

Salomon Manier

List of Publications by Year in descending order

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Version: 2024-02-01

135
papers

4,752
citations

117625

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docs citations

139
times ranked

7190
citing authors

#	ARTICLE	IF	CITATIONS
1	Current state and next-generation CAR-T cells in multiple myeloma. <i>Blood Reviews</i> , 2022, 54, 100929.	5.7	38
2	Bortezomib and high-dose melphalan conditioning regimen in frontline multiple myeloma: an IFM randomized phase 3 study. <i>Blood</i> , 2022, 139, 2747-2757.	1.4	16
3	Progression signature underlies clonal evolution and dissemination of multiple myeloma. <i>Blood</i> , 2021, 137, 2360-2372.	1.4	26
4	International harmonization in performing and reporting minimal residual disease assessment in multiple myeloma trials. <i>Leukemia</i> , 2021, 35, 18-30.	7.2	69
5	Can palliative care consultation increase integration of palliative care for patients with hematologic malignancies?. <i>Blood Advances</i> , 2021, 5, 2123-2127.	5.2	4
6	Antibody-drug conjugate in multiple myeloma. <i>Hematologie</i> , 2021, 27, 26-34.	0.0	0
7	Daratumumab-Based Treatment for Immunoglobulin Light-Chain Amyloidosis. <i>New England Journal of Medicine</i> , 2021, 385, 46-58.	27.0	268
8	Effective anti-BCMA retreatment in multiple myeloma. <i>Blood Advances</i> , 2021, 5, 3016-3020.	5.2	30
9	Can Patient-Reported Ocular Symptoms Guide Dose Modifications in Patients with Relapsed/Refractory Multiple Myeloma Receiving Belantamab Mafodotin?. <i>Blood</i> , 2021, 138, 2746-2746.	1.4	3
10	Bortezomib, Lenalidomide and Dexamethasone (VRd) Followed By Ciltacabtagene Autoleucl Versus Vrd Followed By Lenalidomide and Dexamethasone (Rd) Maintenance in Patients with Newly Diagnosed Multiple Myeloma Not Intended for Transplant: A Randomized, Phase 3 Study (CARTITUDE-5). <i>Blood</i> , 2021, 138, 1835-1835.	1.4	10
11	Real-Life Survival Data after Triple-Exposure to Proteasome Inhibitors (PI), Immunomodulators (IMiD) and Anti-CD38 in Multiple Myeloma Patients in the Emmy Cohort. <i>Blood</i> , 2021, 138, 3764-3764.	1.4	2
12	Single-Cell RNA-Sequencing Identifies Immune Biomarkers of Response to Immunotherapy in Patients with High-Risk Smoldering Myeloma. <i>Blood</i> , 2021, 138, 330-330.	1.4	2
13	In Multiple Myeloma, High-Risk Secondary Genetic Events Observed at Relapse Are Present from the Diagnosis in Tiny Undetectable Subclones. <i>Blood</i> , 2021, 138, 77-77.	1.4	2
14	Ixazomib and Daratumumab without Dexamethasone (I-Dara) in Elderly Frail RRMM Patients. a Multicenter Phase 2 Study (IFM 2018-02) of the Intergroupe Francophone Du Myélome (IFM). <i>Blood</i> , 2021, 138, 83-83.	1.4	4
15	Modified Delphi Method Identifies Consensus Areas for Routine Minimal Residual Disease Testing in Multiple Myeloma. <i>Blood</i> , 2021, 138, 1631-1631.	1.4	0
16	A simplified frailty scale predicts outcomes in transplant-ineligible patients with newly diagnosed multiple myeloma treated in the FIRST (MM-020) trial. <i>Leukemia</i> , 2020, 34, 224-233.	7.2	122
17	On the road to molecular prognostication in SMM. <i>Leukemia</i> , 2020, 34, 331-332.	7.2	1
18	Daratumumab is effective in the relapsed or refractory systemic light-chain amyloidosis but associated with high infection burden in a frail real-life population. <i>British Journal of Haematology</i> , 2020, 188, e24-e27.	2.5	26

