

# Bart Barlogie

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

253  
papers

6,880  
citations

37  
h-index

82  
g-index

256  
ext. papers

7,662  
ext. citations

3.7  
avg, IF

5.02  
L-index

#	Paper	IF	Citations
253	Feasibility of Outpatient Stem Cell Transplantation in Multiple Myeloma and Risk Factors Predictive of Hospital Admission.. <i>Journal of Clinical Medicine</i> , <b>2022</b> , 11,	5.1	3
252	N-Cadherin Stabilizes $\beta$ Catenin and Promotes $\beta$ Catenin/TCF Transcriptional Activation and Cell Adhesion-Mediated Drug Resistance in Multiple Myeloma. <i>Blood</i> , <b>2021</b> , 138, 1572-1572	2.2	
251	High-risk transcriptional profiles in multiple myeloma are an acquired feature that can occur in any subtype and more frequently with each subsequent relapse. <i>British Journal of Haematology</i> , <b>2021</b> , 195, 283-286	4.5	2
250	Timing of Autologous Stem Cell Transplantation for Multiple Myeloma in the Era of Current Therapies. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , <b>2020</b> , 20, e734-e751	2	1
249	Genomic analysis of primary plasma cell leukemia reveals complex structural alterations and high-risk mutational patterns. <i>Blood Cancer Journal</i> , <b>2020</b> , 10, 70	7	16
248	and Mutations Associate with Adverse Outcome in a Long-term Follow-up of Patients with Multiple Myeloma. <i>Clinical Cancer Research</i> , <b>2020</b> , 26, 2422-2432	12.9	17
247	Increased Muscle CXCR4 Expression in the Setting of Rare Muscle-invasive Multiple Myeloma. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , <b>2020</b> , 20, e341-e344	2	0
246	28-Day Metronomic Therapy for Relapsed Refractory Multiple Myeloma. <i>Blood</i> , <b>2020</b> , 136, 13-13	2.2	
245	Bone marrow microenvironments that contribute to patient outcomes in newly diagnosed multiple myeloma: A cohort study of patients in the Total Therapy clinical trials. <i>PLoS Medicine</i> , <b>2020</b> , 17, e1003323	11.6	10
244	Hemophagocytic relapsed intramedullary plasmacytoma. <i>International Journal of Hematology</i> , <b>2020</b> , 111, 888-890	2.3	
243	An acquired high-risk chromosome instability phenotype in multiple myeloma: Jumping 1q Syndrome. <i>Blood Cancer Journal</i> , <b>2019</b> , 9, 62	7	17
242	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?. <i>Theranostics</i> , <b>2019</b> , 9, 4756-4763	12.1	7
241	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. <i>Blood</i> , <b>2019</b> , 134, 4333-4333	2.2	2
240	Long-Term Outcome of Total Therapy Regimens: Impact of Molecular Subgroups. <i>Blood</i> , <b>2019</b> , 134, 3309-3309	2.3	1
239	Stem cell mutations can be detected in myeloma patients years before onset of secondary leukemias. <i>Blood Advances</i> , <b>2019</b> , 3, 3962-3967	7.8	4
238	Mesenchymal stem cells gene signature in high-risk myeloma bone marrow linked to suppression of distinct IGFBP2-expressing small adipocytes. <i>British Journal of Haematology</i> , <b>2019</b> , 184, 578-593	4.5	11
237	Gene Expression Profiling Reveals Aberrant T-cell Marker Expression on Tumor Cells of Waldenström's Macroglobulinemia. <i>Clinical Cancer Research</i> , <b>2019</b> , 25, 201-209	12.9	3

