

Marcos Gonzalez Daz

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.

229
papers

16,896
citations

57
h-index

127
g-index

233
ext. papers

19,096
ext. citations

5.8
avg, IF

5.15
L-index

#	Paper	IF	Citations
229	Overexpression of wild type RRAS2, without oncogenic mutations, drives chronic lymphocytic leukemia.. <i>Molecular Cancer</i> , 2022 , 21, 35	42.1	3
228	MYD88 Mutations: Transforming the Landscape of IgM Monoclonal Gammopathies. <i>International Journal of Molecular Sciences</i> , 2022 , 23, 5570	6.3	1
227	Cellular and humoral immunogenicity of the mRNA-1273 SARS-CoV-2 vaccine in patients with hematologic malignancies. <i>Blood Advances</i> , 2021 ,	7.8	8
226	The Hydropathy Index of the HCDR3 Region of the B-Cell Receptor Identifies Two Subgroups of IGHV-Mutated Chronic Lymphocytic Leukemia Patients With Distinct Outcome. <i>Frontiers in Oncology</i> , 2021 , 11, 723722	5.3	
225	Siglec-6 is a novel target for CAR T-cell therapy in acute myeloid leukemia. <i>Blood</i> , 2021 , 138, 1830-1842	2.2	9
224	Liquid biopsy: a non-invasive approach for Hodgkin lymphoma genotyping. <i>British Journal of Haematology</i> , 2021 , 195, 542-551	4.5	1
223	Genetic complexity impacts the clinical outcome of follicular lymphoma patients. <i>Blood Cancer Journal</i> , 2021 , 11, 11	7	0
222	Impact of measurable residual disease by decentralized flow cytometry: a PETHEMA real-world study in 1076 patients with acute myeloid leukemia. <i>Leukemia</i> , 2021 , 35, 2358-2370	10.7	8
221	Allele and haplotype frequencies of HLA-A, -B, -C, -DRB1, -DQB1 and -DQA1 in Castile and Leon region from North West of Spain. <i>Human Immunology</i> , 2021 , 82, 549-550	2.3	0
220	Genomic mutation profile in progressive chronic lymphocytic leukemia patients prior to first-line chemoimmunotherapy with FCR and rituximab maintenance (REM). <i>PLoS ONE</i> , 2021 , 16, e0257353	3.7	
219	Building a network of TP53 and IGHV testing reference centers across Spain: the Red53 initiative. <i>Annals of Hematology</i> , 2021 , 100, 825-830	3	1
218	Identification of relapse-associated gene mutations by next-generation sequencing in low-risk acute myeloid leukaemia patients. <i>British Journal of Haematology</i> , 2020 , 189, 718-730	4.5	6
217	Biological Features and Prognostic Impact of Bone Marrow Infiltration in Patients with Diffuse Large B-Cell Lymphoma. <i>Cancers</i> , 2020 , 12,	6.6	4
216	Biological and clinical significance of dysplastic hematopoiesis in patients with newly diagnosed multiple myeloma. <i>Blood</i> , 2020 , 135, 2375-2387	2.2	11
215	Characteristics and outcome of adult patients with acute promyelocytic leukemia and increased body mass index treated with the PETHEMA Protocols. <i>European Journal of Haematology</i> , 2020 , 104, 162-169	3.8	2
214	Comparison of next-generation sequencing (NGS) and next-generation flow (NGF) for minimal residual disease (MRD) assessment in multiple myeloma. <i>Blood Cancer Journal</i> , 2020 , 10, 108	7	20
213	Immunoglobulin gene rearrangement IGHV3-48 is a predictive marker of histological transformation into aggressive lymphoma in follicular lymphomas. <i>Blood Cancer Journal</i> , 2019 , 9, 52	7	4

