Gareth J Morgan

List of Publications by Year
in descending order

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595	36,252	3930 88	4012 176
papers	citations	h-index	g-index

607	607	607	25833
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Design and standardization of PCR primers and protocols for detection of clonal immunoglobulin and T-cell receptor gene recombinations in suspect lymphoproliferations: Report of the BIOMED-2 Concerted Action BMH4-CT98-3936. Leukemia, 2003, 17, 2257-2317.	3.3	2,788
2	International Staging System for Multiple Myeloma. Journal of Clinical Oncology, 2005, 23, 3412-3420.	0.8	2,404
3	High-Dose Chemotherapy with Hematopoietic Stem-Cell Rescue for Multiple Myeloma. New England Journal of Medicine, 2003, 348, 1875-1883.	13.9	1,648
4	Revised International Staging System for Multiple Myeloma: A Report From International Myeloma Working Group. Journal of Clinical Oncology, 2015, 33, 2863-2869.	0.8	1,525
5	Thalidomide and immunomodulatory derivatives augment natural killer cell cytotoxicity in multiple myeloma. Blood, 2001, 98, 210-216.	0.6	869
6	Prevention of thalidomide- and lenalidomide-associated thrombosis in myeloma. Leukemia, 2008, 22, 414-423.	3.3	787
7	The genetic architecture of multiple myeloma. Nature Reviews Cancer, 2012, 12, 335-348.	12.8	741
8	Risk of progression and survival in multiple myeloma relapsing after therapy with IMiDs and bortezomib: A multicenter international myeloma working group study. Leukemia, 2012, 26, 149-157.	3.3	664
9	First-line treatment with zoledronic acid as compared with clodronic acid in multiple myeloma (MRC) Tj ETQq1 1 0.784314 rgBT /Ove	6.3	505
10	Mutational Spectrum, Copy Number Changes, and Outcome: Results of a Sequencing Study of Patients With Newly Diagnosed Myeloma. Journal of Clinical Oncology, 2015, 33, 3911-3920.	0.8	463
11	Early Mortality After Diagnosis of Multiple Myeloma: Analysis of Patients Entered Onto the United Kingdom Medical Research Council Trials Between 1980 and 2002â€"Medical Research Council Adult Leukaemia Working Party. Journal of Clinical Oncology, 2005, 23, 9219-9226.	0.8	402
12	Myeloma management guidelines: a consensus report from the Scientific Advisors of the International Myeloma Foundation. The Hematology Journal, 2003, 4, 379-398.	2.0	374
13	Minimal Residual Disease Assessed by Multiparameter Flow Cytometry in Multiple Myeloma: Impact on Outcome in the Medical Research Council Myeloma IX Study. Journal of Clinical Oncology, 2013, 31, 2540-2547.	0.8	372
14	Genetic variation in TNF and IL10 and risk of non-Hodgkin lymphoma: a report from the InterLymph Consortium. Lancet Oncology, The, 2006, 7, 27-38.	5.1	345
15	Identification of novel mutational drivers reveals oncogene dependencies in multiple myeloma. Blood, 2018, 132, 587-597.	0.6	335
16	International Myeloma Working Group Consensus Statement for the Management, Treatment, and Supportive Care of Patients With Myeloma Not Eligible for Standard Autologous Stem-Cell Transplantation. Journal of Clinical Oncology, 2014, 32, 587-600.	0.8	330
17	A compendium of myeloma-associated chromosomal copy number abnormalities and their prognostic value. Blood, 2010, 116, e56-e65.	0.6	315
18	The role of maintenance thalidomide therapy in multiple myeloma: MRC Myeloma IX results and meta-analysis. Blood, 2012, 119, 7-15.	0.6	315

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19	A high-risk, Double-Hit, group of newly diagnosed myeloma identified by genomic analysis. Leukemia, 2019, 33, 159-170.	3.3	313
