Maurizio Zangari

List of Publications by Year in descending order

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187 6,978 36 81 papers citations h-index g-index

188 188 188 6428
all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	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
2	Clinical implications of loss of bone marrow minimal residual disease negativity in multiple myeloma. Blood Advances, 2022, 6, 808-817.	2.5	14
3	Tandem autologous stem cell transplantation in patients with persistent bone marrow minimal residual disease after first transplantation in multiple myeloma. American Journal of Hematology, 2022, 97, .	2.0	O
4	Feasibility of Outpatient Stem Cell Transplantation in Multiple Myeloma and Risk Factors Predictive of Hospital Admission. Journal of Clinical Medicine, 2022, 11, 1640.	1.0	6
5	First―versus secondâ€generation Bruton tyrosine kinase inhibitors in Waldenström's Macroglobulinemia: A systematic review and metaâ€analysis. American Journal of Hematology, 2022, 97, 942-950.	2.0	5
6	Epigenomic translocation of H3K4me3 broad domains over oncogenes following hijacking of super-enhancers. Genome Research, 2022, 32, 1343-1354.	2.4	8
7	Enrollment of Black Participants in Pivotal Clinical Trials Supporting US Food and Drug Administration Approval of Chimeric Antigen Receptor–T Cell Therapy for Hematological Malignant Neoplasms. JAMA Network Open, 2022, 5, e228161.	2.8	22
8	Clinical efficacy of sequencing CD38 targeting monoclonal antibodies in relapsed refractory multiple myeloma: A multiâ€institutional experience. American Journal of Hematology, 2022, 97, .	2.0	4
9	Chimeric Antigen Receptor T-Cell Therapy in Multiple Myeloma—Challenges and Potential Solutions. JAMA Oncology, 2022, 8, 823.	3.4	6
10	Monitoring treatment response and disease progression in myeloma with circulating cellâ€free DNA. European Journal of Haematology, 2021, 106, 230-240.	1.1	21
11	The molecular make up of smoldering myeloma highlights the evolutionary pathways leading to multiple myeloma. Nature Communications, 2021, 12, 293.	5.8	54
12	Bone remineralization of lytic lesions in multiple myeloma – The Arkansas experience. Bone, 2021, 146, 115876.	1.4	4
13	Salvage autologous stem cell transplantation in daratumumab refractory multiple myeloma (MM) Journal of Clinical Oncology, 2021, 39, e20031-e20031.	0.8	1
14	Persistent bone marrow minimal residual disease as a "highâ€riskâ€disease feature in multiple myeloma. American Journal of Hematology, 2021, 96, E341-E344.	2.0	4
15	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
16	PHF19 inhibition as a therapeutic target in multiple myeloma. Current Research in Translational Medicine, 2021, 69, 103290.	1.2	5
17	Salvage Autologous Stem Cell Transplantation in Daratumumab-Refractory Multiple Myeloma. Cancers, 2021, 13, 4019.	1.7	9
18	Predicting risk of progression in relapsed multiple myeloma using traditional risk models, focal lesion assessment with PET-CT and minimal residual disease status. Haematologica, 2021, 106, 0-0.	1.7	2

