Gareth J Morgan

List of Publications by Year
in descending order

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595 papers	36,252 citations	3919 88 h-index	3997 176 g-index
607 all docs	607 docs citations	607 times ranked	25833 citing authors

#	Article	IF	Citations
1	Chromothripsis as a pathogenic driver of multiple myeloma. Seminars in Cell and Developmental Biology, 2022, 123, 115-123.	2.3	22
2	Plasma cells expression from smouldering myeloma to myeloma reveals the importance of the PRC2 complex, cell cycle progression, and the divergent evolutionary pathways within the different molecular subgroups. Leukemia, 2022, 36, 591-595.	3.3	6
3	Inflammation and infection in plasma cell disorders: how pathogens shape the fate of patients. Leukemia, 2022, 36, 613-624.	3.3	11
4	Minimal Residual Disease After Autologous Stem-Cell Transplant for Patients With Myeloma: Prognostic Significance and the Impact of Lenalidomide Maintenance and Molecular Risk. Journal of Clinical Oncology, 2022, 40, 2889-2900.	0.8	29
5	Ixazomib with cyclophosphamide and dexamethasone in relapsed or refractory myeloma: MUKeight phase II randomised controlled trial results. Blood Cancer Journal, 2022, 12, 52.	2.8	8
6	Epigenomic translocation of H3K4me3 broad domains over oncogenes following hijacking of super-enhancers. Genome Research, 2022, 32, 1343-1354.	2.4	8
7	Myeloma Genome Project Panel is a Comprehensive Targeted Genomics Panel for Molecular Profiling of Patients with Multiple Myeloma. Clinical Cancer Research, 2022, 28, 2854-2864.	3.2	6
8	Structural variants shape the genomic landscape and clinical outcome of multiple myeloma. Blood Cancer Journal, 2022, 12, .	2.8	7
9	Perspectives on the Risk-Stratified Treatment of Multiple Myeloma. Blood Cancer Discovery, 2022, 3, 273-284.	2.6	24
10	Genetic subtypes of smoldering multiple myeloma are associated with distinct pathogenic phenotypes and clinical outcomes. Nature Communications, 2022, 13, .	5.8	11
11	Differential RNA splicing as a potentially important driver mechanism in multiple myeloma. Haematologica, 2021, 106, 736-745.	1.7	20
12	Heterogenous mutation spectrum and deregulated cellular pathways in aberrant plasma cells underline molecular pathology of light-chain amyloidosis. Haematologica, 2021, 106, 601-604.	1.7	2
13	Designing Evolutionary-based Interception Strategies to Block the Transition from Precursor Phases to Multiple Myeloma. Clinical Cancer Research, 2021, 27, 15-23.	3.2	20
14	Optimising the value of immunomodulatory drugs during induction and maintenance in transplant ineligible patients with newly diagnosed multiple myeloma: results from Myeloma XI, a multicentre, openâ€label, randomised, Phase III trial. British Journal of Haematology, 2021, 192, 853-868.	1.2	14
15	Carfilzomib, lenalidomide, dexamethasone, and cyclophosphamide (KRdc) as induction therapy for transplant-eligible, newly diagnosed multiple myeloma patients (Myeloma XI+): Interim analysis of an open-label randomised controlled trial. PLoS Medicine, 2021, 18, e1003454.	3.9	18
16	The molecular make up of smoldering myeloma highlights the evolutionary pathways leading to multiple myeloma. Nature Communications, 2021, 12, 293.	5.8	54
17	Positive selection as the unifying force for clonal evolution in multiple myeloma. Leukemia, 2021, 35, 1511-1515.	3.3	10
18	Whole-genome sequencing reveals progressive versus stable myeloma precursor conditions as two distinct entities. Nature Communications, 2021, 12, 1861.	5.8	68

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19	Bortezomib, Vorinostat, and Dexamethasone Combination Therapy in Relapsed Myeloma: Results of the Phase 2 MUK four Trial. Clinical Lymphoma, Myeloma and Leukemia, 2021, 21, 154-161.e3.	0.2	11