20	Personalized therapy in multiple myeloma according to patient age and vulnerability: a report of the European Myeloma Network (EMN). Blood, 2011, 118, 4519-4529.	0.6	309
21	Monoclonal B lymphocytes with the characteristics of â€œindolentâ€•chronic lymphocytic leukemia are present in 3.5% of adults with normal blood counts. Blood, 2002, 100, 635-639.	0.6	305
22	A novel prognostic model in myeloma based on co-segregating adverse FISH lesions and the ISS: analysis of patients treated in the MRC Myeloma IX trial. Leukemia, 2012, 26, 349-355.	3.3	298
23	The Requirement for DNAM-1, NKG2D, and NKp46 in the Natural Killer Cell-Mediated Killing of Myeloma Cells. Cancer Research, 2007, 67, 8444-8449.	0.4	284
24	Antimyeloma activity of heat shock protein-90 inhibition. Blood, 2005, 107, 1092-1100.	0.6	278
25	Spatial genomic heterogeneity in multiple myeloma revealed by multi-region sequencing. Nature Communications, 2017, 8, 268.	5.8	277
26	APOBEC family mutational signatures are associated with poor prognosis translocations in multiple myeloma. Nature Communications, 2015, 6, 6997.	5.8	261
27	Germinal center phenotype and bcl-2 expression combined with the International Prognostic Index improves patient risk stratification in diffuse large B-cell lymphoma. Blood, 2002, 99, 1136-1143.	0.6	252
28	Guidelines for the diagnosis and management of multiple myeloma 2011. British Journal of Haematology, 2011, 154, 32-75.	1.2	252
29	Intraclonal heterogeneity is a critical early event in the development of myeloma and precedes the development of clinical symptoms. Leukemia, 2014, 28, 384-390.	3.3	252
30	Quantitation of minimal disease levels in chronic lymphocytic leukemia using a sensitive flow cytometric assay improves the prediction of outcome and can be used to optimize therapy. Blood, 2001, 98, 29-35.	0.6	249
31	Lenalidomide maintenance versus observation for patients with newly diagnosed multiple myeloma (Myeloma XI): a multicentre, open-label, randomised, phase 3 trial. Lancet Oncology, The, 2019, 20, 57-73.	5.1	245
32	Polymorphism in glutathione S-transferase P1 is associated with susceptibility to chemotherapy-induced leukemia. Proceedings of the National Academy of Sciences of the United States of America, 2001, 98, 11592-11597.	3.3	233
33	Intraclonal heterogeneity and distinct molecular mechanisms characterize the development of t(4;14) and t(11;14) myeloma. Blood, 2012, 120, 1077-1086.	0.6	231
34	Heat shock protein inhibition is associated with activation of the unfolded protein response pathway in myeloma plasma cells. Blood, 2007, 110, 2641-2649.	0.6	219
35	Aberrant global methylation patterns affect the molecular pathogenesis and prognosis of multiple myeloma. Blood, 2011, 117, 553-562.	0.6	217
36	Insights into the multistep transformation of MGUS to myeloma using microarray expression analysis. Blood, 2003, 102, 4504-4511.	0.6	212

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37	Polymorphisms in the thymidylate synthase and serine hydroxymethyltransferase genes and risk of adult acute lymphocytic leukemia. Blood, 2002, 99, 3786-3791.	0.6	210
38	Single-cell genetic analysis reveals the composition of initiating clones and phylogenetic patterns of branching and parallel evolution in myeloma. Leukemia, 2014, 28, 1705-1715.	3.3	207
39	Structure of the Ire1 autophosphorylation complex and implications for the unfolded protein response. EMBO Journal, 2011, 30, 894-905.	3.5	201