236	The Pattern of Mesenchymal Stem Cell Expression Is an Independent Marker of Outcome in Multiple Myeloma. <i>Clinical Cancer Research</i> , <b>2018</b> , 24, 2913-2919	12.9	17
235	Treatment to suppression of focal lesions on positron emission tomography-computed tomography is a therapeutic goal in newly diagnosed multiple myeloma. <i>Haematologica</i> , <b>2018</b> , 103, 1047-1053	6.6	29
234	Lack of a Spleen Signal on Diffusion Weighted MRI Is Associated with High Tumor Burden and Poor Prognosis in Multiple Myeloma. <i>Blood</i> , <b>2018</b> , 132, 4471-4471	2.2	
233	Precision Medicine for Relapsed Multiple Myeloma on the Basis of an Integrative Multiomics Approach. <i>JCO Precision Oncology</i> , <b>2018</b> , 2018,	3.6	10
232	Clinical Presentation and Gene Expression Profiling of Immunoglobulin M Multiple Myeloma Compared With Other Myeloma Subtypes and Waldenström Macroglobulinemia. <i>Journal of Global Oncology</i> , <b>2018</b> , 4, 1-8	2.6	1
231	Walking on myeloma. <i>Blood</i> , <b>2018</b> , 132, 1724	2.2	2
230	Risk stratification of smoldering multiple myeloma: predictive value of free light chains and group-based trajectory modeling. <i>Blood Advances</i> , <b>2018</b> , 2, 1470-1479	7.8	22
229	The prognostic value of the depth of response in multiple myeloma depends on the time of assessment, risk status and molecular subtype. <i>Haematologica</i> , <b>2017</b> , 102, e313-e316	6.6	21
228	The level of deletion 17p and bi-allelic inactivation of has a significant impact on clinical outcome in multiple myeloma. <i>Haematologica</i> , <b>2017</b> , 102, e364-e367	6.6	44
227	Clinical characteristics and prognostic factors in multiple myeloma patients with light chain deposition disease. <i>American Journal of Hematology</i> , <b>2017</b> , 92, 739-745	7.1	21
226	Bortezomib with lenalidomide and dexamethasone versus lenalidomide and dexamethasone alone in patients with newly diagnosed myeloma without intent for immediate autologous stem-cell transplant (SWOG S0777): a randomised, open-label, phase 3 trial. <i>Lancet, The</i> , <b>2017</b> , 389, 519-527	40	496
225	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. <i>Lancet Haematology,the</i> , <b>2017</b> , 4, e443-e451	14.6	16
224	Drug Combinations with Transplantation for Myeloma. <i>New England Journal of Medicine</i> , <b>2017</b> , 377, 91-259.2		1
223	Adverse Metaphase Cytogenetics Can Be Overcome by Adding Bortezomib and Thalidomide to Fractionated Melphalan Transplants. <i>Clinical Cancer Research</i> , <b>2017</b> , 23, 2665-2672	12.9	9
222	Assessment of Total Lesion Glycolysis by F FDG PET/CT Significantly Improves Prognostic Value of GEP and ISS in Myeloma. <i>Clinical Cancer Research</i> , <b>2017</b> , 23, 1981-1987	12.9	57
221	The varied distribution and impact of RAS codon and other key DNA alterations across the translocation cyclin D subgroups in multiple myeloma. <i>Oncotarget</i> , <b>2017</b> , 8, 27854-27867	3.3	19
220	The effect of novel therapies in high-molecular-risk multiple myeloma. <i>Clinical Advances in Hematology and Oncology</i> , <b>2017</b> , 15, 870-879	0.6	6
219	The Spectrum and Clinical Impact of Epigenetic Modifier Mutations in Myeloma. <i>Clinical Cancer Research</i> , <b>2016</b> , 22, 5783-5794	12.9	56

