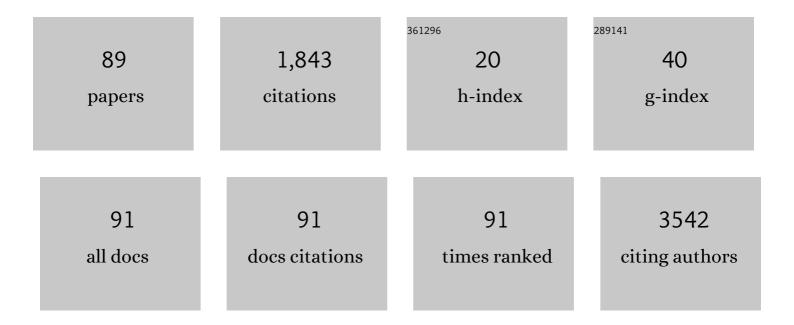
List of Publications by Year in descending order

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ROSA AVALA

#	Article	IF	CITATIONS
1	The Minimal Residual Disease Using Liquid Biopsies in Hematological Malignancies. Cancers, 2022, 14, 1310.	1.7	16
2	Realâ€world analysis of main clinical outcomes in patients with polycythemia vera treated with ruxolitinib or best available therapy after developing resistance/intolerance to hydroxyurea. Cancer, 2022, 128, 2441-2448.	2.0	14
3	Monitoring of clonal evolution of acute myeloid leukemia identifies the leukemia subtype, clinical outcome and potential new drug targets for post-remission strategies or relapse. Haematologica, 2021, 106, 2325-2333.	1.7	18
4	Measurable residual disease in elderly acute myeloid leukemia: results from the PETHEMA-FLUGAZA phase 3 clinical trial. Blood Advances, 2021, 5, 760-770.	2.5	18
5	A typical acute lymphoblastic leukemia JAK2 variant, R683C, causes an aggressive form of familial thrombocytosis when germline. Leukemia, 2021, 35, 3295-3298.	3.3	2
6	The Mutational Landscape of Acute Myeloid Leukaemia Predicts Responses and Outcomes in Elderly Patients from the PETHEMA-FLUGAZA Phase 3 Clinical Trial. Cancers, 2021, 13, 2458.	1.7	7
7	Prognostic heterogeneity of adult Bâ€cell precursor acute lymphoblastic leukaemia patients with t(1;19)(q23;p13)/ TCF3â€PBX1 treated with measurable residual diseaseâ€oriented protocols. British Journal of Haematology, 2021, , .	1.2	2
8	Clonal hematopoiesis-defining mutations have no impact on the development of thrombosis in a cohort of patients with myeloid pathology. Leukemia Research, 2021, 108, 106613.	0.4	0
9	Increased von Willebrand factor antigen and low ADAMTS13 activity are related to poor prognosis in covidâ€19 patients. International Journal of Laboratory Hematology, 2021, 43, O152-O155.	0.7	23
10	Networking for advanced molecular diagnosis in acute myeloid leukemia patients is possible: the PETHEMA NGS-AML project. Haematologica, 2021, 106, 3079-3089.	1.7	15
11	Prognostic significance of FLT3-ITD length in AML patients treated with intensive regimens. Scientific Reports, 2021, 11, 20745.	1.6	11
12	Does RAD21 Co-Mutation Have a Role in DNMT3A Mutated AML? Results of Harmony Alliance AML Database. Blood, 2021, 138, 608-608.	0.6	0
13	Nationwide Laboratory Network for AML Cross-Validated NGS Studies: Results from a Real-Life Cohort of the Pethema Group. Blood, 2021, 138, 1302-1302.	0.6	0
14	Triple Combination of Ruxolutinib, Nilotinib and Prednisone Is Safe and Shows Promising Activity for the Treatment of Myelofibrosis Patients, Results of a Phase Ib Clinical Trial (RUNIC). Blood, 2021, 138, 3655-3655.	0.6	0
15	Potential Utility of Circulating Tumor DNA Monitoring in Primary Mediastinal B-Cell Lymphoma Treated with R-DA-EPOCH. Blood, 2021, 138, 4491-4491.	0.6	0
16	Integrated Multidimensional Flow Cytometry (MFC) and Next-Generation Sequencing (NGS) to Reconstruct Evolutionary Paterns from Dysplasia to Acute Myeloid Leukemia (AML). Blood, 2021, 138, 520-520.	0.6	0
17	Impact of Gender on Molecular AML Subclasses - a Harmony Alliance Study. Blood, 2021, 138, 3438-3438.	0.6	0
18	The Spliceosome As a New Therapeutic Target in Cytarabine-Resistant Acute Myeloid Leukemia. Blood, 2021, 138, 3334-3334.	0.6	0