20	Sex Differences in Multiple Myeloma Biology but not Clinical Outcomes: Results from 3894 Patients in the Myeloma XI Trial. Clinical Lymphoma, Myeloma and Leukemia, 2021, 21, 667-675.	0.2	12
21	From Bench to Bedside. Cancer Journal (Sudbury, Mass), 2021, 27, 213-221.	1.0	1
22	The mutagenic impact of melphalan in multiple myeloma. Leukemia, 2021, 35, 2145-2150.	3.3	32
23	Improving prognostic assignment in older adults with multiple myeloma using acquired genetic features, clonal hemopoiesis and telomere length. Leukemia, 2021, , .	3.3	8
24	Case Report: Two Cases of Cryptosporidiosis in Heavily Pretreated Patients With Myeloma. Clinical Lymphoma, Myeloma and Leukemia, 2021, 21, e545-e547.	0.2	3
25	Highâ€risk transcriptional profiles in multiple myeloma are an acquired feature that can occur in any subtype and more frequently with each subsequent relapse. British Journal of Haematology, 2021, 195, 283-286.	1.2	4
26	Mutations in CRBN and other cereblon pathway genes are infrequently associated with acquired resistance to immunomodulatory drugs. Leukemia, 2021, 35, 3017-3020.	3.3	11
27	Copy number signatures predict chromothripsis and clinical outcomes in newly diagnosed multiple myeloma. Nature Communications, 2021, 12, 5172.	5.8	27
28	Impact of Etiological Cytogenetic Abnormalities on the Depth of Immunoparesis and Survival in Newly Diagnosed Multiple Myeloma. Clinical Lymphoma, Myeloma and Leukemia, 2021, , .	0.2	0
29	Residual Monoclonal Free Light Chain Positivity By Mass Spectrometry Identifies Patients at Increased Risk of Early Relapse Following First-Line Anti-Myeloma Treatment. Blood, 2021, 138, 820-820.	0.6	4
30	Multiomic Mapping of Copy Number and Structural Variation on Chromosome 1 (Chr1) Highlights Multiple Recurrent Disease Drivers. Blood, 2021, 138, 721-721.	0.6	0
31	Insights into high-risk multiple myeloma from an analysis of the role of PHF19 in cancer. Journal of Experimental and Clinical Cancer Research, 2021, 40, 380.	3.5	4
32	Microhomology-mediated end joining drives complex rearrangements and overexpression of MYC and PVT1 in multiple myeloma. Haematologica, 2020, 105, 1055-1066.	1.7	42
33	Role of AID in the temporal pattern of acquisition of driver mutations in multiple myeloma. Leukemia, 2020, 34, 1476-1480.	3.3	39
34	Accelerated single cell seeding in relapsed multiple myeloma. Nature Communications, 2020, 11, 3617.	5.8	41
35	Renal outcome in patients with newly diagnosed multiple myeloma: results from the UK NCRI Myeloma XI trial. Blood Advances, 2020, 4, 5836-5845.	2.5	7
36	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

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37	The functional epigenetic landscape of aberrant gene expression in molecular subgroups of newly diagnosed multiple myeloma. Journal of Hematology and Oncology, 2020, 13, 108.	6.9	20
38	Revealing the Impact of Structural Variants in Multiple Myeloma. Blood Cancer Discovery, 2020, 1, 258-273.	2.6	81
39	Deep sequencing as an approach to understanding the complexity and improving the treatment of multiple myeloma. Expert Review of Precision Medicine and Drug Development, 2020, 5, 363-370.	0.4	0
40	Search for multiple myeloma risk factors using Mendelian randomization. Blood Advances, 2020, 4, 2172-2179.	2.5	27
41	Lenalidomide before and after ASCT for transplant-eligible patients of all ages in the randomized, phase III, Myeloma XI trial. Haematologica, 2020, 106, haematol.2020.247130.	1.7	16
42	Genomic analysis of primary plasma cell leukemia reveals complex structural alterations and high-risk mutational patterns. Blood Cancer Journal, 2020, 10, 70.	2.8	27