218	Genome-wide association study identifies variation at 6q25.1 associated with survival in multiple myeloma. <i>Nature Communications</i> , <b>2016</b> , 7, 10290	17.4	26
217	Disease and Outcome Disparities in Multiple Myeloma (MM): Exploring the Role of Race/Ethnicity and Obesity in Cooperative Group Clinical Trials. <i>Blood</i> , <b>2016</b> , 128, 1192-1192	2.2	2
216	Signatures of Mesenchymal Cell Lineages and Microenvironment Factors Are Dysregulated in High Risk Myeloma. <i>Blood</i> , <b>2016</b> , 128, 2065-2065	2.2	1
215	The 70-Gene MyPRSR prognostic Risk Score Signature Predicts Increased Risk of Progression from MGUS to Multiple Myeloma Requiring Treatment. <i>Blood</i> , <b>2016</b> , 128, 3275-3275	2.2	1
214	Extensive Regional Intra-Clonal Heterogeneity in Multiple Myeloma - Implications for Diagnostics, Risk Stratification and Targeted Treatment. <i>Blood</i> , <b>2016</b> , 128, 3278-3278	2.2	2
213	Mesenchymal Stem Cells Preconditioned with Myeloma Cells from High-Risk Patients Support the Growth of Myeloma Cells from Low-Risk Patients. <i>Blood</i> , <b>2016</b> , 128, 3304-3304	2.2	2
212	Automated Multiparameter Flow Cytometry (MFC) Immunophenotyping for Reproducible Identification of High Risk Smoldering Multiple Myeloma (SMM). <i>Blood</i> , <b>2016</b> , 128, 373-373	2.2	1
211	The Clinical Impact of Macrofocal Disease in Multiple Myeloma Differs Between Presentation and Relapse. <i>Blood</i> , <b>2016</b> , 128, 4431-4431	2.2	7
210	Use of Multiple Myeloma 70-Gene Prognostic Risk Score As a Continuous Predictor of Patient Outcome. <i>Blood</i> , <b>2016</b> , 128, 5614-5614	2.2	1
209	Network Modeling Reveals CDC42BPA and CLEC11A As Novel Driver Genes of t(4; 14) Multiple Myeloma. <i>Blood</i> , <b>2016</b> , 128, 802-802	2.2	1
208	Integrative Network Analysis of Newly Diagnosed Multiple Myeloma Identifies a Novel RNA-Seq Based High Riskgene Signature. <i>Blood</i> , <b>2016</b> , 128, 3285-3285	2.2	0
207	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. <i>Blood</i> , <b>2016</b> , 128, 2064-2064	2.2	
206	High Risk Myeloma Is Characterized By the Bi-Allelic Inactivation of CDKN2C and RB1. <i>Blood</i> , <b>2016</b> , 128, 4416-4416	2.2	0
205	Aberrant a-to-I RNA Editing and Prognostic Impact of Adar in Multiple Myeloma Patients with 1q Amplification. <i>Blood</i> , <b>2016</b> , 128, 357-357	2.2	
204	Mutation Burden in Multiple Myeloma Is Captured By Gene Expression Profiles. <i>Blood</i> , <b>2016</b> , 128, 4450-4450	2.2	
203	CA-125 secreting IgG kappa multiple myeloma. <i>American Journal of Hematology</i> , <b>2016</b> , 91, E457-8	7.1	
202	MAF protein mediates innate resistance to proteasome inhibition therapy in multiple myeloma. <i>Blood</i> , <b>2016</b> , 128, 2919-2930	2.2	36
201	Tight Junction Protein 1 Modulates Proteasome Capacity and Proteasome Inhibitor Sensitivity in Multiple Myeloma via EGFR/JAK1/STAT3 Signaling. <i>Cancer Cell</i> , <b>2016</b> , 29, 639-652	24.3	67