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19	Harmony Alliance Provides a Machine Learning Researching Tool to Predict the Risk of Relapse after First Remission in AML Patients Treated without Allogeneic Haematopoietic Stem Cell Transplantation. Blood, 2021, 138, 4041-4041.	0.6	2
20	MPL S505C enhances driver mutations at W515 in essential thrombocythemia. Blood Cancer Journal, 2021, 11, 188.	2.8	2
21	Impact of COVID-19 in patients with multiple myeloma based on a global data network. Blood Cancer Journal, 2021, 11, 198.	2.8	25
22	Novel DNMT3A Germline Variant in a Patient with Multiple Paragangliomas and Papillary Thyroid Carcinoma. Cancers, 2020, 12, 3304.	1.7	5
23	Clinical course and risk factors for mortality from COVIDâ€19 in patients with haematological malignancies. European Journal of Haematology, 2020, 105, 597-607.	1.1	73
24	Jumping Translocation in a Patient with Acute Leukemia and Fatal Evolution. Case Reports in Oncology, 2020, 13, 1026-1030.	0.3	0
25	High-sensitivity microsatellite instability assessment for the detection of mismatch repair defects in normal tissue of biallelic germline mismatch repair mutation carriers. Journal of Medical Genetics, 2020, 57, 269-273.	1.5	20
26	Improving the prediction of acute myeloid leukaemia outcomes by complementing mutational profiling with <i>ex vivo</i> chemosensitivity. British Journal of Haematology, 2020, 189, 672-683.	1.2	11
27	miR-146a rs2431697 identifies myeloproliferative neoplasm patients with higher secondary myelofibrosis progression risk. Leukemia, 2020, 34, 2648-2659.	3.3	18
28	Analysis of SNP Array Abnormalities in Patients with DE NOVO Acute Myeloid Leukemia with Normal Karyotype. Scientific Reports, 2020, 10, 5904.	1.6	8
29	A novel targeted RNA-Seq panel identifies a subset of adult patients with acute lymphoblastic leukemia with BCR-ABL1-like characteristics. Blood Cancer Journal, 2020, 10, 43.	2.8	10
30	CaracterÃsticas clÃnico-biológicas de los pacientes con mielofibrosis: un análisis de 1.000 casos del Registro Español de Mielofibrosis. Medicina ClÃnica, 2020, 155, 152-158.	0.3	3
31	Measurable Residual Disease (MRD) in Elderly Acute Myeloid Leukemia (AML): Results from the Pethema-Flugaza Phase III Clinical Trial. Blood, 2020, 136, 32-32.	0.6	0
32	Study of the Role of Splicing Factor SRRM2 in Cytarabine Treatment Resistance in Acute Myeloid Leukemia. Blood, 2020, 136, 8-9.	0.6	0
33	Validation of the High-Risk Prognostic Score Defined By the Presence of Mutations in NRAS or TP53 in a Cohort of 497 Patients with Acute Myeloid Leukemia. Blood, 2020, 136, 4-5.	0.6	0
34	Detection of Emerging Resistant Clones in Philadelphia-Positive Leukemia Patients Exposed to Tyrosine Kinase Inhibitors. Correlation of cDNA and Gdna Approaches. Blood, 2020, 136, 6-8.	0.6	1
35	Differences in the Mutational Landscape of Myeloid Malignancies (acute myeloid leukemia,) Tj ETQq1 1 0.78431 136, 41-42.	4 rgBT /O 0.6	verlock 10 Tf 0
36	Minimal Residual Disease Monitoring from Liquid Biopsy By Next Generation Sequencing in Follicular Lymphoma Patients. Blood, 2020, 136, 31-33.	0.6	2

