Marcos Gonzalez DÃ-az

List of Publications by Year
in descending order

230 papers	20,570 citations	20817 60 h-index	11052 137 g-index
233 all docs	233 docs citations	233 times ranked	22082 citing authors

#	Article	IF	Citations
1	Interlaboratory Analytical Validation of a Next-Generation Sequencing Strategy for Clonotypic Assessment and Minimal Residual Disease Monitoring in Multiple Myeloma. Archives of Pathology and Laboratory Medicine, 2022, 146, 862-871.	2.5	7
2	Cellular and humoral immunogenicity of the mRNA-1273 SARS-CoV-2 vaccine in patients with hematologic malignancies. Blood Advances, 2022, 6, 774-784.	5.2	42
3	Overexpression of wild type RRAS2, without oncogenic mutations, drives chronic lymphocytic leukemia. Molecular Cancer, 2022, 21, 35.	19.2	11
4	MYD88 Mutations: Transforming the Landscape of IgM Monoclonal Gammopathies. International Journal of Molecular Sciences, 2022, 23, 5570.	4.1	14
5	Genetic complexity impacts the clinical outcome of follicular lymphoma patients. Blood Cancer Journal, 2021, 11, 11.	6.2	3
6	Impact of measurable residual disease by decentralized flow cytometry: a PETHEMA real-world study in 1076 patients with acute myeloid leukemia. Leukemia, 2021, 35, 2358-2370.	7.2	31
7	Siglec-6 is a novel target for CAR T-cell therapy in acute myeloid leukemia. Blood, 2021, 138, 1830-1842.	1.4	40
8	Liquid biopsy: a nonâ€invasive approach for Hodgkin lymphoma genotyping. British Journal of Haematology, 2021, 195, 542-551.	2.5	14
9	Allele and haplotype frequencies of HLA-A, -B, -C, -DRB1, -DQB1 and -DQA1 in Castile and Leon region from North West of Spain. Human Immunology, 2021, 82, 549-550.	2.4	1
10	Genomic mutation profile in progressive chronic lymphocytic leukemia patients prior to first-line chemoimmunotherapy with FCR and rituximab maintenance (REM). PLoS ONE, 2021, 16, e0257353.	2.5	1
11	Building a network of TP53 and IGHV testing reference centers across Spain: the Red53 initiative. Annals of Hematology, 2021, 100, 825-830.	1.8	2
12	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. Frontiers in Oncology, 2021, 11, 723722.	2.8	0
13	Characteristics and outcome of adult patients with acute promyelocytic leukemia and increased body mass index treated with the PETHEMA Protocols. European Journal of Haematology, 2020, 104, 162-169.	2.2	6
14	Comparison of next-generation sequencing (NGS) and next-generation flow (NGF) for minimal residual disease (MRD) assessment in multiple myeloma. Blood Cancer Journal, 2020, 10, 108.	6.2	60
15	Identification of relapseâ€associated gene mutations by nextâ€generation sequencing in lowâ€risk acute myeloid leukaemia patients. British Journal of Haematology, 2020, 189, 718-730.	2.5	12
16	Biological Features and Prognostic Impact of Bone Marrow Infiltration in Patients with Diffuse Large B-cell Lymphoma. Cancers, 2020, 12, 474.	3.7	10
17	Biological and clinical significance of dysplastic hematopoiesis in patients with newly diagnosed multiple myeloma. Blood, 2020, 135, 2375-2387.	1.4	24
18	Immunoglobulin gene rearrangement IGHV3-48 is a predictive marker of histological transformation into aggressive lymphoma in follicular lymphomas. Blood Cancer Journal, 2019, 9, 52.	6.2	11

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19	Life expectancy of follicular lymphoma patients in complete response at 30Âmonths is similar to that of the Spanish general population. British Journal of Haematology, 2019, 185, 480-491.	2.5	26
20	Blood monitoring of circulating tumor plasma cells by next generation flow in multiple myeloma after therapy. Blood, 2019, 134, 2218-2222.	1.4	66
21	Exportinâ€1 E571K mutation is a common finding in patients with classical Hodgkin lymphoma. Hematological Oncology, 2019, 37, 215-218.	1.7	2