200	Phenotypic and genomic analysis of multiple myeloma minimal residual disease tumor cells: a new model to understand chemoresistance. <i>Blood</i> , <b>2016</b> , 127, 1896-906	2.2	65
199	Clonal selection and double-hit events involving tumor suppressor genes underlie relapse in myeloma. <i>Blood</i> , <b>2016</b> , 128, 1735-44	2.2	129
198	A common genetic variant in 19q13.3 is associated with outcome of multiple myeloma patients treated with Total Therapy 2 and 3. <i>British Journal of Haematology</i> , <b>2016</b> , 174, 991-3	4.5	6
197	Removing batch effects from purified plasma cell gene expression microarrays with modified ComBat. <i>BMC Bioinformatics</i> , <b>2015</b> , 16, 63	3.6	48
196	American Society of Blood and Marrow Transplantation, European Society of Blood and Marrow Transplantation, Blood and Marrow Transplant Clinical Trials Network, and International Myeloma Working Group Consensus Conference on Salvage Hematopoietic Cell Transplantation in Patients with Relapsed Multiple Myeloma. <i>Biology of Blood and Marrow Transplantation</i> , <b>2015</b> , 21, 2039-2051	4.7	114
195	Prospective analysis of antigen-specific immunity, stem-cell antigens, and immune checkpoints in monoclonal gammopathy. <i>Blood</i> , <b>2015</b> , 126, 2475-8	2.2	85
194	Evidence of an epigenetic origin for high-risk 1q21 copy number aberrations in multiple myeloma. <i>Blood</i> , <b>2015</b> , 125, 3756-9	2.2	31
193	Four genes predict high risk of progression from smoldering to symptomatic multiple myeloma (SWOG S0120). <i>Haematologica</i> , <b>2015</b> , 100, 1214-21	6.6	34
192	Allelic mutations in noncoding genomic sequences construct novel transcription factor binding sites that promote gene overexpression. <i>Genes Chromosomes and Cancer</i> , <b>2015</b> , 54, 692-701	5	5
191	Primary myeloma interaction and growth in coculture with healthy donor hematopoietic bone marrow. <i>BMC Cancer</i> , <b>2015</b> , 15, 864	4.8	9
190	Using a latent class model to refine risk stratification in multiple myeloma. <i>Statistics in Medicine</i> , <b>2015</b> , 34, 2971-80	2.3	2
189	Muscular Relapse in a Patient With Multiple Myeloma. <i>Journal of Clinical Oncology</i> , <b>2015</b> , 33, e125-9	2.2	
188	The Composition and Clinical Impact of Focal Lesions and Their Impact on the Microenvironment in Myeloma. <i>Blood</i> , <b>2015</b> , 126, 1806-1806	2.2	1
187	Melphalan Affects Genes Critical for Myeloma Survival, Homing, and Response to Cytokines and Chemokines. <i>Blood</i> , <b>2015</b> , 126, 1808-1808	2.2	1
186	Upfront 28-Day Metronomic Therapy for High-Risk Multiple Myeloma (HRMM). <i>Blood</i> , <b>2015</b> , 126, 1843-1843	2.2	1
185	High Risk Multiple Myeloma Demonstrates Marked Spatial Genomic Heterogeneity Between Focal Lesions and Random Bone Marrow; Implications for Targeted Therapy and Treatment Resistance. <i>Blood</i> , <b>2015</b> , 126, 20-20	2.2	4
184	Bortezomib, Lenalidomide and Dexamethasone Vs. Lenalidomide and Dexamethasone in Patients (Pts) with Previously Untreated Multiple Myeloma without an Intent for Immediate Autologous Stem Cell Transplant (ASCT): Results of the Randomized Phase III Trial SWOG S0777. <i>Blood</i> , <b>2015</b> , 126, 25-25	2.2	39
183	Impact of Minimal Residual Disease in High and Standard Risk Multiple Myeloma. <i>Blood</i> , <b>2015</b> , 126, 2979-2979	2.2	2

182	Comprehensive Genomic Profiling of Multiple Myeloma in the Course of Clinical Care Identifies Targetable and Prognostically Significant Genomic Alterations. <i>Blood</i> , <b>2015</b> , 126, 369-369	2.2	1
181	The Impact of Combination Chemotherapy and Tandem Stem Cell Transplant on Clonal Substructure and Mutational Pattern at Relapse of MM. <i>Blood</i> , <b>2015</b> , 126, 372-372	2.2	1
180	Specific Exosomal microRNA Are Differentially Expressed Between High and Low-Risk Myeloma Suggesting They Are Pathogenically Important. <i>Blood</i> , <b>2015</b> , 126, 4189-4189	2.2	2
179	Outcomes of Autologous Transplantation for Treatment-Related AML and MDS in Previously Treated Multiple Myeloma Patients (pts). <i>Blood</i> , <b>2015</b> , 126, 1997-1997	2.2	
178	Assessment of Total Lesion Glycolysis and Metabolic Tumor Volume Improve the Clinical Value of Focal Lesion Assessment By FDG PET/CT in Myeloma. <i>Blood</i> , <b>2015</b> , 126, 724-724	2.2	
177	Stem Cell-like Characteristics of MM Plasma Cells Vary By ROS Levels: Implications for Targeted Therapy. <i>Blood</i> , <b>2015</b> , 126, 1820-1820	2.2	
176	Deletion of TP53 (17p13) Is Associated with Poor Outcome for Newly Diagnosed High-Risk Multiple Myeloma. <i>Blood</i> , <b>2015</b> , 126, 2982-2982	2.2	
175	Molecular Subtyping and Risk Stratification for the Classification of Myeloma. <i>Blood</i> , <b>2015</b> , 126, 4173-4173	2.2	
174	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. <i>Blood</i> , <b>2015</b> , 126, 3181-3181	2.2	
173	Differential ICAM3 Gene Expression Correlates with Susceptibility to Natural Killer Cell-Mediated Lysis in Multiple Myeloma. <i>Blood</i> , <b>2015</b> , 126, 2990-2990	2.2	
172	Defining the Impact of Tandem Autologous Stem Cell Transplantation in Multiple Myeloma: A Case-Match Analysis in the Total Therapy Trials. <i>Blood</i> , <b>2015</b> , 126, 3182-3182	2.2	
171	Extending Metronomic Therapy to 28 Days (metro28) for Relapsed Refractory Multiple Myeloma (RRMM). <i>Blood</i> , <b>2015</b> , 126, 5395-5395	2.2	
170	Identification of Biomarkers Associated with MAF-Mediated Resistance to Proteasome Inhibitors in t(14;16) Multiple Myeloma. <i>Blood</i> , <b>2015</b> , 126, 3020-3020	2.2	1
169	Identification and Validation of IMiD-14 Model Predictive of IMiD Resistance in Multiple Myeloma. <i>Blood</i> , <b>2015</b> , 126, 4183-4183	2.2	
168	Re-Mineralization of Large Pelvic Lytic Lesions By CT Imaging in Patients with Multiple Myeloma: The Arkansas Experience. <i>Blood</i> , <b>2015</b> , 126, 4193-4193	2.2	
167	47 Genes Define Myeloma Cell Acquired Resistance to Bortezomib and Have Profound Prognostic Implications in Multiple Myeloma. <i>Blood</i> , <b>2015</b> , 126, 499-499	2.2	
166	Gene Expression Profiling of Extramedullary Disease-Related Toward Identification of a Terminal Disease Pathway in Multiple Myeloma. <i>Blood</i> , <b>2015</b> , 126, 1777-1777	2.2	
165	Second primary malignancies with lenalidomide therapy for newly diagnosed myeloma: a meta-analysis of individual patient data. <i>Lancet Oncology</i> , <b>2014</b> , 15, 333-42	21.7	206