212	Life expectancy of follicular lymphoma patients in complete response at 30 months is similar to that of the Spanish general population. <i>British Journal of Haematology</i> , 2019 , 185, 480-491	4.5	14
211	Blood monitoring of circulating tumor plasma cells by next generation flow in multiple myeloma after therapy. <i>Blood</i> , 2019 , 134, 2218-2222	2.2	35
210	Exportin-1 E571K mutation is a common finding in patients with classical Hodgkin lymphoma. <i>Hematological Oncology</i> , 2019 , 37, 215-218	1.3	1
209	Mutations in the RAS-BRAF-MAPK-ERK pathway define a specific subgroup of patients with adverse clinical features and provide new therapeutic options in chronic lymphocytic leukemia. <i>Haematologica</i> , 2019 , 104, 576-586	6.6	28
208	Richter transformation driven by Epstein-Barr virus reactivation during therapy-related immunosuppression in chronic lymphocytic leukaemia. <i>Journal of Pathology</i> , 2018 , 245, 61-73	9.4	15
207	Low-count monoclonal B-cell lymphocytosis persists after seven years of follow up and is associated with a poorer outcome. <i>Haematologica</i> , 2018 , 103, 1198-1208	6.6	19
206	Basophil-lineage commitment in acute promyelocytic leukemia predicts for severe bleeding after starting therapy. <i>Modern Pathology</i> , 2018 , 31, 1318-1331	9.8	8
205	Characterizing patients with multiple chromosomal aberrations detected by FISH in chronic lymphocytic leukemia. <i>Leukemia and Lymphoma</i> , 2018 , 59, 633-642	1.9	6
204	Residual normal B-cell profiles in monoclonal B-cell lymphocytosis versus chronic lymphocytic leukemia. <i>Leukemia</i> , 2018 , 32, 2701-2705	10.7	8
203	A novel high-throughput assay reveals antiproliferative effects of idelalisib and ibrutinib in chronic lymphocytic leukemia. <i>Oncotarget</i> , 2018 , 9, 26019-26031	3.3	5
202	The mutational landscape of small lymphocytic lymphoma compared to non-early stage chronic lymphocytic leukemia. <i>Leukemia and Lymphoma</i> , 2018 , 59, 2318-2326	1.9	2
201	Focal Adhesion Genes Refine the Intermediate-Risk Cytogenetic Classification of Acute Myeloid Leukemia. <i>Cancers</i> , 2018 , 10,	6.6	5
200	A novel predictive approach for GVHD after allogeneic SCT based on clinical variables and cytokine gene polymorphisms. <i>Blood Advances</i> , 2018 , 2, 1719-1737	7.8	20
199	Next Generation Flow for highly sensitive and standardized detection of minimal residual disease in multiple myeloma. <i>Leukemia</i> , 2017 , 31, 2094-2103	10.7	298
198	Update of the Grupo Español de Leucemia Linfocítica Crónica clinical guidelines of the management of chronic lymphocytic leukemia. <i>Medicina Clínica</i> , 2017 , 148, 381.e1-381.e9	1	1
197	Host virus and pneumococcus-specific immune responses in high-count monoclonal B-cell lymphocytosis and chronic lymphocytic leukemia: implications for disease progression. <i>Haematologica</i> , 2017 , 102, 1238-1246	6.6	7
196	Next-generation sequencing and FISH studies reveal the appearance of gene mutations and chromosomal abnormalities in hematopoietic progenitors in chronic lymphocytic leukemia. <i>Journal of Hematology and Oncology</i> , 2017 , 10, 83	22.4	28
195	Donor CTLA-4 Genotype Modulates the Immune Response to Minor Histocompatibility Antigen Mismatches. <i>Biology of Blood and Marrow Transplantation</i> , 2017 , 23, 2042-2047	4.7	4