22	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. Haematologica, 2019, 104, 576-586.	3.5	40
23	Richter transformation driven by Epsteinâ€“Barr virus reactivation during therapyâ€related immunosuppression in chronic lymphocytic leukaemia. Journal of Pathology, 2018, 245, 61-73.	4.5	24
24	Low-count monoclonal B-cell lymphocytosis persists after seven years of follow up and is associated with a poorer outcome. Haematologica, 2018, 103, 1198-1208.	3.5	34
25	Basophil-lineage commitment in acute promyelocytic leukemia predicts for severe bleeding after starting therapy. Modern Pathology, 2018, 31, 1318-1331.	5.5	9
26	Characterizing patients with multiple chromosomal aberrations detected by FISH in chronic lymphocytic leukemia. Leukemia and Lymphoma, 2018, 59, 633-642.	1.3	8
27	The mutational landscape of small lymphocytic lymphoma compared to non-early stage chronic lymphocytic leukemia. Leukemia and Lymphoma, 2018, 59, 2318-2326.	1.3	5
28	Focal Adhesion Genes Refine the Intermediate-Risk Cytogenetic Classification of Acute Myeloid Leukemia. Cancers, 2018, 10, 436.	3.7	8
29	A novel predictive approach for GVHD after allogeneic SCT based on clinical variables and cytokine gene polymorphisms. Blood Advances, 2018, 2, 1719-1737.	5.2	25
30	Residual normal B-cell profiles in monoclonal B-cell lymphocytosis versus chronic lymphocytic leukemia. Leukemia, 2018, 32, 2701-2705.	7.2	19
31	A novel ex vivo high-throughput assay reveals antiproliferative effects of idelalisib and ibrutinib in chronic lymphocytic leukemia. Oncotarget, 2018, 9, 26019-26031.	1.8	8
32	Next Generation Flow for highly sensitive and standardized detection of minimal residual disease in multiple myeloma. Leukemia, 2017, 31, 2094-2103.	7.2	486
33	ActualizaciÃ³n de las guÃas nacionales de consenso del Grupo EspaÃ±ol de Leucemia LinfocÃtica CrÃ³nica para el tratamiento y seguimiento de la leucemia linfocÃtica crÃ³nica. Medicina ClÃnica, 2017, 148, 381.e1-381.e9.	0.6	2
34	Host virus and pneumococcus-specific immune responses in high-count monoclonal B-cell lymphocytosis and chronic lymphocytic leukemia: implications for disease progression. Haematologica, 2017, 102, 1238-1246.	3.5	9
35	Next-generation sequencing and FISH studies reveal the appearance of gene mutations and chromosomal abnormalities in hematopoietic progenitors in chronic lymphocytic leukemia. Journal of Hematology and Oncology, 2017, 10, 83.	17.0	38
36	Donor CTLA-4 Genotype Modulates the Immune Response to Minor Histocompatibility Antigen Mismatches. Biology of Blood and Marrow Transplantation, 2017, 23, 2042-2047.	2.0	13

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37	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. Leukemia and Lymphoma, 2017, 58, 1144-1152.	1.3	28
38	Application of a molecular diagnostic algorithm for haemophilia A and B using next-generation sequencing of entire F8, F9 and VWF genes. Thrombosis and Haemostasis, 2017, 117, 66-74.	3.4	36
39	HLA specificities are associated with prognosis in IGHV-mutated CLL-like high-count monoclonal B cell lymphocytosis. PLoS ONE, 2017, 12, e0172978.	2.5	4
40	Mesenchymal stromal cells (MSC) from JAK2+ myeloproliferative neoplasms differ from normal MSC and contribute to the maintenance of neoplastic hematopoiesis. PLoS ONE, 2017, 12, e0182470.	2.5	19
41	Karyotypic complexity rather than chromosome 8 abnormalities aggravates the outcome of chronic lymphocytic leukemia patients with TP53 aberrations. Oncotarget, 2016, 7, 80916-80924.	1.8	29
42	Microvesicles from Mesenchymal Stromal Cells Are Involved in HPC-Microenvironment Crosstalk in Myelodysplastic Patients. PLoS ONE, 2016, 11, e0146722.	2.5	70
43	Design and application of a 23â€gene panel by nextâ€generation sequencing for inherited coagulation bleeding disorders. Haemophilia, 2016, 22, 590-597.	2.1	43
