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	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
164	Minimal Residual Disease in the Maintenance Setting in Myeloma: Prognostic Significance and Impact of Lenalidomide. Blood, 2017, 130, 904-904.	0.6	12
165	The varied distribution and impact of RAS codon and other key DNA alterations across the translocation cyclin D subgroups in multiple myeloma. Oncotarget, 2017, 8, 27854-27867.	0.8	25
166	Search for rare protein altering variants influencing susceptibility to multiple myeloma. Oncotarget, 2017, 8, 36203-36210.	0.8	11
167	Hyperhaploid karyotypes in multiple myeloma. Oncotarget, 2017, 8, 78259-78260.	0.8	4
168	MYC Rearrangements in Multiple Myeloma Are Complex, Can Involve More Than Five Different Chromosomes, and Correlate with Increased Expression of MYC and a Distinct Downstream Gene Expression Pattern. Blood, 2017, 130, 65-65.	0.6	3
169	Disentangling the microRNA regulatorymilieuin multiple myeloma: integrative genomics analysis outlines mixed miRNA-TF circuits and pathway-derived networks modulated in t(4;14) patients. Oncotarget, 2016, 7, 2367-2378.	0.8	41
170	An Outbreak of Respiratory Syncytial Virus Infections in an Outpatient Cancer Unit: Clinical Characteristics and Molecular Investigations. Open Forum Infectious Diseases, 2016, 3, .	0.4	0
171	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
172	MAF protein mediates innate resistance to proteasome inhibition therapy in multiple myeloma. Blood, 2016, 128, 2919-2930.	0.6	57
173	Minimal residual disease following autologous stem cell transplant in myeloma: impact on outcome is independent of induction regimen. Haematologica, 2016, 101, e69-e71.	1.7	41
174	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
175	Percutaneous Device Closure of Paravalvular Leak. Circulation, 2016, 134, 934-944.	1.6	109
176	Clonal selection and double-hit events involving tumor suppressor genes underlie relapse in myeloma. Blood, 2016, 128, 1735-1744.	0.6	170
177	Multiple myeloma risk variant at 7p15.3 creates an IRF4-binding site and interferes with CDCA7L expression. Nature Communications, 2016, 7, 13656.	5.8	32
178	Genome-wide association study identifies multiple susceptibility loci for multiple myeloma. Nature Communications, 2016, 7, 12050.	5.8	146
179	The Spectrum and Clinical Impact of Epigenetic Modifier Mutations in Myeloma. Clinical Cancer Research, 2016, 22, 5783-5794.	3.2	81
180	Genomewide profiling of copyâ€number alteration in monoclonal gammopathy of undetermined significance. European Journal of Haematology, 2016, 97, 568-575.	1.1	24

#	Article	IF	Citations
181	Genetic factors influencing the risk of multiple myeloma bone disease. Leukemia, 2016, 30, 883-888.	3.3	11
182	Monoclonal antibody therapy in multiple myeloma: where do we stand and where are we going?. Immunotherapy, 2016, 8, 367-384.	1.0	6
183	Impact of Genes Highly Correlated with MMSET Myeloma on the Survival of Non-MMSET Myeloma Patients. Clinical Cancer Research, 2016, 22, 4039-4044.	3.2	14
184	The safety of pomalidomide for the treatment of multiple myeloma. Expert Opinion on Drug Safety, 2016, 15, 535-547.	1.0	6
185	Genome-wide association study identifies variation at 6q25.1 associated with survival in multiple myeloma. Nature Communications, 2016, 7, 10290.	5.8	31
186	Clinical value of molecular subtyping multiple myeloma using gene expression profiling. Leukemia, 2016, 30, 423-430.	3.3	58
187	Risk of multiple myeloma in a caseâ€“spouse study. Leukemia and Lymphoma, 2016, 57, 1450-1459.	0.6	8
188	The Co-Occurrence of MAF Translocations in RAS Mutated Multiple Myeloma Confers Resistance to MEK Inhibition. Blood, 2016, 128, 1138-1138.	0.6	2
189	Lenalidomide Is a Highly Effective Maintenance Therapy in Myeloma Patients of All Ages; Results of the Phase III Myeloma XI Study. Blood, 2016, 128, 1143-1143.	0.6	26
190	Primary IMiD Refractory Myeloma; Results from 3894 Patients Treated in the Phase III Myeloma XI Study. Blood, 2016, 128, 1144-1144.	0.6	5
191	Signatures of Mesenchymal Cell Lineages and Microenvironment Factors Are Dysregulated in High Risk Myeloma. Blood, 2016, 128, 2065-2065.	0.6	1
192	Myeloma-Derived Exosomes and Soluble Factors Suppress Natural Killer Cell Function. Blood, 2016, 128, 2066-2066.	0.6	2
193	Myeloma Exosomes Prime the Microenvironment to Support Survival and Growth of Myeloma Cells. Blood, 2016, 128, 2067-2067.	0.6	2
194	Response Adapted Induction Treatment Improves Outcomes for Myeloma Patients; Results of the Phase III Myeloma XI Study. Blood, 2016, 128, 244-244.	0.6	9
195	Concurrent Amplification of MYC and 1q21 in Multiple Myeloma: Focal and Segmental Jumping Translocations of MYC. Blood, 2016, 128, 3266-3266.	0.6	1
196	The 70-Gene MyPRSR prognostic Risk Score Signature Predicts Increased Risk of Progression from MGUS to Multiple Myeloma Requring Treatment. Blood, 2016, 128, 3275-3275.	0.6	1
197	Extensive Regional Intra-Clonal Heterogeneity in Multiple Myeloma - Implications for Diagnostics, Risk Stratification and Targeted Treatment. Blood, 2016, 128, 3278-3278.	0.6	2
198	Mesenchymal Stem Cells Preconditioned with Myeloma Cells from High-Risk Patients Support the Growth of Myeloma Cells from Low-Risk Patients. Blood, 2016, 128, 3304-3304.	0.6	3

#	Article	IF	Citations
199	A Phase I Dose-Escalation Study of the Class 1 Selective Histone Deacetylase Inhibitor CHR-3996 in Combination with Tosedostat for Patients with Relapsed, Refractory Multiple Myeloma: Results of the Muk Three Trial. Blood, 2016, 128, 3321-3321.	0.6	5
200	Automated Multiparameter Flow Cytometry (MFC) Immunophenotyping for Reproducible Identification of High Risk Smoldering Multiple Myeloma (SMM). Blood, 2016, 128, 373-373.	0.6	1
201	Comparison of MRD Detection By MFC, NGS and PET-CT in Patients at Different Treatment Stages for Multiple Myeloma. Blood, 2016, 128, 377-377.	0.6	1
202	Identifying Ultra-High Risk Myeloma By Integrated Molecular Genetic and Gene Expression Profiling. Blood, 2016, 128, 4407-4407.	0.6	2
203	The Impact of Maintenance Lenalidomide on the Mutational Status of the Myeloma Clone at Relapse in the NCRI Myeloma XI Trial for Newly Diagnosed Multiple Myeloma Patients (NDMM). Blood, 2016, 128, 4412-4412.	0.6	2
204	The Clinical Impact of Macrofocal Disease in Multiple Myeloma Differs Between Presentation and Relapse. Blood, 2016, 128, 4431-4431.	0.6	8
205	Daratumumab Single Agent and Daratumumab Plus Pomalidomide and Dexametasone in Relapsed/Refractory Multiple Myeloma: A Real Life Retrospective Evaluation. Blood, 2016, 128, 4516-4516.	0.6	8
206	A Favorable BCL-2 Family Expression Profile May Explain the Increased Susceptibility of the t(11;14) Multiple Myeloma Subgroup to Single Agent Venetoclax. Blood, 2016, 128, 5613-5613.	0.6	9
207	Use of Multiple Myeloma 70-Gene Prognostic Risk Score As a Continuous Predicitor of Patient Outcome. Blood, 2016, 128, 5614-5614.	0.6	1
208	DNA Methylation Profiling of Myeloma Trial Patients Reveals Specific Epigenetic Changes Associated with Outcome. Blood, 2016, 128, 804-804.	0.6	1
209	A Survey of Fusion Genes in Myeloma Identifies Kinase Domain Activation Which Could be Targeted with Available Treatments. Blood, 2016, 128, 117-117.	0.6	1
210	Next Generation Sequencing (NGS) Based Minimal Residual Disease (MRD) Testing Is Highly Predictive of Overall and Progression Free Survival in the Total Therapy Trials and Shows Different Prognostic Implications in High Vs Standard Risk Multiple Myeloma. Blood, 2016, 128, 2064-2064.	0.6	0
211	The Multiple Myeloma Genome Project: Development of a Molecular Segmentation Strategy for the Clinical Classification of Multiple Myeloma. Blood, 2016, 128, 196-196.	0.6	2
212	Spatial-Temporal Genomic Analyses Reveal a Component of "Big Bang" Kinetics in Multiple Myeloma Evolution. Blood, 2016, 128, 239-239.	0.6	3
213	High Risk Myeloma Is Characterized By the Bi-Allelic Inactivation of CDKN2C and RB1. Blood, 2016, 128, 4416-4416.	0.6	1
214	The Mutational and Signaling Landscape of Multiple Myeloma Varies Dependent upon Translocation Cyclin D (TC) Subgroup. Blood, 2016, 128, 4441-4441.	0.6	0
215	The Metabolic Phenotype of Myeloma Plasma Cells Differs Between Active and Residual Disease States. Blood, 2016, 128, 4438-4438.	0.6	0
216	Translocations and Jumping Rearrangements at 8q24 Result in over-Expression of MYC and are Key Drivers of Disease Progression. Blood, 2016, 128, 115-115.	0.6	2