40	Bortezomib (Velcade?) in the treatment of multiple myeloma. Therapeutics and Clinical Risk Management, 2006, 2, 271-279.	0.9	197
41	Preclinical evaluation of the proteasome inhibitor bortezomib in cancer therapy. Cancer Cell International, 2005, 5, 18.	1.8	196
42	Flow cytometric disease monitoring in multiple myeloma: the relationship between normal and neoplastic plasma cells predicts outcome after transplantation. Blood, 2002, 100, 3095-3100.	0.6	194
43	Curing myeloma at last: defining criteria and providing the evidence. Blood, 2014, 124, 3043-3051.	0.6	194
44	Oral ixazomib maintenance following autologous stem cell transplantation (TOURMALINE-MM3): a double-blind, randomised, placebo-controlled phase 3 trial. Lancet, The, 2019, 393, 253-264.	6.3	187
45	Cyclophosphamide, thalidomide, and dexamethasone (CTD) as initial therapy for patients with multiple myeloma unsuitable for autologous transplantation. Blood, 2011, 118, 1231-1238.	0.6	179
46	Evolutionary biology of high-risk multiple myeloma. Nature Reviews Cancer, 2017, 17, 543-556.	12.8	178
47	Prediction of outcome in newly diagnosed myeloma: a meta-analysis of the molecular profiles of 1905 trial patients. Leukemia, 2018, 32, 102-110.	3.3	177
48	Integration of global SNP-based mapping and expression arrays reveals key regions, mechanisms, and genes important in the pathogenesis of multiple myeloma. Blood, 2006, 108, 1733-1743.	0.6	176
49	Clonal selection and double-hit events involving tumor suppressor genes underlie relapse in myeloma. Blood, 2016, 128, 1735-1744.	0.6	170
50	Circulating plasma cells in multiple myeloma: characterization and correlation with disease stage. British Journal of Haematology, 1997, 97, 46-55.	1.2	165
51	Immunoglobulin gene rearrangements and the pathogenesis of multiple myeloma. Blood, 2007, 110, 3112-3121.	0.6	157
52	Effects of zoledronic acid versus clodronic acid on skeletal morbidity in patients with newly diagnosed multiple myeloma (MRC Myeloma IX): secondary outcomes from a randomised controlled trial. Lancet Oncology, The, 2011, 12, 743-752.	5.1	151
53	Mapping of Chromosome 1p Deletions in Myeloma Identifies FAM46C at 1p12 and CDKN2C at 1p32.3 as Being Genes in Regions Associated with Adverse Survival. Clinical Cancer Research, 2011, 17, 7776-7784.	3.2	147
54	Global methylation analysis identifies prognostically important epigenetically inactivated tumor suppressor genes in multiple myeloma. Blood, 2013, 122, 219-226.	0.6	147

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55	Genome-wide association study identifies multiple susceptibility loci for multiple myeloma. Nature Communications, 2016, 7, 12050.	5.8	146
56	Cyclophosphamide, thalidomide, and dexamethasone as induction therapy for newly diagnosed multiple myeloma patients destined for autologous stem-cell transplantation: MRC Myeloma IX randomized trial results. Haematologica, 2012, 97, 442-450.	1.7	144
57	Safety and efficacy of pomalidomide plus low-dose dexamethasone in STRATUS (MM-010): a phase 3b study in refractory multiple myeloma. Blood, 2016, 128, 497-503.	0.6	144
58	Common variation at 3q26.2, 6p21.33, 17p11.2 and 22q13.1 influences multiple myeloma risk. Nature Genetics, 2013, 45, 1221-1225.	9.4	143
59	Long-term Follow-up of MRC Myeloma IX Trial: Survival Outcomes with Bisphosphonate and Thalidomide Treatment. Clinical Cancer Research, 2013, 19, 6030-6038.	3.2	143
60	Translocations at 8q24 juxtapose MYC with genes that harbor superenhancers resulting in overexpression and poor prognosis in myeloma patients. Blood Cancer Journal, 2014, 4, e191-e191.	2.8	142