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19	Clonal hematopoiesis is associated with adverse outcomes in multiple myeloma patients undergoing transplant. <i>Nature Communications</i> , 2020, 11, 2996.	12.8	98
20	Randomized Trial Comparing Double Versus Triple Bortezomib-Based Regimen in Patients With Multiple Myeloma and Acute Kidney Injury Due to Cast Nephropathy. <i>Journal of Clinical Oncology</i> , 2020, 38, 2647-2657.	1.6	24
21	Genome instability in multiple myeloma. <i>Leukemia</i> , 2020, 34, 2887-2897.	7.2	63
22	Single-cell RNA sequencing reveals compromised immune microenvironment in precursor stages of multiple myeloma. <i>Nature Cancer</i> , 2020, 1, 493-506.	13.2	209
23	Genomic Profiling of Smoldering Multiple Myeloma Identifies Patients at a High Risk of Disease Progression. <i>Journal of Clinical Oncology</i> , 2020, 38, 2380-2389.	1.6	110
24	Antigen excess pitfall for free light chains measurements solved by ELISA assay. <i>American Journal of Hematology</i> , 2019, 94, E120-E122.	4.1	3
25	Daratumumab and dexamethasone is safe and effective for triple refractory myeloma patients: final results of the IFM 2014 (Etoile du Nord) trial. <i>British Journal of Haematology</i> , 2019, 187, 319-327.	2.5	18
26	Response to pneumococcal vaccination in multiple myeloma. <i>Cancer Medicine</i> , 2019, 8, 3822-3830.	2.8	20
27	Immunotherapy in Multiple Myeloma: Accelerating on the Path to the Patient. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , 2019, 19, 332-344.	0.4	16
28	Citron Rho-interacting kinase silencing causes cytokinesis failure and reduces tumor growth in multiple myeloma. <i>Blood Advances</i> , 2019, 3, 995-1002.	5.2	15
29	Progression signature underlies clonal evolution and dissemination of Multiple Myeloma. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , 2019, 19, e19-e20.	0.4	0
30	A Prospective Phase II of Daratumumab in Previously Treated Systemic Light-Chain (AL) Amyloidosis: Updated Results. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , 2019, 19, e40-e41.	0.4	4
31	Single-cell RNA sequencing reveals compromised immune microenvironment in precursor stages of multiple myeloma. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , 2019, 19, e27.	0.4	0
32	Genomic profiling of smoldering multiple myeloma identifies patients at a high risk of disease progression.. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , 2019, 19, e5-e6.	0.4	1
33	In vivo modeling of clonal competition using CRISPR-based gene editing reveals novel fitness variables in multiple myeloma. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , 2019, 19, e110.	0.4	0
34	A High-Throughput Drug Screen Reveals a Novel Compound Class That Significantly Depletes IRF4 Expression in Multiple Myeloma. <i>Blood</i> , 2019, 134, 5545-5545.	1.4	1
35	Maintenance with Weekly Carfilzomib in Elderly Newly Diagnosed Multiple Myeloma (IFM 1212-03). <i>Blood</i> , 2019, 134, 3190-3190.	1.4	0
36	MYC Overexpressing Multiple Myeloma Are Dependent on GLS1. <i>Blood</i> , 2019, 134, 853-853.	1.4	0