43	Multiple Myeloma DREAM Challenge reveals epigenetic regulator PHF19 as marker of aggressive disease. Leukemia, 2020, 34, 1866-1874.	3.3	36
44	Reconstructing the evolutionary history of multiple myeloma. Best Practice and Research in Clinical Haematology, 2020, 33, 101145.	0.7	21
45	Antibody-based targeting of BCMA in multiple myeloma. Lancet Oncology, The, 2020, 21, 186-187.	5.1	2
46	BRAF and DIS3 Mutations Associate with Adverse Outcome in a Long-term Follow-up of Patients with Multiple Myeloma. Clinical Cancer Research, 2020, 26, 2422-2432.	3.2	37
47	Long-term outcomes after autologous stem cell transplantation for multiple myeloma. Blood Advances, 2020, 4, 422-431.	2.5	66
48	Whole-Genome Sequencing Reveals Evidence of Two Biologically and Clinically Distinct Entities: Progressive Versus Stable Myeloma Precursor Disease. Blood, 2020, 136, 47-48.	0.6	2
49	Clinical Development of a Non-Gene-Edited Allogeneic Bcma-Targeting CAR T-Cell Product in Relapsed or Refractory Multiple Myeloma. Blood, 2020, 136, 27-28.	0.6	6
50	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
51	Bone marrow microenvironments that contribute to patient outcomes in newly diagnosed multiple myeloma: A cohort study of patients in the Total Therapy clinical trials. PLoS Medicine, 2020, 17, e1003323.	3.9	33
52	Autologous stem cell transplantation is safe and effective for fit older myeloma patients: exploratory results from the Myeloma XI trial. Haematologica, 2020, Online ahead of print, 0-0.	1.7	16
53	Subclonal evolution in disease progression from MGUS/SMM to multiple myeloma is characterised by clonal stability. Leukemia, 2019, 33, 457-468.	3.3	96
54	A high-risk, Double-Hit, group of newly diagnosed myeloma identified by genomic analysis. Leukemia, 2019, 33, 159-170.	3.3	313

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55	An acquired high-risk chromosome instability phenotype in multiple myeloma: Jumping 1q Syndrome. Blood Cancer Journal, 2019, 9, 62.	2.8	23
56	Lack of Spleen Signal on Diffusion Weighted MRI is associated with High Tumor Burden and Poor Prognosis in Multiple Myeloma: A Link to Extramedullary Hematopoiesis?. Theranostics, 2019, 9, 4756-4763.	4.6	12
57	Response-adapted intensification with cyclophosphamide, bortezomib, and dexamethasone versus no intensification in patients with newly diagnosed multiple myeloma (Myeloma XI): a multicentre, open-label, randomised, phase 3 trial. Lancet Haematology,the, 2019, 6, e616-e629.	2.2	42
58	Targeting both BET and CBP/EP300 proteins with the novel dual inhibitors NEO2734 and NEO1132 leads to anti-tumor activity in Multiple Myeloma. Clinical Lymphoma, Myeloma and Leukemia, 2019, 19, e120-e121.	0.2	1
59	Phenome-wide association analysis of LDL-cholesterol lowering genetic variants in PCSK9. BMC Cardiovascular Disorders, 2019, 19, 240.	0.7	22
60	Transcriptome-wide association study of multiple myeloma identifies candidate susceptibility genes. Human Genomics, 2019, 13, 37.	1.4	14
61	Genome-wide interaction and pathway-based identification of key regulators in multiple myeloma. Communications Biology, 2019, 2, 89.	2.0	14
62	Immunotherapy in Multiple Myeloma: Accelerating on the Path to the Patient. Clinical Lymphoma, Myeloma and Leukemia, 2019, 19, 332-344.	0.2	16
63	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
64	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
65	Stem cell mutations can be detected in myeloma patients years before onset of secondary leukemias. Blood Advances, 2019, 3, 3962-3967.	2.5	12
66	Long-term Analysis Of Multiple Sequential Samples Reveals Patterns Of Progression In Smoldering Myeloma. Clinical Lymphoma, Myeloma and Leukemia, 2019, 19, e59-e60.	0.2	0