61	Deletion of chromosome 13 detected by conventional cytogenetics is a critical prognostic factor in myeloma. Leukemia, 2006, 20, 1610-1617.	3.3	141
62	Common variation at 3p22.1 and 7p15.3 influences multiple myeloma risk. Nature Genetics, 2012, 44, 58-61.	9.4	137
63	Gene mapping and expression analysis of 16q loss of heterozygosity identifies WWOX and CYLD as being important in determining clinical outcome in multiple myeloma. Blood, 2007, 110, 3291-3300.	0.6	133
64	Essential Role of Caveolae in Interleukin-6- and Insulin-like Growth Factor I-triggered Akt-1-mediated Survival of Multiple Myeloma Cells. Journal of Biological Chemistry, 2003, 278, 5794-5801.	1.6	128
65	Characterization of IGH locus breakpoints in multiple myeloma indicates a subset of translocations appear to occur in pregerminal center B cells. Blood, 2013, 121, 3413-3419.	0.6	128
66	Low NAD(P)H:quinone oxidoreductase 1 activity is associated with increased risk of acute leukemia in adults. Blood, 2001, 97, 1422-1426.	0.6	125
67	Homozygous Deletion Mapping in Myeloma Samples Identifies Genes and an Expression Signature Relevant to Pathogenesis and Outcome. Clinical Cancer Research, 2010, 16, 1856-1864.	3.2	124
68	The clinical relevance and management of monoclonal gammopathy of undetermined significance and related disorders: recommendations from the European Myeloma Network. Haematologica, 2014, 99, 984-996.	1.7	124
69	Rearrangement of the BCL6 locus at 3q27 is an independent poor prognostic factor in nodal diffuse large B-cell lymphoma. British Journal of Haematology, 2002, 117, 322-332.	1.2	113
70	Combination of flow cytometry and functional imaging for monitoring of residual disease in myeloma. Leukemia, 2019, 33, 1713-1722.	3.3	112
71	Potent and Selective KDM5 Inhibitor Stops Cellular Demethylation of H3K4me3 at Transcription Start Sites and Proliferation of MM1S Myeloma Cells. Cell Chemical Biology, 2017, 24, 371-380.	2.5	111
72	Percutaneous Device Closure of Paravalvular Leak. Circulation, 2016, 134, 934-944.	1.6	109

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73	Genetic variation in XPD predicts treatment outcome and risk of acute myeloid leukemia following chemotherapy. Blood, 2004, 104, 3872-3877.	0.6	108
74	Expert panel consensus statement on the optimal use of pomalidomide in relapsed and refractory multiple myeloma. Leukemia, 2014, 28, 1573-1585.	3.3	108
75	Trends in autologous hematopoietic cell transplantation for multiple myeloma in Europe: increased use and improved outcomes in elderly patients in recent years. Bone Marrow Transplantation, 2015, 50, 209-215.	1.3	108
76	Results of the MRC pilot study show autografting for younger patients with chronic lymphocytic leukemia is safe and achieves a high percentage of molecular responses. Blood, 2005, 105, 397-404.	0.6	107
77	XBP1s levels are implicated in the biology and outcome of myeloma mediating different clinical outcomes to thalidomide-based treatments. Blood, 2010, 116, 250-253.	0.6	107
78	MMSET deregulation affects cell cycle progression and adhesion regulons in t(4;14) myeloma plasma cells. Haematologica, 2009, 94, 78-86.	1.7	106
79	Prediction of high- and low-risk multiple myeloma based on gene expression and the International Staging System. Blood, 2015, 126, 1996-2004.	0.6	106
80	The impact of attaining a minimal disease state after high-dose melphalan and autologous transplantation for multiple myeloma. British Journal of Haematology, 2001, 112, 814-819.	1.2	103