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37	Targeting MYC in multiple myeloma. <i>Leukemia</i> , 2018, 32, 1295-1306.	7.2	89
38	Antibody-Dependent Cellular Phagocytosis by Macrophages is a Novel Mechanism of Action of Elotuzumab. <i>Molecular Cancer Therapeutics</i> , 2018, 17, 1454-1463.	4.1	70
39	Platelets Enhance Multiple Myeloma Progression via IL-1 β Upregulation. <i>Clinical Cancer Research</i> , 2018, 24, 2430-2439.	7.0	44
40	Final analysis of survival outcomes in the phase 3 FIRST trial of up-front treatment for multiple myeloma. <i>Blood</i> , 2018, 131, 301-310.	1.4	216
41	Pathology and Genetics of Multiple Myeloma. , 2018, , .		0
42	Myeloma MRD by deep sequencing from circulating tumor DNA does not correlate with results obtained in the bone marrow. <i>Blood Advances</i> , 2018, 2, 2811-2813.	5.2	69
43	Profiling of circulating exosomal miRNAs in patients with Waldenström Macroglobulinemia. <i>PLoS ONE</i> , 2018, 13, e0204589.	2.5	17
44	A predictive model for risk of early grade 3 infection in patients with multiple myeloma not eligible for transplant: analysis of the FIRST trial. <i>Leukemia</i> , 2018, 32, 1404-1413.	7.2	53
45	Inhibition of microRNA-138 enhances bone formation in multiple myeloma bone marrow niche. <i>Leukemia</i> , 2018, 32, 1739-1750.	7.2	34
46	Deregulation and Targeting of TP53 Pathway in Multiple Myeloma. <i>Frontiers in Oncology</i> , 2018, 8, 665.	2.8	47
47	Blocking IFNAR1 inhibits multiple myeloma-driven Treg expansion and immunosuppression. <i>Journal of Clinical Investigation</i> , 2018, 128, 2487-2499.	8.2	80
48	Phase II Trial of Combination of Elotuzumab, Lenalidomide, and Dexamethasone in High-Risk Smoldering Multiple Myeloma. <i>Blood</i> , 2018, 132, 154-154.	1.4	19
49	Founding Precision Therapy in 1q-Amplified Multiple Myeloma. <i>Blood</i> , 2018, 132, 1007-1007.	1.4	12
50	Efficacy and Safety of Daratumumab in a Frail Real-Life Relapsed or Refractory Systemic Light-Chain Amyloidosis Population (AL): Report on 15 Cases from the North of France. <i>Blood</i> , 2018, 132, 5660-5660.	1.4	4
51	Deciphering Clonal Evolution and Dissemination of Multiple Myeloma Cells In Vivo. <i>Blood</i> , 2018, 132, 55-55.	1.4	0
52	In Vivo Modeling of Clonal Competition Using CRISPR-Based Gene Editing Reveals Novel Fitness Variables in Multiple Myeloma. <i>Blood</i> , 2018, 132, 57-57.	1.4	0
53	Prognostic role of circulating exosomal miRNAs in multiple myeloma. <i>Blood</i> , 2017, 129, 2429-2436.	1.4	214
54	A novel in vivo model for studying conditional dual loss of BLIMP1 and p53 in B cells, leading to tumor transformation. <i>American Journal of Hematology</i> , 2017, 92, E138-E145.	4.1	3

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55	Inhibiting the oncogenic translation program is an effective therapeutic strategy in multiple myeloma. <i>Science Translational Medicine</i> , 2017, 9, .	12.4	53
56	The Mutational Landscape of Circulating Tumor Cells in Multiple Myeloma. <i>Cell Reports</i> , 2017, 19, 218-224.	6.4	92
57	Proteomic characterization of human multiple myeloma bone marrow extracellular matrix. <i>Leukemia</i> , 2017, 31, 2426-2434.	7.2	72
58	Circulating tumor markers: harmonizing the yin and yang of CTCs and ctDNA for precision medicine. <i>Annals of Oncology</i> , 2017, 28, 468-477.	1.2	62
59	Metformin Affects Cortical Bone Mass and Marrow Adiposity in Diet-Induced Obesity in Male Mice. <i>Endocrinology</i> , 2017, 158, 3369-3385.	2.8	54
60	The LIN28B/let-7 axis is a novel therapeutic pathway in multiple myeloma. <i>Leukemia</i> , 2017, 31, 853-860.	7.2	72
61	Genomic complexity of multiple myeloma and its clinical implications. <i>Nature Reviews Clinical Oncology</i> , 2017, 14, 100-113.	27.6	413
62	Role of IRF4 in resistance to immunomodulatory (IMiD) compounds in Waldenström's macroglobulinemia. <i>Oncotarget</i> , 2017, 8, 112917-112927.	1.8	5
63	Future Directions in the Evaluation and Treatment of Precursor Plasma Cell Disorders. <i>American Society of Clinical Oncology Educational Book / ASCO American Society of Clinical Oncology Meeting</i> , 2016, 35, e400-e406.	3.8	2
64	IgM and IgM Measurements for the Assessment of Patients with Waldenström's Macroglobulinaemia. <i>Clinical Cancer Research</i> , 2016, 22, 5152-5158.	7.0	9
65	Cell autonomous and microenvironmental regulation of tumor progression in precursor states of multiple myeloma. <i>Current Opinion in Hematology</i> , 2016, 23, 426-433.	2.5	33
66	Exome sequencing reveals recurrent germ line variants in patients with familial Waldenström macroglobulinemia. <i>Blood</i> , 2016, 127, 2598-2606.	1.4	22
67	Epigenetics in Multiple Myeloma. <i>Cancer Treatment and Research</i> , 2016, 169, 35-49.	0.5	7
68	Genomic Aberrations in Multiple Myeloma. <i>Cancer Treatment and Research</i> , 2016, 169, 23-34.	0.5	21
69	Targeting vasculogenesis to prevent progression in multiple myeloma. <i>Leukemia</i> , 2016, 30, 1103-1115.	7.2	46
70	Exosomes in Tumor Angiogenesis. <i>Methods in Molecular Biology</i> , 2016, 1464, 25-34.	0.9	32
71	Driver Mutation in Waldenström's Macroglobulinemia and Their Clonal Heterogeneity during Progression and Relapse. <i>Blood</i> , 2016, 128, 1092-1092.	1.4	2
72	In Vivo Genome-Wide Crispr Library Screen in a Xenograft Mouse Model of Tumor Growth and Metastasis of Multiple Myeloma. <i>Blood</i> , 2016, 128, 1137-1137.	1.4	2

