Matthew W Jenner

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73 papers 1,923 citations 21 h-index g-index

73 ext. papers ext. citations 4.6 avg, IF L-index

#	Paper	IF	Citations
73	A compendium of myeloma-associated chromosomal copy number abnormalities and their prognostic value. <i>Blood</i> , 2010 , 116, e56-65	2.2	263
72	Integration of global SNP-based mapping and expression arrays reveals key regions, mechanisms, and genes important in the pathogenesis of multiple myeloma. <i>Blood</i> , 2006 , 108, 1733-43	2.2	163
71	Lenalidomide maintenance versus observation for patients with newly diagnosed multiple myeloma (Myeloma XI): a multicentre, open-label, randomised, phase 3 trial. <i>Lancet Oncology, The</i> , 2019 , 20, 57-73	21.7	154
70	Gene mapping and expression analysis of 16q loss of heterozygosity identifies WWOX and CYLD as being important in determining clinical outcome in multiple myeloma. <i>Blood</i> , 2007 , 110, 3291-300	2.2	121
69	Homozygous deletion mapping in myeloma samples identifies genes and an expression signature relevant to pathogenesis and outcome. <i>Clinical Cancer Research</i> , 2010 , 16, 1856-64	12.9	109
68	Prediction of outcome in newly diagnosed myeloma: a meta-analysis of the molecular profiles of 1905 trial patients. <i>Leukemia</i> , 2018 , 32, 102-110	10.7	108
67	MMSET deregulation affects cell cycle progression and adhesion regulons in t(4;14) myeloma plasma cells. <i>Haematologica</i> , 2009 , 94, 78-86	6.6	94
66	Lenalidomide (Revlimid), in combination with cyclophosphamide and dexamethasone (RCD), is an effective and tolerated regimen for myeloma patients. <i>British Journal of Haematology</i> , 2007 , 137, 268-9	4.5	86
65	A randomized phase III study of carfilzomib vs low-dose corticosteroids with optional cyclophosphamide in relapsed and refractory multiple myeloma (FOCUS). <i>Leukemia</i> , 2017 , 31, 107-114	10.7	81
64	Deletions of CDKN2C in multiple myeloma: biological and clinical implications. <i>Clinical Cancer Research</i> , 2008 , 14, 6033-41	12.9	77
63	The combination of cyclophosphamide, velcade and dexamethasone induces high response rates with comparable toxicity to velcade alone and velcade plus dexamethasone. <i>Haematologica</i> , 2007 , 92, 1149-50	6.6	62
62	The addition of cyclophosphamide to lenalidomide and dexamethasone in multiply relapsed/refractory myeloma patients; a phase I/II study. <i>British Journal of Haematology</i> , 2010 , 150, 326	- 3 3	56
61	Second malignancies in the context of lenalidomide treatment: an analysis of 2732 myeloma patients enrolled to the Myeloma XI trial. <i>Blood Cancer Journal</i> , 2016 , 6, e506	7	49
60	Use of a biosimilar granulocyte colony-stimulating factor for peripheral blood stem cell mobilization: an analysis of mobilization and engraftment. <i>British Journal of Haematology</i> , 2013 , 162, 107-11	4.5	45
59	The combination of cyclophosphomide, thalidomide and dexamethasone is an effective alternative to cyclophosphamide - vincristine - doxorubicin - methylprednisolone as induction chemotherapy prior to autologous transplantation for multiple myeloma: a case-matched analysis. <i>Leukemia and</i>	1.9	38
58	Advances in oral therapy for multiple myeloma. <i>Lancet Oncology, The</i> , 2006 , 7, 316-25	21.7	34
57	The relative importance of factors predicting outcome for myeloma patients at different ages: results from 3894 patients in the Myeloma XI trial. <i>Leukemia</i> , 2020 , 34, 604-612	10.7	32

