Jesus Maria Hernandez Rivas

List of Publications by Year
in descending order

253 papers	13,430 citations	30070 54 h-index	26613 107 g-index

259 all docs	259 docs citations	259 times ranked	15791 citing authors

#	Article	IF	Citations
1	Molecular Dissection of Structural Variations Involved in Antithrombin Deficiency. Journal of Molecular Diagnostics, 2022, 24, 462-475.	2.8	7
2	Validation of the Burkitt Lymphoma International Prognostic Index in patients treated with two prospective chemoimmunotherapy trials in Spain. Leukemia and Lymphoma, 2022, 63, 1993-1996.	1.3	2
3	TRAF3 alterations are frequent in delâ€3â€²<scp>IGH</scp> chronic lymphocytic leukemia patients and define a specific subgroup with adverse clinical features. American Journal of Hematology, 2022, 97, 903-914.	4.1	3
4	Second Revision of the International Staging System (R2-ISS) for Overall Survival in Multiple Myeloma: A European Myeloma Network (EMN) Report Within the HARMONY Project. Journal of Clinical Oncology, 2022, 40, 3406-3418.	1.6	115
5	CRISPR/Cas9-Directed Gene Trap Constitutes a Selection System for Corrected BCR/ABL Leukemic Cells in CML. International Journal of Molecular Sciences, 2022, 23, 6386.	4.1	3
6	Co-occurrence of cohesin complex and Ras signaling mutations during progression from myelodysplastic syndromes to secondary acute myeloid leukemia. Haematologica, 2021, 106, 2215-2223.	3.5	12
7	Chemotherapy or allogeneic transplantation in high-risk Philadelphia chromosomeâ€“negative adult lymphoblastic leukemia. Blood, 2021, 137, 1879-1894.	1.4	48
8	Rearrangements of ATP5Lâ€KMT2A in acute lymphoblastic leukaemia. British Journal of Haematology, 2021, 192, e139-e144.	2.5	3
9	Dissecting the role of TP53 alterations in del(11q) chronic lymphocytic leukemia. Clinical and Translational Medicine, 2021, 11, e304.	4.0	7
10	NEMHESYSâ€”European Perspective on the Implementation of Next-generation Sequencing Into Clinical Diagnostics. HemaSphere, 2021, 5, e541.	2.7	2
11	The Evolving Landscape of Chronic Lymphocytic Leukemia on Diagnosis, Prognosis and Treatment. Diagnostics, 2021, 11, 853.	2.6	15
12	Biological significance of monoallelic and biallelic BIRC3 loss in del(11q) chronic lymphocytic leukemia progression. Blood Cancer Journal, 2021, 11, 127.	6.2	12
13	Vascular target organ damage in patients with Philadelphia negative myeloproliferative syndrome: A propensity score analysis. Medicina ClÃnica, 2021, , .	0.6	1
14	CRISPR-Cas9 Technology as a Tool to Target Gene Drivers in Cancer: Proof of Concept and New Opportunities to Treat Chronic Myeloid Leukemia. CRISPR Journal, 2021, 4, 519-535.	2.9	3
15	Adverse prognostic impact of complex karyotype (â‰¥3 cytogenetic alterations) in adult T-cell acute lymphoblastic leukemia (T-ALL). Leukemia Research, 2021, 109, 106612.	0.8	11
16	Clinical, biological, and prognostic implications of SF3B1 co-occurrence mutations in very low/low- and intermediate-risk MDS patients. Annals of Hematology, 2021, 100, 1995-2004.	1.8	9
17	A novel genetic variant in <scp>PTGS1</scp> affects Nâ€glycosylation of cyclooxygenaseâ€1 causing a dominantâ€negative effect on platelet function and bleeding diathesis. American Journal of Hematology, 2021, 96, E83-E88.	4.1	2
18	Identification By Whole Exome Sequencing of the Molecular Defect in a Novel Gene Related to Glycosylation in Two Unrelated Families with Syndromic Macrothrombocytopenia. Blood, 2021, 138, 588-588.	1.4	0

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19	Spanish Guidelines for the use of targeted deep sequencing in myelodysplastic syndromes and chronic myelomonocytic leukaemia. British Journal of Haematology, 2020, 188, 605-622.	2.5	25
20	Valores de referencia de parÃ¡metros de rigidez arterial y su relaciÃ³n con los factores de riesgo cardiovascular en poblaciÃ³n espaÃ±ola. Estudio EVA. Revista Espanola De Cardiologia, 2020, 73, 43-52.	1.2	20