81	Cancer-Selective Targeting of the NF-Î²B Survival Pathway with GADD45Î²/MKK7 Inhibitors. Cancer Cell, 2014, 26, 495-508.	7.7	99
82	The impact of extramedullary disease at presentation on the outcome of myeloma. Leukemia and Lymphoma, 2009, 50, 230-235.	0.6	97
83	The sialyltransferase ST3GAL6 influences homing and survival in multiple myeloma. Blood, 2014, 124, 1765-1776.	0.6	97
84	Assessment of Total Lesion Glycolysis by 18F FDG PET/CT Significantly Improves Prognostic Value of GEP and ISS in Myeloma. Clinical Cancer Research, 2017, 23, 1981-1987.	3.2	97
85	Lenalidomide (Revlimid), in combination with cyclophosphamide and dexamethasone (RCD), is an effective and tolerated regimen for myeloma patients. British Journal of Haematology, 2007, 137, 268-269.	1.2	96
86	Subclonal evolution in disease progression from MGUS/SMM to multiple myeloma is characterised by clonal stability. Leukemia, 2019, 33, 457-468.	3.3	96
87	Genetic Factors Underlying the Risk of Thalidomide-Related Neuropathy in Patients With Multiple Myeloma. Journal of Clinical Oncology, 2011, 29, 797-804.	0.8	95
88	High-Producer Haplotypes of Tumor Necrosis Factor Alpha and Lymphotoxin Alpha Are Associated With an Increased Risk of Myeloma and Have an Improved Progression-Free Survival After Treatment. Journal of Clinical Oncology, 2000, 18, 2843-2851.	0.8	91
89	The CCND1 c.870G>A polymorphism is a risk factor for t(11;14)(q13;q32) multiple myeloma. Nature Genetics, 2013, 45, 522-525.	9.4	91
90	The impact of intra-clonal heterogeneity on the treatment of multiple myeloma. British Journal of Haematology, 2014, 165, 441-454.	1.2	91

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91	Polymorphic variation in GSTP1 modulates outcome following therapy for multiple myeloma. Blood, 2003, 102, 2345-2350.	0.6	90
92	European Perspective on Multiple Myeloma Treatment Strategies in 2014. Oncologist, 2014, 19, 829-844.	1.9	90
93	Carfilzomib resistance due to ABCB1/MDR1 overexpression is overcome by nelfinavir and lopinavir in multiple myeloma. Leukemia, 2018, 32, 391-401.	3.3	89
94	Deletions of CDKN2C in Multiple Myeloma: Biological and Clinical Implications. Clinical Cancer Research, 2008, 14, 6033-6041.	3.2	88
95	Identification of multiple risk loci and regulatory mechanisms influencing susceptibility to multiple myeloma. Nature Communications, 2018, 9, 3707.	5.8	86
96	Myeloma management guidelines: a consensus report from the Scientific Advisors of the International Myeloma Foundation. The Hematology Journal, 2003, 4, 379-98.	2.0	86
97	Current Multiple Myeloma Treatment Strategies with Novel Agents: A European Perspective. Oncologist, 2010, 15, 6-25.	1.9	85
98	Polymorphic variation within the glutathione S-transferase genes and risk of adult acute leukaemia. Carcinogenesis, 2000, 21, 43-47.	1.3	84
99	Differentiation stage of myeloma plasma cells: biological and clinical significance. Leukemia, 2017, 31, 382-392.	3.3	83
100	The Spectrum and Clinical Impact of Epigenetic Modifier Mutations in Myeloma. Clinical Cancer Research, 2016, 22, 5783-5794.	3.2	81
101	Overexpression of EZH2 in multiple myeloma is associated with poor prognosis and dysregulation of cell cycle control. Blood Cancer Journal, 2017, 7, e549-e549.	2.8	81
102	Revealing the Impact of Structural Variants in Multiple Myeloma. Blood Cancer Discovery, 2020, 1, 258-273.	2.6	81