194	The number of tumor infiltrating T-cell subsets in lymph nodes from patients with Hodgkin lymphoma is associated with the outcome after first line ABVD therapy. <i>Leukemia and Lymphoma</i> , 2017 , 58, 1144-1152	1.9	16
193	Application of a molecular diagnostic algorithm for haemophilia A and B using next-generation sequencing of entire F8, F9 and VWF genes. <i>Thrombosis and Haemostasis</i> , 2017 , 117, 66-74	7	27
192	HLA specificities are associated with prognosis in IGHV-mutated CLL-like high-count monoclonal B cell lymphocytosis. <i>PLoS ONE</i> , 2017 , 12, e0172978	3.7	2
191	Mesenchymal stromal cells (MSC) from JAK2+ myeloproliferative neoplasms differ from normal MSC and contribute to the maintenance of neoplastic hematopoiesis. <i>PLoS ONE</i> , 2017 , 12, e0182470	3.7	9
190	Sarcoma histiocítico en intestino delgado: estudio y discusión de un caso con reordenamiento clonal linfoide B. <i>Revista Espanola De Patologia</i> , 2016 , 49, 254-258	1.2	
189	High-resolution copy number analysis of paired normal-tumor samples from diffuse large B cell lymphoma. <i>Annals of Hematology</i> , 2016 , 95, 253-62	3	14
188	Minimal residual disease evaluation by flow cytometry is a complementary tool to cytogenetics for treatment decisions in acute myeloid leukaemia. <i>Leukemia Research</i> , 2016 , 40, 1-9	2.7	25
187	The Presence of MDS-like Phenotypic Abnormalities (MDS-PA) Identifies Newly Diagnosed Multiple Myeloma (MM) Patients with MDS/AML-Related Somatic Mutations and Inferior Survival. <i>Blood</i> , 2016 , 128, 375-375	2.2	1
186	Karyotypic complexity rather than chromosome 8 abnormalities aggravates the outcome of chronic lymphocytic leukemia patients with TP53 aberrations. <i>Oncotarget</i> , 2016 , 7, 80916-80924	3.3	23
185	Microvesicles from Mesenchymal Stromal Cells Are Involved in HPC-Microenvironment Crosstalk in Myelodysplastic Patients. <i>PLoS ONE</i> , 2016 , 11, e0146722	3.7	56
184	Design and application of a 23-gene panel by next-generation sequencing for inherited coagulation bleeding disorders. <i>Haemophilia</i> , 2016 , 22, 590-7	3.3	34
183	A high proportion of cells carrying trisomy 12 is associated with a worse outcome in patients with chronic lymphocytic leukemia. <i>Hematological Oncology</i> , 2016 , 34, 84-92	1.3	21
182	Multipronged functional proteomics approaches for global identification of altered cell signalling pathways in B-cell chronic lymphocytic leukaemia. <i>Proteomics</i> , 2016 , 16, 1193-203	4.8	12
181	Homeobox NKX2-3 promotes marginal-zone lymphomagenesis by activating B-cell receptor signalling and shaping lymphocyte dynamics. <i>Nature Communications</i> , 2016 , 7, 11889	17.4	32
180	MiRNA expression profile of chronic lymphocytic leukemia patients with 13q deletion. <i>Leukemia Research</i> , 2016 , 46, 30-6	2.7	7
179	Clinical impact of clonal and subclonal TP53, SF3B1, BIRC3, NOTCH1, and ATM mutations in chronic lymphocytic leukemia. <i>Blood</i> , 2016 , 127, 2122-30	2.2	188
178	Origin of Waldenstrom's macroglobulinaemia. <i>Best Practice and Research in Clinical Haematology</i> , 2016 , 29, 136-147	4.2	9
177	Subjects with chronic lymphocytic leukaemia-like B-cell clones with stereotyped B-cell receptors frequently show MDS-associated phenotypes on myeloid cells. <i>British Journal of Haematology</i> , 2015 , 168, 258-67	4.5	5