21	Transcriptomic analysis of patients with immune thrombocytopenia treated with eltrombopag. Platelets, 2020, 31, 993-1000.	2.3	10
22	Comprehensive Custom NGS Panel Validation for the Improvement of the Stratification of B-Acute Lymphoblastic Leukemia Patients. Journal of Personalized Medicine, 2020, 10, 137.	2.5	4
23	Cytogenetic and genomic analysis of a patient with turner syndrome and t(2;12): a case report. Molecular Cytogenetics, 2020, 13, 46.	0.9	4
24	Chronic lymphocytic leukemia patients with <scp>IGH</scp> translocations are characterized by a distinct genetic landscape with prognostic implications. International Journal of Cancer, 2020, 147, 2780-2792.	5.1	19
25	Helpful Criteria When Implementing NGS Panels in Childhood Lymphoblastic Leukemia. Journal of Personalized Medicine, 2020, 10, 244.	2.5	1
26	Core outcome set measurement for future clinical trials in acute myeloid leukemia: the HARMONY study protocol using a multi-stakeholder consensus-based Delphi process and a final consensus meeting. Trials, 2020, 21, 437.	1.6	5
27	Machine learning applied to gene expression analysis of T-lymphocytes in patients with cGVHD. Bone Marrow Transplantation, 2020, 55, 1668-1670.	2.4	3
28	Integrated Genomic Analysis of Chromosomal Alterations and Mutations in B-Cell Acute Lymphoblastic Leukemia Reveals Distinct Genetic Profiles at Relapse. Diagnostics, 2020, 10, 455.	2.6	6
29	A pediatric regimen for adolescents and young adults with Philadelphia chromosomeâ€negative acute lymphoblastic leukemia: Results of the ALLRE08 PETHEMA trial. Cancer Medicine, 2020, 9, 2317-2329.	2.8	13
30	CRISPR/Cas9-generated models uncover therapeutic vulnerabilities of del(11q) CLL cells to dual BCR and PARP inhibition. Leukemia, 2020, 34, 1599-1612.	7.2	21
31	ETV6/RUNX1 Fusion Gene Abrogation Decreases the Oncogenicity of Tumour Cells in a Preclinical Model of Acute Lymphoblastic Leukaemia. Cells, 2020, 9, 215.	4.1	16
32	Genome-wide transcriptomics leads to the identification of deregulated genes after deferasirox therapy in low-risk MDS patients. Pharmacogenomics Journal, 2020, 20, 664-671.	2.0	3
33	Treatment of Frail Older Adults and Elderly Patients With Philadelphia Chromosome-negative Acute Lymphoblastic Leukemia: Results of a Prospective Trial With Minimal Chemotherapy. Clinical Lymphoma, Myeloma and Leukemia, 2020, 20, e513-e522.	0.4	5
34	Phase 3 Study of Lenalidomide (LEN) Vs Placebo in Non-Transfusion Dependent (TD) Low Risk Del(5q) MDS Patients - Interim Analysis of the European Sintra-REV Trial. Blood, 2020, 136, 28-29.	1.4	8
35	Biological Impact of Monoallelic and Biallelic BIRC3 Loss in Del(11q) Chronic Lymphocytic Leukemia Progression. Blood, 2020, 136, 4-4.	1.4	0
36	Chronic graft-versus-host disease could ameliorate the impact of adverse somatic mutations in patients with myelodysplastic syndromes and hematopoietic stem cell transplantation. Annals of Hematology, 2019, 98, 2151-2162.	1.8	2

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37	Molecular profiling refines minimal residual diseaseâ€based prognostic assessment in adults with Philadelphia chromosomeâ€negative Bâ€cell precursor acute lymphoblastic leukemia. Genes Chromosomes and Cancer, 2019, 58, 815-819.	2.8	6
38	Splice donor site sgRNAs enhance CRISPR/Cas9-mediated knockout efficiency. PLoS ONE, 2019, 14, e0216674.	2.5	19
39	Incidence and outcome after first molecular versus overt recurrence in patients with Philadelphia chromosomeâ€"positive acute lymphoblastic leukemia included in the ALL Ph08 trial from the Spanish PETHEMA Group. Cancer, 2019, 125, 2810-2817.	4.1	13
40	Identification of novel variants in ten patients with Hermansky-Pudlak syndrome by high-throughput sequencing. Annals of Medicine, 2019, 51, 141-148.	3.8	11