#	Article	IF	Citations
217	Multiple Myeloma with a Deletion of Chromosome 17p: TP53 Mutations Are Highly Prevalent and Negatively Affect Prognosis. Blood, 2016, 128, 3271-3271.	0.6	2
218	A molecular diagnostic approach able to detect the recurrent genetic prognostic factors typical of presenting myeloma. Genes Chromosomes and Cancer, 2015, 54, 91-98.	1.5	31
219	Implementation of genome-wide complex trait analysis to quantify the heritability in multiple myeloma. Scientific Reports, 2015, 5, 12473.	1.6	16
220	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
221	Evidence of an epigenetic origin for high-risk 1q21 copy number aberrations in multiple myeloma. Blood, 2015, 125, 3756-3759.	0.6	41
222	Four genes predict high risk of progression from smoldering to symptomatic multiple myeloma (SWOG S0120). Haematologica, 2015, 100, 1214-1221.	1.7	44
223	The 7p15.3 (rs4487645) association for multiple myeloma shows strong allele-specific regulation of the MYC-interacting gene CDCA7L in malignant plasma cells. Haematologica, 2015, 100, e110-e113.	1.7	27
224	Bendamustine, thalidomide and dexamethasone combination therapy for relapsed/refractory myeloma patients: results of the MUKone randomized dose selection trial. British Journal of Haematology, 2015, 170, 336-348.	1.2	19
225	Successful mobilization of PBSCs predicts favorable outcomes in multiple myeloma patients treated with novel agents and autologous transplantation. Bone Marrow Transplantation, 2015, 50, 673-678.	1.3	21
226	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
227	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
228	APOBEC family mutational signatures are associated with poor prognosis translocations in multiple myeloma. Nature Communications, 2015, 6, 6997.	5.8	261
229	Removing batch effects from purified plasma cell gene expression microarrays with modified ComBat. BMC Bioinformatics, 2015, 16, 63.	1.2	73
230	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
231	A gene expression based predictor for high risk myeloma treated with intensive therapy and autologous stem cell rescue. Leukemia and Lymphoma, 2015, 56, 594-601.	0.6	7
232	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
233	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
234	Overall survival of relapsed and refractory multiple myeloma patients after adjusting for crossover in the <scp>MM</scp>â€003 trial for pomalidomide plus lowâ€dose dexamethasone. British Journal of Haematology, 2015, 168, 820-823.	1.2	21

#	Article	IF	Citations
235	The Composition and Clinical Impact of Focal Lesions and Their Impact on the Microenvironment in Myeloma. Blood, 2015, 126, 1806-1806.	0.6	2
236	Melphalan Affects Genes Critical for Myeloma Survival, Homing, and Response to Cytokines and Chemokines. Blood, 2015, 126, 1808-1808.	0.6	2
237	Upfront 28-Day Metronomic Therapy for High-Risk Multiple Myeloma (HRMM). Blood, 2015, 126, 1843-1843.	0.6	1
238	Myeloma XI Trial for Newly Diagnosed Multiple Myeloma (NDMM); A Report of Second Primary Malignancy (SPM) Rates and the Importance of Review of Reported Cases. Blood, 2015, 126, 1847-1847.	0.6	1
239	High Risk Multiple Myeloma Demonstrates Marked Spatial Genomic Heterogeneity Between Focal Lesions and Random Bone Marrow; Implications for Targeted Therapy and Treatment Resistance. Blood, 2015, 126, 20-20.	0.6	7
240	Impact of Minimal Residual Disease in High and Standard Risk Multiple Myeloma. Blood, 2015, 126, 2979-2979.	0.6	2
241	Comprehensive Genomic Profiling of Multiple Myeloma in the Course of Clinical Care Identifies Targetable and Prognostically Significant Genomic Alterations. Blood, 2015, 126, 369-369.	0.6	1
242	The Impact of Combination Chemotherapy and Tandem Stem Cell Transplant on Clonal Substructure and Mutational Pattern at Relapse of MM. Blood, 2015, 126, 372-372.	0.6	1
243	Specific Exosomal microRNA Are Differentially Expressed Between High and Low-Risk Myeloma Suggesting They Are Pathogenically Important. Blood, 2015, 126, 4189-4189.	0.6	2
244	The combination of HDAC and aminopeptidase inhibitors is highly synergistic in myeloma and leads to disruption of the NFÎºB signalling pathway. Oncotarget, 2015, 6, 17314-17327.	0.8	9
245	Integrated analysis of microRNAs, transcription factors and target genes expression discloses a specific molecular architecture of hyperdiploid multiple myeloma. Oncotarget, 2015, 6, 19132-19147.	0.8	41
246	Outcomes of Autologous Transplantation for Treatment-Related AML and MDS in Previously Treated Multiple Myeloma Patients (pts). Blood, 2015, 126, 1997-1997.	0.6	0
247	Assessment of Total Lesion Glycolysis and Metabolic Tumor Volume Improve the Clinical Value of Focal Lesion Assessment By FDG PET/CT in Myeloma. Blood, 2015, 126, 724-724.	0.6	0
248	Stem Cell-like Characteristics of MM Plasma Cells Vary By ROS Levels: Implications for Targeted Therapy. Blood, 2015, 126, 1820-1820.	0.6	1
249	Deletion of TP53 (17p13) Is Associated with Poor Outcome for Newly Diagnosed High-Risk Multiple Myeloma. Blood, 2015, 126, 2982-2982.	0.6	0
250	Molecular Subtyping and Risk Stratification for the Classification of Myeloma. Blood, 2015, 126, 4173-4173.	0.6	0
251	A Prognostic 51-Gene Signature Linked to Abnormal Metaphase Cytogenetics Identifies Myeloma Patients Who Benefit from Fractionated Melphalan Dosing and Added Bortezomib, Thalidomide and Dexamethasone As Conditioning for Autologous Stem Cell Transplant. Blood, 2015, 126, 3181-3181.	0.6	2
252	Identifying Targets for Therapy in High Risk t(4;14) Myeloma Using Multi-Level Molecular and Phenotypic Analysis of Isogenic MMSET and MMSET Knock out Cell Lines. Blood, 2015, 126, 1792-1792.	0.6	0

#	Article	IF	Citations
253	Differential ICAM3 Gene Expression Correlates with Susceptibility to Natural Killer Cell-Mediated Lysis in Multiple Myeloma. Blood, 2015, 126, 2990-2990.	0.6	0
254	Defining the Impact of Tandem Autologous Stem Cell Transplantation in Multiple Myeloma: A Case-Match Analysis in the Total Therapy Trials. Blood, 2015, 126, 3182-3182.	0.6	1
255	Profound Impact of Sample Processing Delay on Gene Expression of Multiple Myeloma Plasma Cells. Blood, 2015, 126, 2996-2996.	0.6	0
256	Extending Metronomic Therapy to 28 Days (metro28) for Relapsed Refractory Multiple Myeloma (RRMM). Blood, 2015, 126, 5395-5395.	0.6	0
257	Identification of Biomarkers Associated with MAF-Mediated Resistance to Proteasome Inhibitors in t(14;16) Multiple Myeloma. Blood, 2015, 126, 3020-3020.	0.6	1
258	Re-Mineralization of Large Pelvic Lytic Lesions By CT Imaging in Patients with Multiple Myeloma: The Arkansas Experience. Blood, 2015, 126, 4193-4193.	0.6	0
259	Gene Expression Profiling of Extramedullary Disease-Related Toward Identification of a Terminal Disease Pathway in Multiple Myeloma. Blood, 2015, 126, 1777-1777.	0.6	23
260	B-cell malignancies: capture-sequencing strategies for identification of gene rearrangements and translocations into immunoglobulin gene loci. Blood and Lymphatic Cancer: Targets and Therapy, 2014, , 107.	1.2	0
261	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
262	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
263	Expert panel consensus statement on the optimal use of pomalidomide in relapsed and refractory multiple myeloma. Leukemia, 2014, 28, 1573-1585.	3.3	108
264	Biology and Treatment of Myeloma. Clinical Lymphoma, Myeloma and Leukemia, 2014, 14, S65-S70.	0.2	15
265	Current and potential epigenetic targets in multiple myeloma. Epigenomics, 2014, 6, 215-228.	1.0	11
266	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
267	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
268	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
269	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
270	Biologic Frontiers in Multiple Myeloma: From Biomarker Identification to Clinical Practice. Clinical Cancer Research, 2014, 20, 804-813.	3.2	29

