

# Frederik Damm

## List of Publications by Year in descending order

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Version: 2024-02-01

59  
papers

3,896  
citations

218677

26  
h-index

149698

56  
g-index

60  
all docs

60  
docs citations

60  
times ranked

5470  
citing authors

#	ARTICLE	IF	CITATIONS
1	Incidence and Prognostic Influence of <i>DNMT3A</i> Mutations in Acute Myeloid Leukemia. <i>Journal of Clinical Oncology</i> , 2011, 29, 2889-2896.	1.6	351
2	Prognostic impact, concurrent genetic mutations, and gene expression features of AML with CEBPA mutations in a cohort of 1182 cytogenetically normal AML patients: further evidence for CEBPA double mutant AML as a distinctive disease entity. <i>Blood</i> , 2011, 117, 2469-2475.	1.4	341
3	Prognostic Significance of <i>ASXL1</i> Mutations in Patients With Myelodysplastic Syndromes. <i>Journal of Clinical Oncology</i> , 2011, 29, 2499-2506.	1.6	258
4	Impact of <i>IDH1</i> R132 Mutations and an <i>IDH1</i> Single Nucleotide Polymorphism in Cytogenetically Normal Acute Myeloid Leukemia: SNP rs11554137 Is an Adverse Prognostic Factor. <i>Journal of Clinical Oncology</i> , 2010, 28, 2356-2364.	1.6	229
5	Mutations affecting mRNA splicing define distinct clinical phenotypes and correlate with patient outcome in myelodysplastic syndromes. <i>Blood</i> , 2012, 119, 3211-3218.	1.4	220
6	Acquired Initiating Mutations in Early Hematopoietic Cells of CLL Patients. <i>Cancer Discovery</i> , 2014, 4, 1088-1101.	9.4	213
7	<i>IDH1</i> mutations in patients with myelodysplastic syndromes are associated with an unfavorable prognosis. <i>Haematologica</i> , 2010, 95, 1668-1674.	3.5	177
8	Prognostic impact of <i>IDH2</i> mutations in cytogenetically normal acute myeloid leukemia. <i>Blood</i> , 2010, 116, 614-616.	1.4	170
9	<i>BCOR</i> and <i>BCORL1</i> mutations in myelodysplastic syndromes and related disorders. <i>Blood</i> , 2013, 122, 3169-3177.	1.4	169
10	Mutations in the cohesin complex in acute myeloid leukemia: clinical and prognostic implications. <i>Blood</i> , 2014, 123, 914-920.	1.4	167
11	Role of Donor Clonal Hematopoiesis in Allogeneic Hematopoietic Stem-Cell Transplantation. <i>Journal of Clinical Oncology</i> , 2019, 37, 375-385.	1.6	163
12	Hematopoietic lineage distribution and evolutionary dynamics of clonal hematopoiesis. <i>Leukemia</i> , 2018, 32, 1908-1919.	7.2	137
13	Single Nucleotide Polymorphism in the Mutational Hotspot of <i>WT1</i> Predicts a Favorable Outcome in Patients With Cytogenetically Normal Acute Myeloid Leukemia. <i>Journal of Clinical Oncology</i> , 2010, 28, 578-585.	1.6	119
14	Next-generation sequencing for minimal residual disease monitoring in acute myeloid leukemia patients with <i>FLT3</i> ITD or <i>NPM1</i> mutations. <i>Genes Chromosomes and Cancer</i> , 2012, 51, 689-695.	2.8	114
15	Integrative prognostic risk score in acute myeloid leukemia with normal karyotype. <i>Blood</i> , 2011, 117, 4561-4568.	1.4	99
16	Genomic landscape and clonal evolution of acute myeloid leukemia with t(8;21): an international study on 331 patients. <i>Blood</i> , 2019, 133, 1140-1151.	1.4	96
17	Frequent <i>NFKBIE</i> deletions are associated with poor outcome in primary mediastinal B-cell lymphoma. <i>Blood</i> , 2016, 128, 2666-2670.	1.4	82
18	Rare occurrence of <i>DNMT3A</i> mutations in myelodysplastic syndromes. <i>Haematologica</i> , 2011, 96, 1870-1873.	3.5	67

