

# Jan Stuchly

## List of Publications by Year in descending order

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Version: 2024-02-01

32  
papers

887  
citations

430874

18  
h-index

477307

29  
g-index

34  
all docs

34  
docs citations

34  
times ranked

1887  
citing authors

#	ARTICLE	IF	CITATIONS
1	<i>DUX4</i>, <i>ZNF384</i> and <i>PAX5</i>-P80R mutated B-cell precursor acute lymphoblastic leukemia frequently undergo monocytic switch. <i>Haematologica</i> , 2021, 106, 2066-2075.	3.5	29
2	A novel class of ZNF384 aberrations in acute leukemia. <i>Blood Advances</i> , 2021, 5, 4393-4397.	5.2	11
3	A homozygous deletion in the SLC19A1 gene as a cause of folate-dependent recurrent megaloblastic anemia. <i>Blood</i> , 2020, 135, 2427-2431.	1.4	13
4	Single-Cell Profiling of Signal Transduction Pathways in Pediatric T-Cell Acute Lymphoblastic Leukemia By Mass Cytometry: Dissecting JAK/STAT and PI3K/Akt/mTOR Active Signalling. <i>Blood</i> , 2020, 136, 38-39.	1.4	0
5	Molecular Basis of Cisplatin Resistance in Testicular Germ Cell Tumors. <i>Cancers</i> , 2019, 11, 1316.	3.7	12
6	<i>ERG</i> deletions in childhood acute lymphoblastic leukemia with <i>DUX4</i> rearrangements are mostly polyclonal, prognostically relevant and their detection rate strongly depends on screening method sensitivity. <i>Haematologica</i> , 2019, 104, 1407-1416.	3.5	34
7	Genomic landscape of pediatric B-other acute lymphoblastic leukemia in a consecutive European cohort. <i>Haematologica</i> , 2019, 104, 1396-1406.	3.5	78
8	The EuroFlow PID Orientation Tube for Flow Cytometric Diagnostic Screening of Primary Immunodeficiencies of the Lymphoid System. <i>Frontiers in Immunology</i> , 2019, 10, 246.	4.8	100
9	A high-throughput pipeline for validation of antibodies. <i>Nature Methods</i> , 2018, 15, 909-912.	19.0	52
10	Appearance of cytomegalovirus-specific T cells predicts fast resolution of viremia post hematopoietic stem cell transplantation. <i>Cytometry Part B - Clinical Cytometry</i> , 2017, 92, 380-388.	1.5	18
11	Common Variable Immunodeficiency patients with a phenotypic profile of immunosenescence present with thrombocytopenia. <i>Scientific Reports</i> , 2017, 7, 39710.	3.3	31
12	<i>ETV6/RUNX1</i>-like acute lymphoblastic leukemia: A novel B-cell precursor leukemia subtype associated with the CD27/CD44 immunophenotype. <i>Genes Chromosomes and Cancer</i> , 2017, 56, 608-616.	2.8	63
13	An activating mutation of GNB1 is associated with resistance to tyrosine kinase inhibitors in ETV6-ABL1-positive leukemia. <i>Oncogene</i> , 2017, 36, 5985-5994.	5.9	21
14	Lymphocyte enrichment using CD81-targeted immunoaffinity matrix. <i>Cytometry Part A: the Journal of the International Society for Analytical Cytology</i> , 2017, 91, 62-72.	1.5	9
15	Wilms tumor gene 1 (WT1), TP53, RAS/BRAF and KIT aberrations in testicular germ cell tumors. <i>Cancer Letters</i> , 2016, 376, 367-376.	7.2	16
16	MetaMass, a tool for meta-analysis of subcellular proteomics data. <i>Nature Methods</i> , 2016, 13, 837-840.	19.0	30
17	Characterization of leukemias with ETV6-ABL1 fusion. <i>Haematologica</i> , 2016, 101, 1082-1093.	3.5	66
18	Distinct bilineal leukemia immunophenotypes are not genetically determined. <i>Blood</i> , 2016, 128, 2263-2266.	1.4	23

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19	High-resolution Antibody Array Analysis of Childhood Acute Leukemia Cells. <i>Molecular and Cellular Proteomics</i> , 2016, 15, 1246-1261.	3.8	10
20	GNB1 K89M Drives TKI Resistance in ETV6-ABL1-Positive Leukemia. <i>Blood</i> , 2016, 128, 751-751.	1.4	0
21	The predictive strength of next-generation sequencing MRD detection for relapse compared with current methods in childhood ALL. <i>Blood</i> , 2015, 126, 1045-1047.	1.4	82
22	L-Asparaginase More Effectively Targets Leukemic Cells with Low Glycolytic Activity. <i>Blood</i> , 2015, 126, 1285-1285.	1.4	3
23	Characterization of Leukemias with ETV6-ABL1 Fusion. <i>Blood</i> , 2015, 126, 84-84.	1.4	1
24	Analyses of large flow cytometry datasets. <i>Cytometry Part A: the Journal of the International Society for Analytical Cytology</i> , 2014, 85, 203-205.	1.5	6
25	Next Generation Amplicon Sequencing of Immunoglobulin Heavy Chain Gene Rearrangements for Minimal Residual Disease (MRD) Stratification in Childhood Acute Lymphoblastic Leukemia (ALL): A Comparison with Classical qPCR-Based Technique. <i>Blood</i> , 2014, 124, 2395-2395.	1.4	0
26	Novel Flow Cytometry-Based Method Of Affinity Proteomics Revealing Expression, Post-Translational Modification and Proteolysis In Primary Childhood Acute Leukemias. <i>Blood</i> , 2013, 122, 2553-2553.	1.4	0
27	Real-time PCR quantification of major Wilms's tumor gene 1 (WT1) isoforms in acute myeloid leukemia, their characteristic expression patterns and possible functional consequences. <i>Leukemia</i> , 2012, 26, 2086-2095.	7.2	31
28	An automated analysis of highly complex flow cytometry-based proteomic data. <i>Cytometry Part A: the Journal of the International Society for Analytical Cytology</i> , 2012, 81A, 120-129.	1.5	13
29	Signature profiles of CMV-specific T-cells in patients with CMV reactivation after hematopoietic SCT. <i>Bone Marrow Transplantation</i> , 2011, 46, 1089-1098.	2.4	43
30	Plasma EBV-DNA monitoring in Epstein-Barr virus-positive Hodgkin lymphoma patients. <i>Apmis</i> , 2011, 119, 10-16.	2.0	34
31	Multiplexed immunoprecipitation with 1725 commercially available antibodies to cellular proteins. <i>Proteomics</i> , 2011, 11, 4578-4582.	2.2	27
32	Profiling of polychromatic flow cytometry data on B-cells reveals patients' clusters in common variable immunodeficiency. <i>Cytometry Part A: the Journal of the International Society for Analytical Cytology</i> , 2009, 75A, 902-909.	1.5	24