#	Article	IF	Citations
271	Cancer-Selective Targeting of the NF-Î²B Survival Pathway with GADD45Î²/MKK7 Inhibitors. Cancer Cell, 2014, 26, 495-508.	7.7	99
272	Optimizing the treatment of multiple myeloma. Nature Reviews Clinical Oncology, 2014, 11, 686-688.	12.5	0
273	The impact of intra-clonal heterogeneity on the treatment of multiple myeloma. British Journal of Haematology, 2014, 165, 441-454.	1.2	91
274	European Perspective on Multiple Myeloma Treatment Strategies in 2014. Oncologist, 2014, 19, 829-844.	1.9	90
275	Inherited genetic susceptibility to multiple myeloma. Leukemia, 2014, 28, 518-524.	3.3	60
276	Reply to M. Roschewski et al. Journal of Clinical Oncology, 2014, 32, 476-477.	0.8	0
277	Phase II Study of Vincristine Sulfate Liposome Injection (Marqibo) and Rituximab for Patients With Relapsed and Refractory Diffuse Large B-Cell Lymphoma or Mantle Cell Lymphoma in Need of Palliative Therapy. Clinical Lymphoma, Myeloma and Leukemia, 2014, 14, 37-42.	0.2	31
278	Understanding the multiple biological aspects leading to myeloma. Haematologica, 2014, 99, 605-612.	1.7	25
279	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
280	The sialyltransferase ST3GAL6 influences homing and survival in multiple myeloma. Blood, 2014, 124, 1765-1776.	0.6	97
281	Curing myeloma at last: defining criteria and providing the evidence. Blood, 2014, 124, 3043-3051.	0.6	194
282	Curing Multiple Myeloma (MM) with Total Therapy (TT). Blood, 2014, 124, 195-195.	0.6	3
283	Further Evolution of Metronomic Therapy Extended to 28 Days (Metro28) for Relapsed Refractory Multiple Myeloma (RRMM). Blood, 2014, 124, 2128-2128.	0.6	1
284	The Spectrum of Epigenetic Mutations in Myeloma and Their Clinical Impact. Blood, 2014, 124, 2194-2194.	0.6	2
285	Somatic Mutation Spectrum in Monoclonal Gammopathy of Undetermined Significance Compared to Multiple Myeloma. Blood, 2014, 124, 3346-3346.	0.6	1
286	Higher Expressions of PTH Receptor Type 1 and/or 2 in Bone Marrow Is Associated to Longer Survival in Newly Diagnosed Myeloma Patients Enrolled in Total Therapy 3. Blood, 2014, 124, 3409-3409.	0.6	5
287	Minimal Residual Disease (MRD) in Myeloma: Independent Outcome Prediction and Sequential Survival Benefits per Log Tumour Reduction. Blood, 2014, 124, 3416-3416.	0.6	1
288	Characterization of the Mutational Landscape of Multiple Myeloma Using Comprehensive Genomic Profiling. Blood, 2014, 124, 3418-3418.	0.6	3

#	Article	IF	Citations
289	Multiple Myeloma Cells Express Functional E-Selectin Ligands Which Can be Inhibited Both in-Vitro and in-Vivo Leading to Prolongation of Survival in a Murine Transplant Model. Blood, 2014, 124, 4718-4718.	0.6	4
290	Targeted MEK Inhibition in Patients with Previously Treated Multiple Myeloma. Blood, 2014, 124, 4775-4775.	0.6	4
291	Safety and Efficacy in the Stratus (MM-010) Trial, a Single-Arm Phase 3b Study Evaluating Pomalidomide + Low-Dose Dexamethasone in Patients with Refractory or Relapsed and Refractory Multiple Myeloma. Blood, 2014, 124, 80-80.	0.6	2
292	A Novel Functional Role for MMSET in RNA Processing Based on the Link Between the REIIBP Isoform and Its Interaction with the SMN Complex. PLoS ONE, 2014, 9, e99493.	1.1	5
293	Flow Cytometry Defined Cytoplasmic Immunoglobulin Index Is a Major Prognostic Factor for Progression of Asymptomatic Monoclonal Gammopathies to Clinical Multiple Myeloma. Blood, 2014, 124, 2079-2079.	0.6	0
294	Identifying a Gene Expression (GEP)-Based Model Predicting for Progression from AMM to Cmm Requiring Therapy in S0120 Patients Treated at Mirt. Blood, 2014, 124, 2078-2078.	0.6	0
295	PET-CT Defined Focal Lesions at Baseline and Day 7 Predict Outcome in GEP 70 Defined High Risk Multiple Myeloma Patients. Blood, 2014, 124, 3407-3407.	0.6	0
296	High Resolution Genome Wide DNA Methylation Analysis in a Large Trial Group Reveals a Novel Epigenetically Defined Subgroup of Myeloma Patients Characterized By Developmental Gene Hypermethylation. Blood, 2014, 124, 2189-2189.	0.6	0
297	Prediction of High and Low-Risk Multiple Myeloma Based on the EMC92 Gene Expression Signature and the International Staging System. Blood, 2014, 124, 3358-3358.	0.6	0
298	Mafb Protein Confers Primary Resistance of Myeloma to Proteasome Inhibitors. Blood, 2014, 124, 2091-2091.	0.6	1
299	Apobec Family Mutational Signatures Are Associated with Poor Prognosis Translocations in Multiple Myeloma. Blood, 2014, 124, 723-723.	0.6	0
300	Defining Risk of MGUS and AMM Progression to Myeloma By Ig Heavy-Chain FISH. Blood, 2014, 124, 3408-3408.	0.6	0
301	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
302	Role of thalidomide in the treatment of patients with multiple myeloma. Critical Reviews in Oncology/Hematology, 2013, 88, S14-S22.	2.0	22
303	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
304	Identification of a novel t(7;14) translocation in multiple myeloma resulting in overexpression of EGFR. Genes Chromosomes and Cancer, 2013, 52, 817-822.	1.5	7
305	Global methylation analysis identifies prognostically important epigenetically inactivated tumor suppressor genes in multiple myeloma. Blood, 2013, 122, 219-226.	0.6	147
306	Hsp70 inhibition induces myeloma cell death via the intracellular accumulation of immunoglobulin and the generation of proteotoxic stress. Cancer Letters, 2013, 339, 49-59.	3.2	31

#	Article	IF	Citations
307	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
308	Identification of Autophosphorylation Inhibitors of the Inositol-Requiring Enzyme 1 Alpha (IRE1Î±) by High-Throughput Screening Using a DELFIA Assay. Journal of Biomolecular Screening, 2013, 18, 298-308.	2.6	13
309	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
310	Response and biological subtype of myeloma are independent prognostic factors and combine to define outcome after highâ€dose therapy. British Journal of Haematology, 2013, 161, 291-294.	1.2	4
311	A TC classification-based predictor for multiple myeloma using multiplexed real-time quantitative PCR. Leukemia, 2013, 27, 1754-1757.	3.3	36
312	Understanding the interplay between the proteasome pathway and autophagy in response to dual PI3K/mTOR inhibition in myeloma cells is essential for their effective clinical application. Leukemia, 2013, 27, 2397-2403.	3.3	22
313	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
314	Transplants for the elderly in myeloma. Blood, 2013, 122, 1332-1334.	0.6	7
315	Proof of the Concept to Use a Malignant B Cell Line Drug Screen Strategy for Identification and Weight of Melphalan Resistance Genes in Multiple Myeloma. PLoS ONE, 2013, 8, e83252.	1.1	13
316	Low Expression Of The FUCA1 Gene Is An Adverse Prognostic Factor In Myeloma and Combined With High Sialyltransferase Gene Expression Identifies Patients At Increased Risk Of Early Disease Progression and Death. Blood, 2013, 122, 1864-1864.	0.6	4
317	The Value Of Serum Free Light Chain Monitoring Compared To Urinary Bence-Jones Measurement In Light Chain Only Myeloma. Blood, 2013, 122, 1895-1895.	0.6	3
318	MYC Translocations In Multiple Myeloma Involve Recruitment Of Enhancer Elements Resulting In Over-Expression and Decreased Overall Survival. Blood, 2013, 122, 274-274.	0.6	2
319	Single-Cell Genetic Analysis Reveals The Genetic Composition Of Founder Clones, Phylogenetic Patterns Of Branching and Parallel Evolution, and Clonal Fluctuations Following Patient Treatment In Multiple Myeloma. Blood, 2013, 122, 398-398.	0.6	0
320	Myeloma:diagnosis complications and supportive care. Hematology, 2012, 17, s109-s111.	0.7	2
321	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
322	The role of maintenance thalidomide therapy in multiple myeloma: MRC Myeloma IX results and meta-analysis. Blood, 2012, 119, 7-15.	0.6	315
323	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
324	Common variation at 3p22.1 and 7p15.3 influences multiple myeloma risk. Nature Genetics, 2012, 44, 58-61.	9.4	137