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19	Eight-Color Flow Cytometry Phenotypic Markers and Disease Progression in Monoclonal Gammopathy of Unknown Significance. Blood, 2021, 138, 2713-2713.	0.6	O
20	N-Cadherin Stabilizes \hat{l}^2 -Catenin and Promotes \hat{l}^2 -Catenin/TCF Transcriptional Activation and Cell Adhesion-Mediated Drug Resistance in Multiple Myeloma. Blood, 2021, 138, 1572-1572.	0.6	0
21	Enrollment of Black Americans in Pivotal Clinical Trials Supporting Food and Drug Administration (FDA) Chimeric Antigen Receptor (CAR)-T Cell Therapy Approval in Hematological Malignancies. Blood, 2021, 138, 566-566.	0.6	4
22	Concomitant Deletion of Short Arm (del 1p) and Amplification or Gain (1q21) of Chromosome 1 By Fluorescence in Situ Hybridization (FISH) Is Associated with Poor Clinical Outcome. Blood, 2021, 138, 1627-1627.	0.6	0
23	Ethnic Disparities in AL Amyloidosis Outcomes Among Hospitalized Patients in the United States. Blood, 2021, 138, 4110-4110.	0.6	1
24	Predicting risk of progression in relapsed multiple myeloma using traditional risk models, focal lesion assessment with PET-CT and minimal residual disease status. Haematologica, 2021, , .	1.7	0
25	Daratumumab in highâ€risk relapsed/refractory multiple myeloma patients: adverse effect of chromosome 1q21 gain/amplification and GEP70 status on outcome. British Journal of Haematology, 2020, 189, 67-71.	1.2	35
26	Innate Biomineralization. International Journal of Molecular Sciences, 2020, 21, 4820.	1.8	4
27	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
28	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
29	Primary Plasma Cell Neoplasm of the Kidney Without Formation of a Mass and Its Renal Manifestations: An Interstitial Variant of Renal Plasmacytoma?. Clinical Lymphoma, Myeloma and Leukemia, 2020, 20, e551-e555.	0.2	0
30	<i>BRAF</i> and <i>DIS3</i> 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
31	Long-term outcomes after autologous stem cell transplantation for multiple myeloma. Blood Advances, 2020, 4, 422-431.	2.5	66
32	Late Relapsing Multiple Myeloma \hat{a} %¥ 10 Years after Treatment on Total Therapy Protocols Are Associated with Good Outcome. Blood, 2020, 136, 11-12.	0.6	3
33	A meta-analysis of genome-wide association studies of multiple myeloma among men and women of African ancestry. Blood Advances, 2020, 4, 181-190.	2.5	16
34	Clinical implications of loss of minimal residual disease (MRD) negativity in multiple myeloma Journal of Clinical Oncology, 2020, 38, 8514-8514.	0.8	2
35	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
36	Feasibility of Outpatient Autologous Stem Cell Transplantation in Multiple Myeloma and Risk Factors Predicting Hospital Admission. Blood, 2020, 136, 44-44.	0.6	2

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37	Iron Trafficking through Macrophages Regulates Signaling Pathways in Myeloma. Blood, 2020, 136, 2-2.	0.6	O
38	Predicting Risk of Progression in Relapsed Multiple Myeloma Using Minimal Residual Disease Status and Focal Lesion Assessment with PET-CT. Blood, 2020, 136, 24-24.	0.6	0
39	An Improved Animal Model of Multiple Myeloma Bone Disease. Blood, 2020, 136, 31-31.	0.6	O
40	CST6 Is a Small Autocrine Molecule That Targets Myeloma Growth and Bone Destruction. Blood, 2020, 136, 21-21.	0.6	0
41	An acquired high-risk chromosome instability phenotype in multiple myeloma: Jumping 1q Syndrome. Blood Cancer Journal, 2019, 9, 62.	2.8	23
42	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
43	A Phase 1 Study of Intravenous Busulfan as a Conditioning Regimen for Multiple Myeloma. Cell Transplantation, 2019, 28, 1624-1631.	1.2	0
44	Bacteremias following autologous stem cell transplantation for multiple myeloma: Risk factors and outcomes. Transplant Infectious Disease, 2019, 21, e13052.	0.7	6
45	FRAX is a robust predictor of baseline vertebral fractures in multiple myeloma patients. Bone, 2019, 121, 134-138.	1.4	3
46	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
47	Effect of lowâ€dose bortezomib on bone formation in smouldering multiple myeloma. British Journal of Haematology, 2019, 184, 308-310.	1.2	2
48	Analysis of the Sub-Clonal Structure of Smoldering Myeloma over Time Provides a New Means of Disease Monitoring and Highlights Evolutionary Trajectories Leading to Myeloma. Blood, 2019, 134, 4333-4333.	0.6	2
49	EARLY Results of TOTAL Therapy 7 (TT7): High Response Rates of NEWLY Diagnosed High Risk Myeloma to Daratumumab. Blood, 2019, 134, 4569-4569.	0.6	4
50	The mTOR Component, Rictor, Is Regulated By the Microenvironment to Control Dormancy and Proliferative States in Myeloma Cells. Blood, 2019, 134, 4412-4412.	0.6	0
51	Long-Term Outcome of Total Therapy Regimens: Impact of Molecular Subgroups. Blood, 2019, 134, 3309-3309.	0.6	2
52	The Role of PHF19 As a Promoter of Tumorigenicity and Therapeutic Target in Multiple Myeloma. Blood, 2019, 134, 508-508.	0.6	0
53	Comprehensive Investigation of White Blood Cell and Gene Expression Profiles As Risk Factors for Multiple Myeloma in African Americans. Blood, 2019, 134, 4379-4379.	0.6	0
54	The Translational Switch of MYC Protein Aliases in Myeloma Tumor Cells. Blood, 2019, 134, 4390-4390.	0.6	O