176	Effect of mismatching for mHA UTA2-1 on clinical outcome after HLA-identical sibling donor allo-SCT. <i>Bone Marrow Transplantation</i> , 2015 , 50, 298-300	4.4	1
175	Non-coding recurrent mutations in chronic lymphocytic leukaemia. <i>Nature</i> , 2015 , 526, 519-24	50.4	565
174	The predominant myeloma clone at diagnosis, CDR3 defined, is constantly detectable across all stages of disease evolution. <i>Leukemia</i> , 2015 , 29, 1435-7	10.7	14
173	MicroRNA-223 is a novel negative regulator of HSP90B1 in CLL. <i>BMC Cancer</i> , 2015 , 15, 238	4.8	12
172	A B-cell epigenetic signature defines three biologic subgroups of chronic lymphocytic leukemia with clinical impact. <i>Leukemia</i> , 2015 , 29, 598-605	10.7	92
171	CXCR4 expression enhances diffuse large B cell lymphoma dissemination and decreases patient survival. <i>Journal of Pathology</i> , 2015 , 235, 445-55	9.4	55
170	Detection of chromothripsis-like patterns with a custom array platform for chronic lymphocytic leukemia. <i>Genes Chromosomes and Cancer</i> , 2015 , 54, 668-80	5	18
169	Panobinostat as part of induction and maintenance for elderly patients with newly diagnosed acute myeloid leukemia: phase Ib/II panobidara study. <i>Haematologica</i> , 2015 , 100, 1294-300	6.6	23
168	CIP2A high expression is a poor prognostic factor in normal karyotype acute myeloid leukemia. <i>Haematologica</i> , 2015 , 100, e183-5	6.6	15
167	The Genotype of the Donor for the (GT) _n Polymorphism in the Promoter/Enhancer of FOXP3 Is Associated with the Development of Severe Acute GVHD but Does Not Affect the GVL Effect after Myeloablative HLA-Identical Allogeneic Stem Cell Transplantation. <i>PLoS ONE</i> , 2015 , 10, e0140454	3.7	8
166	A Low Frequency of Losses in 11q Chromosome Is Associated with Better Outcome and Lower Rate of Genomic Mutations in Patients with Chronic Lymphocytic Leukemia. <i>PLoS ONE</i> , 2015 , 10, e0143073	3.7	23
165	Do endothelial cells belong to the primitive stem leukemic clone in CML? Role of extracellular vesicles. <i>Leukemia Research</i> , 2015 , 39, 921-4	2.7	10
164	Extranodal and nodal diffuse large B cell lymphoma of the head and neck: two different entities?. <i>Annals of Hematology</i> , 2015 , 94, 609-16	3	7
163	Role of minimal residual disease and chimerism after reduced-intensity and myeloablative allo-transplantation in acute myeloid leukemia and high-risk myelodysplastic syndrome. <i>Leukemia Research</i> , 2014 , 38, 551-6	2.7	8
162	Hairy cell leukemia treated initially with purine analogs: a retrospective study of 107 patients from the Spanish Cooperative Group on Chronic Lymphocytic Leukemia (GELLC). <i>Leukemia and Lymphoma</i> , 2014 , 55, 1007-12	1.9	14
161	Mutations in TLR/MYD88 pathway identify a subset of young chronic lymphocytic leukemia patients with favorable outcome. <i>Blood</i> , 2014 , 123, 3790-6	2.2	82
160	Genomic complexity and IGHV mutational status are key predictors of outcome of chronic lymphocytic leukemia patients with TP53 disruption. <i>Haematologica</i> , 2014 , 99, e231-4	6.6	27
159	Prognostic value of deep sequencing method for minimal residual disease detection in multiple myeloma. <i>Blood</i> , 2014 , 123, 3073-9	2.2	306