#	Article	IF	Citations
325	Understanding the molecular biology of myeloma and its therapeutic implications. Expert Review of Hematology, 2012, 5, 603-617.	1.0	14
326	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
327	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
328	What is the evidence for the use of bisphosphonate therapy in newly diagnosed multiple myeloma patients lacking bone disease?. Hematology American Society of Hematology Education Program, 2012, 2012, 350-353.	0.9	4
329	How to use new biology to guide therapy in multiple myeloma. Hematology American Society of Hematology Education Program, 2012, 2012, 342-349.	0.9	25
330	The genetic architecture of multiple myeloma. Nature Reviews Cancer, 2012, 12, 335-348.	12.8	741
331	Targeting Bone in Myeloma. Recent Results in Cancer Research, 2012, 192, 127-143.	1.8	4
332	Sequential Immunomodulatory Drug (IMiD) and Proteosome Inhibitor Therapy Improves Response Rates in Newly Diagnosed Multiple Myeloma: Preliminary Results From the Myeloma XI Trial. Blood, 2012, 120, 335-335.	0.6	5
333	Efficacy and side-effect profile of long-term bisphosphonate therapy in patients (pts) with multiple myeloma (MM): MRC myeloma IX study results.. Journal of Clinical Oncology, 2012, 30, 8015-8015.	0.8	1
334	Regional differences in the treatment approaches for relapsed multiple myeloma: An IMF study.. Journal of Clinical Oncology, 2012, 30, 8095-8095.	0.8	9
335	A Modified Method for Whole Exome Resequencing from Minimal Amounts of Starting DNA. PLoS ONE, 2012, 7, e32617.	1.1	24
336	The Natural History of Myeloma Patients Who Relapse Early After High Dose Therapy.. Blood, 2012, 120, 3125-3125.	0.6	0
337	Base-Pair Resolution Mapping of IGH Translocations in Multiple Myeloma Using Targeted Capture and Massively Parallel Sequencing. Blood, 2012, 120, 3490-3490.	0.6	0
338	Adjusting for Patient Crossover in Clinical Trials Using External Data: A Case Study of Lenalidomide for Advanced Multiple Myeloma. Value in Health, 2011, 14, 672-678.	0.1	9
339	Treatment of relapsed and refractory multiple myeloma in the era of novel agents. Cancer Treatment Reviews, 2011, 37, 266-283.	3.4	66
340	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
341	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
342	Aberrant global methylation patterns affect the molecular pathogenesis and prognosis of multiple myeloma. Blood, 2011, 117, 553-562.	0.6	217

#	Article	IF	Citations
343	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
344	Guidelines for the diagnosis and management of multiple myeloma 2011. British Journal of Haematology, 2011, 154, 32-75.	1.2	252
345	Structure of the Ire1 autophosphorylation complex and implications for the unfolded protein response. EMBO Journal, 2011, 30, 894-905.	3.5	201
346	Targeting Bone as a Therapy for Myeloma. Cancer Microenvironment, 2011, 4, 299-311.	3.1	3
347	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
348	Genetic factors underlying the risk of bortezomib induced peripheral neuropathy in multiple myeloma patients. Haematologica, 2011, 96, 1728-1732.	1.7	48
349	A Gene Expressionâ€“Based Predictor for Myeloma Patients at High Risk of Developing Bone Disease on Bisphosphonate Treatment. Clinical Cancer Research, 2011, 17, 6347-6355.	3.2	27
350	Gender Disparities in the Tumor Genetics and Clinical Outcome of Multiple Myeloma. Cancer Epidemiology Biomarkers and Prevention, 2011, 20, 1703-1707.	1.1	39
351	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
352	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
353	Could DNA methylation become a useful measure for multiple myeloma prognoses?. Expert Review of Hematology, 2011, 4, 125-127.	1.0	3
354	The outcome of high-dose chemotherapy and auto-SCT in patients with multiple myeloma: a UK/Ireland and European benchmarking comparative analysis. Bone Marrow Transplantation, 2011, 46, 1210-1218.	1.3	7
355	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
356	Novel Drugs in Myeloma: Harnessing Tumour Biology to Treat Myeloma. Recent Results in Cancer Research, 2011, 183, 151-187.	1.8	7
357	MRC Myeloma IX, 6 Year Median Follow-up (FU) Highlights the Importance of Long-Term FU in Myeloma Clinical Trials and Differential Effects of Thalidomide in High- and Low-Risk Disease. Blood, 2011, 118, 993-993.	0.6	2
358	High Absolute T Regulatory Cell Counts and Resistance of CD8 Central Memory Cells to Killing by Fludarabine Predicts for Poor Responses to DLI in the Context of a Pre-DLI Lymphoreduction Strategy. Blood, 2011, 118, 1912-1912.	0.6	16
359	Whole Genome Sequencing Illuminates the Genetic and Biological Features Underlying the Transition of SMM to MM. Blood, 2011, 118, 296-296.	0.6	1
360	The Interaction of Response and FISH-Based Risk Stratification to Better Define Clinical Outcome in Myeloma. Blood, 2011, 118, 1823-1823.	0.6	0

#	Article	IF	Citations
361	A Novel Mouse Model of Multiple Myeloma Representative of Human Disease and Its Use in Preclinical Therapeutic Assessment. Blood, 2011, 118, 2907-2907.	0.6	0
362	High Readmission Rates Are Associated with a Significant Economic Burden and Poor Outcome in Patients with Grade 3/4 Acute GvHD. Blood, 2011, 118, 2061-2061.	0.6	0
363	Differences in Patterns of Treatment and Outcome Among Patients with Relapsed Refractory Myeloma From United States, Europe and Asia,. Blood, 2011, 118, 3989-3989.	0.6	0
364	Exome Sequencing of the t(4;14) and t(11;14) Translocation Specific Subgroups of MM. Blood, 2011, 118, 1817-1817.	0.6	0
365	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
366	A compendium of myeloma-associated chromosomal copy number abnormalities and their prognostic value. Blood, 2010, 116, e56-e65.	0.6	315
367	p53 protein overexpression in bone marrow biopsies from chronic lymphocytic leukaemia is associated with TP53 deletion and resistance to fludarabine. Journal of Hematopathology, 2010, 3, 61-68.	0.2	3
368	Antitumor Effects and Anticancer Applications of Bisphosphonates. Seminars in Oncology, 2010, 37, S30-S40.	0.8	57
369	Nonâ€Homologous Endâ€Joining Gene Profiling Reveals Distinct Expression Patterns Associated with Lymphoma and Multiple Myeloma. British Journal of Haematology, 2010, 149, 258-262.	1.2	15
370	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
371	Targeting heat shock protein 72 enhances Hsp90 inhibitor-induced apoptosis in myeloma. Leukemia, 2010, 24, 1804-1807.	3.3	71
372	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
373	Current Multiple Myeloma Treatment Strategies with Novel Agents: A European Perspective. Oncologist, 2010, 15, 6-25.	1.9	85
374	High expression levels of the mammalian target of rapamycin inhibitorDEPTORare predictive of response to thalidomide in myeloma. Leukemia and Lymphoma, 2010, 51, 2126-2129.	0.6	26
375	Transfection of siRNAs in Multiple Myeloma Cell Lines. Methods in Molecular Biology, 2010, 623, 299-309.	0.4	6
376	First-line treatment with zoledronic acid as compared with clodronic acid in multiple myeloma (MRC) Tj ETQq0 0 0 rgBT /Overlock 10 Tf	6.3	505
377	Efficacy and outcome of autologous transplantation in rare myelomas. Haematologica, 2010, 95, 2126-2133.	1.7	51
378	Optimising Bone Disease In Myeloma; Zoledronic Acid Plus Thalidomide Combinations Improves Survival and Bone Endpoints: Results of the MRC Myeloma IX Trial. Blood, 2010, 116, 311-311.	0.6	3