67	Enrichment for copy number alterations and a unique pattern of gene mutations characterize multiple myeloma in elderly patients. Clinical Lymphoma, Myeloma and Leukemia, 2019, 19, e81-e82.	0.2	0
68	Large deletions (>10.9 MB) in 17p and bi-allelic TP53 inactivation events in newly-diagnosed multiple myeloma are associated with higher clonal cell fraction and poor prognosis. Clinical Lymphoma, Myeloma and Leukemia, 2019, 19, e81.	0.2	0
69	Sequential minimal residual disease (MRD) monitoring: Results from the UK Myeloma XI trial. Clinical Lymphoma, Myeloma and Leukemia, 2019, 19, e45-e46.	0.2	6
70	Circulating cell free DNA is a biomarker for GEP70 risk score and tumor burden in myeloma. Clinical Lymphoma, Myeloma and Leukemia, 2019, 19, e62.	0.2	0
71	Quadruplet KCRD (Carfilzomib, Cyclophosphamide, Lenalidomide and Dexamethasone) Induction for Newly Diagnosed Myeloma Patients. Clinical Lymphoma, Myeloma and Leukemia, 2019, 19, e2.	0.2	1
72	A detailed exploration of using RNA-Seq data in established multiple myeloma gene expression profile microarray based risk scores. Clinical Lymphoma, Myeloma and Leukemia, 2019, 19, e57-e58.	0.2	1

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73	Preclinical evaluation of the new GPRC5DxCD3 (JNJ-7564) bispecific antibody for the treatment of multiple myeloma. Clinical Lymphoma, Myeloma and Leukemia, 2019, 19, e122-e123.	0.2	3
74	FRAX is a robust predictor of baseline vertebral fractures in multiple myeloma patients. Bone, 2019, 121, 134-138.	1.4	3
75	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
76	Combination of flow cytometry and functional imaging for monitoring of residual disease in myeloma. Leukemia, 2019, 33, 1713-1722.	3.3	112
77	Mesenchymal stem cells gene signature in highâ€risk myeloma bone marrow linked to suppression of distinct IGFBP2â€expressing small adipocytes. British Journal of Haematology, 2019, 184, 578-593.	1.2	18
78	Genetic correlation between multiple myeloma and chronic lymphocytic leukaemia provides evidence for shared aetiology. Blood Cancer Journal, 2019, 9, 1.	2.8	40
79	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
80	The Spectrum of Exomic Mutation in Elderly Myeloma Differs Substantially from Patients at Younger Ages Consistent with a Different Evolutionary Trajectory to Full Blown Disease Based on Age of Onset. Blood, 2019, 134, 4346-4346.	0.6	2
81	Chromoplexy and Chromothripsis Are Important Prognostically in Myeloma and Deregulate Gene Function By a Range of Mechanisms. Blood, 2019, 134, 3767-3767.	0.6	5
82	Analysis of Intestinal Microbiome in Multiple Myeloma Reveals Progressive Dysbiosis Compared to MGUS and Healthy Individuals. Blood, 2019, 134, 3076-3076.	0.6	10
83	Poor overall survival in hyperhaploid multiple myeloma is defined by double-hit bi-allelic inactivation of TP53. Oncotarget, 2019, 10, 732-737.	0.8	13
84	Genetic Segmentation and Targeted Therapeutics for Multiple Myeloma. Oncology & Hematology Review, 2019, 15, 87.	0.2	2
85	Kinase domain activation through gene rearrangement in multiple myeloma. Leukemia, 2018, 32, 2435-2444.	3.3	26
86	Loss of heterozygosity as a marker of homologous repair deficiency in multiple myeloma: a role for PARP inhibition?. Leukemia, 2018, 32, 1561-1566.	3.3	39
87	HSF1 Is Essential for Myeloma Cell Survival and A Promising Therapeutic Target. Clinical Cancer Research, 2018, 24, 2395-2407.	3.2	46
88	The multiple myeloma risk allele at 5q15 lowers ELL2 expression and increases ribosomal gene expression. Nature Communications, 2018, 9, 1649.	5.8	22
89	Thymic PTH Increases After Thyroparathyroidectomy in C57BL/KaLwRij Mice. Endocrinology, 2018, 159, 1561-1569.	1.4	4
90	The Pattern of Mesenchymal Stem Cell Expression Is an Independent Marker of Outcome in Multiple Myeloma. Clinical Cancer Research, 2018, 24, 2913-2919.	3.2	30