103	Gastric marginal zone lymphoma is associated with polymorphisms in genes involved in inflammatory response and antioxidative capacity. Blood, 2003, 102, 1007-1011.	0.6	79
104	Non-Hodgkin's lymphoma, obesity and energy homeostasis polymorphisms. British Journal of Cancer, 2005, 93, 811-816.	2.9	79
105	The interleukin-6 receptor alpha-chain (CD126) is expressed by neoplastic but not normal plasma cells. Blood, 2000, 96, 3880-3886.	0.6	78
106	Genetic variants of NHEJ DNA ligase IV can affect the risk of developing multiple myeloma, a tumour characterised by aberrant class switch recombination. Journal of Medical Genetics, 2002, 39, 900-905.	1.5	77
107	Untangling the unfolded protein response. Cell Cycle, 2008, 7, 865-869.	1.3	76
108	Genetic abnormalities during transition from Helicobacter-pylori-associated gastritis to low-grade MALToma. Lancet, The, 1995, 345, 26-27.	6.3	75

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109	Bâ€lymphocyte suppression in multiple myeloma is a reversible phenomenon specific to normal Bâ€cell progenitors and plasma cell precursors. British Journal of Haematology, 1998, 100, 176-183.	1.2	74
110	Age has a profound effect on the incidence and significance of chromosome abnormalities in myeloma. Leukemia, 2005, 19, 1634-1642.	3.3	73
111	Removing batch effects from purified plasma cell gene expression microarrays with modified ComBat. BMC Bioinformatics, 2015, 16, 63.	1.2	73
112	Myeloma aetiology and epidemiology. Biomedicine and Pharmacotherapy, 2002, 56, 223-234.	2.5	72
113	Risk of Non-Hodgkin Lymphoma Associated with Polymorphisms in Folate-Metabolizing Genes. Cancer Epidemiology Biomarkers and Prevention, 2005, 14, 2999-3003.	1.1	72
114	The combination of cyclophosphamide, velcade and dexamethasone (CVD) induces high response rates with comparable toxicity to velcade alone (V) and velcade plus dexamethasone (VD). Haematologica, 2007, 92, 1149-1150.	1.7	71
115	Targeting heat shock protein 72 enhances Hsp90 inhibitor-induced apoptosis in myeloma. Leukemia, 2010, 24, 1804-1807.	3.3	71
116	The spectrum of somatic mutations in monoclonal gammopathy of undetermined significance indicates a less complex genomic landscape than that in multiple myeloma. Haematologica, 2017, 102, 1617-1625.	1.7	71
117	A clinical prediction model for outcome and therapy delivery in transplant-ineligible patients with myeloma (UK Myeloma Research Alliance Risk Profile): a development and validation study. Lancet Haematology,the, 2019, 6, e154-e166.	2.2	71
118	Serum free immunoglobulin light chain evaluation as a marker of impact from intraclonal heterogeneity on myeloma outcome. Blood, 2014, 123, 3414-3419.	0.6	68
119	Second malignancies in the context of lenalidomide treatment: an analysis of 2732 myeloma patients enrolled to the Myeloma XI trial. Blood Cancer Journal, 2016, 6, e506-e506.	2.8	68
120	Whole-genome sequencing reveals progressive versus stable myeloma precursor conditions as two distinct entities. Nature Communications, 2021, 12, 1861.	5.8	68
121	Poor metabolizers at the cytochrome P450 2D6 and 2C19 loci are at increased risk of developing adult acute leukaemia. Pharmacogenetics and Genomics, 2000, 10, 605-615.	5.7	67
122	Maintenance Treatment and Survival in Patients With Myeloma. JAMA Oncology, 2018, 4, 1389.	3.4	67
123	Clonal evolution in myeloma: the impact of maintenance lenalidomide and depth of response on the genetics and sub-clonal structure of relapsed disease in uniformly treated newly diagnosed patients. Haematologica, 2019, 104, 1440-1450.	1.7	67