#	Article	IF	Citations
379	Hypermethylation Is A Key Feature of the Transition of Multiple Myeloma to Plasma Cell Leukemia. Blood, 2010, 116, 535-535.	0.6	1
380	Thalidomide Maintenance Significantly Improves Progression-Free Survival (PFS) and Overall Survival (OS) of Myeloma Patients When Effective Relapse Treatments Are Used: MRC Myeloma IX Results. Blood, 2010, 116, 623-623.	0.6	13
381	Defining Myeloma Patients at High Risk of Developing Bone Disease While on Bisphosphonate Treatment. Blood, 2010, 116, 782-782.	0.6	3
382	Identification of Single Nucleotide Polymorphism Interactions Associated with Survival and Risk In Multiple Myeloma Using Novel Data Mining Methods. Blood, 2010, 116, 2973-2973.	0.6	0
383	Expression Profile and up-Regulation of Telomere-Associated Proteins In Multiple Myeloma. Blood, 2010, 116, 4050-4050.	0.6	0
384	Deletion 13, Detected by Metaphase Analysis, Is Not a Significant Prognostic Indicator In Myeloma. Blood, 2010, 116, 2980-2980.	0.6	0
385	The Introduction of Novel Agents Improves Outcomes of Young Patients with Myeloma (MM) Treated with Autologous Stem Cell Transplant (ASCT). Blood, 2010, 116, 1348-1348.	0.6	0
386	Cost-Effectiveness of Zoledronic Acid Versus Clodronate In Patients with Multiple Myeloma From a Canadian Healthcare System Perspective.. Blood, 2010, 116, 3812-3812.	0.6	0
387	Defining High Risk Myeloma Using Co-Segregating FISH Variables; Results of MRC Myeloma IX. Blood, 2010, 116, 1907-1907.	0.6	1
388	Autophagy Is a Key Myeloma Survival Pathway That Can Be Manipulated Therapeutically to Enhance Apoptosis. Blood, 2010, 116, 4083-4083.	0.6	1
389	Results of a Prospective Clinical Trial of Pre-DLI Lymphoreduction Using Oral Fludarabine In Patients with Mixed Chimerism Post Allogeneic Transplant.. Blood, 2010, 116, 1299-1299.	0.6	26
390	Genetic polymorphisms of EPHX1, Gsk3Î², TNFSF8 and myeloma cell DKK-1 expression linked to bone disease in myeloma. Leukemia, 2009, 23, 1913-1919.	3.3	17
391	The impact of extramedullary disease at presentation on the outcome of myeloma. Leukemia and Lymphoma, 2009, 50, 230-235.	0.6	97
392	Aminopeptidase inhibition as a targeted treatment strategy in myeloma. Molecular Cancer Therapeutics, 2009, 8, 762-770.	1.9	44
393	MMSET deregulation affects cell cycle progression and adhesion regulons in t(4;14) myeloma plasma cells. Haematologica, 2009, 94, 78-86.	1.7	106
394	Genetic Associations with Bortezomib Mediated Neuropathy in Multiple Myeloma.. Blood, 2009, 114, 1794-1794.	0.6	18
395	Developing a SNP Classifier for Predicting Peripheral Neuropathy by Bortezomib in Multiple Myeloma Patients.. Blood, 2009, 114, 1800-1800.	0.6	1
396	Natural History of Multiple Myeloma Relapsing After Therapy with IMiDs and Bortezomib: A Multicenter International Myeloma Working Group Study.. Blood, 2009, 114, 2878-2878.	0.6	18

#	Article	IF	Citations
397	The Addition of Thalidomide to the Induction Treatment of Newly Presenting Myeloma Patients Increases the CR Rate Which Is Likely to Translate Into Improved PFS and OS.. Blood, 2009, 114, 352-352.	0.6	14
398	Genetic Variations Associated with Overall and Progression-Free Survival in Multiple Myeloma Patients Treated with Thalidomide Combinations.. Blood, 2009, 114, 426-426.	0.6	2
399	Abnormal Serum IgA Kappa / IgA Lambda Ratios at Maximum Response Predict Poor Progression Free Survival in Myeloma Patients.. Blood, 2009, 114, 4879-4879.	0.6	1
400	Gene Expression Profiling Classifies Splenic Marginal Zone Lymphoma and Hairy Cell Leukemia-Variant as Related Diseases That Are Distinct From Typical Hairy Cell Leukemia.. Blood, 2009, 114, 3467-3467.	0.6	0
401	Global Methylation Array Analysis of Multiple Myeloma Samples Indicate An Alteration of Epigenetics During the Transition From MGUS to Myeloma and An Increased Frequency of Gene Methylation in t(4;14) Myeloma.. Blood, 2009, 114, 121-121.	0.6	0
402	Inhibition of HDACs and Aminopeptidases Is Highly Synergistic in Myeloma Cells Resulting in Cell Death Via the Upregulation of BIRC3, a Key Mediator of NF-KappaB Signalling.. Blood, 2009, 114, 607-607.	0.6	1
403	Molecular Characterization of Human Multiple Myeloma Cell Lines by Genome-Wide Profiling.. Blood, 2009, 114, 1793-1793.	0.6	0
404	UTX, a Histone Demethylase, Is Inactivated through Homozygous Deletion, Mutation, and DNA Methylation in Multiple Myeloma.. Blood, 2009, 114, 1798-1798.	0.6	0
405	Serum Free Light Chain and Serum Heavy / Light Chain Ratios Are Independently Prognostic in Multiple Myeloma Patients.. Blood, 2009, 114, 4880-4880.	0.6	0
406	Non-Cycling Cells From Acute Myeloid Leukemia Patients Harbor the FLT3-ITD Mutation and Are Insensitive to TKI258, a Potent FLT3-Directed Inhibitor, in Vitro.. Blood, 2009, 114, 479-479.	0.6	1
407	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
408	Prevention of thalidomide- and lenalidomide-associated thrombosis in myeloma. Leukemia, 2008, 22, 414-423.	3.3	787
409	Position statement on the use of bortezomib in multiple myeloma. International Journal of Laboratory Hematology, 2008, 30, 1-10.	0.7	15
410	Survival and outcome of blastoid variant myeloma following treatment with the novel thalidomide containing regime DTâ€PACE. European Journal of Haematology, 2008, 81, 432-436.	1.1	16
411	Streptolysin-O reversible permeabilisation is an effective method to transfect siRNAs into myeloma cells. Journal of Immunological Methods, 2008, 333, 147-155.	0.6	19
412	Lenalidomide: A new therapy for multiple myeloma. Cancer Treatment Reviews, 2008, 34, 283-291.	3.4	39
413	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
414	Characterising the TP53-deleted subgroup of chronic lymphocytic leukemia: an analysis of additional cytogenetic abnormalities detected by interphase fluorescencein situhybridisation and array-based comparative genomic hybridisation. Leukemia and Lymphoma, 2008, 49, 1879-1886.	0.6	12

#	Article	IF	Citations
415	Outcome of high-dose cytarabine-based induction therapy followed by hematopoietic stem cell transplantation in acute myeloid leukemia: influence of karyotype. Leukemia and Lymphoma, 2008, 49, 2284-2290.	0.6	2
416	Untangling the unfolded protein response. Cell Cycle, 2008, 7, 865-869.	1.3	76
417	An update on drug combinations for treatment of myeloma. Expert Opinion on Investigational Drugs, 2008, 17, 1-12.	1.9	10
418	Deletions of CDKN2C in Multiple Myeloma: Biological and Clinical Implications. Clinical Cancer Research, 2008, 14, 6033-6041.	3.2	88
419	Genetic associations with thalidomide mediated venous thrombotic events in myeloma identified using targeted genotyping. Blood, 2008, 112, 4924-4934.	0.6	65
420	Genetic Variation in ADME Genes Is Associated with Thalidomide Related Peripheral Neuropathy in Multiple Myeloma Patients.. Blood, 2008, 112, 1675-1675.	0.6	3
421	Economic Evaluation of Lenalidomide Combined with Dexamethasone for the Treatment of Multiple Myeloma in the UK. Blood, 2008, 112, 2400-2400.	0.6	7
422	CRD: A Phase 1 Dose Escalation Study to Determine the Maximum Tolerated Dose of Cyclophosphamide in Combination with Lenalidomide and Dexamethasone in Relapsed/Refractory Myeloma. Blood, 2008, 112, 3707-3707.	0.6	4
423	Maintenance Thalidomide May Improve Progression Free but Not Overall Survival; Results from the Myeloma IX Maintenance Randomisation. Blood, 2008, 112, 656-656.	0.6	35
424	An Early CT-Diagnosis Based Treatment Strategy for Invasive Fungal Infection in Allogeneic Transplant Recipients Using Caspofungin First Line: An Effective Strategy with Low Mortality.. Blood, 2008, 112, 1149-1149.	0.6	0
425	Autologous Transplantation Is the Optimum Approach to the Management of Myeloma Patients with Extramedullary Disease at Presentation. Blood, 2008, 112, 3313-3313.	0.6	0
426	The Impact of Constitutional Copy Number Variants in Myeloma. Blood, 2008, 112, 496-496.	0.6	1
427	Molecular Characterization of Human Multiple Myeloma Cell Lines by Genome-Wide Profiling Identifies Kinase Pathway Alterations.. Blood, 2008, 112, 1694-1694.	0.6	0
428	Analytical Approaches for the BOAC SNP Panel Association with Progression Free Survival in Myeloma. Blood, 2008, 112, 2715-2715.	0.6	0
429	High Resolution Genomic Profiling Using Single Nucleotide Polymorphism Microarrays Identifies Multiple Novel Genomic Minimally Deleted Regions in Multiple Myeloma. Blood, 2008, 112, 625-625.	0.6	0
430	High Resolution Genomic Profiling Using Single Nucleotide Polymorphism Microarrays Reveals Novel Genomic Lesions in Hairy Cell Leukaemia and Hairy-Cell Leukaemia Variant. Blood, 2008, 112, 3136-3136.	0.6	0
431	A Single Institutionâ€™s Experience with Allogeneic Haematopoietic Stem Cell Transplantation for Acute Lymphoblastic Leukaemia: Analysis of Factors Leading to Improved Outcome. Blood, 2008, 112, 4322-4322.	0.6	0
432	XBP1 Expression Is An Important Prognostic Factor for Newly Diagnosed Myeloma Patients.. Blood, 2008, 112, 1686-1686.	0.6	0