41	Molecular Diagnosis of Inherited Coagulation and Bleeding Disorders. Seminars in Thrombosis and Hemostasis, 2019, 45, 695-707.	2.7	32
42	The poor prognosis of low hypodiploidy in adults with Bâ€cell precursor acute lymphoblastic leukaemia is restricted to older adults and elderly patients. British Journal of Haematology, 2019, 186, 263-268.	2.5	6
43	1q23.1 homozygous deletion and downregulation of Fc receptor-like family genes confer poor prognosis in chronic lymphocytic leukemia. Clinical and Experimental Medicine, 2019, 19, 261-267.	3.6	4
44	DNA damage response-related alterations define the genetic background of patients with chronic lymphocytic leukemia and chromosomal gains. Experimental Hematology, 2019, 72, 9-13.	0.4	9
45	Clinical, cytogenetic, and molecular findings in a patient with ring chromosome 4: case report and literature review. BMC Medical Genomics, 2019, 12, 167.	1.5	2
46	CLL cells cumulate genetic aberrations prior to the first therapy even in outwardly inactive disease phase. Leukemia, 2019, 33, 518-558.	7.2	15
47	Increased survival due to lower toxicity for highâ€risk Tâ€cell acute lymphoblastic leukemia patients in two consecutive pediatricâ€inspired PETHEMA trials. European Journal of Haematology, 2019, 102, 79-86.	2.2	14
48	Sitosterolemia: Diagnosis, Metabolic and Hematological Abnormalities, Cardiovascular Disease and Management. Current Medicinal Chemistry, 2019, 26, 6766-6775.	2.4	21
49	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
50	Azacitidine improves outcome in higherâ€risk <scp>MDS</scp> patients with chromosome 7 abnormalities: a retrospective comparison of <scp>GESMD</scp> and <scp>GFM</scp> registries. British Journal of Haematology, 2018, 181, 350-359.	2.5	11
51	Frequency and prognostic significance of additional cytogenetic abnormalities to the Philadelphia chromosome in young and older adults with acute lymphoblastic leukemia. Leukemia and Lymphoma, 2018, 59, 146-154.	1.3	17
52	Characterizing patients with multiple chromosomal aberrations detected by FISH in chronic lymphocytic leukemia. Leukemia and Lymphoma, 2018, 59, 633-642.	1.3	8
53	Prospective randomized trial of 5 days azacitidine versus supportive care in patients with lower-risk myelodysplastic syndromes without 5q deletion and transfusion-dependent anemia. Leukemia and Lymphoma, 2018, 59, 1095-1104.	1.3	15
54	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

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55	Introducing high-throughput sequencing into mainstream genetic diagnosis practice in inherited platelet disorders. Haematologica, 2018, 103, 148-162.	3.5	96
56	Identification of two novel mutations in RASGRP2 affecting platelet CalDAG-GEFI expression and function in patients with bleeding diathesis. Platelets, 2018, 29, 192-195.	2.3	26
57	A Modern Approach to the Molecular Diagnosis of Inherited Bleeding Disorders. Journal of Molecular and Genetic Medicine: an International Journal of Biomedical Research, 2018, 12, .	0.1	2
58	The International Prognostic Index for Patients with Chronic Lymphocytic Leukemia Has the Higher Value in Predicting Overall Outcome Compared with the Barcelona-Brno Biomarkers Only Prognostic Model and the MD Anderson Cancer Center Prognostic Index. BioMed Research International, 2018, 2018, 1-8.	1.9	18
59	New Challenges in Targeting Signaling Pathways in Acute Lymphoblastic Leukemia by NGS Approaches: An Update. Cancers, 2018, 10, 110.	3.7	35
60	Targeted genome editing in acute lymphoblastic leukemia: a review. BMC Biotechnology, 2018, 18, 45.	3.3	13
61	Genomic Instability and a Preferential Involvement of Ras Pathway in the Myelodysplastic Syndromes Evolution to Secondary Acute Myeloid Leukemia. Blood, 2018, 132, 3082-3082.	1.4	0
62	Nuevos mÃ©todos diagnÃ³sticos en los trastornos plaquetarios hereditarios. Medicina ClÃ¬nica, 2017, 148, 71-77.	0.6	6