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55	Eltrombopag Following Chemotherapy and G-CSF+/- Plerixafor for Mobilization and Collection of Hematopoietic Progenitor Cells (HPC) in Lymphoma and Myeloma Patients. Blood, 2019, 134, 5638-5638.	0.6	O
56	Kinase domain activation through gene rearrangement in multiple myeloma. Leukemia, 2018, 32, 2435-2444.	3.3	26
57	Surgical thyroparathyroidectomy prevents progression of 5TGM1 murine multiple myeloma in vivo. Journal of Bone Oncology, 2018, 12, 19-22.	1.0	3
58	Thymic PTH Increases After Thyroparathyroidectomy in C57BL/KaLwRij Mice. Endocrinology, 2018, 159, 1561-1569.	1.4	4
59	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
60	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
61	The presence of large focal lesions is a strong independent prognostic factor in multiple myeloma. Blood, 2018, 132, 59-66.	0.6	75
62	Visualizing collagen proteolysis by peptide hybridization: From 3D cell culture to in vivo imaging. Biomaterials, 2018, 183, 67-76.	5.7	49
63	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
64	Baseline and on-Treatment Bone Marrow Microenvironments Predict Myeloma Patient Outcomes and Inform Potential Intervention Strategies. Blood, 2018, 132, 1882-1882.	0.6	3
65	The Mutational Landscape of Primary Plasma Cell Leukemia. Blood, 2018, 132, 114-114.	0.6	2
66	Chromothripsis and Chromoplexy Are Associated with DNA Instability and Adverse Clinical Outcome in Multiple Myeloma. Blood, 2018, 132, 408-408.	0.6	3
67	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
68	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
69	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
70	Myeloma Patient-Derived Bone Marrow Serum Negatively Regulates Natural Killer Cell Activity. Blood, 2018, 132, 4468-4468.	0.6	0
71	Combination of Flow Cytometry and Functional Imaging for Monitoring of Residual Disease in Myeloma. Blood, 2018, 132, 3185-3185.	0.6	0
72	Extracting Prognostic Molecular Information from PET-CT Imaging of Multiple Myeloma Using Radiomic Approaches. Blood, 2018, 132, 1906-1906.	0.6	1