#	Article	IF	Citations
433	Large Scale Evaluation of Genetic Variation and the Risk of Multiple Myeloma.. Blood, 2008, 112, 1679-1679.	0.6	0
434	Homozygous Deletions Can Be Used to Define a Cell Death Specific Gene Expression Signature Able to Predict Outcome in Myeloma. Blood, 2008, 112, 2725-2725.	0.6	0
435	Genome-Wide Profiling of DNA Copy Number Variation in CLL Cases Lacking 17p- (TP53) or 11q- (ATM) Abnormalities Selected from the CLL4 Study. Blood, 2008, 112, 3140-3140.	0.6	0
436	Inhibitors of Heat Shock Protein 72 Induction Enhance Apoptosis Induced by Hsp90 Inhibitors. Blood, 2008, 112, 2625-2625.	0.6	1
437	Collection of peripheral blood stem cells in new patients with myeloma receiving minimal or no prior cytoreductive therapy. Hematology, 2007, 12, 113-115.	0.7	3
438	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
439	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
440	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
441	Immunoglobulin gene rearrangements and the pathogenesis of multiple myeloma. Blood, 2007, 110, 3112-3121.	0.6	157
442	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
443	Demonstration of changes in plasma cell subsets in multiple myeloma. Haematologica, 2007, 92, 1135-1138.	1.7	31
444	Non-Hodgkin Lymphoma Secondary to Cancer Chemotherapy: Table 1.. Cancer Epidemiology Biomarkers and Prevention, 2007, 16, 377-380.	1.1	65
445	Haplotypic structure across the ÎºBÎ± gene (NFKBIA) and association with multiple myeloma. Cancer Letters, 2007, 246, 92-99.	3.2	32
446	Peripheral blood or bone marrow cells in reduced-intensity or myeloablative conditioning allogeneic HLA identical sibling donor transplantation for multiple myeloma. Haematologica, 2007, 92, 1513-1518.	1.7	34
447	Haplotype uncertainty in association studies. Genetic Epidemiology, 2007, 31, 348-357.	0.6	13
448	RAD51 homologous recombination repair gene haplotypes and risk of acute myeloid leukaemia. Leukemia Research, 2007, 31, 169-174.	0.4	33
449	A single-tube six-colour flow cytometry screening assay for the detection of minimal residual disease in myeloma. Leukemia, 2007, 21, 2046-2049.	3.3	39
450	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

#	Article	IF	Citations
451	Proteomic evaluation of pathways associated with dexamethasone-mediated apoptosis and resistance in multiple myeloma. British Journal of Haematology, 2007, 139, 559-567.	1.2	34
452	Different IGH Rearrangement and Somatic Hypermutation Patterns in Hairy-Cell Leukemia, Hairy-Cell Leukemia Variant and Splenic Marginal Zone Lymphoma.. Blood, 2007, 110, 2082-2082.	0.6	2
453	Aminopeptidase Inhibition as a Targeted Treatment Strategy in Myeloma.. Blood, 2007, 110, 2505-2505.	0.6	2
454	Thalidomide Combinations Improve Response Rates; Results from the MRC IX Study.. Blood, 2007, 110, 3593-3593.	0.6	33
455	Integration of Gene Mapping and Expression Arrays Identifies Mechanisms by Which Genes Are Dysregulated as a Result of Copy Number Loss and Gain Associated with IgH Translocations in Multiple Myeloma.. Blood, 2007, 110, 395-395.	0.6	1
456	The Combination of Velcade, Idarubicin and Melphalan (VIM) Demonstrates Significant Clinical Activity in Relapsed/Refractory Myeloma Patients.. Blood, 2007, 110, 2727-2727.	0.6	8
457	High Dose Methylprednisolone and Rituximab Is an Effective Therapy in Advanced Refractory Chronic Lymphocytic Leukaemia Resistant to Fludarabine Therapy.. Blood, 2007, 110, 3125-3125.	0.6	0
458	The Tyrosine Kinase Inhibitor TK258 Differentially Targets Proliferating Human Hematopoietic Progenitor Cells but Does Not Eliminate the Quiescent Fraction.. Blood, 2007, 110, 3356-3356.	0.6	0
459	Immunoglobulin Gene Translocations Are Frequent in Unclassifiable CD5+ Small B-Cell Leukemias and in Particular, a Novel Subtype Characterised by t(14;19)(q32;q13) and Deregulation of BCL3 Is Described.. Blood, 2007, 110, 4230-4230.	0.6	0
460	Thalidomide in Combination with Idarubicin, Dexamethasone and Etoposide (TIDE) Is an Effective Oral Combination in Heavily Pre-Treated Myeloma Patients.. Blood, 2007, 110, 4841-4841.	0.6	0
461	Correlation between TP53 Deletion and Protein Overexpression in Chronic Lymphocytic Leukaemia and Its Impact on Prognostic.. Blood, 2007, 110, 2076-2076.	0.6	0
462	A Phase I Study of CHR-2797, an Orally Active Aminopeptidase Inhibitor in Elderly and/or Treatment Refractory Patients with Acute Myeloid Leukemia or Multiple Myeloma.. Blood, 2007, 110, 443-443.	0.6	2
463	Genome-Wide Identification of Gene Expression Networks Affected by Genomic Changes in Multiple Myeloma.. Blood, 2007, 110, 2494-2494.	0.6	4
464	Mutation and Methylation Analysis of WWOX and CYLD on 16q; Potential Tumor Suppressor Genes in Myeloma.. Blood, 2007, 110, 2473-2473.	0.6	0
465	Maintenance Chemotherapy (MC) after Autotransplantation in First-Remission Acute Lymphoblastic Leukemia (ALL): Single-Center Experience of 100 Patients.. Blood, 2007, 110, 2874-2874.	0.6	0
466	Screening of Homozygous Deletions Identifies Key Deregulated Genes and Pathways in Multiple Myeloma.. Blood, 2007, 110, 2474-2474.	0.6	0
467	An Integrated Pharmacogenomic Strategy for the Definition of Thalidomide Response Signatures in Presenting Cases of Multiple Myeloma.. Blood, 2007, 110, 2493-2493.	0.6	0
468	Defining Complete Response in Multiple Myeloma: Role of the Serum Free Light Chain Assay and Multiparameter Flow Cytometry.. Blood, 2007, 110, 1479-1479.	0.6	1