63	A two-step approach for sequencing spliceosome-related genes as a complementary diagnostic assay in MDS patients with ringed sideroblasts. Leukemia Research, 2017, 56, 82-87.	0.8	4
64	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
65	Clinical characteristics of patients with central nervous system relapse in BCR-ABL1-positive acute lymphoblastic leukemia: the importance of characterizing ABL1 mutations in cerebrospinal fluid. Annals of Hematology, 2017, 96, 1069-1075.	1.8	21
66	Mutations in TP53 and JAK2 are independent prognostic biomarkers in B-cell precursor acute lymphoblastic leukaemia. British Journal of Cancer, 2017, 117, 256-265.	6.4	34
67	Multidimensional assessment of patient condition and mutational analysis in peripheral blood, as tools to improve outcome prediction in myelodysplastic syndromes: A prospective study of the Spanish MDS group. American Journal of Hematology, 2017, 92, E534-E541.	4.1	6
68	Clinical and biological significance of isolated Y chromosome loss in myelodysplastic syndromes and chronic myelomonocytic leukemia. A report from the Spanish MDS Group. Leukemia Research, 2017, 63, 85-89.	0.8	9
69	Hyperdiploidy as a rare event that accompanies poor prognosis markers in <scp>CLL</scp>. European Journal of Haematology, 2017, 98, 142-148.	2.2	4
70	Wiskottâ€“Aldrich syndrome in a child presenting with macrothrombocytopenia. Platelets, 2017, 28, 417-420.	2.3	19
71	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
72	Next-generation sequencing in chronic lymphocytic leukemia: recent findings and new horizons. Oncotarget, 2017, 8, 71234-71248.	1.8	25

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73	The CRISPR/Cas9 system efficiently reverts the tumorigenic ability of BCR/ABL in vitro and in a xenograft model of chronic myeloid leukemia. Oncotarget, 2017, 8, 26027-26040.	1.8	30
74	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
75	Bone marrow fibrosis in myelodysplastic syndromes: a prospective evaluation including mutational analysis. Oncotarget, 2016, 7, 30492-30503.	1.8	41
76	Genome-Wide DNA Copy Number Analysis of Acute Lymphoblastic Leukemia Identifies New Genetic Markers Associated with Clinical Outcome. PLoS ONE, 2016, 11, e0148972.	2.5	40
77	Chromothripsis Is a Recurrent Genomic Abnormality in High-Risk Myelodysplastic Syndromes. PLoS ONE, 2016, 11, e0164370.	2.5	28
78	FISHing in the dark: How the combination of FISH and conventional karyotyping improves the diagnostic yield in CpGâ€stimulated chronic lymphocytic leukemia. American Journal of Hematology, 2016, 91, 978-983.	4.1	14
79	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
80	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
81	Association between different risk factors and vascular accelerated ageing (EVA study): study protocol for a cross-sectional, descriptive observational study. BMJ Open, 2016, 6, e011031.	1.9	37
82	Identification of expression patterns in the progression of disease stages by integration of transcriptomic data. BMC Bioinformatics, 2016, 17, 432.	2.6	15
83	Copy number alterations in adult patients with mature B acute lymphoblastic leukemia treated with specific immunochemotherapy. Medicina ClÃnica (English Edition), 2016, 147, 488-491.	0.2	0
84	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
85	MiRNA expression profile of chronic lymphocytic leukemia patients with 13q deletion. Leukemia Research, 2016, 46, 30-36.	0.8	8
86	Novel mutations in RASGRP2, which encodes CalDAG-GEFI, abrogate Rap1 activation, causing platelet dysfunction. Blood, 2016, 128, 1282-1289.	1.4	68
87	Uniparental disomy causes deficiencies of vitamin Kâ€dependent proteins. Journal of Thrombosis and Haemostasis, 2016, 14, 2410-2418.	3.8	6
88	Population pharmacokinetics of doxorubicin and doxorubicinol in patients diagnosed with nonâ€Hodgkin's lymphoma. British Journal of Clinical Pharmacology, 2016, 82, 1517-1527.	2.4	24