44	A high proportion of cells carrying trisomy 12 is associated with a worse outcome in patients with chronic lymphocytic leukemia. Hematological Oncology, 2016, 34, 84-92.	1.7	26
45	Multipronged functional proteomics approaches for global identification of altered cell signalling pathways in Bâ€cell chronic lymphocytic leukaemia. Proteomics, 2016, 16, 1193-1203.	2.2	15
46	Homeobox NKX2-3 promotes marginal-zone lymphomagenesis by activating B-cell receptor signalling and shaping lymphocyte dynamics. Nature Communications, 2016, 7, 11889.	12.8	42
47	MiRNA expression profile of chronic lymphocytic leukemia patients with 13q deletion. Leukemia Research, 2016, 46, 30-36.	0.8	8
48	Clinical impact of clonal and subclonal TP53, SF3B1, BIRC3, NOTCH1, and ATM mutations in chronic lymphocytic leukemia. Blood, 2016, 127, 2122-2130.	1.4	260
49	Origin of Waldenstrom's macroglobulinaemia. Best Practice and Research in Clinical Haematology, 2016, 29, 136-147.	1.7	17
50	Sarcoma histiocÃtico en intestino delgado: estudio y discusiÃ³n de un caso con reordenamiento clonal linfoide B. Revista Espanola De Patologia, 2016, 49, 254-258.	0.2	0
51	High-resolution copy number analysis of paired normal-tumor samples from diffuse large B cell lymphoma. Annals of Hematology, 2016, 95, 253-262.	1.8	19
52	Minimal residual disease evaluation by flow cytometry is a complementary tool to cytogenetics for treatment decisions in acute myeloid leukaemia. Leukemia Research, 2016, 40, 1-9.	0.8	29
53	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. Blood, 2016, 128, 375-375.	1.4	1
54	Detection of chromothripsisâ€like patterns with a custom array platform for chronic lymphocytic leukemia. Genes Chromosomes and Cancer, 2015, 54, 668-680.	2.8	23

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55	Panobinostat as part of induction and maintenance for elderly patients with newly diagnosed acute myeloid leukemia: phase Ib/II panobidara study. Haematologica, 2015, 100, 1294-1300.	3.5	27
56	CIP2A high expression is a poor prognostic factor in normal karyotype acute myeloid leukemia. Haematologica, 2015, 100, e183-e185.	3.5	20
57	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. PLoS ONE, 2015, 10, e0140454.	2.5	11
58	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. PLoS ONE, 2015, 10, e0143073.	2.5	24
59	Do endothelial cells belong to the primitive stem leukemic clone in CML? Role of extracellular vesicles. Leukemia Research, 2015, 39, 921-924.	0.8	12
60	Extranodal and nodal diffuse large B cell lymphoma of the head and neck: two different entities?. Annals of Hematology, 2015, 94, 609-616.	1.8	7
61	Subjects with chronic lymphocytic leukaemiaâ€like <scp>B</scp>â€cell clones with stereotyped <scp>B</scp>â€cell receptors frequently show <scp>MDS</scp>â€associated phenotypes on myeloid cells. British Journal of Haematology, 2015, 168, 258-267.	2.5	5
62	Effect of mismatching for mHA UTA2-1 on clinical outcome after HLA-identical sibling donor allo-SCT. Bone Marrow Transplantation, 2015, 50, 298-300.	2.4	1
63	Non-coding recurrent mutations in chronic lymphocytic leukaemia. Nature, 2015, 526, 519-524.	27.8	749
64	The predominant myeloma clone at diagnosis, CDR3 defined, is constantly detectable across all stages of disease evolution. Leukemia, 2015, 29, 1435-1437.	7.2	17
65	MicroRNA-223 is a novel negative regulator of HSP90B1 in CLL. BMC Cancer, 2015, 15, 238.	2.6	16
66	A B-cell epigenetic signature defines three biologic subgroups of chronic lymphocytic leukemia with clinical impact. Leukemia, 2015, 29, 598-605.	7.2	129
67	<scp>CXCR4</scp> expression enhances diffuse large B cell lymphoma dissemination and decreases patient survival. Journal of Pathology, 2015, 235, 445-455.	4.5	71