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19	Clonal hematopoiesis in patients with anti-neutrophil cytoplasmic antibody-associated vasculitis. <i>Haematologica</i> , 2020, 105, e264-e267.	3.5	56
20	Prognostic Importance of Histone Methyltransferase <i>MLL5</i> Expression in Acute Myeloid Leukemia. <i>Journal of Clinical Oncology</i> , 2011, 29, 682-689.	1.6	53
21	STAT3 mutations identified in human hematologic neoplasms induce myeloid malignancies in a mouse bone marrow transplantation model. <i>Haematologica</i> , 2013, 98, 1748-1752.	3.5	50
22	Prognostic significance of expression levels of stem cell regulators MSI2 and NUMB in acute myeloid leukemia. <i>Annals of Hematology</i> , 2013, 92, 315-323.	1.8	48
23	Single-cell analysis based dissection of clonality in myelofibrosis. <i>Nature Communications</i> , 2020, 11, 73.	12.8	46
24	FLT3-internal tandem duplication and age are the major prognostic factors in patients with relapsed acute myeloid leukemia with normal karyotype. <i>Haematologica</i> , 2011, 96, 681-686.	3.5	45
25	Prognostic significance of combined MN1, ERG, BAALC, and EVI1 (MEBE) expression in patients with myelodysplastic syndromes. <i>Annals of Hematology</i> , 2012, 91, 1221-1233.	1.8	37
26	DNMT3A mutations are rare in childhood acute myeloid leukemia. <i>Haematologica</i> , 2011, 96, 1238-1240.	3.5	34
27	A Recurrent Activating Missense Mutation in Waldenström Macroglobulinemia Affects the DNA Binding of the ETS Transcription Factor SPI1 and Enhances Proliferation. <i>Cancer Discovery</i> , 2019, 9, 796-811.	9.4	30
28	<i>TP53</i> Mutations Predict Sensitivity to Adjuvant Gemcitabine in Patients with Pancreatic Ductal Adenocarcinoma: Next-Generation Sequencing Results from the CONKO-001 Trial. <i>Clinical Cancer Research</i> , 2020, 26, 3732-3739.	7.0	28
29	<i>TET2</i> mutations in cytogenetically normal acute myeloid leukemia: Clinical implications and evolutionary patterns. <i>Genes Chromosomes and Cancer</i> , 2014, 53, 824-832.	2.8	27
30	Impact of <i>MLL5</i> expression on decitabine efficacy and DNA methylation in acute myeloid leukemia. <i>Haematologica</i> , 2014, 99, 1456-1464.	3.5	26
31	14q deletions are associated with trisomy 12, <i>NOTCH1</i> mutations and unmutated <i>IGHV</i> genes in chronic lymphocytic leukemia and small lymphocytic lymphoma. <i>Genes Chromosomes and Cancer</i> , 2014, 53, 657-666.	2.8	25
32	Clonal hematopoiesis and its emerging effects on cellular therapies. <i>Leukemia</i> , 2021, 35, 2752-2758.	7.2	21
33	Clonal hematopoiesis is associated with improved survival in patients with metastatic colorectal cancer from the FIRE-3 trial. <i>Blood</i> , 2022, 139, 1593-1597.	1.4	21
34	Spliceosome mutations in myelodysplastic syndromes and chronic myelomonocytic leukemia. <i>Oncotarget</i> , 2012, 3, 1284-1293.	1.8	19
35	Single-cell omics: Overview, analysis, and application in biomedical science. <i>Journal of Cellular Biochemistry</i> , 2021, 122, 1571-1578.	2.6	18
36	The clinical implications of clonal hematopoiesis in hematopoietic cell transplantation. <i>Blood Reviews</i> , 2021, 46, 100744.	5.7	16