89	Prognostic impact of chromosomal translocations in myelodysplastic syndromes and chronic myelomonocytic leukemia patients. A study by the spanish group of myelodysplastic syndromes. Genes Chromosomes and Cancer, 2016, 55, 322-327.	2.8	7
90	The presence of genomic imbalances is associated with poor outcome in patients with burkitt lymphoma treated with doseâ€intensive chemotherapy including rituximab. British Journal of Haematology, 2016, 172, 428-438.	2.5	20

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91	Pharmacogenetics and pharmacogenomics as tools in cancer therapy. Drug Metabolism and Personalized Therapy, 2016, 31, 25-34.	0.6	23
92	Current opinion and consensus statement regarding the diagnosis, prognosis, and treatment of patients with essential thrombocythemia: a survey of the Spanish Group of Ph-negative Myeloproliferative Neoplasms (GEMFIN) using the Delphi method. Annals of Hematology, 2016, 95, 719-732.	1.8	5
93	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
94	A rare but recurrent t(8;13)(q24;q14) translocation in Bâ€cell chronic lymphocytic leukaemia causing <scp>MYC</scp> upâ€regulation and concomitant loss of <scp>PVT</scp>1, miRâ€15/16 and <scp>DLEU</scp>7. British Journal of Haematology, 2016, 172, 296-299.	2.5	7
95	DNA copy number profiling reveals different patterns of chromosomal instability within colorectal cancer according to the age of onset. Molecular Carcinogenesis, 2016, 55, 705-716.	2.7	30
96	Incidence, clinical and biological characteristics and outcome of secondary acute lymphoblastic leukemia after solid organ or hematologic malignancy. Leukemia and Lymphoma, 2016, 57, 86-91.	1.3	14
97	Integrative analysis of DNA copy number, DNA methylation and gene expression in multiple myeloma reveals alterations related to relapse. Oncotarget, 2016, 7, 80664-80679.	1.8	11
98	Analyse multiple disease subtypes and build associated gene networks using genome-wide expression profiles. BMC Genomics, 2015, 16, S3.	2.8	17
99	Prognostic significance of copy number alterations in adolescent and adult patients with precursor <scp>B</scp> acute lymphoblastic leukemia enrolled in <scp>PETHEMA</scp> protocols. Cancer, 2015, 121, 3809-3817.	4.1	43
100	Deregulation of Genes Related to Iron and Mitochondrial Metabolism in Refractory Anemia with Ring Sideroblasts. PLoS ONE, 2015, 10, e0126555.	2.5	21
101	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
102	Correlation of myelodysplastic syndromes with i(17)(q10) and <scp>TP</scp>53 and <scp>SETBP</scp>1 mutations. British Journal of Haematology, 2015, 171, 137-141.	2.5	11
103	Non-coding recurrent mutations in chronic lymphocytic leukaemia. Nature, 2015, 526, 519-524.	27.8	749
104	MicroRNA-223 is a novel negative regulator of HSP90B1 in CLL. BMC Cancer, 2015, 15, 238.	2.6	16
105	Fluorescencein situhybridization analysis does not increase detection rate for trisomy 8 in chronic myelomonocytic leukemia. Leukemia and Lymphoma, 2015, 56, 242-243.	1.3	1
106	Impact of transplant eligibility and availability of a human leukocyte antigen-identical matched related donor on outcome of older patients with acute lymphoblastic leukemia. Leukemia and Lymphoma, 2015, 56, 2812-2818.	1.3	5
107	A robust estimation of exon expression to identify alternative spliced genes applied to human tissues and cancer samples. BMC Genomics, 2014, 15, 879.	2.8	12
108	Transcriptome analysis reveals molecular profiles associated with evolving steps of monoclonal gammopathies. Haematologica, 2014, 99, 1365-1372.	3.5	65

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109	TET2Overexpression in Chronic Lymphocytic Leukemia Is Unrelated to the Presence ofTET2Variations. BioMed Research International, 2014, 2014, 1-6.	1.9	12