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73	Whole-Exome Sequencing and Targeted Deep Sequencing of cfDNA Enables a Comprehensive Mutational Profiling of Multiple Myeloma. <i>Blood</i> , 2016, 128, 197-197.	1.4	8
74	Future Directions in the Evaluation and Treatment of Precursor Plasma Cell Disorders. American Society of Clinical Oncology Educational Book / ASCO American Society of Clinical Oncology Meeting, 2016, 36, e400-e406.	3.8	4
75	Whole Exome Sequencing and Targeted Sequencing Reveal the Heterogeneity of Genomic Evolution and Mutational Profile in Smoldering Multiple Myeloma. <i>Blood</i> , 2016, 128, 237-237.	1.4	0
76	Microrna-138 Regulates Osteogenic Differentiation and Its Inhibition Presents a Novel Therapeutic Line to Prevent Bone Lytic Lesions in Multiple Myeloma. <i>Blood</i> , 2016, 128, 4483-4483.	1.4	0
77	Dual Conditional Loss of BLIMP-1 and p53 in B-Cells Drives B-Cell Lymphomagenesis. <i>Blood</i> , 2016, 128, 4169-4169.	1.4	0
78	In Vivo Analysis of Clonal Evolution of Multiple Myeloma. <i>Blood</i> , 2016, 128, 799-799.	1.4	0
79	Profiling of Circulating Exosomes in Patients with Waldenström Macroglobulinemia. <i>Blood</i> , 2016, 128, 2940-2940.	1.4	1
80	CXCR4 Regulates Extra-Medullary Myeloma through Epithelial-Mesenchymal-Transition-like Transcriptional Activation. <i>Cell Reports</i> , 2015, 12, 622-635.	6.4	123
81	The cancer glycome: Carbohydrates as mediators of metastasis. <i>Blood Reviews</i> , 2015, 29, 269-279.	5.7	91
82	Targeting the bone marrow microenvironment in multiple myeloma. <i>Immunological Reviews</i> , 2015, 263, 160-172.	6.0	323
83	Mutational Profile and Prognostic Relevance of Circulating Tumor Cells in Multiple Myeloma. <i>Blood</i> , 2015, 126, 23-23.	1.4	37
84	Characterization of the Role of Regulatory T Cells (Tregs) in Inducing Progression of Multiple Myeloma. <i>Blood</i> , 2015, 126, 502-502.	1.4	4
85	Can Assessment of Patient Monoclonal Immunoglobulins By Heavy/Light Analysis be Used to Assign Patient Responses Analogous to IMWG Response Criteria?. <i>Blood</i> , 2015, 126, 3042-3042.	1.4	0
86	MYC Regulation Via the LIN28B/Let-7 Axis in Multiple Myeloma. <i>Blood</i> , 2015, 126, 1755-1755.	1.4	0
87	Circulating Exosomal microRNAs Are Prognostic Markers in Multiple Myeloma. <i>Blood</i> , 2015, 126, 1770-1770.	1.4	4
88	Platelets/Megakaryocytes Are Critical Regulators of Tumor Progression in Multiple Myeloma. <i>Blood</i> , 2015, 126, 1793-1793.	1.4	1
89	Global Epigenetic Regulation of MicroRNAs in Multiple Myeloma. <i>PLoS ONE</i> , 2014, 9, e110973.	2.5	29
90	The Role of miRNAs in Plasma Cell Dyscrasias. <i>MicroRNA (Sharqah, United Arab Emirates)</i> , 2014, 2, 165-173.	1.2	3