124	Aetiology of bone disease and the role of bisphosphonates in multiple myeloma. Lancet Oncology, The, 2003, 4, 284-292.	5.1	66
125	Treatment of relapsed and refractory multiple myeloma in the era of novel agents. Cancer Treatment Reviews, 2011, 37, 266-283.	3.4	66
126	Long-term outcomes after autologous stem cell transplantation for multiple myeloma. Blood Advances, 2020, 4, 422-431.	2.5	66

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127	Non-Hodgkin Lymphoma Secondary to Cancer Chemotherapy: Table 1.. Cancer Epidemiology Biomarkers and Prevention, 2007, 16, 377-380.	1.1	65
128	Genetic associations with thalidomide mediated venous thrombotic events in myeloma identified using targeted genotyping. Blood, 2008, 112, 4924-4934.	0.6	65
129	The role of second autografts in the management of myeloma at first relapse. Haematologica, 2006, 91, 141-2.	1.7	62
130	Inherited genetic susceptibility to multiple myeloma. Leukemia, 2014, 28, 518-524.	3.3	60
131	Karyotype and age in acute myeloid leukemia.. Cancer Genetics and Cytogenetics, 2001, 126, 155-161.	1.0	59
132	Factors Influencing the Outcome of a Second Autologous Stem Cell Transplant (ASCT) in Relapsed Multiple Myeloma: A Study from the British Society ofÂBlood and Marrow Transplantation Registry. Biology of Blood and Marrow Transplantation, 2011, 17, 1638-1645.	2.0	59
133	The clinical impact and molecular biology of del(17p) in multiple myeloma treated with conventional or thalidomideâ€based therapy. Genes Chromosomes and Cancer, 2011, 50, 765-774.	1.5	59
134	Epigenetic consequences of AML1-ETO action at the human c-FMS locus. EMBO Journal, 2003, 22, 2798-2809.	3.5	58
135	Clinical value of molecular subtyping multiple myeloma using gene expression profiling. Leukemia, 2016, 30, 423-430.	3.3	58
136	Thrombosis in patients with myeloma treated in the Myeloma IX and Myeloma XI phase 3 randomized controlled trials. Blood, 2020, 136, 1091-1104.	0.6	58
137	Antitumor Effects and Anticancer Applications of Bisphosphonates. Seminars in Oncology, 2010, 37, S30-S40.	0.8	57
138	The addition of cyclophosphamide to lenalidomide and dexamethasone in multiply relapsed/refractory myeloma patients; a phase I/II study. British Journal of Haematology, 2010, 150, 326-333.	1.2	57
139	Lenalidomide-induced diarrhea in patients with myeloma is caused by bile acid malabsorption that responds to treatment. Blood, 2014, 124, 2467-2468.	0.6	57
140	MAF protein mediates innate resistance to proteasome inhibition therapy in multiple myeloma. Blood, 2016, 128, 2919-2930.	0.6	57
141	The level of deletion 17p and bi-allelic inactivation of TP53 has a significant impact on clinical outcome in multiple myeloma. Haematologica, 2017, 102, e364-e367.	1.7	57
142	Tobacco and Alcohol Consumption and the Risk of Non-Hodgkin Lymphoma. Cancer Causes and Control, 2004, 15, 771-780.	0.8	55
143	Assessing myeloma bone disease with whole-body diffusion-weighted imaging: comparison with x-ray skeletal survey by regionÂand relationship with laboratory estimatesÂof disease burden. Clinical Radiology, 2015, 70, 614-621.	0.5	54
144	The molecular make up of smoldering myeloma highlights the evolutionary pathways leading to multiple myeloma. Nature Communications, 2021, 12, 293.	5.8	54

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145	Mutations of theAML1 gene in acute myeloid leukemia of FAB types M0 and M7. Genes Chromosomes and Cancer, 2002, 34, 24-32.	1.5	53