164	CYR61/CCN1 overexpression in the myeloma microenvironment is associated with superior survival and reduced bone disease. <i>Blood</i> , <b>2014</b> , 124, 2051-60	2.2	23
163	The future of autologous stem cell transplantation in myeloma. <i>Blood</i> , <b>2014</b> , 124, 328-33	2.2	35
162	Treatment recommendations for patients with Waldenström macroglobulinemia (WM) and related disorders: IWWM-7 consensus. <i>Blood</i> , <b>2014</b> , 124, 1404-11	2.2	107
161	Curing myeloma at last: defining criteria and providing the evidence. <i>Blood</i> , <b>2014</b> , 124, 3043-51	2.2	167
160	Patterns of central nervous system involvement in relapsed and refractory multiple myeloma. <i>Clinical Lymphoma, Myeloma and Leukemia</i> , <b>2014</b> , 14, 211-4	2	38
159	Can autologous bone marrow transplantation improve systolic function in patients with multiple myeloma related cardiac amyloidosis?. <i>International Journal of Cardiology</i> , <b>2014</b> , 172, 265-6	3.2	3
158	Total Therapy 4 (TT4) for GEP70-Defined Low Risk Clinical Multiple Myeloma (CMM): Results of Patients Randomized to a Standard v Light Arm (S-TT4 v L-TT4). <i>Blood</i> , <b>2014</b> , 124, 1199-1199	2.2	3
157	Curing Multiple Myeloma (MM) with Total Therapy (TT). <i>Blood</i> , <b>2014</b> , 124, 195-195	2.2	3
156	Further Evolution of Metronomic Therapy Extended to 28 Days (Metro28) for Relapsed Refractory Multiple Myeloma (RRMM). <i>Blood</i> , <b>2014</b> , 124, 2128-2128	2.2	0
155	Validation of a Predictive Formula for Collection of Hematopoietic Progenitor Cells (HPC) By Leukapheresis at 2 Institutions Using 4 Different Machine Protocols. <i>Blood</i> , <b>2014</b> , 124, 2458-2458	2.2	2
154	Waldenstrom's Macroglobulinemia Associated Bone Disease the UAMS Experience. <i>Blood</i> , <b>2014</b> , 124, 2999-2999	2.2	2
153	Modified Combat Removes Batch Effects from Myeloma Cell GEP-derived Risk Scores and Molecular Subgroup Assignment. <i>Blood</i> , <b>2014</b> , 124, 3355-3355	2.2	1
152	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. <i>Blood</i> , <b>2014</b> , 124, 3409-3409	2.2	1
151	Characterization of the Mutational Landscape of Multiple Myeloma Using Comprehensive Genomic Profiling. <i>Blood</i> , <b>2014</b> , 124, 3418-3418	2.2	3
150	Targeted MEK Inhibition in Patients with Previously Treated Multiple Myeloma. <i>Blood</i> , <b>2014</b> , 124, 4775-4775	2.2	4
149	Evidence of an Epigenetic Origin for High-Risk 1q21 Copy Number Aberrations in Multiple Myeloma. <i>Blood</i> , <b>2014</b> , 124, 725-725	2.2	1
148	Sustained Growth of Primary Myeloma Cells in Coculture with Whole Donor Bone Marrow Is Associated with Induced Secretion of the Microenvironmental Mediator of Cytokinesis, Hemicentin-1. <i>Blood</i> , <b>2014</b> , 124, 3403-3403	2.2	
147	Studies of the Proteasome Inhibitor Sensitivity Modulator Tight Junction Protein 1 Highlight a Role for Signaling through the Epidermal Growth Factor Receptor in Determining Proteasome Capacity. <i>Blood</i> , <b>2014</b> , 124, 3414-3414	2.2	