110	Mutation Status and Immunoglobulin Gene Rearrangements in Patients from Northwest and Central Region of Spain with Chronic Lymphocytic Leukemia. BioMed Research International, 2014, 2014, 1-8.	1.9	14
111	Validation and clinical evaluation of a UHPLC method with fluorescence detector for plasma quantification of doxorubicin and doxorubicinol in haematological patients. Journal of Chromatography B: Analytical Technologies in the Biomedical and Life Sciences, 2014, 955-956, 93-97.	2.3	17
112	Exome sequencing reveals novel and recurrent mutations with clinical impact in blastic plasmacytoid dendritic cell neoplasm. Leukemia, 2014, 28, 823-829.	7.2	148
113	Interstitial 13q14 deletions detected in the karyotype and translocations with concomitant deletion at 13q14 in chronic lymphocytic leukemia: Different genetic mechanisms but equivalent poorer clinical outcome. Genes Chromosomes and Cancer, 2014, 53, 788-797.	2.8	15
114	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
115	Prognostic significance of complex karyotype and monosomal karyotype in adult patients with acute lymphoblastic leukemia treated with riskâ€adapted protocols. Cancer, 2014, 120, 3958-3964.	4.1	24
116	Treatment of High-Risk Philadelphia Chromosomeâ€“Negative Acute Lymphoblastic Leukemia in Adolescents and Adults According to Early Cytologic Response and Minimal Residual Disease After Consolidation Assessed by Flow Cytometry: Final Results of the PETHEMA ALL-AR-03 Trial. Journal of Clinical Oncology, 2014, 32, 1595-1604.	1.6	227
117	Genomic analysis of clonal eosinophils by <scp>CGH</scp> arrays reveals new genetic regions involved in chronic eosinophilia. European Journal of Haematology, 2014, 93, 422-428.	2.2	1
118	Prognostic impact and landscape of NOTCH1 mutations in chronic lymphocytic leukemia (CLL): a study on 852 patients. Leukemia, 2013, 27, 2393-2396.	7.2	65
119	Prognostic impact of the number of methylated genes in myelodysplastic syndromes and acute myeloid leukemias treated with azacytidine. Annals of Hematology, 2013, 92, 1543-1552.	1.8	16
120	Reciprocal translocations in myelodysplastic syndromes and chronic myelomonocytic leukemias: Review of 5,654 patients with an evaluable karyotype. Genes Chromosomes and Cancer, 2013, 52, 753-763.	2.8	15
121	Genome-wide profiling of methylation identifies novel targets with aberrant hypermethylation and reduced expression in low-risk myelodysplastic syndromes. Leukemia, 2013, 27, 610-618.	7.2	31
122	Application of FISH 7q in MDS patients without monosomy 7 or 7q deletion by conventional G-banding cytogenetics: Does â˜7/7qâ˜ detection by FISH have prognostic value?. Leukemia Research, 2013, 37, 416-421.	0.8	16
123	Clinical Applications of BAC Array-CGH to the Study of Diffuse Large B-Cell Lymphomas. Methods in Molecular Biology, 2013, 973, 121-145.	0.9	1
124	Alteration in Endoglin-Related Angiogenesis in Refractory Cytopenia with Multilineage Dysplasia. PLoS ONE, 2013, 8, e53624.	2.5	3
125	Chronic lymphocytic leukemia: a clinical and molecular heterogenous disease. Cancer Genetics, 2013, 206, 49-62.	0.4	63
126	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

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127	Response to lenalidomide in myelodysplastic syndromes with del(5q): influence of cytogenetics and mutations. British Journal of Haematology, 2013, 162, 74-86.	2.5	73
128	Landscape of somatic mutations and clonal evolution in mantle cell lymphoma. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 18250-18255.	7.1	488
129	Single nucleotide polymorphism array karyotyping: A diagnostic and prognostic tool in myelodysplastic syndromes with unsuccessful conventional cytogenetic testing. Genes Chromosomes and Cancer, 2013, 52, 1167-1177.	2.8	44