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91	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
92	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
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	Distinct promoter methylation profile reveals spatial epigenetic heterogeneity in 2 myeloma patients with multifocal extramedullary relapses. Clinical Epigenetics, 2018, 10, 158.	1.8	2
95	The genomic landscape of plasma cells in systemic light chain amyloidosis. Blood, 2018, 132, 2775-2777.	0.6	12
96	Subclonal TP53 copy number is associated with prognosis in multiple myeloma. Blood, 2018, 132, 2465-2469.	0.6	29
97	Identification of multiple risk loci and regulatory mechanisms influencing susceptibility to multiple myeloma. Nature Communications, 2018, 9, 3707.	5.8	86
98	A multiple myeloma classification system that associates normal B-cell subset phenotypes with prognosis. Blood Advances, 2018, 2, 2400-2411.	2.5	5
99	Maintaining therapeutic progress in multiple myeloma by integrating genetic and biological advances into the clinic. Expert Review of Hematology, 2018, 11, 513-523.	1.0	8
100	Serum free light chain levels and renal function at diagnosis in patients with multiple myeloma. BMC Nephrology, 2018, 19, 178.	0.8	24
101	MAFb protein confers intrinsic resistance to proteasome inhibitors in multiple myeloma. BMC Cancer, 2018, 18, 724.	1.1	26
102	Maintenance Treatment and Survival in Patients With Myeloma. JAMA Oncology, 2018, 4, 1389.	3.4	67
103	Identification of novel mutational drivers reveals oncogene dependencies in multiple myeloma. Blood, 2018, 132, 587-597.	0.6	335
104	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
105	Maintenance Therapy with the Oral Proteasome Inhibitor (PI) Ixazomib Significantly Prolongs Progression-Free Survival (PFS) Following Autologous Stem Cell Transplantation (ASCT) in Patients with Newly Diagnosed Multiple Myeloma (NDMM): Phase 3 Tourmaline-MM3 Trial. Blood, 2018, 132, 301-301.	0.6	9
106	Deep Immunoprofiling of the Bone Marrow Microenvironmental Changes Underlying the Multistep Progression of Multiple Myeloma. Blood, 2018, 132, 243-243.	0.6	1
107	Long-Term Follow-up Identifies Double Hit and Key Mutations As Impacting Progression Free and Overall Survival in Multiple Myeloma. Blood, 2018, 132, 110-110.	0.6	1
108	Baseline and on-Treatment Bone Marrow Microenvironments Predict Myeloma Patient Outcomes and Inform Potential Intervention Strategies. Blood, 2018, 132, 1882-1882.	0.6	3

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109	A Quadruplet Regimen Comprising Carfilzomib, Cyclophosphamide, Lenalidomide, Dexamethasone (KCRD) Vs an Immunomodulatory Agent Containing Triplet (CTD/CRD) Induction Therapy Prior to Autologous Stem Cell Transplant: Results of the Myeloma XI Study. Blood, 2018, 132, 302-302.	0.6	6
110	The Mutational Landscape of Primary Plasma Cell Leukemia. Blood, 2018, 132, 114-114.	0.6	2
111	Phase 2 Study of Venetoclax Plus Carfilzomib and Dexamethasone in Patients with Relapsed/Refractory Multiple Myeloma. Blood, 2018, 132, 303-303.	0.6	15
112	A High-Risk Multiple Myeloma Group Identified By Integrative Multi-Omics Segmentation of Newly Diagnosed Patients. Blood, 2018, 132, 3165-3165.	0.6	2
113	Chromothripsis and Chromoplexy Are Associated with DNA Instability and Adverse Clinical Outcome in Multiple Myeloma. Blood, 2018, 132, 408-408.	0.6	3
114	The genomic features associated with high-risk multiple myeloma. Oncotarget, 2018, 9, 35478-35479.	0.8	6
115	Clinical Application of Epigenetic Modifier Mutations in Myeloma. Blood, 2018, 132, SCI-39-SCI-39.	0.6	0