- 146 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 2.2
- 145 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 2.2
- 144 ATRA Upregulates Cell Surface CD1D on Myeloma Cells and Sensitizes Them to iNKT Cell-Mediated Lysis. *Blood*, **2014**, 124, 2102-2102 2.2
- 143 Low BCL11A Expression in the Myeloma Microenvironment at Diagnosis Is Associated with Early Development of MDS Cytogenetic Abnormalities and Poor Overall Survival. *Blood*, **2014**, 124, 2012-2012<sup>2,2</sup>
- 142 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 2.2
- 141 Low-Dose 28-Day Metronomically Scheduled Therapy (METRO) for Newly Diagnosed High-Risk Multiple Myeloma: A Pilot Study. *Blood*, **2014**, 124, 5770-5770 2.2
- 140 Advanced Osteolytic Lesions (OL), Mobilization and Collection of Hematopoietic Progenitor Cells (HPC) in Multiple Myeloma (MM). *Blood*, **2014**, 124, 3858-3858 2.2
- 139 Exomic microRNA Profiling of Bone Marrow Aspirate Plasma and Comparison with mRNA Profiles Used in the Clinical Management of Multiple Myeloma. *Blood*, **2014**, 124, 5681-5681 2.2
- 138 Mafk Protein Confers Primary Resistance of Myeloma to Proteasome Inhibitors. *Blood*, **2014**, 124, 2091-2091
- 137 Defining Risk of MGUS and AMM Progression to Myeloma By Ig Heavy-Chain FISH. *Blood*, **2014**, 124, 3408-3408
- 136 The Antimalarial Agent Artesunate Exerts Its Antimyeloma Activity By Affecting The Mitochondria and The Reactive Oxygen Status Of The Myeloma Cells and Its Efficacy Depends On Intracellular Bivalent Iron Levels. *Blood*, **2013**, 122, 4444-4444 2.2 1
- 135 Role Of Cytogenetic Abnormalities At Baseline and During 5-Year Follow-Up In Multiple Myeloma Patients Treated On The Total Therapy 3 Protocol. *Blood*, **2013**, 122, 3137-3137 2.2
- 134 MAF Protein Elicits Innate Resistance To Bortezomib In Multiple Myeloma. *Blood*, **2013**, 122, 281-281 2.2 1
- 133 Non-Producing Multiple Myeloma (MM) Is a Distinct Subset Of Non-Secretory MM Characterized By High Cyclin D1 Expression and Decreased Progression Free Survival. *Blood*, **2013**, 122, 1911-1911 2.2 1
- 132 Impact Of Elotuzumab Therapy On Circulating and Ex Vivo Activated/Expanded Autologous Natural Killer (Auto-ENK) Cell Activity. *Blood*, **2013**, 122, 5389-5389 2.2
- 131 Healthy Donor Whole Bone Marrow Cells Preconditioned With Myeloma Patient Serum Support Long-Term Survival Of Primary Myeloma and Reveal Altered Microenvironmental Pathways. *Blood*, **2013**, 122, 3118-3118 2.2
- 130 Inhibition Of BTK Activity In Myeloma Cells Within a Supportive Microenvironment Promotes Their Growth But Suppresses Metastasis. *Blood*, **2013**, 122, 4432-4432 2.2
- 129 Macrophages Activation By ICAM1 Antibody Combined With Lenalidomide Has Enhanced Anti-Myeloma Activity In a Supportive Microenvironment In Vivo and In Vitro. *Blood*, **2013**, 122, 1926-1926<sup>2,2</sup> 1



