

The genetic architecture of multiple myeloma

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Synthetic miR-34a Mimics as a Novel Therapeutic Agent for Multiple Myeloma: <i>In Vitro</i> and <i>In Vivo</i> Evidence. <i>Clinical Cancer Research</i> , 2012, 18, 6260-6270.	3.2	213
2	Heat Shock Protein 90 and Role of Its Chemical Inhibitors in Treatment of Hematologic Malignancies. <i>Pharmaceuticals</i> , 2012, 5, 779-801.	1.7	14
3	Effect of autoimmune diseases on incidence and survival in subsequent multiple myeloma. <i>Journal of Hematology and Oncology</i> , 2012, 5, 59.	6.9	38
4	Understanding the molecular biology of myeloma and its therapeutic implications. <i>Expert Review of Hematology</i> , 2012, 5, 603-617.	1.0	14
5	Cancer heterogeneity: origins and implications for genetic association studies. <i>Trends in Genetics</i> , 2012, 28, 538-543.	2.9	28
6	The expression pattern of small nucleolar and small Cajal body-specific RNAs characterizes distinct molecular subtypes of multiple myeloma. <i>Blood Cancer Journal</i> , 2012, 2, e96-e96.	2.8	70
7	Genomic instability in multiple myeloma: mechanisms and therapeutic implications. <i>Expert Opinion on Biological Therapy</i> , 2013, 13, S69-S82.	1.4	35
8	Gene expression-based prediction of myeloma cell sensitivity to histone deacetylase inhibitors. <i>British Journal of Cancer</i> , 2013, 109, 676-685.	2.9	50
9	A multiparameter flow cytometry immunophenotypic algorithm for the identification of newly diagnosed symptomatic myeloma with an MGUS-like signature and long-term disease control. <i>Leukemia</i> , 2013, 27, 2056-2061.	3.3	78
10	Georgia on my mind: multiple myeloma highlights at ASH 2012. <i>Memo - Magazine of European Medical Oncology</i> , 2013, 6, 189-192.	0.3	1
11	Molecular profiling of multiple myeloma: from gene expression analysis to next-generation sequencing. <i>Expert Opinion on Biological Therapy</i> , 2013, 13, S55-S68.	1.4	20
12	The CCND1 c.870G>A polymorphism is a risk factor for t(11;14)(q13;q32) multiple myeloma. <i>Nature Genetics</i> , 2013, 45, 522-525.	9.4	91
13	Myeloma: Classification and Risk Assessment. <i>Seminars in Oncology</i> , 2013, 40, 554-566.	0.8	24
15	Novel players in multiple myeloma pathogenesis: Role of protein kinases CK2 and GSK3. <i>Leukemia Research</i> , 2013, 37, 221-227.	0.4	28
16	Incorporating Novel Agents in the Management of Elderly Myeloma Patients. <i>Current Hematologic Malignancy Reports</i> , 2013, 8, 261-269.	1.2	4
18	Identification of a novel t(7;14) translocation in multiple myeloma resulting in overexpression of <i>EGFR</i> . <i>Genes Chromosomes and Cancer</i> , 2013, 52, 817-822.	1.5	7
19	Global methylation analysis identifies prognostically important epigenetically inactivated tumor suppressor genes in multiple myeloma. <i>Blood</i> , 2013, 122, 219-226.	0.6	147
20	Clinical drug resistance linked to interconvertible phenotypic and functional states of tumor-propagating cells in multiple myeloma. <i>Blood</i> , 2013, 121, 318-328.	0.6	112

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21	MAGE-C1/CT7 and MAGE-C2/CT10 are frequently expressed in multiple myeloma and can be explored in combined immunotherapy for this malignancy. <i>Cancer Immunology, Immunotherapy</i> , 2013, 62, 191-195.	2.0	23
22	Genome-wide analysis of primary plasma cell leukemia identifies recurrent imbalances associated with changes in transcriptional profiles. <i>American Journal of Hematology</i> , 2013, 88, 16-23.	2.0	60
23	Cell cycle genes co-expression in multiple myeloma and plasma cell leukemia. <i>Genomics</i> , 2013, 102, 243-249.	1.3	15
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37	Progression in Smoldering Myeloma Is Independently Determined by the Chromosomal Abnormalities del(17p), t(4;14), Gain 1q, Hyperdiploidy, and Tumor Load. <i>Journal of Clinical Oncology</i> , 2013, 31, 4325-4332.	0.8	200
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39	The Translocation t(4;14) Can Be Present Only in Minor Subclones in Multiple Myeloma. <i>Clinical Cancer Research</i> , 2013, 19, 4634-4637.	3.2	12
40	A Method for Measurement of Drug Sensitivity of Myeloma Cells Co-Cultured with Bone Marrow Stromal Cells. <i>Journal of Biomolecular Screening</i> , 2013, 18, 637-646.	2.6	21
41	DNA repair pathways in human multiple myeloma. <i>Cell Cycle</i> , 2013, 12, 2760-2773.	1.3	52
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50	Characterization of IGH locus breakpoints in multiple myeloma indicates a subset of translocations appear to occur in pregerminal center B cells. <i>Blood</i> , 2013, 121, 3413-3419.	0.6	128
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87	The Road to Treating Smoldering Multiple Myeloma. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , 2014, 14, S59-S64.	0.2	2
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