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91	Distinct roles of class I PI3K isoforms in multiple myeloma cell survival and dissemination. Blood Cancer Journal, 2014, 4, e204-e204.	6.2	11
92	IgA kappa/IgA lambda heavy/light chain assessment in the management of patients with IgA myeloma. Cancer, 2014, 120, 3952-3957.	4.1	29
93	Targeting survival and cell trafficking in multiple myeloma and Waldenstrom macroglobulinemia using pan-class I PI3K inhibitor, buparlisib. American Journal of Hematology, 2014, 89, 1030-1036.	4.1	14
94	Investigating osteogenic differentiation in multiple myeloma using a novel 3D bone marrow niche model. Blood, 2014, 124, 3250-3259.	1.4	109
95	Role of endothelial progenitor cells in cancer progression. Biochimica Et Biophysica Acta: Reviews on Cancer, 2014, 1846, 26-39.	7.4	70
96	Regulation of microRNAs in cancer metastasis. Biochimica Et Biophysica Acta: Reviews on Cancer, 2014, 1845, 255-265.	7.4	132
97	Engineered nanomedicine for myeloma and bone microenvironment targeting. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 10287-10292.	7.1	234
98	The sialyltransferase ST3GAL6 influences homing and survival in multiple myeloma. Blood, 2014, 124, 1765-1776.	1.4	97
99	Pyk2 promotes tumor progression in multiple myeloma. Blood, 2014, 124, 2675-2686.	1.4	51
100	Clonal-Heterogeneity and Propensity for Bone Metastasis in Multiple Myeloma. Blood, 2014, 124, 3370-3370.	1.4	1
101	Dissecting the Mechanisms of Activity of SLAMF7 and the Targeting Antibody Elotuzumab in Multiple Myeloma. Blood, 2014, 124, 3431-3431.	1.4	8
102	Proteomic Characterization of the Multiple Myeloma Bone Marrow Extracellular Matrix. Blood, 2014, 124, 2051-2051.	1.4	1
103	Citron Rho-Interacting Serine/Threonine kinase (CIT) Is a Novel Therapeutic Target in Multiple Myeloma Cells. Blood, 2014, 124, 3430-3430.	1.4	0
104	Early Trafficking of Bone Marrow Derived-Endothelial Progenitor Cells Promotes Multiple Myeloma Progression. Blood, 2014, 124, 4719-4719.	1.4	0
105	Comparison of Waldenstrom Macroglobulinemia Responses Using Immunoglobulin Heavy / Light Chain Analysis and Conventional Electrophoresis Techniques. Blood, 2014, 124, 2978-2978.	1.4	1
106	Prognostic Value of Circulating Exosomal microRNAs in 112 Patients with Multiple Myeloma. Blood, 2014, 124, 2056-2056.	1.4	0
107	microRNA Aberrations in Waldenstrom Macroglobulinemia. Clinical Lymphoma, Myeloma and Leukemia, 2013, 13, 205-207.	0.4	14
108	Genomic Studies Have Identified Multiple Mechanisms of Genetic Changes in Waldenstrom Macroglobulinemia. Clinical Lymphoma, Myeloma and Leukemia, 2013, 13, 202-204.	0.4	8