158	Lineage-specific function of Engrailed-2 in the progression of chronic myelogenous leukemia to T-cell blast crisis. <i>Cell Cycle</i> , 2014 , 13, 1717-26	4.7	7
157	Circulating clonotypic B cells in multiple myeloma and monoclonal gammopathy of undetermined significance. <i>Haematologica</i> , 2014 , 99, 155-62	6.6	23
156	Molecular and cytogenetic characterization of expanded B-cell clones from multiclonal versus monoclonal B-cell chronic lymphoproliferative disorders. <i>Haematologica</i> , 2014 , 99, 897-907	6.6	17
155	Late differentiation syndrome in acute promyelocytic leukemia: a challenging diagnosis. <i>Hematology Reports</i> , 2014 , 6, 5654	0.9	1
154	TET2 overexpression in chronic lymphocytic leukemia is unrelated to the presence of TET2 variations. <i>BioMed Research International</i> , 2014 , 2014, 814294	3	11
153	Detection of MYD88 L265P mutation by real-time allele-specific oligonucleotide polymerase chain reaction. <i>Applied Immunohistochemistry and Molecular Morphology</i> , 2014 , 22, 768-73	1.9	22
152	Critical evaluation of ASO RQ-PCR for minimal residual disease evaluation in multiple myeloma. A comparative analysis with flow cytometry. <i>Leukemia</i> , 2014 , 28, 391-7	10.7	129
151	The use of CD138 positively selected marrow samples increases the applicability of minimal residual disease assessment by PCR in patients with multiple myeloma. <i>Annals of Hematology</i> , 2013 , 92, 97-100	3	16
150	Pediatric primary follicular mucinosis: further evidence of its relationship with mycosis fungoides. <i>Pediatric Dermatology</i> , 2013 , 30, e218-20	1.9	8
149	Chronic lymphocytic leukemia: a clinical and molecular heterogeneous disease. <i>Cancer Genetics</i> , 2013 , 206, 49-62	2.3	51
148	NOTCH1 mutations identify a genetic subgroup of chronic lymphocytic leukemia patients with high risk of transformation and poor outcome. <i>Leukemia</i> , 2013 , 27, 1100-6	10.7	135
147	MYD88 L265P is a marker highly characteristic of, but not restricted to, Waldenström macroglobulinemia. <i>Leukemia</i> , 2013 , 27, 1722-8	10.7	179
146	Clinical practice guidelines for diagnosis, treatment, and follow-up of patients with mantle cell lymphoma. Recommendations from the GEL/TAMO Spanish Cooperative Group. <i>Annals of Hematology</i> , 2013 , 92, 1151-79	3	18
145	Rituximab maintenance after first-line therapy with rituximab, fludarabine, cyclophosphamide, and mitoxantrone (R-FCM) for chronic lymphocytic leukemia. <i>Blood</i> , 2013 , 122, 3951-9	2.2	47
144	Evaluating gene expression profiling by quantitative polymerase chain reaction to develop a clinically feasible test for outcome prediction in multiple myeloma. <i>British Journal of Haematology</i> , 2013 , 163, 223-34	4.5	5
143	HLA specificities are related to development and prognosis of diffuse large B-cell lymphoma. <i>Blood</i> , 2013 , 122, 1448-54	2.2	18
142	Combined patterns of IGHV repertoire and cytogenetic/molecular alterations in monoclonal B lymphocytosis versus chronic lymphocytic leukemia. <i>PLoS ONE</i> , 2013 , 8, e67751	3.7	21
141	Risk of placenta-mediated pregnancy complications or pregnancy-related VTE in VTE-asymptomatic families of probands with VTE and heterozygosity for factor V Leiden or G20210 prothrombin mutation. <i>European Journal of Haematology</i> , 2012 , 89, 250-5	3.8	4

