) Tj ETQq1	1 0 <b>1728</b> 431	4 rgBT /Overl
68	Cancer-Germline Antigen Expression Discriminates Clinical Outcome to CTLA-4 Blockade. Cell, 2018, 173, 624-633.e8.	13.5	113
69	Absence of Secondary Malignant Neoplasms in Children With High-Risk Acute Lymphoblastic Leukemia Treated With Dexrazoxane. Journal of Clinical Oncology, 2008, 26, 1106-1111.	0.8	111
70	Mechanisms of PD-L1/PD-1–mediated CD8 T-cell dysfunction in the context of aging-related immune defects in the Eµ-TCL1 CLL mouse model. Blood, 2015, 126, 212-221.	0.6	111
71	Targeting protein disulfide isomerase with the flavonoid isoquercetin to improve hypercoagulability in advanced cancer. JCl Insight, 2019, 4, .	2.3	110
72	Sensitive Detection of Minimal Residual Disease in Patients Treated for Early-Stage Breast Cancer. Clinical Cancer Research, 2020, 26, 2556-2564.	3.2	109

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73	Two familial ALS proteins function in prevention/repair of transcription-associated DNA damage. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E7701-E7709.	3.3	105
74	Pten mediates Myc oncogene dependence in a conditional zebrafish model of T cell acute lymphoblastic leukemia. Journal of Experimental Medicine, 2011, 208, 1595-1603.	4.2	104
75	High-level ROR1 associates with accelerated disease progression in chronic lymphocytic leukemia. Blood, 2016, 128, 2931-2940.	0.6	102
76	The low incidence of secondary acute myelogenous leukaemia in children and adolescents treated with dexrazoxane for acute lymphoblastic leukaemia: A report from the Dana-Farber Cancer Institute ALL Consortium. European Journal of Cancer, 2011, 47, 1373-1379.	1.3	99
77	Endogenous Glucocorticoid Signaling Regulates CD8+ T Cell Differentiation and Development of Dysfunction in the Tumor Microenvironment. Immunity, 2020, 53, 658-671.e6.	6.6	98
78	Clonal hematopoiesis is associated with adverse outcomes in multiple myeloma patients undergoing transplant. Nature Communications, 2020, 11, 2996.	5.8	98
79	Indications and Results of HLA-Identical Sibling Hematopoietic Cell Transplantation for Sickle Cell Disease. Biology of Blood and Marrow Transplantation, 2016, 22, 207-211.	2.0	97
80	Protein disulfide isomerase inhibition blocks thrombin generation in humans by interfering with platelet factor V activation. JCI Insight, 2017, 2, e89373.	2.3	96
81	Memory B cell repertoire for recognition of evolving SARS-CoV-2 spike. Cell, 2021, 184, 4969-4980.e15.	13.5	94
82	Characterization of T cell repertoire in patients with graft-versus-leukemia after donor lymphocyte infusion Journal of Clinical Investigation, 1997, 100, 855-866.	3.9	94
83	Absence of Biallelic <i>TCR</i> γ Deletion Predicts Early Treatment Failure in Pediatric T-Cell Acute Lymphoblastic Leukemia. Journal of Clinical Oncology, 2010, 28, 3816-3823.	0.8	93
84	A peripheral immune signature of responsiveness to PD-1 blockade in patients with classical Hodgkin lymphoma. Nature Medicine, 2020, 26, 1468-1479.	15.2	87
85	Growth dynamics in naturally progressing chronic lymphocytic leukaemia. Nature, 2019, 570, 474-479.	13.7	86
86	Somatic Mutations in MDS Patients Are Associated with Clinical Features and Predict Prognosis Independent of the IPSS-R: Analysis of Combined Datasets from the International Working Group for Prognosis in MDS-Molecular Committee. Blood, 2015, 126, 907-907.	0.6	85
