

# Jan Jacob Schuringa

## List of Publications by Year in descending order

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91  
papers

3,366  
citations

136740

32  
h-index

155451

55  
g-index

96  
all docs

96  
docs citations

96  
times ranked

5810  
citing authors

#	ARTICLE	IF	CITATIONS
1	The Glycolytic Gatekeeper PDK1 defines different metabolic states between genetically distinct subtypes of human acute myeloid leukemia. <i>Nature Communications</i> , 2022, 13, 1105.	5.8	14
2	Inhibition of the succinyl dehydrogenase complex in acute myeloid leukemia leads to a lactate-fueled respiratory metabolic vulnerability. <i>Nature Communications</i> , 2022, 13, 2013.	5.8	22
3	Monocytosis and its association with clonal hematopoiesis in community-dwelling individuals. <i>Blood Advances</i> , 2022, 6, 4174-4184.	2.5	8
4	The Expression of NTAL and Its Protein Interactors Is Associated With Clinical Outcomes in Acute Myeloid Leukemia. <i>Molecular and Cellular Proteomics</i> , 2021, 20, 100091.	2.5	1
5	Pretransplantation MRD in Older Patients With AML After Treatment With Decitabine or Conventional Chemotherapy. <i>Transplantation and Cellular Therapy</i> , 2021, 27, 246-252.	0.6	9
6	MLL5 improves ATRA driven differentiation and promotes xenotransplant engraftment in acute promyelocytic leukemia model. <i>Cell Death and Disease</i> , 2021, 12, 371.	2.7	5
7	Prevalence, predictors, and outcomes of clonal hematopoiesis in individuals aged ≥80 years. <i>Blood Advances</i> , 2021, 5, 2115-2122.	2.5	44
8	The USP7-TRIM27 axis mediates non-canonical PRC1.1 function and is a druggable target in leukemia. <i>IScience</i> , 2021, 24, 102435.	1.9	19
9	Peripheral blood cytopenias in the aging general population and risk of incident hematological disease and mortality. <i>Blood Advances</i> , 2021, 5, 3266-3278.	2.5	6
10	CombiFlow: Combinatorial AML-specific plasma membrane expression profiles allow longitudinal tracking of clones. <i>Blood Advances</i> , 2021, , .	2.5	4
11	The Combination of Gefitinib With ATRA and ATO Induces Myeloid Differentiation in Acute Promyelocytic Leukemia Resistant Cells. <i>Frontiers in Oncology</i> , 2021, 11, 686445.	1.3	8
12	CombiFlow: Flow cytometry-based identification and characterization of genetically and functionally distinct AML subclones. <i>STAR Protocols</i> , 2021, 2, 100864.	0.5	1
13	The IL1-IL1RAP axis plays an important role in the inflammatory leukemic niche that favors acute myeloid leukemia proliferation over normal hematopoiesis. <i>Haematologica</i> , 2021, 106, 3067-3078.	1.7	18
14	The EHA Research Roadmap: Normal Hematopoiesis. <i>HemaSphere</i> , 2021, 5, e669.	1.2	1
15	HUWE1 cooperates with RAS activation to control leukemia cell proliferation and human hematopoietic stem cells differentiation fate. <i>Cancer Gene Therapy</i> , 2020, 27, 830-833.	2.2	4
16	Dissecting Clonal Heterogeneity in AML. <i>Cancer Cell</i> , 2020, 38, 782-784.	7.7	16
17	Reduced SLIT2 is Associated with Increased Cell Proliferation and Arsenic Trioxide Resistance in Acute Promyelocytic Leukemia. <i>Cancers</i> , 2020, 12, 3134.	1.7	7
18	Innovations, challenges, and minimal information for standardization of humanized mice. <i>EMBO Molecular Medicine</i> , 2020, 12, e8662.	3.3	82