130	Biallelic losses of 13q do not confer a poorer outcome in chronic lymphocytic leukaemia: analysis of 627 patients with isolated 13q deletion. British Journal of Haematology, 2013, 163, 47-54.	2.5	13
131	Doseâ€intensive chemotherapy including rituximab in Burkitt's leukemia or lymphoma regardless of human immunodeficiency virus infection status. Cancer, 2013, 119, 1660-1668.	4.1	63
132	Comparative Genomics with Multi-agent Systems. Advances in Intelligent Systems and Computing, 2013, , 175-181.	0.6	0
133	SNP-based mapping arrays reveal high genomic complexity in monoclonal gammopathies, from MGUS to myeloma status. Leukemia, 2012, 26, 2521-2529.	7.2	100
134	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
135	Automatic knowledge extraction in sequencing analysis with multiagent system and grid computing. Journal of Integrative Bioinformatics, 2012, 9, 93-104.	1.5	2
136	Molecular Characterization of Chronic Lymphocytic Leukemia Patients with a High Number of Losses in 13q14. PLoS ONE, 2012, 7, e48485.	2.5	37
137	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
138	Prognostic value of trisomy 8 as a single anomaly and the influence of additional cytogenetic aberrations in primary myelodysplastic syndromes. British Journal of Haematology, 2012, 159, 311-321.	2.5	25
139	Highâ€throughput sequencing analysis of the chromosome 7q32 deletion reveals <scp>IRF</scp>5 as a potential tumour suppressor in splenic marginalâ€zone lymphoma. British Journal of Haematology, 2012, 158, 712-726.	2.5	45
140	Treatment of young patients with <scp>P</scp>hiladelphia chromosomeâ€positive acute lymphoblastic leukaemia using increased dose of imatinib and deintensified chemotherapy before allogeneic stem cell transplantation. British Journal of Haematology, 2012, 159, 78-81.	2.5	43
141	Expression of VAV1 in the tumour microenvironment of glioblastoma multiforme. Journal of Neuro-Oncology, 2012, 110, 69-77.	2.9	12
142	Response to imatinib mesylate in patients with hypereosinophilic syndrome. International Journal of Hematology, 2012, 96, 320-326.	1.6	16
143	Imatinib therapy of chronic myeloid leukemia restores the expression levels of key genes for DNA damage and cell-cycle progression. Pharmacogenetics and Genomics, 2012, 22, 381-388.	1.5	12
144	Exome sequencing identifies recurrent mutations of the splicing factor SF3B1 gene in chronic lymphocytic leukemia. Nature Genetics, 2012, 44, 47-52.	21.4	893

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145	Incidence and clinical characteristics of myeloproliferative neoplasms displaying a <scp>PDGFRB</scp> rearrangement. European Journal of Haematology, 2012, 89, 37-41.	2.2	42
146	Integration of Global Spectral Karyotyping, CGH Arrays, and Expression Arrays Reveals Important Genes in the Pathogenesis of Glioblastoma Multiforme. Annals of Surgical Oncology, 2012, 19, 2367-2379.	1.5	9
147	SAMasGC: Sequencing Analysis with a Multiagent System and Grid Computing. Advances in Intelligent and Soft Computing, 2012, , 209-216.	0.2	0
148	Visual Analysis Tool in Comparative Genomics. Advances in Intelligent and Soft Computing, 2012, , 121-127.	0.2	0
149	Automatic knowledge extraction in sequencing analysis with multiagent system and grid computing. Journal of Integrative Bioinformatics, 2012, 9, 206.	1.5	2
150	Cytogenetic risk stratification in chronic myelomonocytic leukemia. Haematologica, 2011, 96, 375-383.	3.5	226
151	Whole-genome sequencing identifies recurrent mutations in chronic lymphocytic leukaemia. Nature, 2011, 475, 101-105.	27.8	1,364
152	Evaluation of Spanish Gaucher disease patients after a 6-month imiglucerase shortage. Blood Cells, Molecules, and Diseases, 2011, 46, 115-118.	1.4	32
153	Molecular Characterization of the Region 7q22.1 in Splenic Marginal Zone Lymphomas. PLoS ONE, 2011, 6, e24939.	2.5	23
154	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
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