#	Article	IF	Citations
469	Proteasome inhibition and multiple myeloma. Current Opinion in Investigational Drugs, 2007, 8, 447-51.	2.3	6
470	The combination of cyclophosphomide, thalidomide and dexamethasone is an effective alternative to cyclophosphamide â€“ vincristine â€“ doxorubicin â€“ methylprednisolone as induction chemotherapy prior to autologous transplantation for multiple myeloma: a case-matched analysis. Leukemia and Lymphoma, 2006, 47, 2335-2338.	0.6	43
471	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
472	Advances in oral therapy for multiple myeloma. Lancet Oncology, The, 2006, 7, 316-325.	5.1	42
473	Neuron-specific enolase expression in multiple myeloma. Lancet Oncology, The, 2006, 7, 960.	5.1	3
474	Use of Single Nucleotide Polymorphismâ€“Based Mapping Arrays to Detect Copy Number Changes and Loss of Heterozygosity in Multiple Myeloma. Clinical Lymphoma and Myeloma, 2006, 7, 186-192.	1.4	17
475	Haplotypic variation in MRE11, RAD50 and NBS1 and risk of non-Hodgkin's lymphoma. Leukemia and Lymphoma, 2006, 47, 2567-2583.	0.6	13
476	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
477	Association between non-Hodgkin lymphoma and haplotypes in the TNF region. British Journal of Haematology, 2006, 133, 293-300.	1.2	28
478	Deletion of chromosome 13 detected by conventional cytogenetics is a critical prognostic factor in myeloma. Leukemia, 2006, 20, 1610-1617.	3.3	141
479	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
480	New Developments in the Epidemiology of Cancer Prognosis: Traditional and Molecular Predictors of Treatment Response and Survival. Cancer Epidemiology Biomarkers and Prevention, 2006, 15, 2042-2046.	1.1	15
481	Fine Mapping and Expression Analysis of Chromosome 1 with the Aim of Defining Critically Deregulated Genes Important in the Pathogenesis of Myeloma.. Blood, 2006, 108, 112-112.	0.6	2
482	The Combination of Cyclophosphamide, Velcade and Dexamethasone (CVD) Induces High Response Rates with Minimal Toxicity Compared to Velcade Alone (V) and Velcade Plus Dexamethasone (VD).. Blood, 2006, 108, 3537-3537.	0.6	6
483	Lenolidamide (Revlimid), in Combination with Cyclophosphamide and Dexamethasone (CRD) Is an Effective Regimen for Heavily Pre-Treated Myeloma Patients.. Blood, 2006, 108, 3555-3555.	0.6	2
484	Bortezomib (Velcade?) in the treatment of multiple myeloma. Therapeutics and Clinical Risk Management, 2006, 2, 271-279.	0.9	197
485	Genetic variation in Th1/Th2 cytokines and the risk of developing childhood leukaemia. FASEB Journal, 2006, 20, LB123.	0.2	0
486	Lymphocyte Count on Day 56 Predicts for Reduced Risk of Infection Related Mortality Following Non-Myeloablative Allogeneic Haematopoietic Stem Cell Transplantation.. Blood, 2006, 108, 5324-5324.	0.6	0

#	Article	IF	Citations
487	Prophylactic Defibrotide in Allogeneic Stem Cell Transplantation: Low Morbidity and Zero Mortality from Veno-Occlusive Disease.. Blood, 2006, 108, 2968-2968.	0.6	0
488	Abnormalities of 16q in Multiple Myeloma Are Associated with Poor Prognosis: 500K Gene Mapping and Expression Correlations Identify Two Potential Tumor Suppressor Genes, WWOX and CYLD.. Blood, 2006, 108, 110-110.	0.6	1
489	The Genetic Contribution to the Aetiology of Thalidomide Associated VTE.. Blood, 2006, 108, 246-246.	0.6	9
490	Incidence of Second Malignant Neoplasms (SMN) after Haematopoetic Stem Cell Transplant.. Blood, 2006, 108, 551-551.	0.6	3
491	Sub-Classification of Hyperdiploid Myeloma Using Global Gene Expression Profiling and SNP-Based Mapping Arrays.. Blood, 2006, 108, 3390-3390.	0.6	0
492	Pregnancy Outcome Following Haematopoietic Stem Cell Transplantation.. Blood, 2006, 108, 3367-3367.	0.6	0
493	The role of second autografts in the management of myeloma at first relapse. Haematologica, 2006, 91, 141-2.	1.7	62
494	The bone marrow microenvironment influences the differential chemokine receptor expression of normal and neoplastic plasma cells. Blood, 2005, 105, 4895-4896.	0.6	5
495	Antimyeloma activity of heat shock protein-90 inhibition. Blood, 2005, 107, 1092-1100.	0.6	278
496	Response to Goodman et al.: Amyloidosis, not myeloma. British Journal of Haematology, 2005, 129, 159-160.	1.2	0
497	Haplotypes in the tumour necrosis factor region and myeloma. British Journal of Haematology, 2005, 129, 358-365.	1.2	42
498	Polymorphisms in cytochrome P450 17A1 and risk of non-Hodgkin lymphoma. British Journal of Haematology, 2005, 129, 618-621.	1.2	20
499	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
500	Evolving treatment strategies for myeloma. British Journal of Cancer, 2005, 92, 217-221.	2.9	18
501	Non-Hodgkin's lymphoma, obesity and energy homeostasis polymorphisms. British Journal of Cancer, 2005, 93, 811-816.	2.9	79
502	Age has a profound effect on the incidence and significance of chromosome abnormalities in myeloma. Leukemia, 2005, 19, 1634-1642.	3.3	73
503	Benzene and the hemopoietic stem cell. Chemico-Biological Interactions, 2005, 153-154, 217-222.	1.7	26
504	Preclinical evaluation of the proteasome inhibitor bortezomib in cancer therapy. Cancer Cell International, 2005, 5, 18.	1.8	196

#	Article	IF	Citations
505	Risk of Non-Hodgkin Lymphoma Associated with Polymorphisms in Folate-Metabolizing Genes. Cancer Epidemiology Biomarkers and Prevention, 2005, 14, 2999-3003.	1.1	72
506	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
507	International Staging System for Multiple Myeloma. Journal of Clinical Oncology, 2005, 23, 3412-3420.	0.8	2,404
508	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
509	A Phase I Trial of CHIR-258, a Multitargeted RTK Inhibitor, in Acute Myeloid Leukemia (AML).. Blood, 2005, 106, 2794-2794.	0.6	1
510	Comparison of IgH Gene Rearrangement Configuration between Hairy Cell Leukemia (HCL) and Hairy Cell Leukemia Variant (HCL-v).. Blood, 2005, 106, 4997-4997.	0.6	0
511	A Comparative Analysis of Outcomes in Primary Myelodysplastic Syndrome (MDS) and Therapy-Related MDS Reveals Two Subgroups with Differing Risk Profiles: Implications for the Application of a Prognostic Classification.. Blood, 2005, 106, 2532-2532.	0.6	1
512	Insights into the Basis of Chromosomal Imbalances during the Clonal Evolution of Multiple Myeloma Using SNP Array Analysis.. Blood, 2005, 106, 621-621.	0.6	0
513	SNP Genotypes as Prognostic Factors Associated with Staging in Myeloma: The First Combined SNP Association Analysis of SWOG, ECOG, and MRC Clinical Trials from the Bank on a Cure.. Blood, 2005, 106, 3489-3489.	0.6	0
514	Development of a Panel of 3,500 SNP Based Functional Genetic Variants Relevant to the Etiology and Outcome in Multiple Myeloma.. Blood, 2005, 106, 620-620.	0.6	0
515	Ultra-Rapid, High-Throughput Molecular Diagnostics in Hemato-Oncology.. Blood, 2005, 106, 3270-3270.	0.6	0
516	Combinations of ZAP-70, CD38 and IgVH Mutational Status as Predictors of Time to First Treatment in CLL.. Blood, 2005, 106, 711-711.	0.6	0
517	B- Cell Chronic Lymphocytic Leukaemia Complicated by Aggressive T-Cell Lymphoma: Clinical and Molecular Analysis of a Rare Variant of Richterâ€™s Syndrome.. Blood, 2005, 106, 4999-4999.	0.6	0
518	Identification of Collaborating Oncogeneic Events Leading to Disease Progression in Myeloma Cases with a t(4;14) and t(11;14) Using SNP and Gene Expression Arrays.. Blood, 2005, 106, 1542-1542.	0.6	0
519	Status of Chromosome 13 in Multiple Myeloma: Integrated Approach Using SNP Mapping Array and Gene Expression Array.. Blood, 2005, 106, 1563-1563.	0.6	0
520	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
521	DNA mismatch repair pathway defects in the pathogenesis and evolution of myeloma. Carcinogenesis, 2004, 25, 1795-1803.	1.3	29
522	Tobacco and Alcohol Consumption and the Risk of Non-Hodgkin Lymphoma. Cancer Causes and Control, 2004, 15, 771-780.	0.8	55