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37	A genetically defined signature of responsiveness to erlotinib in early-stage pancreatic cancer patients: Results from the CONKO-005 trial. <i>EBioMedicine</i> , 2021, 66, 103327.	6.1	16
38	Modeling clonal hematopoiesis in umbilical cord blood cells by CRISPR/Cas9. <i>Leukemia</i> , 2022, 36, 1102-1110.	7.2	14
39	The landscape of genetic aberrations in myxofibrosarcoma. <i>International Journal of Cancer</i> , 2022, 151, 565-577.	5.1	13
40	Comprehensive CRISPR-Cas9 screens identify genetic determinants of drug responsiveness in multiple myeloma. <i>Blood Advances</i> , 2021, 5, 2391-2402.	5.2	10
41	<i>CD117</i> expression associates with other molecular markers and is not an independent prognostic factor in cytogenetically normal acute myeloid leukaemia. <i>British Journal of Haematology</i> , 2012, 158, 208-215.	2.5	9
42	Clonal hematopoiesis of indeterminate potential-related epigenetic age acceleration correlates with clonal hematopoiesis of indeterminate potential clone size in patients with high morbidity. <i>Haematologica</i> , 2022, 107, 1703-1708.	3.5	8
43	The hypomorphic TERT A1062T variant is associated with increased treatment-related toxicity in acute myeloid leukemia. <i>Annals of Hematology</i> , 2017, 96, 895-904.	1.8	7
44	<i>Nfkbie</i> -deficiency leads to increased susceptibility to develop B-cell lymphoproliferative disorders in aged mice. <i>Blood Cancer Journal</i> , 2020, 10, 38.	6.2	7
45	Lack of noncanonical RAS mutations in cytogenetically normal acute myeloid leukemia. <i>Annals of Hematology</i> , 2014, 93, 977-982.	1.8	6
46	Haematopoietic ageing through the lens of single-cell technologies. <i>DMM Disease Models and Mechanisms</i> , 2021, 14, .	2.4	6
47	Refining AML Treatment: The Role of Genetics in Response and Resistance Evaluation to New Agents. <i>Cancers</i> , 2022, 14, 1689.	3.7	6
48	Phosphoinositide Phospholipase C $\beta$ 1 ( <i>PI-PLC<math>\beta</math>1</i> ) Gene in Myelodysplastic Syndromes and Cytogenetically Normal Acute Myeloid Leukemia: Not a Deletion, but Increased <i>PI-PLC<math>\beta</math>1</i> Expression Is an Independent Prognostic Factor. <i>Journal of Clinical Oncology</i> , 2010, 28, e384-e387.	1.6	5
49	Reply to I.H.I.M. Hollink et al. <i>Journal of Clinical Oncology</i> , 2010, 28, e527-e528.	1.6	4
50	<i>NADH</i> dehydrogenase subunit 4 variant sequences in childhood acute myeloid leukaemia. <i>British Journal of Haematology</i> , 2013, 161, 891-895.	2.5	3
51	Reply to S. Masuda. <i>Journal of Clinical Oncology</i> , 2011, 29, 4593-4594.	1.6	2
52	Clinical Impact of TERT A1062T Mutations in Younger Patients with Acute Myeloblastic Leukemia. <i>Blood</i> , 2012, 120, 1381-1381.	1.4	2
53	Acute left ventricular insufficiency in a Burkitt Lymphoma patient with myocardial involvement and extensive local tumor cell lysis: a case report. <i>BMC Cardiovascular Disorders</i> , 2022, 22, 31.	1.7	2
54	Genetic changes of miR-182 G106A: rather a polymorphism than a somatic mutation. <i>Annals of Hematology</i> , 2011, 90, 1107-1109.	1.8	1

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55	FLT3-ITD and Age Are the Major Prognostic Factors In Relapsed AML with Normal Karyotype. Blood, 2010, 116, 1719-1719.	1.4	1
56	Genomic Landscape and Clonal Evolution of AML. Hematologic Malignancies, 2021, , 103-118.	0.2	0
57	ID1 Expression Correlates with CEBPA Mutational Status and Is Not An Independent Risk Factor in Cytogenetically Normal AML,. Blood, 2011, 118, 3554-3554.	1.4	0
58	Next Generation Sequencing for Minimal Residual Disease Monitoring in AML Patients with FLT3-ITD,. Blood, 2011, 118, 3548-3548.	1.4	0
59	The Clinical and Prognostic Influence Of Mutations In The Cohesin Complex In Acute Myeloid Leukemia. Blood, 2013, 122, 1314-1314.	1.4	0