68	Lineage-specific function of Engrailed-2 in the progression of chronic myelogenous leukemia to T-cell blast crisis. Cell Cycle, 2014, 13, 1717-1726.	2.6	7
69	Circulating clonotypic B cells in multiple myeloma and monoclonal gammopathy of undetermined significance. Haematologica, 2014, 99, 155-162.	3.5	23
70	Molecular and cytogenetic characterization of expanded B-cell clones from multiclonal versus monoclonal B-cell chronic lymphoproliferative disorders. Haematologica, 2014, 99, 897-907.	3.5	22
71	Late differentiation syndrome in acute promyelocytic leukemia: a challenging diagnosis. Hematology Reports, 2014, 6, 5654.	0.8	4
72	TET2Overexpression in Chronic Lymphocytic Leukemia Is Unrelated to the Presence ofTET2Variations. BioMed Research International, 2014, 2014, 1-6.	1.9	12

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73	Detection of MYD88 L265P Mutation by Real-Time Allele-Specific Oligonucleotide Polymerase Chain Reaction. Applied Immunohistochemistry and Molecular Morphology, 2014, 22, 768-773.	1.2	28
74	Critical evaluation of ASO RQ-PCR for minimal residual disease evaluation in multiple myeloma. A comparative analysis with flow cytometry. Leukemia, 2014, 28, 391-397.	7.2	155
75	Role of minimal residual disease and chimerism after reduced-intensity and myeloablative allo-transplantation in acute myeloid leukemia and high-risk myelodysplastic syndrome. Leukemia Research, 2014, 38, 551-556.	0.8	11
76	Hairy cell leukemia treated initially with purine analogs: a retrospective study of 107 patients from the Spanish Cooperative Group on Chronic Lymphocytic Leukemia (GELLC). Leukemia and Lymphoma, 2014, 55, 1007-1012.	1.3	20
77	Mutations in TLR/MYD88 pathway identify a subset of young chronic lymphocytic leukemia patients with favorable outcome. Blood, 2014, 123, 3790-3796.	1.4	97
78	Genomic complexity and IGHV mutational status are key predictors of outcome of chronic lymphocytic leukemia patients with TP53 disruption. Haematologica, 2014, 99, e231-e234.	3.5	33
79	Prognostic value of deep sequencing method for minimal residual disease detection in multiple myeloma. Blood, 2014, 123, 3073-3079.	1.4	380
80	The use of CD138 positively selected marrow samples increases the applicability of minimal residual disease assessment by PCR in patients with multiple myeloma. Annals of Hematology, 2013, 92, 97-100.	1.8	18
81	Pediatric Primary Follicular Mucinosis: Further Evidence of its Relationship with Mycosis Fungoides. Pediatric Dermatology, 2013, 30, e218-20.	0.9	10
82	Chronic lymphocytic leukemia: a clinical and molecular heterogenous disease. Cancer Genetics, 2013, 206, 49-62.	0.4	63
83	NOTCH1 mutations identify a genetic subgroup of chronic lymphocytic leukemia patients with high risk of transformation and poor outcome. Leukemia, 2013, 27, 1100-1106.	7.2	167
84	MYD88 L265P is a marker highly characteristic of, but not restricted to, WaldenstrÃ¶mâ€™s macroglobulinemia. Leukemia, 2013, 27, 1722-1728.	7.2	238
85	Clinical practice guidelines for diagnosis, treatment, and follow-up of patients with mantle cell lymphoma. Recommendations from the GEL/TAMO Spanish Cooperative Group. Annals of Hematology, 2013, 92, 1151-1179.	1.8	22
86	Rituximab maintenance after first-line therapy with rituximab, fludarabine, cyclophosphamide, and mitoxantrone (R-FCM) for chronic lymphocytic leukemia. Blood, 2013, 122, 3951-3959.	1.4	55
87	Evaluating gene expression profiling by quantitative polymerase chain reaction to develop a clinically feasible test for outcome prediction in multiple myeloma. British Journal of Haematology, 2013, 163, 223-234.	2.5	7
88	HLA specificities are related to development and prognosis of diffuse large B-cell lymphoma. Blood, 2013, 122, 1448-1454.	1.4	23