128	Hyperhaploid Multiple Myeloma (MM): A Rare Karyotypic Subgroup Retaining Disomy 18 and 1q12~23 Amplification. <i>Blood</i> , <b>2012</b> , 120, 3983-3983	2.2	1
127	Phase II Study of Pomalidomide (Pom) in Genomically Defined High Risk Relapsed and Refractory Multiple Myeloma (RRMM). <i>Blood</i> , <b>2012</b> , 120, 4083-4083	2.2	2
126	Fresh Ex Vivo Expanded Natural Killer Cells Demonstrate Robust Proliferation in Vivo in High-Risk Relapsed Multiple Myeloma (MM) Patients. <i>Blood</i> , <b>2012</b> , 120, 579-579	2.2	2
125	Myeloma Can Modulate Expanded Natural Killer Cell Function Through Multiple Mechanisms. <i>Blood</i> , <b>2012</b> , 120, 4020-4020	2.2	
124	Fulminant Onset of Acute Leukemia (FOAL) After Total Therapies (TT) for Multiple Myeloma (MM): Absence of MDS Pathological Criteria within 3 Months of Prior MM Follow-up. <i>Blood</i> , <b>2012</b> , 120, 1458-1458	2.2	2
123	The Antimalarial Agent Artesunate Overcomes Bortezomib Resistance in Myeloma Cell Lines Through Non-Caspase Mediated Apoptosis. <i>Blood</i> , <b>2012</b> , 120, 4015-4015	2.2	
122	Gene Expression Profiling (GEP) in MGUS and AMM: Predictors of Progression.. <i>Blood</i> , <b>2012</b> , 120, 2933-2933	2.2	
121	Identifying the Outliers Among Gene Expression Profiling (GEP)-Defined Low-Risk Myeloma Patients Treated with Total Therapy 2 and 3 (TT2, TT3). <i>Blood</i> , <b>2012</b> , 120, 195-195	2.2	
120	FISH and GEP Based Prediction of Chromosomal Translocation Identifies Myeloma Patients Who Do Not Benefit From Bortezomib. <i>Blood</i> , <b>2012</b> , 120, 1814-1814	2.2	
119	Prognostic Significance of DNA/Cig Flow Cytometry Assay in the Era of Novel Therapies in Multiple Myeloma (MM).. <i>Blood</i> , <b>2012</b> , 120, 2918-2918	2.2	
118	Renal Function Impairment (creatinine $\geq$ 2mg/dL) Limits Progress Noted with the Transition From Total Therapies TT1 to TT2 to TT3 Across Age Groups. <i>Blood</i> , <b>2012</b> , 120, 1962-1962	2.2	
117	Hematopoietic Progenitor Cell (HPC) Collection Is Feasible in Previously Transplanted Multiple Myeloma Patients and Plerixafor Improves Collection. <i>Blood</i> , <b>2012</b> , 120, 4127-4127	2.2	
116	Lenalidomide Suppression of Multiple Myeloma Cell Proliferation Is Associated with Downregulation of LEF/TCF Activity. <i>Blood</i> , <b>2012</b> , 120, 5014-5014	2.2	
115	The use of molecular-based risk stratification and pharmacogenomics for outcome prediction and personalized therapeutic management of multiple myeloma. <i>International Journal of Hematology</i> , <b>2011</b> , 94, 321-333	2.3	24
114	Could CR mean cure?. <i>Blood</i> , <b>2011</b> , 118, 483	2.2	8
113	Prognostic factor analyses of myeloma survival with intergroup trial S9321 (INT 0141): examining whether different variables govern different time segments of survival. <i>Annals of Hematology</i> , <b>2011</b> , 90, 423-8	3	22
112	Secreted Frizzled-Related Protein-3 (sFRP3) Is Produced by Myeloma Cells and Augments Wnt3a-Induced Differentiation of Mesenchymal Stem Cells and OPG Production in Osteoblasts. <i>Blood</i> , <b>2011</b> , 118, 808-808	2.2	1
111	Inducible Heme Oxygenase 1 (HMOX1) Promotes Osteoblastogenesis, and Inhibits Osteoclastogenesis and Myeloma-Induced Bone Disease. <i>Blood</i> , <b>2011</b> , 118, 627-627	2.2	