146	Cleavage of BLOC1S1 mRNA by IRE1 Is Sequence Specific, Temporally Separate from XBP1 Splicing, and Dispensable for Cell Viability under Acute Endoplasmic Reticulum Stress. Molecular and Cellular Biology, 2015, 35, 2186-2202.	1.1	53
147	A Global Expression-based Analysis of the Consequences of the t(4;14) Translocation in Myeloma. Clinical Cancer Research, 2004, 10, 5692-5701.	3.2	51
148	t(3;14)(p14;q32) Results in aberrant expression ofFOXP1 in a case of diffuse large B-cell lymphoma. Genes Chromosomes and Cancer, 2006, 45, 164-168.	1.5	51
149	Efficacy and outcome of autologous transplantation in rare myelomas. Haematologica, 2010, 95, 2126-2133.	1.7	51
150	Characterisation of immunoparesis in newly diagnosed myeloma and its impact on progression-free and overall survival in both old and recent myeloma trials. Leukemia, 2018, 32, 1727-1738.	3.3	50
151	Combinations of ZAP-70, CD38 and IGHV mutational status as predictors of time to first treatment in CLL. Leukemia and Lymphoma, 2008, 49, 2108-2115.	0.6	48
152	Genetic factors underlying the risk of bortezomib induced peripheral neuropathy in multiple myeloma patients. Haematologica, 2011, 96, 1728-1732.	1.7	48
153	Bi-allelic inactivation is more prevalent at relapse in multiple myeloma, identifying RB1 as an independent prognostic marker. Blood Cancer Journal, 2017, 7, e535-e535.	2.8	48
154	Treatment to suppression of focal lesions on positron emission tomography-computed tomography is a therapeutic goal in newly diagnosed multiple myeloma. Haematologica, 2018, 103, 1047-1053.	1.7	47
155	Immunologic approaches for the treatment of multiple myeloma. Cancer Treatment Reviews, 2017, 55, 190-199.	3.4	46
156	HSF1 Is Essential for Myeloma Cell Survival and A Promising Therapeutic Target. Clinical Cancer Research, 2018, 24, 2395-2407.	3.2	46
157	COVID-19 Infections and Clinical Outcomes in Patients with Multiple Myeloma in New York City: A Cohort Study from Five Academic Centers. Blood Cancer Discovery, 2020, 1, 234-243.	2.6	46
158	Residual disease detection using fluorescent polymerase chain reaction at 20 weeks of therapy predicts clinical outcome in childhood acute lymphoblastic leukemia.. Journal of Clinical Oncology, 1998, 16, 3616-3627.	0.8	45
159	Long-term outcomes of previously untreated myeloma patients: responses to induction chemotherapy and high-dose melphalan incorporated within a risk stratification model can help to direct the use of novel treatments. British Journal of Haematology, 2005, 129, 607-614.	1.2	45
160	Genomic variation in myeloma: design, content, and initial application of the Bank On A Cure SNP Panel to detect associations with progression-free survival. BMC Medicine, 2008, 6, 26.	2.3	45
161	Assessment of IgH PCR strategies in multiple myeloma.. Journal of Clinical Pathology, 1996, 49, 672-675.	1.0	44
162	Minimal residual disease monitoring in multiple myeloma. Best Practice and Research in Clinical Haematology, 2002, 15, 197-222.	0.7	44

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163	High-dose cyclophosphamide with stem cell rescue for severe rheumatoid arthritis: Short-term efficacy correlates with reduction of macroscopic and histologic synovitis. Arthritis and Rheumatism, 2002, 46, 837-839.	6.7	44
164	Aminopeptidase inhibition as a targeted treatment strategy in myeloma. Molecular Cancer Therapeutics, 2009, 8, 762-770.	1.9	44
165	Four genes predict high risk of progression from smoldering to symptomatic multiple myeloma (SWOG S0120). Haematologica, 2015, 100, 1214-1221.	1.7	44
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