#	Article	IF	Citations
523	Translocation t(11;14) in multiple myeloma: Analysis of translocation breakpoints on der(11) and der(14) chromosomes suggests complex molecular mechanisms of recombination. Genes Chromosomes and Cancer, 2004, 39, 151-155.	1.5	13
524	Association of metabolic gene polymorphisms with tobacco consumption in healthy controls. International Journal of Cancer, 2004, 110, 266-270.	2.3	21
525	Genetic variation in XPD predicts treatment outcome and risk of acute myeloid leukemia following chemotherapy. Blood, 2004, 104, 3872-3877.	0.6	108
526	Economic Evaluation of Bortezomib (VELCADE) for Relapsed and Refractory Multiple Myeloma.. Blood, 2004, 104, 268-268.	0.6	5
527	Innovative approaches for diagnosis and monitoring. , 2004, , 164-178.		0
528	Responses to Induction Chemotherapy and High Dose Melphalan Can Be Incorporated within a Risk Stratification Model and Help to Identify Myeloma Patients Who May Benefit from Novel Therapies.. Blood, 2004, 104, 2394-2394.	0.6	0
529	Proteomic Evaluation of Pathways Associated with Dexamethasone-Induced Apoptosis and Resistance in Multiple Myeloma.. Blood, 2004, 104, 642-642.	0.6	5
530	Newly Diagnosed Myeloma Pateints Are at Risk of Venous Thrombotic Events - High Risk Patients Need To Be Identified and Recieve Thromboprophylaxis: The MRC Experience.. Blood, 2004, 104, 2395-2395.	0.6	1
531	The Bone Marrow Microenvironment Influences the Differential Chemokine Receptor Expression of Normal and Neoplastic Plasma Cells.. Blood, 2004, 104, 2353-2353.	0.6	12
532	IgH Rearrangements and Mutational Status in Class-Switched IgG B-CLL.. Blood, 2004, 104, 2793-2793.	0.6	1
533	Follow up Analysis for MRC Myeloma VII Trial.. Blood, 2004, 104, 927-927.	0.6	1
534	Epigenetic consequences of AML1-ETO action at the human c-FMS locus. EMBO Journal, 2003, 22, 2798-2809.	3.5	58
535	Poor metabolizer status at the cytochrome p450 2c19 and 2d6 loci does not modulate susceptibility to therapy-related acute myeloid leukaemia. British Journal of Haematology, 2003, 121, 192-194.	1.2	8
536	The use of genetic microarray analysis to classify and predict prognosis in haematological malignancies. International Journal of Laboratory Hematology, 2003, 25, 209-220.	0.2	14
537	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
538	Genomic characterization of the chromosomal breakpoints of t(4;14) of multiple myeloma suggests more than one possible aetiological mechanism. Oncogene, 2003, 22, 1103-1113.	2.6	21
539	The Detection of t(14;18) in Archival Lymph Nodes. Journal of Molecular Diagnostics, 2003, 5, 168-175.	1.2	39
540	Aetiology of bone disease and the role of bisphosphonates in multiple myeloma. Lancet Oncology, The, 2003, 4, 284-292.	5.1	66

#	Article	IF	Citations
541	High-Dose Chemotherapy with Hematopoietic Stem-Cell Rescue for Multiple Myeloma. New England Journal of Medicine, 2003, 348, 1875-1883.	13.9	1,648
542	Differential transcription factor occupancy but evolutionarily conserved chromatin features at the human and mouse M-CSF (CSF-1) receptor loci. Nucleic Acids Research, 2003, 31, 5805-5816.	6.5	17
543	Insights into the multistep transformation of MGUS to myeloma using microarray expression analysis. Blood, 2003, 102, 4504-4511.	0.6	212
544	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
545	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
546	CYP1A1*2B (Val) allele is overrepresented in a subgroup of acute myeloid leukemia patients with poor-risk karyotype associated with NRAS mutation, but not associated withFLT3 internal tandem duplication. Blood, 2003, 101, 2770-2774.	0.6	41
547	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
548	The Leukaemia-Associated AML1-ETO Fusion Protein Acts as a Sequence Specific Transcriptional Repressor in vivo by Altering the Chromatin Structure of Target Genes. Clinical Science, 2003, 104, 30P-30P.	0.0	0
549	Polymorphic variation in GSTP1 modulates outcome following therapy for multiple myeloma. Blood, 2003, 102, 2345-2350.	0.6	90
550	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
551	Polymorphisms in the thymidylate synthase and serine hydroxymethyltransferase genes and risk of adult acute lymphocytic leukemia. Blood, 2002, 99, 3786-3791.	0.6	210
552	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
553	Long Distance Vectorette PCR (LDV PCR). , 2002, 192, 275-284.		1
554	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
555	Follicular lymphoma with a novel t(14;18) breakpoint involving the immunoglobulin heavy chain switch mu region indicates an origin from germinal center B cells. Blood, 2002, 99, 716-718.	0.6	24
556	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
557	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
558	Minimal residual disease monitoring in multiple myeloma. Best Practice and Research in Clinical Haematology, 2002, 15, 197-222.	0.7	44

#	Article	IF	Citations
559	Metabolic Enzyme Polymorphisms and Susceptibility to Acute Leukemia in Adults. Molecular Diagnosis and Therapy, 2002, 2, 79-92.	3.3	26
560	Myeloma aetiology and epidemiology. Biomedicine and Pharmacotherapy, 2002, 56, 223-234.	2.5	72
561	High dose immunosuppressive therapy and stem cell transplantation in autoimmune and inflammatory diseases. International Immunopharmacology, 2002, 2, 399-414.	1.7	7
562	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
563	Isotype class switching and the pathogenenesis of multiple myeloma. Hematological Oncology, 2002, 20, 75-85.	0.8	25
564	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
565	Genetic polymorphisms in microsomal epoxide hydrolase â€˜and susceptibility to adult acute myeloid leukaemia â€˜with defined cytogenetic abnormalities. British Journal of Haematology, 2002, 116, 587-594.	1.2	21
566	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
567	Allele and haplotype frequency at human leucocyte antigen class I/II and immunomodulatory cytokine loci in patients with myelodysplasia and acute myeloid leukaemia: in search of an autoimmune aetiology. British Journal of Haematology, 2002, 117, 541-545.	1.2	13
568	A molecular study of the t(4;14) in multiple myeloma. British Journal of Haematology, 2002, 118, 514-520.	1.2	29
569	Patients entered into MRC AML trials are biologically representative of the totality of the disease in the UK. International Journal of Laboratory Hematology, 2002, 24, 263-265.	0.2	5
570	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
571	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
572	Thalidomide and immunomodulatory derivatives augment natural killer cell cytotoxicity in multiple myeloma. Blood, 2001, 98, 210-216.	0.6	869
573	Interleukin 6, tumour necrosis factor Î± and lymphotoxin Î± polymorphisms in monoclonal gammopathy of uncertain significance and multiple myeloma. British Journal of Haematology, 2001, 112, 249-250.	1.2	13
574	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
575	Autologous stem cell transplantation for rapidly progressive jo-1-positive polymyositis with long-term follow-up. British Journal of Haematology, 2001, 113, 840-841.	1.2	25
576	Karyotype and age in acute myeloid leukemia.. Cancer Genetics and Cytogenetics, 2001, 126, 155-161.	1.0	59

#	Article	IF	Citations
577	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
578	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
579	The interleukin-6 receptor alpha-chain (CD126) is expressed by neoplastic but not normal plasma cells. Blood, 2000, 96, 3880-3886.	0.6	78
580	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
581	Polymorphic variation within the glutathione S-transferase genes and risk of adult acute leukaemia. Carcinogenesis, 2000, 21, 43-47.	1.3	84
582	Variability of polymerase chain reaction detection of the bcl-2-IgH translocation in an international multicentre study. Annals of Oncology, 1999, 10, 1349-1354.	0.6	38
583	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
584	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
585	TRUE SPECTRUM OF 14q32 TRANSLOCATIONS IN MULTIPLE MYELOMA. British Journal of Haematology, 1998, 103, 1209-1210.	1.2	7
586	p53 gene mutations in multiple myeloma.. Journal of Clinical Pathology, 1997, 50, 18-20.	2.1	16
587	Comparison of fluorescent consensus IgH PCR and alleleâ€specific oligonucleotide probing in the detection of minimal residual disease in childhood ALL. British Journal of Haematology, 1997, 97, 457-459.	1.2	23
588	Circulating plasma cells in multiple myeloma: characterization and correlation with disease stage. British Journal of Haematology, 1997, 97, 46-55.	1.2	165
589	A phase 1 study to address the safety and efficacy of granulocyte colony-stimulating factor for the mobilization of hematopoietic progenitor cells in active rheumatoid arthritis. Arthritis and Rheumatism, 1997, 40, 1838-1842.	6.7	25
590	Assessment of IgH PCR strategies in multiple myeloma.. Journal of Clinical Pathology, 1996, 49, 672-675.	1.0	44
591	Allele Imbalance at Tumour Suppressor Loci During the Indolent Phase of Follicle Centre Cell Lymphoma. Leukemia and Lymphoma, 1996, 22, 113-117.	0.6	1
592	Peripheral blood stem cell transplantation in myeloma using CD34 selected cells. Bone Marrow Transplantation, 1996, 17, 723-7.	1.3	20
593	Genetic abnormalities during transition from Helicobacter-pylori-associated gastritis to low-grade MALToma. Lancet, The, 1995, 345, 26-27.	6.3	75
594	The genetic and epigenetic mechanisms underlying the behavior of myeloma. , 0, , 48-63.		0

#	Article	IF	Citations
595	The evolving role and utility of off-label drug use in multiple myeloma. Exploration of Targeted Anti-tumor Therapy, 0, , .	0.5	0