89	Combined Patterns of IGHV Repertoire and Cytogenetic/Molecular Alterations in Monoclonal B Lymphocytosis versus Chronic Lymphocytic Leukemia. PLoS ONE, 2013, 8, e67751.	2.5	27
90	SNP-based mapping arrays reveal high genomic complexity in monoclonal gammopathies, from MGUS to myeloma status. Leukemia, 2012, 26, 2521-2529.	7.2	100

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91	Expression of MALT1 oncogene in hematopoietic stem/progenitor cells recapitulates the pathogenesis of human lymphoma in mice. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 10534-10539.	7.1	73
92	Impaired expression of DICER, DROSHA, SBDS and some microRNAs in mesenchymal stromal cells from myelodysplastic syndrome patients. Haematologica, 2012, 97, 1218-1224.	3.5	83
93	Molecular Characterization of Chronic Lymphocytic Leukemia Patients with a High Number of Losses in 13q14. PLoS ONE, 2012, 7, e48485.	2.5	37
94	Identification of a novel recurrent gain on 20q13 in chronic lymphocytic leukemia by array CGH and gene expression profiling. Annals of Oncology, 2012, 23, 2138-2146.	1.2	13
95	Molecular Characterization of Immunoglobulin Gene Rearrangements in Diffuse Large B-Cell Lymphoma. American Journal of Pathology, 2012, 181, 1879-1888.	3.8	31
96	Multiple cranial neuropathy and intracranial hypertension associated with all-trans retinoic acid treatment in a young adult patient with acute promyelocytic leukemia. International Journal of Hematology, 2012, 96, 383-385.	1.6	10
97	Prognostic significance of FLT3 mutational status and expression levels in MLL-AF4+ and MLL-germline acute lymphoblastic leukemia. Leukemia, 2012, 26, 2360-2366.	7.2	55
98	Exome sequencing identifies recurrent mutations of the splicing factor SF3B1 gene in chronic lymphocytic leukemia. Nature Genetics, 2012, 44, 47-52.	21.4	893
99	The EuroChimerism concept for a standardized approach to chimerism analysis after allogeneic stem cell transplantation. Leukemia, 2012, 26, 1821-1828.	7.2	83
100	Risk of placentaâ€mediated pregnancy complications or pregnancyâ€related <scp>VTE</scp> in <scp>VTE</scp>â€asymptomatic families of probands with <scp>VTE</scp> and heterozygosity for factor V <scp>L</scp>eiden or <scp>G</scp>20210 prothrombin mutation. European Journal of Haematology, 2012, 89, 250-255.	2.2	5
101	Chronic lymphocytic leukaemia with 17p deletion: a retrospective analysis of prognostic factors and therapy results. British Journal of Haematology, 2012, 157, 67-74.	2.5	39
102	Common Infectious Agents and Monoclonal B-Cell Lymphocytosis: A Cross-Sectional Epidemiological Study among Healthy Adults. PLoS ONE, 2012, 7, e52808.	2.5	32
103	Whole-genome sequencing identifies recurrent mutations in chronic lymphocytic leukaemia. Nature, 2011, 475, 101-105.	27.8	1,364
104	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. Haematologica, 2011, 96, 468-471.	3.5	29
105	Prognostic value of FLT3 mutations in patients with acute promyelocytic leukemia treated with all-trans retinoic acid and anthracycline monochemotherapy. Haematologica, 2011, 96, 1470-1477.	3.5	59
106	Clinical significance of CD56 expression in patients with acute promyelocytic leukemia treated with all-trans retinoic acid and anthracycline-based regimens. Blood, 2011, 117, 1799-1805.	1.4	112
107	Frequency of HLAâ€A, â€B and â€DRB1 specificities and haplotypic associations in the population of Castilla y LeÃ³n (northwestâ€central Spain). Tissue Antigens, 2011, 78, 249-255.	1.0	14
108	Sustained complete remission with single agent rituximab in relapsed follicular lymphoma as transformed disease after unrelated reduced intensity conditioning allogeneic stem cell transplantation. Annals of Hematology, 2011, 90, 227-229.	1.8	1