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19	Chromatin-Based Classification of Genetically Heterogeneous AMLs into Two Distinct Subtypes with Diverse Stemness Phenotypes. <i>Cell Reports</i> , 2019, 26, 1059-1069.e6.	2.9	33
20	Elevated VMP1 expression in acute myeloid leukemia amplifies autophagy and is protective against venetoclax-induced apoptosis. <i>Cell Death and Disease</i> , 2019, 10, 421.	2.7	27
21	Not type of induction therapy but consolidation with allogeneic hematopoietic cell transplantation determines outcome in older AML patients: A single center experience of 355 consecutive patients. <i>Leukemia Research</i> , 2019, 80, 33-39.	0.4	11
22	RUNX1 mutations enhance self-renewal and block granulocytic differentiation in human in vitro models and primary AMLs. <i>Blood Advances</i> , 2019, 3, 320-332.	2.5	27
23	HIF1/2-exerted control over glycolytic gene expression is not functionally relevant for glycolysis in human leukemic stem/progenitor cells. <i>Cancer &amp; Metabolism</i> , 2019, 7, 11.	2.4	46
24	Protein quality control in the nucleolus safeguards recovery of epigenetic regulators after heat shock. <i>ELife</i> , 2019, 8, .	2.8	46
25	Prospective Isolation and Characterization of Genetically and Functionally Distinct AML Subclones. <i>Cancer Cell</i> , 2018, 34, 674-689.e8.	7.7	71
26	Genetically engineered mesenchymal stromal cells produce IL-3 and TPO to further improve human scaffold-based xenograft models. <i>Experimental Hematology</i> , 2017, 51, 36-46.	0.2	19
27	Smart niche usage: release its fat and burn it!. <i>Blood</i> , 2017, 129, 1239-1240.	0.6	1
28	Inhibition of autophagy as a treatment strategy for p53 wild-type acute myeloid leukemia. <i>Cell Death and Disease</i> , 2017, 8, e2927-e2927.	2.7	72
29	BRD3/4 inhibition and FLT3-ligand deprivation target pathways that are essential for the survival of human MLL-AF9+ leukemic cells. <i>PLoS ONE</i> , 2017, 12, e0189102.	1.1	10
30	Modeling of Chronic Myeloid Leukemia: An Overview of <i>In Vivo</i> Murine and Human Xenograft Models. <i>Stem Cells International</i> , 2016, 2016, 1-12.	1.2	8
31	Hypoxia-Like Signatures Induced by BCR-ABL Potentially Alter the Glutamine Uptake for Maintaining Oxidative Phosphorylation. <i>PLoS ONE</i> , 2016, 11, e0153226.	1.1	16
32	Establishing human leukemia xenograft mouse models by implanting human bone marrow-like scaffold-based niches. <i>Blood</i> , 2016, 128, 2949-2959.	0.6	65
33	Loss of ASXL1 triggers an apoptotic response in human hematopoietic stem and progenitor cells. <i>Experimental Hematology</i> , 2016, 44, 1188-1196.e6.	0.2	11
34	Autophagy Proteins ATG5 and ATG7 Are Essential for the Maintenance of Human CD34+ Hematopoietic Stem-Progenitor Cells. <i>Stem Cells</i> , 2016, 34, 1651-1663.	1.4	67
35	Depletion of SAM50 Specifically Targets BCR-ABL-Expressing Leukemic Stem and Progenitor Cells by Interfering with Mitochondrial Functions. <i>Stem Cells and Development</i> , 2016, 25, 427-437.	1.1	10
36	Non-canonical PRC1.1 Targets Active Genes Independent of H3K27me3 and Is Essential for Leukemogenesis. <i>Cell Reports</i> , 2016, 14, 332-346.	2.9	126