87	Gene expression–based discovery of atovaquone as a STAT3 inhibitor and anticancer agent. Blood, 2016, 128, 1845-1853.	0.6	83
88	LMO1 Synergizes with MYCN to Promote Neuroblastoma Initiation and Metastasis. Cancer Cell, 2017, 32, 310-323.e5.	7.7	80
89	Predicting the higher rate of intracranial hemorrhage in glioma patients receiving therapeutic enoxaparin. Blood, 2017, 129, 3379-3385.	0.6	77
90	Integrative Genomic Analysis Implicates Gain of <i>PIK3CA</i> at 3q26 and <i>MYC</i> at 8q24 in Chronic Lymphocytic Leukemia. Clinical Cancer Research, 2012, 18, 3791-3802.	3.2	76

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91	Toxicity and efficacy of defined doses of CD4(+) donor lymphocytes for treatment of relapse after allogeneic bone marrow transplant. Blood, 1998, 91, 3671-80.	0.6	75
92	Refining risk classification in childhood B acute lymphoblastic leukemia: results of DFCI ALL Consortium Protocol 05-001. Blood Advances, 2018, 2, 1449-1458.	2.5	73
93	Pediatricâ€inspired therapy compared to allografting for <scp>P</scp> hiladelphia chromosomeâ€negative adult ALL in first complete remission. American Journal of Hematology, 2016, 91, 322-329.	2.0	72
94	A Murine Model of Chronic Lymphocytic Leukemia Based on B Cell-Restricted Expression of Sf3b1 Mutation and Atm Deletion. Cancer Cell, 2019, 35, 283-296.e5.	7.7	71
95	Synthetic Lethality of Wnt Pathway Activation and Asparaginase in Drug-Resistant Acute Leukemias. Cancer Cell, 2019, 35, 664-676.e7.	7.7	70
96	Landscape of helper and regulatory antitumour CD4+ T cells in melanoma. Nature, 2022, 605, 532-538.	13.7	70
97	Impact of Socioeconomic Status on Timing of Relapse and Overall Survival for Children Treated on Danaâ€Farber Cancer Institute ALL Consortium Protocols (2000–2010). Pediatric Blood and Cancer, 2016, 63, 1012-1018.	0.8	69
98	Integrated single-cell genetic and transcriptional analysis suggests novel drivers of chronic lymphocytic leukemia. Genome Research, 2017, 27, 1300-1311.	2.4	67
99	The Dohner fluorescence <i>inÂsitu</i> hybridization prognostic classification of chronic lymphocytic leukaemia ( <scp>CLL</scp> ): the <scp>CLL</scp> Research Consortium experience. British Journal of Haematology, 2016, 173, 105-113.	1.2	66
100	Somatic mutation as a mechanism of Wnt/ $\hat{I}^2$ -catenin pathway activation in CLL. Blood, 2014, 124, 1089-1098.	0.6	65
101	Notch signaling expands a pre-malignant pool of T-cell acute lymphoblastic leukemia clones without affecting leukemia-propagating cell frequency. Leukemia, 2012, 26, 2069-2078.	3.3	64
102	Trisomy 12 chronic lymphocytic leukemia cells exhibit upregulation of integrin signaling that is modulated by NOTCH1 mutations. Blood, 2014, 123, 4101-4110.	0.6	63
103	NRAS mutations with low allele burden have independent prognostic significance for patients with lower risk myelodysplastic syndromes. Leukemia, 2013, 27, 2077-2081.	3.3	57
104	Polymorphisms in Genes Related to Oxidative Stress Are Associated With Inferior Cognitive Function After Therapy for Childhood Acute Lymphoblastic Leukemia. Journal of Clinical Oncology, 2015, 33, 2205-2211.	0.8	57
105	Phase III Open-Label Randomized Study of Cytarabine in Combination With Amonafide L-Malate or Daunorubicin As Induction Therapy for Patients With Secondary Acute Myeloid Leukemia. Journal of Clinical Oncology, 2015, 33, 1252-1257.	0.8	57