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109	Molecular and flow cytometry characterization during the followâ€up of three simultaneous lymphoproliferative disorders: Hairy cell leukemia, monoclonal Bâ€cell lymphocytosis, and CD4⁺⁺/CD8^{+/â¯dim} Tâ€large granular lymphocytosisâ€"A case report. Cytometry Part B - Clinical Cytometry, 2011, 80B, 195-200.	1.5	6
110	Early intervention during imatinib therapy in patients with newly diagnosed chronic-phase chronic myeloid leukemia: a study of the Spanish PETHEMA group. Haematologica, 2010, 95, 1317-1324.	3.5	53
111	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. Blood, 2010, 116, 5650-5659.	1.4	33
112	Additional chromosome abnormalities in patients with acute promyelocytic leukemia treated with all-trans retinoic acid and chemotherapy. Haematologica, 2010, 95, 424-431.	3.5	84
113	BAALC is an important predictor of refractoriness to chemotherapy and poor survival in intermediate-risk acute myeloid leukemia (AML). Annals of Hematology, 2010, 89, 453-458.	1.8	40
114	Deregulation of microRNA expression in the different genetic subtypes of multiple myeloma and correlation with gene expression profiling. Leukemia, 2010, 24, 629-637.	7.2	188
115	International network of cancer genome projects. Nature, 2010, 464, 993-998.	27.8	2,114
116	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. Haematologica, 2010, 95, 87-95.	3.5	164
117	Long FLT3 internal tandem duplications and reduced PML-RARÂ expression at diagnosis characterize a high-risk subgroup of acute promyelocytic leukemia patients. Haematologica, 2010, 95, 745-751.	3.5	47
118	Mapping of Genetic Abnormalities of Primary Tumours from Metastatic CRC by High-Resolution SNP Arrays. PLoS ONE, 2010, 5, e13752.	2.5	22
119	Rituximab, Fludarabine, Cyclophosphamide, and Mitoxantrone: A New, Highly Active Chemoimmunotherapy Regimen for Chronic Lymphocytic Leukemia. Journal of Clinical Oncology, 2009, 27, 4578-4584.	1.6	116
120	Array comparative genomic hybridization identifies genetic regions associated with outcome in aggressive diffuse large Bâ€cell lymphomas. Cancer, 2009, 115, 3728-3737.	4.1	31
121	Multitargeted sequential therapy with MK-0457 and dasatinib followed by stem cell transplantation for T315I mutated chronic myeloid leukemia. Leukemia Research, 2009, 33, e20-e22.	0.8	5
122	High FOXO3a expression is associated with a poorer prognosis in AML with normal cytogenetics. Leukemia Research, 2009, 33, 1706-1709.	0.8	49
123	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. Haematologica, 2009, 94, 364-371.	3.5	59
124	Influence ofMBL-2mutations in the infection risk of patients with follicular lymphoma treated with rituximab, fludarabine, and cyclophosphamide. Leukemia and Lymphoma, 2009, 50, 1283-1289.	1.3	6
125	Molecular and Epidemiologic Findings of Childhood Acute Leukemia in Costa Rica. Journal of Pediatric Hematology/Oncology, 2009, 31, 131-135.	0.6	12
126	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. Haematologica, 2009, 94, 1242-1249.	3.5	93

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127	Differentiation syndrome in patients with acute promyelocytic leukemia treated with all-trans retinoic acid and anthracycline chemotherapy: characteristics, outcome, and prognostic factors. Blood, 2009, 113, 775-783.	1.4	279
128	Molecular stratification model for prognosis in cytogenetically normal acute myeloid leukemia. Blood, 2009, 114, 148-152.	1.4	78
129	Increased frequency (12%) of circulating chronic lymphocytic leukemiaâ€“like B-cell clones in healthy subjects using a highly sensitive multicolor flow cytometry approach. Blood, 2009, 114, 33-37.	1.4	183
130	Inhibition of proteasome by bortezomib causes intracellular aggregation of hepatic serpins and increases the latent circulating form of antithrombin. Laboratory Investigation, 2008, 88, 306-317.	3.7	11