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37	Constitutive NF- $\kappa$ B activation in AML: Causes and treatment strategies. <i>Critical Reviews in Oncology/Hematology</i> , 2016, 98, 35-44.	2.0	60
38	RUNX1 Mutations Cause a Myeloid Differentiation Block Leading to the Formation of a Long Term Expanding CD34+/CD33+/CD45RA+/CD123+ Cell Population. <i>Blood</i> , 2016, 128, 1979-1979.	0.6	9
39	Mitochondrial Dysfunction in Human Leukemic Stem/Progenitor Cells upon Loss of RAC2. <i>PLoS ONE</i> , 2015, 10, e0128585.	1.1	15
40	Loss of ASXL1 Triggers an Apoptotic Response in Human Hematopoietic Stem and Progenitor Cells. <i>Blood</i> , 2015, 126, 4107-4107.	0.6	0
41	Convergence of Hypoxia and TGF $\beta$ 2 Pathways on Cell Cycle Regulation in Human Hematopoietic Stem/Progenitor Cells. <i>PLoS ONE</i> , 2014, 9, e93494.	1.1	49
42	ELMO1 Is Upregulated in AML CD34+ Stem/Progenitor Cells, Mediates Chemotaxis and Predicts Poor Prognosis in Normal Karyotype AML. <i>PLoS ONE</i> , 2014, 9, e111568.	1.1	12
43	The TAK1-NF- $\kappa$ B axis as therapeutic target for AML. <i>Blood</i> , 2014, 124, 3130-3140.	0.6	47
44	Ageing Impairs Long-Term Hematopoietic Regeneration after Autologous Stem Cell Transplantation. <i>Biology of Blood and Marrow Transplantation</i> , 2014, 20, 865-871.	2.0	28
45	Ex Vivo Assays to Study Self-Renewal, Long-Term Expansion, and Leukemic Transformation of Genetically Modified Human Hematopoietic and Patient-Derived Leukemic Stem Cells. <i>Methods in Molecular Biology</i> , 2014, 1185, 195-210.	0.4	9
46	Loss of ASXL1 Triggers an Apoptotic Response in Human Hematopoietic Stem and Progenitor Cells. <i>Blood</i> , 2014, 124, 4619-4619.	0.6	0
47	A Proteomics and Transcriptomics Approach to Identify Leukemic Stem Cell (LSC) Markers. <i>Molecular and Cellular Proteomics</i> , 2013, 12, 626-637.	2.5	79
48	Nonredundant and locus-specific gene repression functions of PRC1 paralog family members in human hematopoietic stem/progenitor cells. <i>Blood</i> , 2013, 121, 2452-2461.	0.6	54
49	Establishing Human Niche Xenograft Models For Myeloid and Lymphoid Leukemia Driven By MLL-AF9. <i>Blood</i> , 2013, 122, 1646-1646.	0.6	1
50	Differential Localization Of RAC1 and RAC2 Reflects Their Specific Functions In Normal and Leukemic Human Hematopoietic Stem/Progenitor Cells. <i>Blood</i> , 2013, 122, 2892-2892.	0.6	0
51	Decreased PU.1 and Enhanced CITED2 Cooperate To Maintain Self-Renewal In Hematopoietic Stem/Progenitors. <i>Blood</i> , 2013, 122, 2411-2411.	0.6	0
52	Convergence Of Hypoxia and TGF $\beta$ 2 Pathways On Cell Cycle Regulation In Human Hematopoietic Stem/Progenitor Cells. <i>Blood</i> , 2013, 122, 3694-3694.	0.6	1
53	Mouse Versus Human Extrinsic Cues Dictate Transformation Potential In BCR-ABL/BMI1-Induced Leukemia In Humanized Xenograft Models. <i>Blood</i> , 2013, 122, 515-515.	0.6	29
54	STAT5-mediated self-renewal of normal hematopoietic and leukemic stem cells. <i>Jak-stat</i> , 2012, 1, 13-25.	2.2	22