140	Chronic lymphocytic leukaemia with 17p deletion: a retrospective analysis of prognostic factors and therapy results. <i>British Journal of Haematology</i> , 2012 , 157, 67-74	4.5	36
139	Impaired expression of DICER, DROSHA, SBDS and some microRNAs in mesenchymal stromal cells from myelodysplastic syndrome patients. <i>Haematologica</i> , 2012 , 97, 1218-24	6.6	74
138	Molecular characterization of chronic lymphocytic leukemia patients with a high number of losses in 13q14. <i>PLoS ONE</i> , 2012 , 7, e48485	3.7	29
137	Identification of a novel recurrent gain on 20q13 in chronic lymphocytic leukemia by array CGH and gene expression profiling. <i>Annals of Oncology</i> , 2012 , 23, 2138-2146	10.3	12
136	Molecular characterization of immunoglobulin gene rearrangements in diffuse large B-cell lymphoma: antigen-driven origin and IGHV4-34 as a particular subgroup of the non-GCB subtype. <i>American Journal of Pathology</i> , 2012 , 181, 1879-88	5.8	24
135	Multiple cranial neuropathy and intracranial hypertension associated with all-trans retinoic acid treatment in a young adult patient with acute promyelocytic leukemia. <i>International Journal of Hematology</i> , 2012 , 96, 383-5	2.3	7
134	Prognostic significance of FLT3 mutational status and expression levels in MLL-AF4+ and MLL-germline acute lymphoblastic leukemia. <i>Leukemia</i> , 2012 , 26, 2360-6	10.7	45
133	The EuroChimerism concept for a standardized approach to chimerism analysis after allogeneic stem cell transplantation. <i>Leukemia</i> , 2012 , 26, 1821-8	10.7	67
132	SNP-based mapping arrays reveal high genomic complexity in monoclonal gammopathies, from MGUS to myeloma status. <i>Leukemia</i> , 2012 , 26, 2521-9	10.7	81
131	Expression of MALT1 oncogene in hematopoietic stem/progenitor cells recapitulates the pathogenesis of human lymphoma in mice. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, 10534-9	11.5	58
130	Common infectious agents and monoclonal B-cell lymphocytosis: a cross-sectional epidemiological study among healthy adults. <i>PLoS ONE</i> , 2012 , 7, e52808	3.7	25
129	Exome sequencing identifies recurrent mutations of the splicing factor SF3B1 gene in chronic lymphocytic leukemia. <i>Nature Genetics</i> , 2011 , 44, 47-52	36.3	752
128	Upregulation of Dicer is more frequent in monoclonal gammopathies of undetermined significance than in multiple myeloma patients and is associated with longer survival in symptomatic myeloma patients. <i>Haematologica</i> , 2011 , 96, 468-71	6.6	28
127	Prognostic value of FLT3 mutations in patients with acute promyelocytic leukemia treated with all-trans retinoic acid and anthracycline monochemotherapy. <i>Haematologica</i> , 2011 , 96, 1470-7	6.6	48
126	Clinical significance of CD56 expression in patients with acute promyelocytic leukemia treated with all-trans retinoic acid and anthracycline-based regimens. <i>Blood</i> , 2011 , 117, 1799-805	2.2	95
125	Frequency of HLA-A, -B and -DRB1 specificities and haplotypic associations in the population of Castilla y Leń (northwest-central Spain). <i>Tissue Antigens</i> , 2011 , 78, 249-55		13
124	Sustained complete remission with single agent rituximab in relapsed follicular lymphoma as transformed disease after unrelated reduced intensity conditioning allogeneic stem cell transplantation. <i>Annals of Hematology</i> , 2011 , 90, 227-9	3	0
123	Molecular and flow cytometry characterization during the follow-up of three simultaneous lymphoproliferative disorders: hairy cell leukemia, monoclonal B-cell lymphocytosis, and CD4(++)/CD8(+/- dim) T-large granular lymphocytosis--a case report. <i>Cytometry Part B - Clinical Cytometry</i> , 2011 , 80, 185-200	3.4	4