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73	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	O
74	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
75	An Acquired High-Risk Chromosome Instability Phenotype in Multiple Myeloma: Jumping 1q Syndrome. Blood, 2018, 132, 4489-4489.	0.6	1
76	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
77	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
78	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
79	The level of deletion 17p and bi-allelic inactivation of <i>TP53</i> has a significant impact on clinical outcome in multiple myeloma. Haematologica, 2017, 102, e364-e367.	1.7	57
80	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
81	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
82	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
83	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
84	Clonal selection and double-hit events involving tumor suppressor genes underlie relapse in myeloma. Blood, 2016, 128, 1735-1744.	0.6	170
85	Monoclonal antibody therapy in multiple myeloma: where do we stand and where are we going?. Immunotherapy, 2016, 8, 367-384.	1.0	6
86	The effects of proteasome inhibitors on bone remodeling in multiple myeloma. Bone, 2016, 86, 131-138.	1.4	39
87	Signatures of Mesenchymal Cell Lineages and Microenvironment Factors Are Dysregulated in High Risk Myeloma. Blood, 2016, 128, 2065-2065.	0.6	1
88	Concurrent Amplification of MYC and 1q21 in Multiple Myeloma: Focal and Segmental Jumping Translocations of MYC. Blood, 2016, 128, 3266-3266.	0.6	1
89	Extensive Regional Intra-Clonal Heterogeneity in Multiple Myeloma - Implications for Diagnostics, Risk Stratification and Targeted Treatment. Blood, 2016, 128, 3278-3278.	0.6	2
90	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

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91	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
92	The Clinical Impact of Macrofocal Disease in Multiple Myeloma Differs Between Presentation and Relapse. Blood, 2016, 128, 4431-4431.	0.6	8
93	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
94	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
95	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
96	High Risk Myeloma Is Characterized By the Bi-Allelic Inactivation of CDKN2C and RB1. Blood, 2016, 128, 4416-4416.	0.6	1
97	Rigosertib, a Pan RAS Inhibitor, Decreases Mouse and Human Myeloma Cell Growth in Preclinical Models. Blood, 2016, 128, 5664-5664.	0.6	1
98	The Metabolic Phenotype of Myeloma Plasma Cells Differs Between Active and Residual Disease States. Blood, 2016, 128, 4438-4438.	0.6	0
99	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
100	Evidence of an epigenetic origin for high-risk 1q21 copy number aberrations in multiple myeloma. Blood, 2015, 125, 3756-3759.	0.6	41
101	Four genes predict high risk of progression from smoldering to symptomatic multiple myeloma (SWOG S0120). Haematologica, 2015, 100, 1214-1221.	1.7	44
102	Drug Resistance in Hematologic Malignancies: Induction Mechanisms, Genetics, and Therapeutics. BioMed Research International, 2015, 2015, 1-2.	0.9	2
103	The Composition and Clinical Impact of Focal Lesions and Their Impact on the Microenvironment in Myeloma. Blood, 2015, 126, 1806-1806.	0.6	2
104	Melphalan Affects Genes Critical for Myeloma Survival, Homing, and Response to Cytokines and Chemokines. Blood, 2015, 126, 1808-1808.	0.6	2
105	Upfront 28-Day Metronomic Therapy for High-Risk Multiple Myeloma (HRMM). Blood, 2015, 126, 1843-1843.	0.6	1
106	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
107	Impact of Minimal Residual Disease in High and Standard Risk Multiple Myeloma. Blood, 2015, 126, 2979-2979.	0.6	2
108	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