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55	Cancer stem cell definitions and terminology: the devil is in the details. <i>Nature Reviews Cancer</i> , 2012, 12, 767-775.	12.8	599
56	Reconstructing the human hematopoietic niche in immunodeficient mice: opportunities for studying primary multiple myeloma. <i>Blood</i> , 2012, 120, e9-e16.	0.6	104
57	Differential Localization of RAC1 and RAC2 Reflects Their Specific Functions in Normal and Leukemic Human Hematopoietic Stem/Progenitor Cells.. <i>Blood</i> , 2012, 120, 2302-2302.	0.6	0
58	Identification of HIF2 $\alpha$ as an important STAT5 target gene in human hematopoietic stem cells. <i>Blood</i> , 2011, 117, 3320-3330.	0.6	63
59	KRASG12V Enhances Proliferation and Initiates Myelomonocytic Differentiation in Human Stem/Progenitor Cells via Intrinsic and Extrinsic Pathways. <i>Journal of Biological Chemistry</i> , 2011, 286, 6061-6070.	1.6	21
60	Down-regulation of GATA1 uncouples STAT5-induced erythroid differentiation from stem/progenitor cell proliferation. <i>Blood</i> , 2010, 115, 4367-4376.	0.6	22
61	BMI1 collaborates with BCR-ABL in leukemic transformation of human CD34+ cells. <i>Blood</i> , 2010, 116, 4621-4630.	0.6	72
62	Single-Cell STAT5 Signal Transduction Profiling in Normal and Leukemic Stem and Progenitor Cell Populations Reveals Highly Distinct Cytokine Responses. <i>PLoS ONE</i> , 2009, 4, e7989.	1.1	28
63	Repression of BMI1 in normal and leukemic human CD34+ cells impairs self-renewal and induces apoptosis. <i>Blood</i> , 2009, 114, 1498-1505.	0.6	127
64	Ex Vivo Assays to Study Self-Renewal and Long-Term Expansion of Genetically Modified Primary Human Acute Myeloid Leukemia Stem Cells. <i>Methods in Molecular Biology</i> , 2009, 538, 287-300.	0.4	29
65	Autologous Stem Cell Transplantation Induces a Phenotypical Shift From CMP to GMP Progenitors, Reduces Clonogenic Potential and Enhances in Vitro and In Vivo Cycling Activity Defined by 18f-FLT PET Scan.. <i>Blood</i> , 2009, 114, 4473-4473.	0.6	0
66	Overexpression of Oncogenic KRAS G12V in Human Stem and Progenitor Cells Enhances Proliferation and Initiates Monocytic Differentiation Via Intrinsic and Extrinsic Pathways.. <i>Blood</i> , 2009, 114, 3975-3975.	0.6	0
67	Single-Cell STAT5 Signal Transduction Profiling in Normal and Leukemic Stem and Progenitor Cell Populations Reveals Highly Distinct Cytokine Responses.. <i>Blood</i> , 2009, 114, 2510-2510.	0.6	0
68	Identification of Human Hematopoietic Stem Cell-Specific STAT5 Target Genes Involved in Self-Renewal and Transformation.. <i>Blood</i> , 2009, 114, 568-568.	0.6	0
69	Mucin1 expression is enriched in the human stem cell fraction of cord blood and is upregulated in majority of the AML cases. <i>Experimental Hematology</i> , 2008, 36, 1254-1265.	0.2	35
70	Maximal STAT5-Induced Proliferation and Self-Renewal at Intermediate STAT5 Activity Levels. <i>Molecular and Cellular Biology</i> , 2008, 28, 6668-6680.	1.1	76
71	Reduced activation of protein kinase B, Rac, and F-actin polymerization contributes to an impairment of stromal cell-derived factor-1-induced migration of CD34+ cells from patients with myelodysplasia. <i>Blood</i> , 2008, 111, 359-368.	0.6	43
72	Long-term maintenance of human hematopoietic stem/progenitor cells by expression of BMI1. <i>Blood</i> , 2008, 111, 2621-2630.	0.6	98