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109	Implications of Serial Magnetic Resonance Imaging and Positron Emission Tomography Scanning for Survival of Untreated Myeloma Patients Treated with Total Therapy 3. <i>Blood</i> , <b>2011</b> , 118, 3082-3082	2.2	
108	Jumping Translocations 1q12 Contribute to Copy Number (CN) Alterations in Multiple Myeloma (MM): Unexpected Focal Amplifications of Receptor Chromosomes (RC). <i>Blood</i> , <b>2011</b> , 118, 298-298	2.2	
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106	Autologous Expanded Natural Killer Cells As a New Therapeutic Option for High-Risk Myeloma. <i>Blood</i> , <b>2011</b> , 118, 2918-2918	2.2	1
105	Gene Expression Profiling (GEP) Analysis of Plasma Cells (PC) Obtained From MRI-Defined Focal Lesions (FL) Under CT-Guided Fine-Needle Aspiration Provides Better Risk Stratification in Patients with Multiple Myeloma. <i>Blood</i> , <b>2011</b> , 118, 2896-2896	2.2	
104	Reiterative survival analyses of total therapy 2 for multiple myeloma elucidate follow-up time dependency of prognostic variables and treatment arms. <i>Journal of Clinical Oncology</i> , <b>2010</b> , 28, 3023-7	2.2	32
103	Reply to J.C. Regelin et al. <i>Journal of Clinical Oncology</i> , <b>2010</b> , 28, e744-e745	2.2	
102	Long-term follow-up of autotransplantation trials for multiple myeloma: update of protocols conducted by the intergroupe francophone du myelome, southwest oncology group, and university of arkansas for medical sciences. <i>Journal of Clinical Oncology</i> , <b>2010</b> , 28, 1209-14	2.2	125
101	Superior results of Total Therapy 3 (2003-33) in gene expression profiling-defined low-risk multiple myeloma confirmed in subsequent trial 2006-66 with VRD maintenance. <i>Blood</i> , <b>2010</b> , 115, 4168-73	2.2	175
100	Comparing Toxicities and Survival Outcomes with Total Therapy 4 (TT4) for 70-Gene (R70)-Defined Low-Risk Multiple Myeloma (MM) to Results Obtained with Total Therapy 3 Protocols TT3A and TT3B. <i>Blood</i> , <b>2010</b> , 116, 368-368	2.2	4
99	Pacmed Salvage Therapy for Advanced High-Risk Multiple Myeloma (AHRMM). <i>Blood</i> , <b>2010</b> , 116, 1969-1969	2.2	
98	Total Therapy 2 (TT2) for Multiple Myeloma (MM): Contributions to Survival Outcomes of Dosing of Thalidomide (T), Dexamethasone (D) and Interferon (I) Maintenance Components.. <i>Blood</i> , <b>2010</b> , 116, 1356-1356	2.2	
97	Aldehyde Dehydrogenase (ALDH) Vs CD34 for Predicting Engraftment After Autologous Hematopoietic Progenitor Cell (autoHPC) Transplant. <i>Blood</i> , <b>2010</b> , 116, 4441-4441	2.2	
96	Modeling for Cure with Total Therapy (TT) Trials for Newly Diagnosed Multiple Myeloma (MM): Let the Math Speak.. <i>Blood</i> , <b>2009</b> , 114, 744-744	2.2	1
95	A 15 Hour Dosing-Collection Interval for Plerixafor Is at Least as Effective as the Standard 10 Hour Interval.. <i>Blood</i> , <b>2009</b> , 114, 2152-2152	2.2	1
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54	Serum Free-Lite Chain (sFLC) Assay in Multiple Myeloma (MM): Clinical Correlates and Prognostic Implications in Newly Diagnosed MM Patients Treated with Total Therapy 2 or 3 (TT2/3).. <i>Blood</i> , <b>2005</b> , 106, 3490-3490	2.2	2
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