106	Bone marrow transplantation for adolescents and young adults with sickle cell disease: Results of a prospective multicenter pilot study. American Journal of Hematology, 2019, 94, 446-454.	2.0	56
107	CDK6 Antagonizes p53-Induced Responses during Tumorigenesis. Cancer Discovery, 2018, 8, 884-897.	7.7	53
108	Polymorphisms of Asparaginase Pathway and Asparaginase-Related Complications in Children with Acute Lymphoblastic Leukemia. Clinical Cancer Research, 2015, 21, 329-334.	3.2	52

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109	Multifunctional barcoding with ClonMapper enables high-resolution study of clonal dynamics during tumor evolution and treatment. Nature Cancer, 2021, 2, 758-772.	5.7	52
110	Inhibition of calcineurin phosphatase activity in adult bone marrow transplant patients treated with cyclosporine A. Blood, 1994, 84, 3974-3979.	0.6	51
111	Emergence, Involution, and Progression to Carcinoma of Mutant Clones in Normal Endometrial Tissues. Cancer Research, 2014, 74, 2796-2802.	0.4	48
112	Pathological glycogenesis through glycogen synthase 1 and suppression of excessive AMP kinase activity in myeloid leukemia cells. Leukemia, 2015, 29, 1555-1563.	3.3	48
113	Hedgehog pathway mutations drive oncogenic transformation in high-risk T-cell acute lymphoblastic leukemia. Leukemia, 2018, 32, 2126-2137.	3.3	48
114	Phase I trial of the mTOR inhibitor everolimus in combination with multiâ€agent chemotherapy in relapsed childhood acute lymphoblastic leukemia. Pediatric Blood and Cancer, 2018, 65, e27062.	0.8	48
115	Pattern of Frequent But Nontargeted Pharmacologic Thromboprophylaxis for Hospitalized Patients With Cancer at Academic Medical Centers: A Prospective, Cross-Sectional, Multicenter Study. Journal of Clinical Oncology, 2014, 32, 1792-1796.	0.8	45
116	Targeted BMI1 inhibition impairs tumor growth in lung adenocarcinomas with low CEBPα expression. Science Translational Medicine, 2016, 8, 350ra104.	5.8	45
117	A cloning and expression system to probe T-cell receptor specificity and assess functional avidity to neoantigens. Blood, 2018, 132, 1911-1921.	0.6	44
118	Microbial symbionts regulate the primary Ig repertoire. Journal of Experimental Medicine, 2018, 215, 1397-1415.	4.2	43
119	Sequence intrinsic somatic mutation mechanisms contribute to affinity maturation of VRC01-class HIV-1 broadly neutralizing antibodies. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 8614-8619.	3.3	42
120	Immune recall improves antibody durability and breadth to SARS-CoV-2 variants. Science Immunology, 2022, 7, eabp8328.	5.6	40
121	Splicing modulation sensitizes chronic lymphocytic leukemia cells to venetoclax by remodeling mitochondrial apoptotic dependencies. JCI Insight, 2018, 3, .	2.3	39
122	Randomized phase 2 trial of regadenoson for treatment of acute vaso-occlusive crises in sickle cell disease. Blood Advances, 2017, 1, 1645-1649.	2.5	38
123	A Multicenter Phase II Study Using a Dose Intensified Pegylated-Asparaginase Pediatric Regimen in Adults with Untreated Acute Lymphoblastic Leukemia: A DFCI ALL Consortium Trial. Blood, 2015, 126, 80-80.	0.6	38
124	PRC2 loss induces chemoresistance by repressing apoptosis in T cell acute lymphoblastic leukemia. Journal of Experimental Medicine, 2018, 215, 3094-3114.	4.2	37
125	Mapping the evolution of TÂcell states during response and resistance to adoptive cellular therapy. Cell Reports, 2021, 37, 109992.	2.9	37