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109	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
110	Thymus-Derived PTH (TPTH) Is Increased after Thyroparathyroidectomy in C57BL6/Kalwrij Mice and Modulates Mouse Sensitivity to 5TGM1 Myeloma Cell Line. Blood, 2015, 126, 5335-5335.	0.6	1
111	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
112	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
113	Deletion of TP53 (17p13) Is Associated with Poor Outcome for Newly Diagnosed High-Risk Multiple Myeloma. Blood, 2015, 126, 2982-2982.	0.6	0
114	Molecular Subtyping and Risk Stratification for the Classification of Myeloma. Blood, 2015, 126, 4173-4173.	0.6	0
115	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
116	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
117	Extending Metronomic Therapy to 28 Days (metro28) for Relapsed Refractory Multiple Myeloma (RRMM). Blood, 2015, 126, 5395-5395.	0.6	0
118	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
119	Parathyroid hormone receptor mediates the anti-myeloma effect of proteasome inhibitors. Bone, 2014, 61, 39-43.	1.4	11
120	Jumping translocations of $1q12$ in multiple myeloma: a novel mechanism for deletion of $17p$ in cytogenetically defined high-risk disease. Blood, 2014, 123, 2504-2512.	0.6	45
121	Curing Multiple Myeloma (MM) with Total Therapy (TT). Blood, 2014, 124, 195-195.	0.6	3
122	Further Evolution of Metronomic Therapy Extended to 28 Days (Metro28) for Relapsed Refractory Multiple Myeloma (RRMM). Blood, 2014, 124, 2128-2128.	0.6	1
123	Waldenstrom's Macroglobulinemia Associated Bone Disease the UAMS Experience. Blood, 2014, 124, 2999-2999.	0.6	3
124	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
125	Characterization of the Mutational Landscape of Multiple Myeloma Using Comprehensive Genomic Profiling. Blood, 2014, 124, 3418-3418.	0.6	3
126	Targeted MEK Inhibition in Patients with Previously Treated Multiple Myeloma. Blood, 2014, 124, 4775-4775.	0.6	4

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127	Evidence of an Epigenetic Origin for High-Risk 1q21 Copy Number Aberrations in Multiple Myeloma. Blood, 2014, 124, 725-725.	0.6	1
128	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
129	Surgical Control/Cure of 5TGM1 Murine Multiple Myeloma Model By Thyroparathyroidectomy. Blood, 2014, 124, 3387-3387.	0.6	O
130	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
131	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
132	Low-Dose 28-Day Metronomically Scheduled Therapy (METRO) for Newly Diagnosed High-Risk Multiple Myeloma: A Pilot Study. Blood, 2014, 124, 5770-5770.	0.6	0
133	NEK2 Induces Drug Resistance Mainly through Activation of Efflux Drug Pumps and Is Associated with Poor Prognosis in Myeloma and Other Cancers. Cancer Cell, 2013, 23, 48-62.	7.7	232
134	Could hypoxia increase the prevalence of thrombotic complications in polycythemia vera?. Blood Coagulation and Fibrinolysis, 2013, 24, 311-316.	0.5	22
135	Mechanisms of Thrombosis in Paraproteinemias: The Effects of Immunomodulatory Drugs. Seminars in Thrombosis and Hemostasis, 2012, 38, 768-779.	1.5	11
136	Impact of bortezomib on bone health in myeloma: A review of current evidence. Cancer Treatment Reviews, 2012, 38, 968-980.	3.4	44
137	Phase I Exploratory Study of IV Formulation of Panobinostat in Combination with Bortezomib in Relapsed/Refractory Multiple Myeloma Patients: Effect On Serum PTH and Gene Expression Profiling (GEP) Studies. Blood, 2012, 120, 4073-4073.	0.6	1
138	Low Venous Thromboembolic Risk With Bortezomib in Multiple Myeloma and Potential Protective Effect With Thalidomide/Lenalidomide-based Therapy: Review of Data From Phase 3 Trials and Studies of Novel Combination Regimens. Clinical Lymphoma, Myeloma and Leukemia, 2011, 11, 228-236.	0.2	63
139	Activated protein C resistance as measured by residual factor V after Russell \hat{E}^{1} 4s Viper Venom and activated protein C treatment analyzed as a continuous variable in multiple myeloma and normal controls. Blood Coagulation and Fibrinolysis, 2011, 22, 420-423.	0.5	4
140	A prospective evaluation of the biochemical, metabolic, hormonal and structural bone changes associated with bortezomib response in multiple myeloma patients. Haematologica, 2011, 96, 333-336.	1.7	52
141	Alkaline phosphatase variation during carfilzomib treatment is associated with best response in multiple myeloma patients. European Journal of Haematology, 2011, 86, 484-487.	1.1	38
142	Survival Effect of Venous Thromboembolism in Patients With Multiple Myeloma Treated With Lenalidomide and High-Dose Dexamethasone. Journal of Clinical Oncology, 2010, 28, 132-135.	0.8	58
143	Thrombotic Events in Patients With Cancer Receiving Antiangiogenesis Agents. Journal of Clinical Oncology, 2009, 27, 4865-4873.	0.8	200
144	The proteasome inhibitor, bortezomib suppresses primary myeloma and stimulates bone formation in myelomatous and nonmyelomatous bones in vivo. American Journal of Hematology, 2009, 84, 6-14.	2.0	132