116	Global Expression Changes of Malignant Plasma Cells over Time Reveals the Evolutionary Development of Signatures of Aggressive Clinical Behavior. Blood, 2018, 132, 4457-4457.	0.6	0
117	Poor Overall Survival in Hyperhaploid Multiple Myeloma Is Defined By Double-Hit Bi-Allelic Inactivation of TP53. Blood, 2018, 132, 4441-4441.	0.6	0
118	Sequential Improvements in the Outcome of Autologous Stem Cell Transplantation for Multiple Myeloma over Time. Blood, 2018, 132, 3168-3168.	0.6	0
119	Expression Signature of Myeloma Residual Cells Is Characterized By Genes Associated with Proliferation, Epigenetic Modification, and Stem Cell Maintenance. Blood, 2018, 132, 4465-4465.	0.6	1
120	Myeloma Patient-Derived Bone Marrow Serum Negatively Regulates Natural Killer Cell Activity. Blood, 2018, 132, 4468-4468.	0.6	0
121	Mutations and Copy Number Changes Predict Progression from Smoldering Myeloma to Symptomatic Myeloma in the Era of Novel IMWG Criteria. Blood, 2018, 132, 4456-4456.	0.6	0
122	Global 3D-Epigenetic Dysregulation of Cyclin D1 and D2 Actively Controls Their Expression Pattern in Multiple Myeloma. Blood, 2018, 132, 3904-3904.	0.6	0
123	Combination of Flow Cytometry and Functional Imaging for Monitoring of Residual Disease in Myeloma. Blood, 2018, 132, 3185-3185.	0.6	0
124	Extracting Prognostic Molecular Information from PET-CT Imaging of Multiple Myeloma Using Radiomic Approaches. Blood, 2018, 132, 1906-1906.	0.6	1
125	Lack of a Spleen Signal on Diffusion Weighted MRI Is Associated with High Tumor Burden and Poor Prognosis in Multiple Myeloma. Blood, 2018, 132, 4471-4471.	0.6	0
126	Hotspot Mutations in SF3B1 Result in Increased Alternative Splicing in Multiple Myeloma and Activation of Key Cellular Pathways. Blood, 2018, 132, 4454-4454.	0.6	0

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127	Mesenchymal Stem Cells Gene Signature in High-Risk Myeloma Bone Marrow Linked to Suppression of Distinct IGFBP2-Expressing Small Adipocytes. Blood, 2018, 132, 4448-4448.	0.6	0
128	Characterisation of Long-Term Responders to First-Line Myeloma Therapy - Results from the UK Myeloma IX and XI Trials. Blood, 2018, 132, 2000-2000.	0.6	0
129	High Levels of APOBEC3B Gene Expression Contribute to Poor Prognosis in Multiple Myeloma Patients. Blood, 2018, 132, 3897-3897.	0.6	0
130	Mutant KRAS and Brafs Upregulate Stress Granules and Mediate Drug Resistance, Which Can be Modulated By Cox2 Inhibition in Multiple Myeloma. Blood, 2018, 132, 3166-3166.	0.6	0
131	An Acquired High-Risk Chromosome Instability Phenotype in Multiple Myeloma: Jumping 1q Syndrome. Blood, 2018, 132, 4489-4489.	0.6	1
132	Maximizing Pre-Transplant Response Is Associated with Improved Outcome for Myeloma Patients: Exploratory Analysis of the Myeloma XI Trial. Blood, 2018, 132, 3280-3280.	0.6	2
133	Characterization of the Immune Impact of Daratumumab By Mass Cytometry in Multiple Myeloma. Blood, 2018, 132, 4466-4466.	0.6	0
134	Proliferation and Molecular Risk Score of Low Risk Myeloma Cells Are Increased in High Risk Microenvironment Via Augmented Bioavailability of Growth Factors. Blood, 2018, 132, 1929-1929.	0.6	0
135	Genome-wide association analysis of chronic lymphocytic leukaemia, Hodgkin lymphoma and multiple myeloma identifies pleiotropic risk loci. Scientific Reports, 2017, 7, 41071.	1.6	31
136	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
137	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
138	Extensive Remineralization of Large Pelvic Lytic Lesions Following Total Therapy Treatment in Patients With Multiple Myeloma. Journal of Bone and Mineral Research, 2017, 32, 1261-1266.	3.1	9
139	Immunologic approaches for the treatment of multiple myeloma. Cancer Treatment Reviews, 2017, 55, 190-199.	3.4	46