122	Whole-genome sequencing identifies recurrent mutations in chronic lymphocytic leukaemia. <i>Nature</i> , 2011 , 475, 101-5	50.4	1206
121	Deregulation of microRNA expression in the different genetic subtypes of multiple myeloma and correlation with gene expression profiling. <i>Leukemia</i> , 2010 , 24, 629-37	10.7	173
120	International network of cancer genome projects. <i>Nature</i> , 2010 , 464, 993-8	50.4	1613
119	Concurrent intensive chemotherapy and imatinib before and after stem cell transplantation in newly diagnosed Philadelphia chromosome-positive acute lymphoblastic leukemia. Final results of the CSTIBES02 trial. <i>Haematologica</i> , 2010 , 95, 87-95	6.6	136
118	Long FLT3 internal tandem duplications and reduced PML-RAR α expression at diagnosis characterize a high-risk subgroup of acute promyelocytic leukemia patients. <i>Haematologica</i> , 2010 , 95, 745-51	6.6	42
117	Early intervention during imatinib therapy in patients with newly diagnosed chronic-phase chronic myeloid leukemia: a study of the Spanish PETHEMA group. <i>Haematologica</i> , 2010 , 95, 1317-24	6.6	41
116	Does microgranular variant morphology of acute promyelocytic leukemia independently predict a less favorable outcome compared with classical M3 APL? A joint study of the North American Intergroup and the PETHEMA Group. <i>Blood</i> , 2010 , 116, 5650-9	2.2	29
115	Additional chromosome abnormalities in patients with acute promyelocytic leukemia treated with all-trans retinoic acid and chemotherapy. <i>Haematologica</i> , 2010 , 95, 424-31	6.6	72
114	BAALC is an important predictor of refractoriness to chemotherapy and poor survival in intermediate-risk acute myeloid leukemia (AML). <i>Annals of Hematology</i> , 2010 , 89, 453-8	3	33
113	Mapping of genetic abnormalities of primary tumours from metastatic CRC by high-resolution SNP arrays. <i>PLoS ONE</i> , 2010 , 5, e13752	3.7	20
112	Rituximab, fludarabine, cyclophosphamide, and mitoxantrone: a new, highly active chemoimmunotherapy regimen for chronic lymphocytic leukemia. <i>Journal of Clinical Oncology</i> , 2009 , 27, 4578-84	2.2	111
111	Array comparative genomic hybridization identifies genetic regions associated with outcome in aggressive diffuse large B-cell lymphomas. <i>Cancer</i> , 2009 , 115, 3728-37	6.4	29
110	Multitargeted sequential therapy with MK-0457 and dasatinib followed by stem cell transplantation for T315I mutated chronic myeloid leukemia. <i>Leukemia Research</i> , 2009 , 33, e20-2	2.7	4
109	High FOXO3a expression is associated with a poorer prognosis in AML with normal cytogenetics. <i>Leukemia Research</i> , 2009 , 33, 1706-9	2.7	44
108	A high number of losses in 13q14 chromosome band is associated with a worse outcome and biological differences in patients with B-cell chronic lymphoid leukemia. <i>Haematologica</i> , 2009 , 94, 364-71	6.6	50
107	Influence of MBL-2 mutations in the infection risk of patients with follicular lymphoma treated with rituximab, fludarabine, and cyclophosphamide. <i>Leukemia and Lymphoma</i> , 2009 , 50, 1283-9	1.9	5
106	Molecular and epidemiologic findings of childhood acute leukemia in Costa Rica. <i>Journal of Pediatric Hematology/Oncology</i> , 2009 , 31, 131-5	1.2	9
105	Central nervous system involvement at first relapse in patients with acute promyelocytic leukemia treated with all-trans retinoic acid and anthracycline monochemotherapy without intrathecal prophylaxis. <i>Haematologica</i> , 2009 , 94, 1242-9	6.6	78