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145	Bleeding Disorders Associated with Cancer Dysproteinemias. Cancer Treatment and Research, 2009, 148, 295-304.	0.2	5
146	Eight-year median survival in multiple myeloma after total therapy 2: roles of thalidomide and consolidation chemotherapy in the context of total therapy 1. British Journal of Haematology, 2008, 141, 433-444.	1,2	33
147	Hemostatic Dysfunction in Paraproteinemias and Amyloidosis. Seminars in Thrombosis and Hemostasis, 2007, 33, 339-349.	1.5	64
148	A validated gene expression model of high-risk multiple myeloma is defined by deregulated expression of genes mapping to chromosome 1. Blood, 2007, 109, 2276-2284.	0.6	831
149	Effect on survival of treatment-associated venous thromboembolism in newly diagnosed multiple myeloma patients. Blood Coagulation and Fibrinolysis, 2007, 18, 595-598.	0.5	25
150	Testing standard and genetic parameters in 220 patients with multiple myeloma with complete data sets: superiority of molecular genetics. British Journal of Haematology, 2007, 137, 530-536.	1.2	44
151	Incorporating bortezomib into upfront treatment for multiple myeloma: early results of total therapy 3. British Journal of Haematology, 2007, 138, 176-185.	1.2	304
152	Thrombosis in multiple myeloma. Expert Review of Anticancer Therapy, 2007, 7, 307-315.	1.1	57
153	Exploitation of Novel Hyperdiploid and Nonhyperdiploid Myeloma Cell Lines for Studying Innovative Interventions for Myeloma and Its Associated Bone Disease Blood, 2007, 110, 548-548.	0.6	2
154	Stroke in a Multiple Myeloma Cohort Blood, 2007, 110, 4772-4772.	0.6	0
155	The molecular classification of multiple myeloma. Blood, 2006, 108, 2020-2028.	0.6	997
156	Response to Bortezomib and Activation of Osteoblasts in Multiple Myeloma. Clinical Lymphoma and Myeloma, 2006, 7, 109-114.	1.4	46
157	Farnesyltransferase Inhibitors and Rapamycin in the Treatment of Multiple Myeloma. Current Pharmaceutical Biotechnology, 2006, 7, 449-453.	0.9	5
158	A Validated Gene Expression Signature of High Risk Multiple Myeloma Is Defined by Deregulated Expression of Genes Mapping to Chromosome 1 Blood, 2006, 108, 111-111.	0.6	5
159	A Gene Expression Signature of Benign Monoclonal Gammopathy Evident in Multiple Myeloma Is Linked to Good Prognosis Blood, 2006, 108, 3393-3393.	0.6	1
160	Inhibitory effects of osteoblasts and increased bone formation on myeloma in novel culture systems and a myelomatous mouse model. Haematologica, 2006, 91, 192-9.	1.7	127
161	Anticoagulation regimens for thalidomide and lenalidomide. Clinical Advances in Hematology and Oncology, 2006, 4, 658-9.	0.3	1
162	Response to bortezomib is associated to osteoblastic activation in patients with multiple myeloma. British Journal of Haematology, 2005, 131, 71-73.	1.2	180

#	Article	IF	Citations
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