131	The presence of DRB1*01 allele in multiple myeloma patients is associated with an indolent disease. Tissue Antigens, 2008, 71, 548-551.	1.0	3
132	Low expression of ZHX2, but not RCBTB2 or RAN, is associated with poor outcome in multiple myeloma. British Journal of Haematology, 2008, 141, 212-215.	2.5	29
133	Chromosome 14q32 translocations involving the immunoglobulin heavy chain locus in chronic lymphocytic leukaemia identify a disease subset with poor prognosis. British Journal of Haematology, 2008, 142, 529-537.	2.5	78
134	Fludarabine, Cyclophosphamide, and Mitoxantrone as Initial Therapy of Chronic Lymphocytic Leukemia: High Response Rate and Disease Eradication. Clinical Cancer Research, 2008, 14, 155-161.	7.0	117
135	Causes and prognostic factors of remission induction failure in patients with acute promyelocytic leukemia treated with all-trans retinoic acid and idarubicin. Blood, 2008, 111, 3395-3402.	1.4	303
136	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. Blood, 2008, 111, 5130-5141.	1.4	22
137	Expanded cells in monoclonal TCR-Î±Î²+/CD4+/NKa+/CD8â^'/+dim T-LGL lymphocytosis recognize hCMV antigens. Blood, 2008, 112, 4609-4616.	1.4	54
138	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. Blood, 2008, 112, 3130-3134.	1.4	154
139	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. Blood, 2008, 112, 2709-2712.	1.4	213
140	The relevance of preferentially expressed antigen of melanoma (PRAME) as a marker of disease activity and prognosis in acute promyelocytic leukemia. Haematologica, 2008, 93, 1797-1805.	3.5	41
141	Clinical and Prognostic Value of Discrepancies in Microsatellite DNA Regions Between Recipient and Donor in Human Leukocyte Antigen-Identical Allogeneic Transplantation Setting. Transplantation, 2008, 86, 983-990.	1.0	4
142	Molecular characterization of heavy chain immunoglobulin gene rearrangements in Waldenstrom's macroglobulinemia and IgM monoclonal gammopathy of undetermined significance. Haematologica, 2007, 92, 635-642.	3.5	57
143	Monoclonal TCR-VÎ²13.1+/CD4+/NKa+/CD8â^'/+dim T-LGL lymphocytosis: evidence for an antigen-driven chronic T-cell stimulation origin. Blood, 2007, 109, 4890-4898.	1.4	72
144	Antithrombin Cambridge II (A384S): an underestimated genetic risk factor for venous thrombosis. Blood, 2007, 109, 4258-4263.	1.4	104

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145	CTLA-4 polymorphisms and clinical outcome after allogeneic stem cell transplantation from HLA-identical sibling donors.. Blood, 2007, 110, 461-467.	1.4	82
146	Immunoglobulin gene rearrangements and the pathogenesis of multiple myeloma. Blood, 2007, 110, 3112-3121.	1.4	157
147	Hyperhomocysteinemia is a risk factor of recurrent coronary event in young patients irrespective to the MTHFR C677T polymorphism. Thrombosis Research, 2007, 119, 691-698.	1.7	7
148	Molecular Characterization of Complete and Incomplete Immunoglobulin Heavy Chain Gene Rearrangements in Hairy Cell Leukemia. Clinical Lymphoma and Myeloma, 2007, 7, 573-579.	1.4	9
149	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. Leukemia, 2007, 21, 446-452.	7.2	124
150	Role of MTHFR (677, 1298) haplotype in the risk of developing secondary leukemia after treatment of breast cancer and hematological malignancies. Leukemia, 2007, 21, 1413-1422.	7.2	45
151	Characterization of a reference material for BCR-ABL (M-BCR) mRNA quantitation by real-time amplification assays: towards new standards for gene expression measurements. Leukemia, 2007, 21, 1481-1487.	7.2	29
152	Functional class switch recombination may occur â€˜in vivoâ€™ in WaldenstrÃ¶m macroglobulinaemia. British Journal of Haematology, 2007, 136, 114-116.	2.5	18
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