104	Differentiation syndrome in patients with acute promyelocytic leukemia treated with all-trans retinoic acid and anthracycline chemotherapy: characteristics, outcome, and prognostic factors. <i>Blood</i> , 2009 , 113, 775-83	2.2	232
103	Molecular stratification model for prognosis in cytogenetically normal acute myeloid leukemia. <i>Blood</i> , 2009 , 114, 148-52	2.2	68
102	Increased frequency (12%) of circulating chronic lymphocytic leukemia-like B-cell clones in healthy subjects using a highly sensitive multicolor flow cytometry approach. <i>Blood</i> , 2009 , 114, 33-7	2.2	164
101	Inhibition of proteasome by bortezomib causes intracellular aggregation of hepatic serpins and increases the latent circulating form of antithrombin. <i>Laboratory Investigation</i> , 2008 , 88, 306-17	5.9	11
100	The presence of DRB1*01 allele in multiple myeloma patients is associated with an indolent disease. <i>Tissue Antigens</i> , 2008 , 71, 548-51		3
99	Low expression of ZHX2, but not RCBTB2 or RAN, is associated with poor outcome in multiple myeloma. <i>British Journal of Haematology</i> , 2008 , 141, 212-5	4.5	22
98	Chromosome 14q32 translocations involving the immunoglobulin heavy chain locus in chronic lymphocytic leukaemia identify a disease subset with poor prognosis. <i>British Journal of Haematology</i> , 2008 , 142, 529-37	4.5	69
97	Fludarabine, cyclophosphamide, and mitoxantrone as initial therapy of chronic lymphocytic leukemia: high response rate and disease eradication. <i>Clinical Cancer Research</i> , 2008 , 14, 155-61	12.9	103
96	Causes and prognostic factors of remission induction failure in patients with acute promyelocytic leukemia treated with all-trans retinoic acid and idarubicin. <i>Blood</i> , 2008 , 111, 3395-402	2.2	258
95	Association between the proliferative rate of neoplastic B cells, their maturation stage, and underlying cytogenetic abnormalities in B-cell chronic lymphoproliferative disorders: analysis of a series of 432 patients. <i>Blood</i> , 2008 , 111, 5130-41	2.2	17
94	Expanded cells in monoclonal TCR-alpha/beta+/CD4+/NKa+/CD8-/dim T-LGL lymphocytosis recognize hCMV antigens. <i>Blood</i> , 2008 , 112, 4609-16	2.2	48
93	Risk-adapted treatment of acute promyelocytic leukemia with all-trans retinoic acid and anthracycline monochemotherapy: long-term outcome of the LPA 99 multicenter study by the PETHEMA Group. <i>Blood</i> , 2008 , 112, 3130-4	2.2	129
92	Bisphosphonate-related osteonecrosis of the jaw is associated with polymorphisms of the cytochrome P450 CYP2C8 in multiple myeloma: a genome-wide single nucleotide polymorphism analysis. <i>Blood</i> , 2008 , 112, 2709-12	2.2	184
91	The relevance of preferentially expressed antigen of melanoma (PRAME) as a marker of disease activity and prognosis in acute promyelocytic leukemia. <i>Haematologica</i> , 2008 , 93, 1797-805	6.6	33
90	Clinical and prognostic value of discrepancies in microsatellite DNA regions between recipient and donor in human leukocyte antigen-identical allogeneic transplantation setting. <i>Transplantation</i> , 2008 , 86, 983-90	1.8	4
89	Molecular characterization of complete and incomplete immunoglobulin heavy chain gene rearrangements in hairy cell leukemia. <i>Clinical Lymphoma and Myeloma</i> , 2007 , 7, 573-9		8
88	Outcome of patients with acute promyelocytic leukemia failing to front-line treatment with all-trans retinoic acid and anthracycline-based chemotherapy (PETHEMA protocols LPA96 and LPA99): benefit of an early intervention. <i>Leukemia</i> , 2007 , 21, 446-52	10.7	110
87	Role of MTHFR (677, 1298) haplotype in the risk of developing secondary leukemia after treatment of breast cancer and hematological malignancies. <i>Leukemia</i> , 2007 , 21, 1413-22	10.7	39

86	Characterization of a reference material for BCR-ABL (M-BCR) mRNA quantitation by real-time amplification assays: towards new standards for gene expression measurements. <i>Leukemia</i> , 2007 , 21, 1481-7	10.7	25
85	Functional class switch recombination may occur <i>in vivo</i> in Waldenström macroglobulinaemia. <i>British Journal of Haematology</i> , 2007 , 136, 114-6	4.5	12
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