

Stefan Deneberg

List of Publications by Year in descending order

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papers

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687220

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#	ARTICLE	IF	CITATIONS
1	A risk score based on real-world data to predict early death in acute promyelocytic leukemia. <i>Haematologica</i> , 2022, 107, 1528-1537.	1.7	12
2	Safety and efficacy of talacotuzumab plus decitabine or decitabine alone in patients with acute myeloid leukemia not eligible for chemotherapy: results from a multicenter, randomized, phase 2/3 study. <i>Leukemia</i> , 2021, 35, 62-74.	3.3	63
3	Clinical and genomic characterization of patients diagnosed with the provisional entity acute myeloid leukemia with <i>BCR-ABL1</i> , a Swedish population-based study. <i>Genes Chromosomes and Cancer</i> , 2021, 60, 426-433.	1.5	7
4	Molecular status 36 months after TKI discontinuation in CML is highly predictive for subsequent loss of MMR-final report from AFTER-SKI. <i>Leukemia</i> , 2021, 35, 2416-2418.	3.3	13
5	Is there an impact of measurable residual disease as assessed by multiparameter flow cytometry on survival of AML patients treated in clinical practice? A population-based study. <i>Leukemia and Lymphoma</i> , 2021, 62, 1973-1981.	0.6	4
6	Decreasing early mortality in acute myeloid leukaemia in Sweden 1997-2014: improving performance status is a major contributing factor. <i>British Journal of Haematology</i> , 2020, 188, 187-191.	1.2	7
7	AML displays increased CTCF occupancy associated with aberrant gene expression and transcription factor binding. <i>Blood</i> , 2020, 136, 339-352.	0.6	17
8	Real-world data on treatment patterns and outcomes of hypomethylating therapy in patients with newly diagnosed acute myeloid leukaemia aged 60 years. <i>British Journal of Haematology</i> , 2020, 189, e13-e16.	1.2	10
9	The prognostic impact of FLT3-ITD and NPM1 mutation in adult AML is age-dependent in the population-based setting. <i>Blood Advances</i> , 2020, 4, 1094-1101.	2.5	44
10	Improved survival of men 50 to 75 years old with acute myeloid leukemia over a 20-year period. <i>Blood</i> , 2019, 134, 1558-1561.	0.6	38
11	Acute myeloid leukemia in very old patients. <i>Haematologica</i> , 2018, 103, e578-e580.	1.7	17
12	Incidence and prognostic significance of isolated trisomies in adult acute myeloid leukemia: A population-based study from the Swedish AML registry. <i>European Journal of Haematology</i> , 2017, 98, 493-500.	1.1	14
13	Reasons for Decreasing Early Mortality in Acute Myeloid Leukemia: An Epidemiological Study from the Swedish Acute Leukemia Registry. <i>Blood</i> , 2015, 126, 3748-3748.	0.6	0
14	Phenotypic and Functional Alterations of Bone Marrow Mesenchymal Stem and Progenitor Cells in Chronic Myeloid Leukemia. <i>Blood</i> , 2015, 126, 2398-2398.	0.6	0
15	<i>microRNA-34b/c</i> on chromosome 11q23 is aberrantly methylated in chronic lymphocytic leukemia. <i>Epigenetics</i> , 2014, 9, 910-917.	1.3	43
16	Differential methylation in CN-AML preferentially targets non-CGI regions and is dictated by <i>DNMT3A</i> mutational status and associated with predominant hypomethylation of HOX genes. <i>Epigenetics</i> , 2014, 9, 1108-1119.	1.3	74
17	A Combination Regimen of Bortezomib, Cyclophosphamide and Betamethasone Gives Quicker, Better and More Durable Response than VAD/CyBet Regimens: Results from a Swedish Retrospective Analysis. <i>Acta Haematologica</i> , 2013, 130, 7-15.	0.7	5
18	Epigenetics in Myeloid Malignancies. <i>Methods in Molecular Biology</i> , 2012, 863, 119-137.	0.4	7

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19	Genome-Wide DNA Methylation Analysis Shows Enrichment of Differential Methylation in "Open Seas" and Enhancers and Reveals Hypomethylation in DNMT3A Mutated Cytogenetically Normal AML (CN-AML). <i>Blood</i> , 2012, 120, 653-653.	0.6	3
20	Prognostic DNA methylation patterns in cytogenetically normal acute myeloid leukemia are predefined by stem cell chromatin marks. <i>Blood</i> , 2011, 118, 5573-5582.	0.6	67
21	APR-246 exhibits anti-leukemic activity and synergism with conventional chemotherapeutic drugs in acute myeloid leukemia cells. <i>European Journal of Haematology</i> , 2011, 86, 206-215.	1.1	61
22	Gene-specific and global methylation patterns predict outcome in patients with acute myeloid leukemia. <i>Leukemia</i> , 2010, 24, 932-941.	3.3	113
23	Global and HOX Gene DNA Methylation In Normal Karyotype Acute Myeloid Leukemia: Clinical Implications and Molecular Correlations. <i>Blood</i> , 2010, 116, 231-231.	0.6	0
24	Impact of chromosome 13 deletion and plasma cell load on long-term survival of patients with multiple myeloma undergoing autologous transplantation. <i>Oncology Reports</i> , 2009, 22, 137-42.	1.2	7
25	Low p14ARF expression in de novo acute myeloid leukemia with normal karyotype is associated with poor survival. <i>Leukemia and Lymphoma</i> , 2009, 50, 1512-1518.	0.6	4
26	Single nucleotide polymorphism genomic arrays analysis of t(8;21) acute myeloid leukemia cells. <i>Haematologica</i> , 2009, 94, 1301-1306.	1.7	16
27	In Vitro and Ex Vivo Studies On Cell Lines and Primary Human Leukemia Cells of the Effects of APR-246 Alone and in Combination with Conventional Chemotherapeutic Drugs.. <i>Blood</i> , 2009, 114, 2751-2751.	0.6	0
28	The FLT3 inhibitor PKC412 in combination with cytostatic drugs in vitro in acute myeloid leukemia. <i>Cancer Chemotherapy and Pharmacology</i> , 2008, 62, 439-448.	1.1	32
29	Different Incidence and Implications of DNA Hypermethylation in De Novo AML Compared to High-Risk MDS and AML Following MDS.. <i>Blood</i> , 2008, 112, 3337-3337.	0.6	0
30	Relapse of preB-ALL after rituximab treatment for chronic graft versus host disease. Implications for its use?. <i>Medical Oncology</i> , 2007, 24, 354-356.	1.2	5
31	Expression of p14ARF in De Novo AML with Normal Karyotype. Implication on Drug Resistance and Survival.. <i>Blood</i> , 2007, 110, 4261-4261.	0.6	6
32	Effects of low-dose prednisolone on endothelial function, atherosclerosis, and traditional risk factors for atherosclerosis in patients with rheumatoid arthritis—a randomized study. <i>Journal of Rheumatology</i> , 2007, 34, 1810-6.	1.0	85