140	Assessing the effect of obesity-related traits on multiple myeloma using a Mendelian randomisation approach. Blood Cancer Journal, 2017, 7, e573-e573.	2.8	12
141	The prognostic value of the depth of response in multiple myeloma depends on the time of assessment, risk status and molecular subtype. Haematologica, 2017, 102, e313-e316.	1.7	26
142	Diagnosis and monitoring for light chain only and oligosecretory myeloma using serum free light chain tests. British Journal of Haematology, 2017, 178, 220-230.	1.2	34
143	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
144	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

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145	Where are we now with the treatment of multiple myeloma?. Nature Reviews Clinical Oncology, 2017, 14, 461-462.	12.5	11
146	Clinical characteristics and prognostic factors in multiple myeloma patients with light chain deposition disease. American Journal of Hematology, 2017, 92, 739-745.	2.0	36
147	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
148	Genome-wide association study of immunoglobulin light chain amyloidosis in three patient cohorts: comparison with myeloma. Leukemia, 2017, 31, 1735-1742.	3.3	32
149	Evolutionary biology of high-risk multiple myeloma. Nature Reviews Cancer, 2017, 17, 543-556.	12.8	178
150	Neutral tumor evolution in myeloma is associated with poor prognosis. Blood, 2017, 130, 1639-1643.	0.6	20
151	Investigation of a gene signature to predict response to immunomodulatory derivatives for patients with multiple myeloma: an exploratory, retrospective study using microarray datasets from prospective clinical trials. Lancet Haematology,the, 2017, 4, e443-e451.	2.2	20
152	Genetic Predisposition to Multiple Myeloma at 5q15 Is Mediated by an ELL2 Enhancer Polymorphism. Cell Reports, 2017, 20, 2556-2564.	2.9	17
153	Spatial genomic heterogeneity in multiple myeloma revealed by multi-region sequencing. Nature Communications, 2017, 8, 268.	5.8	277
154	Active multiple myeloma suppresses and typically eliminates coexisting MGUS. British Journal of Cancer, 2017, 117, 835-839.	2.9	2
155	Genome-wide association study of clinical parameters in immunoglobulin light chain amyloidosis in three patient cohorts. Haematologica, 2017, 102, e411-e414.	1.7	7
156	Adverse Metaphase Cytogenetics Can Be Overcome by Adding Bortezomib and Thalidomide to Fractionated Melphalan Transplants. Clinical Cancer Research, 2017, 23, 2665-2672.	3.2	13
157	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
158	Differentiation stage of myeloma plasma cells: biological and clinical significance. Leukemia, 2017, 31, 382-392.	3.3	83
159	Hyperhaploidy is a novel high-risk cytogenetic subgroup in multiple myeloma. Leukemia, 2017, 31, 637-644.	3.3	27
160	The efficacy and tolerability of pomalidomide in relapsed/refractory myeloma patients in a â€œreal-worldâ€•study: the Royal Marsden Hospital experience. Leukemia and Lymphoma, 2017, 58, 494-497.	0.6	14
161	Response comparison of multiple myeloma and monoclonal gammopathy of undetermined significance to the same anti-myeloma therapy: a retrospective cohort study. Lancet Haematology,the, 2017, 4, e584-e594.	2.2	6
162	Integration of Genomics Into Treatment: Are We There Yet?. American Society of Clinical Oncology Educational Book / ASCO American Society of Clinical Oncology Meeting, 2017, 37, 569-574.	1.8	2

#	Article	IF	Citations
163	Carfilzomib, Cyclophosphamide and Dexamethasone (KCD) Versus Bortezomib, Cyclophosphamide and Dexamethasone (VCD) for Treatment of First Relapse or Primary Refractory Multiple Myeloma (MM): First Final Analysis of the Phase 2 Muk Five Study. Blood, 2017, 130, 835-835.	0.6	6
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