126	Obatoclax in combination with fludarabine and rituximab is well-tolerated and shows promising clinical activity in relapsed chronic lymphocytic leukemia. Leukemia and Lymphoma, 2015, 56, 3336-3342.	0.6	36

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127	Wnt5a induces ROR1 to recruit DOCK2 to activate Rac1/2 in chronic lymphocytic leukemia. Blood, 2018, 132, 170-178.	0.6	36
128	Efficacy and Toxicity of Pegaspargase and Calaspargase Pegol in Childhood Acute Lymphoblastic Leukemia: Results of DFCI 11-001. Journal of Clinical Oncology, 2021, 39, 3496-3505.	0.8	36
129	Biologic Activity of Autologous, Granulocyte–Macrophage Colony-Stimulating Factor Secreting Alveolar Soft-Part Sarcoma and Clear Cell Sarcoma Vaccines. Clinical Cancer Research, 2015, 21, 3178-3186.	3.2	34
130	A strategy to improve treatmentâ€related mortality and abandonment of therapy for childhood ALL in a developing country reveals the impact of treatment delays. Pediatric Blood and Cancer, 2015, 62, 1395-1402.	0.8	34
131	Impaired mitochondrial function is abrogated by dexrazoxane in doxorubicinâ€ŧreated childhood acute lymphoblastic leukemia survivors. Cancer, 2016, 122, 946-953.	2.0	34
132	Progression in patients with low- and intermediate-1-risk del(5q) myelodysplastic syndromes is predicted by a limited subset of mutations. Haematologica, 2017, 102, 498-508.	1.7	34
133	Prognostic impact of kinase-activating fusions and IKZF1 deletions in pediatric high-risk B-lineage acute lymphoblastic leukemia. Blood Advances, 2018, 2, 529-533.	2.5	34
134	Phase I study of the aurora A kinase inhibitor alisertib with induction chemotherapy in patients with acute myeloid leukemia. Haematologica, 2017, 102, 719-727.	1.7	33
135	Whole-exome sequencing identified genetic risk factors for asparaginase-related complications in childhood ALL patients. Oncotarget, 2017, 8, 43752-43767.	0.8	33
136	Safety Data from a First-in-Human Phase 1 Trial of NKG2D Chimeric Antigen Receptor-T Cells in AML/MDS and Multiple Myeloma. Blood, 2016, 128, 4052-4052.	0.6	32
137	Reproducibility and prognostic significance of morphologic dysplasia in de novo acute myeloid leukemia. Modern Pathology, 2015, 28, 965-976.	2.9	31
138	The use of prophylactic anticoagulation during induction and consolidation chemotherapy in adults with acute lymphoblastic leukemia. Journal of Thrombosis and Thrombolysis, 2018, 45, 306-314.	1.0	31
139	Effectiveness of antibacterial prophylaxis during induction chemotherapy in children with acute lymphoblastic leukemia. Pediatric Blood and Cancer, 2018, 65, e26952.	0.8	31
140	Longitudinal Single-Cell Dynamics of Chromatin Accessibility and Mitochondrial Mutations in Chronic Lymphocytic Leukemia Mirror Disease History. Cancer Discovery, 2021, 11, 3048-3063.	7.7	31
141	A Multicenter Phase I Study Combining Venetoclax with Mini-Hyper-CVD in Older Adults with Untreated and Relapsed/Refractory Acute Lymphoblastic Leukemia. Blood, 2019, 134, 3867-3867.	0.6	30
142	MYC Immunohistochemistry to Identify MYC-Driven B-Cell Lymphomas in Clinical Practice. American Journal of Clinical Pathology, 2016, 145, 166-179.	0.4	29
143	A hotspot mutation in transcription factor IKZF3 drives B cell neoplasia via transcriptional dysregulation. Cancer Cell, 2021, 39, 380-393.e8.	7.7	27