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37	A novel deep targeted sequencing method for minimal residual disease monitoring in acute myeloid leukemia. Haematologica, 2019, 104, 288-296.	1.7	36
38	Protein Carbonylation in Patients with Myelodysplastic Syndrome: An Opportunity for Deferasirox Therapy. Antioxidants, 2019, 8, 508.	2.2	4
39	Minimal Residual Disease Monitoring with Next-Generation Sequencing Methodologies in Hematological Malignancies. International Journal of Molecular Sciences, 2019, 20, 2832.	1.8	42
40	MEK inhibition enhances the response to tyrosine kinase inhibitors in acute myeloid leukemia. Scientific Reports, 2019, 9, 18630.	1.6	24
41	Clinical Utility of a Next-Generation Sequencing Panel for Acute Myeloid Leukemia Diagnostics. Journal of Molecular Diagnostics, 2019, 21, 228-240.	1.2	24
42	Different Clinical Implications of Kinase Domain BCR-ABL1 Variants Detected in Chronic Myeloid Leukemia and Acute Lymphoblastic Leukemia Patients. Blood, 2019, 134, 5368-5368.	0.6	0
43	Multicenter, Open-Label, Single Arm, Phase II Exploratory Study to Evaluate the Effect of a One-Year Consolidation Treatment with Ponatinib 15 Mg on Treatment Free-Remission Rate in Patients with Philadelphia-Positive Chronic Myeloid Leukemia, Who Had Previously Achieved a Deep Molecular Response with Imatinib (PonaZero) study), Blood, 2019, 134, 5904-5904.	0.6	0
44	Wilms Tumor 1 gene expression levels improve risk stratification in <scp>AML</scp> patients. Results of a multicentre study within the Spanish Group for Molecular Biology in Haematology. British Journal of Haematology, 2018, 181, 542-546.	1.2	4
45	Drug-to-drug interactions of tyrosine kinase inhibitors in chronic myeloid leukemia patients. Is it a real problem?. Annals of Hematology, 2018, 97, 2089-2098.	0.8	18
46	Mutational screening of newly diagnosed multiple myeloma patients by deep targeted sequencing. Haematologica, 2018, 103, e544-e548.	1.7	13
47	rs2431697, a Polymorphism of Mir-146a, Is a Precozing Marker of Progression to Secondary Myelofibrosis: New Epigenetic Regulation of Jak/Stat3 Signaling. Blood, 2018, 132, 3072-3072.	0.6	0
48	Study of the Clinical Significance of the Length of FLT3-ITD in Acute Myeloid Leukemia Patients. Blood, 2018, 132, 5278-5278.	0.6	0
49	Protein Carbonylation Pattern Is Altered in Myelodisplastic Syndromes. Blood, 2018, 132, 5502-5502.	0.6	0
50	Analytical and clinical validation of a novel in-house deep-sequencing method for minimal residual disease monitoring in a phase II trial for multiple myeloma. Leukemia, 2017, 31, 1446-1449.	3.3	44
51	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.	0.8	21
52	PTCH1 is a reliable marker for predicting imatinib response in chronic myeloid leukemia patients in chronic phase. PLoS ONE, 2017, 12, e0181366.	1.1	8
53	Mutations in the DNA methylation pathway and number of driver mutations predict response to azacitidine in myelodysplastic syndromes. Oncotarget, 2017, 8, 106948-106961.	0.8	38
54	Ocena minimalnej choroby resztkowej w szpiczaku plazmocytowym. Hematologia, 2017, 8, 219-227.	0.0	0

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55	Circulating Immune Complexes of IgA Bound to Beta 2 Glycoprotein are Strongly Associated with the Occurrence of Acute Thrombotic Events. Journal of Atherosclerosis and Thrombosis, 2016, 23, 1242-1253.	0.9	32
56	Antiplatelet therapy versus observation in low-risk essential thrombocythemia with a CALR mutation. Haematologica, 2016, 101, 926-931.	1.7	118
57	Leucemia mieloide crónica en España: sus caracterÃsticas de presentación han cambiado. Sección española del registro poblacional EUTOS. Revista Clinica Espanola, 2016, 216, 293-300.	0.2	9
58	<scp>CALR</scp> mutations screening should not be studied in splanchnic vein thrombosis. British Journal of Haematology, 2015, 170, 588-589.	1.2	15
59	Oral anticoagulation to prevent thrombosis recurrence in polycythemia vera and essential thrombocythemia. Annals of Hematology, 2015, 94, 911-918.	0.8	49
60	Myeloproliferative neoplasm in a thalassaemic patient: response to treatment with a JAK inhibitor. Annals of Hematology, 2015, 94, 1237-1239.	0.8	5
61	JAK2 exon 12 mutations were not found in liver transplant recipients with or without pretransplant portal vein thrombosis. European Journal of Gastroenterology and Hepatology, 2014, 26, 362-363.	0.8	3
62	Exome sequencing reveals novel and recurrent mutations with clinical impact in blastic plasmacytoid dendritic cell neoplasm. Leukemia, 2014, 28, 823-829.	3.3	148
63	Prognostic value of deep sequencing method for minimal residual disease detection in multiple myeloma. Blood, 2014, 123, 3073-3079.	0.6	380
64	Proteomic analysis reveals heat shock protein 70 has a key role in polycythemia Vera. Molecular Cancer, 2013, 12, 142.	7.9	20
65	Clinical applicability and prognostic significance of molecular response assessed by fluorescentâ€ <scp>PCR</scp> of immunoglobulin genes in multiple myeloma. Results from a <scp>GEM</scp> / <scp>PETHEMA</scp> study. British Journal of Haematology, 2013, 163, 581-589.	1.2	27
66	Inhibition of related JAK/STAT pathways with molecular targeted drugs shows strong synergy with ruxolitinib in chronic myeloproliferative neoplasm. British Journal of Haematology, 2013, 161, 667-676.	1.2	20
67	Correlation of WT1 expression with the burden of total and residual leukemic blasts in bone marrow samples of acute myeloid leukemia patients. Cancer Genetics, 2012, 205, 190-191.	0.2	14
68	Obesity is an independent risk factor for pre-transplant portal vein thrombosis in liver recipients. BMC Gastroenterology, 2012, 12, 114.	0.8	50
69	Acute myeloid leukemia and transcription factors: role of erythroid Krüppel-like factor (EKLF). Cancer Cell International, 2012, 12, 25.	1.8	5
70	Use of <scp>S</scp> orafenib as an effective treatment in an <scp>AML</scp> patient carrying a new point mutation affecting the <scp>J</scp> uxtamembrane domain of <i><scp>FLT</scp>3</i> . British Journal of Haematology, 2012, 158, 555-558.	1.2	5
71	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.	1.7	59
72	Recipient and donor thrombophilia and the risk of portal venous thrombosis and hepatic artery thrombosis in liver recipients. BMC Gastroenterology, 2011, 11, 130.	0.8	31