Daratumumab monotherapy for patients with intermediate-risk or high-risk smoldering multiple 56 myeloma: a randomized, open-label, multicenter, phase 2 study (CENTAURUS). Leukemia, **2020**, 34, 1840 $^{10.7}$ 2 30 Thalidomide Combinations Improve Response Rates; Results from the MRC IX Study.. Blood, 2007, 2.2 29 55 110, 3593-3593 Response-adapted intensification with cyclophosphamide, bortezomib, and dexamethasone versus no intensification in patients with newly diagnosed multiple myeloma (Myeloma XI): a multicentre, 14.6 26 54 open-label, randomised, phase 3 trial. Lancet Haematology, the, 2019, 6, e616-e629 Thrombosis in patients with myeloma treated in the Myeloma IX and Myeloma XI phase 3 53 2.2 23 randomized controlled trials. Blood, 2020, 136, 1091-1104 Idiotypic DNA vaccination for the treatment of multiple myeloma: safety and immunogenicity in a 52 7.4 21 phase I clinical study. Cancer Immunology, Immunotherapy, 2015, 64, 1021-32 Subclonal copy number is associated with prognosis in multiple myeloma. Blood, 2018, 132, 2465-2469 2.2 51 21 Real-world use of pomalidomide and dexamethasone in double refractory multiple myeloma suggests benefit in renal impairment and adverse genetics: a multi-centre UK experience. British 50 19 4.5 Journal of Haematology, **2017**, 176, 908-917 Isatuximab as monotherapy and combined with dexamethasone in patients with 18 49 2.2 relapsed/refractory multiple myeloma. Blood, 2021, 137, 1154-1165 Genome instability is a consequence of transcription deficiency in patients with bone marrow failure harboring biallelic variants. Proceedings of the National Academy of Sciences of the United 48 11.5 17 States of America, 2018, 115, 7777-7782 Updates to the guidelines for the diagnosis and management of multiple myeloma. British Journal 47 4.5 14 of Haematology, **2014**, 167, 131-3 Survival and outcome of blastoid variant myeloma following treatment with the novel thalidomide 46 3.8 14 containing regime DT-PACE. European Journal of Haematology, 2008, 81, 432-6 Predicting ultrahigh risk multiple myeloma by molecular profiling: an analysis of newly diagnosed 10.7 45 13 transplant eligible myeloma XI trial patients. Leukemia, 2020, 34, 3091-3096 Patient-Reported Outcome Results From the Open-Label, Randomized Phase III Myeloma X Trial Evaluating Salvage Autologous Stem-Cell Transplantation in Relapsed Multiple Myeloma. Journal of 44 2.2 12 Clinical Oncology, 2019, 37, 1617-1628 Early relapse after high-dose melphalan autologous stem cell transplant predicts inferior survival and is associated with high disease burden and genetically high-risk disease in multiple myeloma. 43 4.5 11 British Journal of Haematology, 2021, 193, 551-555 Copy number evolution and its relationship with patient outcome-an analysis of 178 matched 42 10.7 10 presentation-relapse tumor pairs from the Myeloma XI trial. Leukemia, 2021, 35, 2043-2053 Updated Results from the Phase 2 Centaurus Study of Daratumumab (DARA) Monotherapy in Patients with Intermediate-Risk or High-Risk Smoldering Multiple Myeloma (SMM). Blood, 2018, 41 2.2 7 132, 1994-1994 Lenalidomide before and after autologous stem cell transplantation for transplant-eligible patients 6.6 6 40 of all ages in the randomized, phase III, Myeloma XI trial. Haematologica, 2021, 106, 1957-1967 The Combination of Cyclophosphamide, Velcade and Dexamethasone (CVD) Induces High Response Rates with Minimal Toxicity Compared to Velcade Alone (V) and Velcade Plus Dexamethasone 39 2.2 5 (VD).. Blood, 2006, 108, 3537-3537

38	MUK OPTIMUM protocol: a screening study to identify high-risk patients with multiple myeloma suitable for novel treatment approaches combined with a phase II study evaluating optimised combination of biological therapy in newly diagnosed high-risk multiple myeloma and plasma cell	3	5
37	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, open-label, randomised, Phase III trial. <i>British Journal of Haematology</i> , 2021 , 192, 853-868	4.5	5
36	Minimal Residual Disease After Autologous Stem-Cell Transplant for Patients With Myeloma: Prognostic Significance and the Impact of Lenalidomide Maintenance and Molecular Risk <i>Journal of Clinical Oncology</i> , 2022 , JCO2102228	2.2	5
35	Pomalidomide + Bortezomib + Low-Dose Dexamethasone Vs Bortezomib + Low-Dose Dexamethasone As Second-Line Treatment in Patients with Lenalidomide-Pretreated Multiple Myeloma: A Subgroup Analysis of the Phase 3 Optimismm Trial. <i>Blood</i> , 2018 , 132, 3278-3278	2.2	4
34	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. <i>Blood</i> , 2008 , 112, 3707-3707	2.2	4
33	Quadruplet Vs Sequential Triplet Induction Therapy Approaches to Maximise Response for Newly Diagnosed, Transplant Eligible, Myeloma Patients. <i>Blood</i> , 2015 , 126, 189-189	2.2	4
32	Autologous stem cell transplantation is safe and effective for fit older myeloma patients: exploratory results from the Myeloma XI trial. <i>Haematologica</i> , 2020 , Online ahead of print,	6.6	4
31	Bortezomib, Vorinostat, and Dexamethasone Combination Therapy in Relapsed Myeloma: Results of the Phase 2 MUK four Trial. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , 2021 , 21, 154-161.e3	2	4
30	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. <i>PLoS Medicine</i> , 2021 , 18, e1003454	11.6	4
29	Liposomal cytarabine in cerebrospinal fluid. British Journal of Haematology, 2009, 145, 679	4.5	3
28	A Phase 1 Study of Carfilzomib-Thalidomide-Dexamethasone in Patients with Relapsed/Refractory AL Amyloidosis - Catalyst Trial Results. <i>Blood</i> , 2019 , 134, 1890-1890	2.2	3
27	Lenolidamide (Revlimid), in Combination with Cyclophosphamide and Dexamethasone (CRD) Is an Effective Regimen for Heavily Pre-Treated Myeloma Patients <i>Blood</i> , 2006 , 108, 3555-3555	2.2	2
26	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). <i>Blood</i> , 2016 , 128, 4412-4412	2.2	2
25	Title - Myeloma XI Trial for Newly Diagnosed Multiple Myeloma (NDMM); Long Term Second Primary Malignancy (SPM) Incidence in the Context of Lenalidomide Maintenance. <i>Blood</i> , 2019 , 134, 31	3 2-3 13	32 ¹
24	Fine Mapping and Expression Analysis of Chromosome 1 with the Aim of Defining Critically Deregulated Genes Important in the Pathogenesis of Myeloma <i>Blood</i> , 2006 , 108, 112-112	2.2	1
23	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 <i>Blood</i> , 2007 , 110, 395-395	2.2	1
22	Sex Differences in Multiple Myeloma Biology but not Clinical Outcomes: Results from 3894 Patients in the Myeloma XI Trial. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , 2021 , 21, 667-675	2	1
21	Pomalidomide, bortezomib, and dexamethasone at first relapse in lenalidomide-pretreated myeloma: A subanalysis of OPTIMISMM by clinical characteristics. <i>European Journal of Haematology</i> , 2022 , 108, 73-83	3.8	1