144	The clinical and functional effects of <i>TERT</i> variants in myelodysplastic syndrome. Blood, 2021, 138, 898-911.	0.6	27

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145	Acute myeloid leukemia cells require 6-phosphogluconate dehydrogenase for cell growth and NADPH-dependent metabolic reprogramming. Oncotarget, 2017, 8, 67639-67650.	0.8	26
146	Outcome of children and adolescents with Down syndrome treated on Danaâ€Farber Cancer Institute Acute Lymphoblastic Leukemia Consortium protocols 00–001 and 05â€001. Pediatric Blood and Cancer, 2018, 65, e27256.	0.8	26
147	Identification of prognostic factors in childhood Tâ€cell acute lymphoblastic leukemia: Results from DFCI ALL Consortium Protocols 05â€001 and 11â€001. Pediatric Blood and Cancer, 2021, 68, e28719.	0.8	26
148	Genomic profiling of a randomized trial of interferon-α vs hydroxyurea in MPN reveals mutation-specific responses. Blood Advances, 2022, 6, 2107-2119.	2.5	26
149	Molecular Classification of MYC-Driven B-Cell Lymphomas by Targeted Gene Expression Profiling of Fixed Biopsy Specimens. Journal of Molecular Diagnostics, 2015, 17, 19-30.	1.2	25
150	Short telomere length predicts nonrelapse mortality after stem cell transplantation for myelodysplastic syndrome. Blood, 2020, 136, 3070-3081.	0.6	25
151	Biased estimation of thrombosis rates in cancer studies using the method of Kaplan and Meier. Journal of Thrombosis and Haemostasis, 2012, 10, 1449-1451.	1.9	24
152	A phase II study of the EGFR inhibitor gefitinib in patients with acute myeloid leukemia. Leukemia Research, 2014, 38, 430-434.	0.4	23
153	Cabozantinib is well tolerated in acute myeloid leukemia and effectively inhibits the resistanceâ€conferring FLT3/tyrosine kinase domain/F691 mutation. Cancer, 2018, 124, 306-314.	2.0	23
154	An investigation of toxicities and survival in Hispanic children and adolescents with ALL: Results from the Danaâ€Farber Cancer Institute ALL Consortium protocol 05â€001. Pediatric Blood and Cancer, 2018, 65, e26871.	0.8	23
155	CXCR4 upregulation is an indicator of sensitivity to B-cell receptor/PI3K blockade and a potential resistance mechanism in B-cell receptor-dependent diffuse large B-cell lymphomas. Haematologica, 2020, 105, 1361-1368.	1.7	23
156	Amino acid–insensitive mTORC1 regulation enables nutritional stress resilience in hematopoietic stem cells. Journal of Clinical Investigation, 2017, 127, 1405-1413.	3.9	23
157	JDP2: An oncogenic bZIP transcription factor in T cell acute lymphoblastic leukemia. Journal of Experimental Medicine, 2018, 215, 1929-1945.	4.2	22
158	Dose-adjusted enoxaparin thromboprophylaxis in hospitalized cancer patients: a randomized, double-blinded multicenter phase 2 trial. Blood Advances, 2020, 4, 2254-2260.	2.5	22
159	Clonal Evolution In Patients With Chronic Lymphocytic Leukemia (CLL) Developing Resistance To BTK Inhibition. Blood, 2013, 122, 866-866.	0.6	22
160	PPARÎ <sup>3</sup> Contributes to Immunity Induced by Cancer Cell Vaccines That Secrete GM-CSF. Cancer Immunology Research, 2018, 6, 723-732.	1.6	21
161	Automated Flow Synthesis of Tumor Neoantigen Peptides for Personalized Immunotherapy. Scientific Reports, 2020, 10, 723.	1.6	21
162	Phase 1/2 trial of vorinostat in patients with sickle cell disease who have not benefitted from hydroxyurea. Blood, 2015, 125, 3668-3669.	0.6	20

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