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73	Epigenomic profiling in polycythaemia vera and essential thrombocythaemia shows low levels of aberrant DNA methylation. Journal of Clinical Pathology, 2011, 64, 1010-1013.	1.0	20
74	Differential expression of JAK2 and Src kinase genes in response to hydroxyurea treatment in polycythemia vera and essential thrombocythemia. Annals of Hematology, 2011, 90, 939-946.	0.8	7
75	Simplifying the detection of MUTYH mutations by high resolution melting analysis. BMC Cancer, 2010, 10, 408.	1.1	5
76	Realâ€ŧime PCR quantification of haematopoietic chimerism after transplantation: a comparison between TaqMan and hybridization probes technologies. International Journal of Laboratory Hematology, 2010, 32, e17-25.	0.7	6
77	Post-Transplant Myelodysplastic Syndromes In Pediatric Liver Transplantation Recipients: a Report of Two Cases. Blood, 2010, 116, 4978-4978.	0.6	2
78	Long-term follow-up of donor chimerism and tolerance after human liver transplantation. Liver Transplantation, 2009, 15, 581-591.	1.3	19
79	Living Donor Liver Transplantation: Usefulness of Hemostatic and Prothrombotic Screening in Potential Donors. Transplantation Proceedings, 2009, 41, 3791-3795.	0.3	16
80	High Resolution Melting Analysis for JAK2 Exon 14 and Exon 12 Mutations. Journal of Molecular Diagnostics, 2009, 11, 155-161.	1.2	48
81	Long-Term Follow-up of Donor Chimerism and Tolerance After Human Liver Transplantation Blood, 2009, 114, 3534-3534.	0.6	0
82	Validity test study of JAK2 V617F and allele burden quantification in the diagnosis of myeloproliferative diseases. Annals of Hematology, 2008, 87, 741-749.	0.8	43
83	Evaluation of minimal residual disease in multiple myeloma patients by fluorescentâ€polymerase chain reaction: the prognostic impact of achieving molecular response. British Journal of Haematology, 2008, 142, 766-774.	1.2	52
84	Importance of JAK2 V617F Allele Burden in the Diagnosis of Myeloproliferative Diseases and Its Association to Age Blood, 2007, 110, 4654-4654.	0.6	0
85	Application of Self-Quenched JH Consensus Primers for Real-Time Quantitative PCR of IGH Gene to Minimal Residual Disease Evaluation in Multiple Myeloma. Journal of Molecular Diagnostics, 2006, 8, 364-370.	1.2	1
86	Breast Cancer–Specific mRNA Transcripts Presence in Peripheral Blood After Adjuvant Chemotherapy Predicts Poor Survival Among High-Risk Breast Cancer Patients Treated With High-Dose Chemotherapy With Peripheral Blood Stem Cell Support. Journal of Clinical Oncology, 2006, 24, 3611-3618.	0.8	36
87	The use of fluorescent molecular beacons in real time PCR of IgH gene rearrangements for quantitative evaluation of multiple myeloma. International Journal of Laboratory Hematology, 2004, 26, 31-35.	0.2	6
88	Grupos de riesgo citogenético en la leucemia mieloide aguda: comparación de los modelos adoptados por los grupos MRC (Medical Research Council, del Reino Unido) y SWOG (Southwest Oncology) Tj ETQq0 0 0 rg	gB <b>D/</b> Øverlo	oc <b>b</b> 10 Tf 50

89	Clinical Course and Risk Factors for Mortality from COVID-19 in Patients with Hematological Malignancies. SSRN Electronic Journal, 0, , .	0.4	0
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