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109	Genome wide SNP array identified multiple mechanisms of genetic changes in Waldenstrom macroglobulinemia. American Journal of Hematology, 2013, 88, 948-954.	4.1	45
110	Lin28B/Let-7 Axis Regulates Multiple Myeloma Proliferation By Enhancing c-Myc and Ras Survival Pathways. Blood, 2013, 122, 273-273.	1.4	3
111	Bone Marrow Mobilization Of Endothelial Progenitor Cells Represents An Early Pathogenic Event During Multiple Myeloma Progression. Blood, 2013, 122, 680-680.	1.4	4
112	Myeloma, IMiDs and thrombosis. Hematologie, 2013, 19, 33-40.	0.0	1
113	Mirna Expression Profiling and Proteomic Analysis Of Circulating Exosomes From Multiple Myeloma Patients. Blood, 2013, 122, 3086-3086.	1.4	2
114	Extramedullary Disease In Waldenstrom's Macroglobulinemia. Blood, 2013, 122, 1773-1773.	1.4	0
115	Silencing The Sialyltransferase Gene ST3GAL6 Inhibits Adhesion and Migration Of Myeloma Cells In Vitro and Reduces The Homing and Proliferation Of Tumor Cells In Vivo. Blood, 2013, 122, 275-275.	1.4	0
116	Microrna-Dependent Modulation Of Osteogenesis In a 3D In Vitro Bone Marrow Model System Of Multiple Myeloma. Blood, 2013, 122, 3093-3093.	1.4	0
117	Reduced steady state-based peripheral blood stem cell harvest rate in multiple myeloma treated with bortezomib-based induction regimens. Leukemia, 2012, 26, 2552-2554.	7.2	1
118	Mechanisms of Activity of the TORC1 Inhibitor Everolimus in Waldenstrom Macroglobulinemia. Clinical Cancer Research, 2012, 18, 6609-6622.	7.0	14
119	Immunomodulator drug-based therapy in myeloma and the occurrence of thrombosis. Expert Review of Hematology, 2012, 5, 619-629.	2.2	15
120	Prognostic value of PINI index in patients with multiple myeloma. European Journal of Haematology, 2012, 88, 306-313.	2.2	22
121	Comparative miRNA Expression Profiling of Circulating Exosomes From MGUS and Smoldering Multiple Myeloma Patients. Blood, 2012, 120, 3975-3975.	1.4	2
122	In Vivo Targeting of Stromal-Derived Factor-1 As a Strategy to Prevent Myeloma Cell Dissemination to Distant Bone Marrow Niches. Blood, 2012, 120, 440-440.	1.4	4
123	90-Yttrium Ibritumomab Tiuxetan (Zevalin) and BEAM Chemotherapy (Z-BEAM) Vs BEAM for Autologous Stem Cell Transplantation in Lymphoma: Toxicity and Long Term Outcome From a Retrospective Multicentric Study of 123 Patients.. Blood, 2012, 120, 2726-2726.	1.4	0
124	Multiple Myeloma-Derived Bone-Marrow Mesenchymal Stem Cells: Microrna-, Gene Expression-Profiling and Functional Characterization. Blood, 2012, 120, 1837-1837.	1.4	0
125	IgA Hevylite® Test As a Surrogate to Serum Protein Electrophoresis (SPEP) or Nephelometry in the Management of IgA Myeloma. Blood, 2012, 120, 3970-3970.	1.4	1
126	CXCR4 Monoclonal Antibody, BMS-936564 (MDX-1338), Modulates Epithelial to Mesenchymal Transition (EMT) in Multiple Myeloma Cells. Blood, 2012, 120, 4009-4009.	1.4	0

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127	Metabolomic Profiling Identifies Mechanisms Regulating Hypoxia-Induced Drug Resistance in Multiple Myeloma. Blood, 2012, 120, 3944-3944.	1.4	0
128	Let-7 MicroRNA Family Members Regulate Cell Proliferation in Multiple Myeloma. Blood, 2012, 120, 570-570.	1.4	0
129	High-Throughput Genomic Analysis in Waldenström's Macroglobulinemia. Clinical Lymphoma, Myeloma and Leukemia, 2011, 11, 106-108.	0.4	19
130	Novel M-Component Based Biomarkers in Waldenström's Macroglobulinemia. Clinical Lymphoma, Myeloma and Leukemia, 2011, 11, 164-167.	0.4	23
131	Hevylite, a Novel M-Component Based Biomarkers of Response to Therapy and Survival in Waldenstrom Macroglobulinemia. Blood, 2011, 118, 2667-2667.	1.4	2
132	The EOS® System for the Detection of Bone Lesions in Patients with Multiple Myeloma,. Blood, 2011, 118, 3921-3921.	1.4	1
133	A20 Gene Deregulation In Waldenstrom's Macroglobulinemia.. Blood, 2010, 116, 3628-3628.	1.4	0
134	Hevylite®, a New Marker of Tumor Measurement In Waldenstrom Macroglobulinemia. Blood, 2010, 116, 5076-5076.	1.4	3
135	SOCS1 and SOCS3 Genes Are Not Methylated In Waldenstrom Macroglobulinemia. Blood, 2010, 116, 2481-2481.	1.4	0