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73	Distinct Gene Expression Profiling in AML in Elderly Versus Younger Patients. <i>Blood</i> , 2008, 112, 2546-2546.	0.6	8
74	The Polycomb Gene BMI1 Collaborates with BCR-ABL in Leukemic Transformation of Human Cord Blood CD34+ Cells. <i>Blood</i> , 2008, 112, 1350-1350.	0.6	0
75	Inhibition of Long-Term Expansion in a Subgroup of Acute Myeloid Leukemia Samples by Dasatinib. <i>Blood</i> , 2008, 112, 4016-4016.	0.6	0
76	STAT5 is required for long-term maintenance of normal and leukemic human stem/progenitor cells. <i>Blood</i> , 2007, 110, 2880-2888.	0.6	91
77	Reintroduction of C/EBP $\beta$ in leukemic CD34+ stem/progenitor cells impairs self-renewal and partially restores myelopoiesis. <i>Blood</i> , 2007, 110, 1317-1325.	0.6	41
78	Expansion of normal and leukemic human hematopoietic stem/progenitor cells requires Rac-mediated interaction with stromal cells. <i>Experimental Hematology</i> , 2007, 35, 782-792.	0.2	22
79	Establishing long-term cultures with self-renewing acute myeloid leukemia stem/progenitor cells. <i>Experimental Hematology</i> , 2007, 35, 1538-1549.	0.2	80
80	AML1/RUNX1, One of the Most Common Targets of Aberration in Acute Myeloid Leukemia as a Transcriptional Regulator of Vascular Endothelial Growth Factor (VEGFA). <i>Blood</i> , 2007, 110, 1618-1618.	0.6	0
81	Transcription Factor Dosage: Maximal STAT5-Induced Proliferation and Stem Cell Self-Renewal at Intermediate STAT5 Activity Levels. <i>Blood</i> , 2007, 110, 2242-2242.	0.6	27
82	Signaling pathways in self-renewing hematopoietic and leukemic stem cells: do all stem cells need a niche?. <i>Human Molecular Genetics</i> , 2006, 15, R210-R219.	1.4	102
83	STAT5-induced self-renewal and impaired myelopoiesis of human hematopoietic stem/progenitor cells involves down-modulation of C/EBP $\beta$ . <i>Blood</i> , 2006, 107, 4326-4333.	0.6	40
84	Enforced Expression of NUP98-HOXA9 in Human CD34+ Cells Enhances Stem Cell Proliferation. <i>Cancer Research</i> , 2006, 66, 11781-11791.	0.4	73
85	Ex-Vivo Expansion of Human Cord Blood CD34+ Cells by Overexpression of Bmi-1. <i>Blood</i> , 2006, 108, 1329-1329.	0.6	0
86	Dose Dependent Effects of STAT5 on Proliferation, Differentiation and Self Renewal of Hematopoietic Stem/Progenitor Cells. <i>Blood</i> , 2006, 108, 1321-1321.	0.6	0
87	Enforced expression of an Flt3 internal tandem duplication in human CD34+ cells confers properties of self-renewal and enhanced erythropoiesis. <i>Blood</i> , 2005, 105, 77-84.	0.6	51
88	Expansion of Normal and Leukemic Human Hematopoietic Stem/Progenitor Cells Requires Rac-Mediated Interaction with Stromal Cells. <i>Blood</i> , 2005, 106, 1398-1398.	0.6	0
89	STAT5-Induced Self-Renewal and Impaired Myelopoiesis of Human Hematopoietic Stem/Progenitor Cells Involves Downmodulation of C/EBP $\beta$ . <i>Blood</i> , 2005, 106, 268-268.	0.6	0
90	Constitutive Activation of STAT5A Promotes Human Hematopoietic Stem Cell Self-Renewal and Erythroid Differentiation. <i>Journal of Experimental Medicine</i> , 2004, 200, 623-635.	4.2	115

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91	Enforced Activation of STAT5A Facilitates the Generation of Embryonic Stem-Derived Hematopoietic Stem Cells That Contribute to Hematopoiesis In Vivo. <i>Stem Cells</i> , 2004, 22, 1191-1204.	1.4	45