(2008-2021)

20	Evaluation of Cardiac Repolarization in the Randomized Phase 2 Study of Intermediate- or High-Risk Smoldering Multiple Myeloma Patients Treated with Daratumumab Monotherapy. <i>Advances in Therapy</i> , 2021 , 38, 1328-1341	4.1	1
19	The Combination of Velcade, Idarubicin and Melphalan (VIM) Demonstrates Significant Clinical Activity in Relapsed/Refractory Myeloma Patients <i>Blood</i> , 2007 , 110, 2727-2727	2.2	O
18	Frailty-adjusted therapy in Transplant Non-Eligible patients with newly diagnosed Multiple Myeloma (FiTNEss (UK-MRA Myeloma XIV Trial)): a study protocol for a randomised phase III trial. <i>BMJ Open</i> , 2022 , 12, e056147	3	0
17	Insights into the Basis of Chromosomal Imbalances during the Clonal Evolution of Multiple Myeloma Using SNP Array Analysis <i>Blood</i> , 2005 , 106, 621-621	2.2	
16	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 <i>Blood</i> , 2005 , 106, 1542-1542	2.2	
15	Status of Chromosome 13 in Multiple Myeloma: Integrated Approach Using SNP Mapping Array and Gene Expression Array <i>Blood</i> , 2005 , 106, 1563-1563	2.2	
14	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 <i>Blood</i> , 2006 , 108, 110-110	2.2	
13	Sub-Classification of Hyperdiploid Myeloma Using Global Gene Expression Profiling and SNP-Based Mapping Arrays <i>Blood</i> , 2006 , 108, 3390-3390	2.2	
12	Thalidomide in Combination with Idarubicin, Dexamethasone and Etoposide (TIDE) Is an Effective Oral Combination in Heavily Pre-Treated Myeloma Patients <i>Blood</i> , 2007 , 110, 4841-4841	2.2	
11	Genome-Wide Identification of Gene Expression Networks Affected by Genomic Changes in Multiple Myeloma <i>Blood</i> , 2007 , 110, 2494-2494	2.2	
10	Mutation and Methylation Analysis of WWOX and CYLD on 16q; Potential Tumor Suppressor Genes in Myeloma <i>Blood</i> , 2007 , 110, 2473-2473	2.2	
9	Screening of Homozygous Deletions Identifies Key Deregulated Genes and Pathways in Multiple Myeloma <i>Blood</i> , 2007 , 110, 2474-2474	2.2	
8	An Integrated Pharmacogenomic Strategy for the Definition of Thalidomide Response Signatures in Presenting Cases of Multiple Myeloma <i>Blood</i> , 2007 , 110, 2493-2493	2.2	
7	Pomalidomide + Bortezomib + Low-Dose Dexamethasone after 1 Prior Line of Therapy in Patients with Lenalidomide-Pretreated Multiple Myeloma: Subanalysis of the Phase 3 Optimismm Trial By Patient Age and Prior Stem Cell Transplant. <i>Blood</i> , 2019 , 134, 3120-3120	2.2	
6	Velcade, Vorinostat and Dexamethasone (V2 D) in Relapsed Myeloma: Results of the Phase 2 Muk Four Trial. <i>Blood</i> , 2015 , 126, 1852-1852	2.2	
5	The Impact of Constitutional Copy Number Variants in Myeloma. <i>Blood</i> , 2008 , 112, 496-496	2.2	
4	Molecular Characterization of Human Multiple Myeloma Cell Lines by Genome-Wide Profiling Identifies Kinase Pathway Alterations <i>Blood</i> , 2008 , 112, 1694-1694	2.2	
3	High Resolution Genomic Profiling Using Single Nucleotide Polymorphism Microarrays Identifies Multiple Novel Genomic Minimally Deleted Regions in Multiple Myeloma. <i>Blood</i> , 2008 , 112, 625-625	2.2	

2	XBP1 Expression Is An Important Prognostic Factor for Newly Diagnosed Myeloma Patients <i>Blood</i> ,
	2008 , 112, 1686-1686

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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

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