

# Jasmin Straube

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3232473/publications.pdf>

Version: 2024-02-01

26  
papers

1,255  
citations

623734

14  
h-index

713466

21  
g-index

28  
all docs

28  
docs citations

28  
times ranked

2769  
citing authors

#	ARTICLE	IF	CITATIONS
1	Expression of CD49f defines subsets of human regulatory T cells with divergent transcriptional landscape and function that correlate with ulcerative colitis disease activity. <i>Clinical and Translational Immunology</i> , 2021, 10, e1334.	3.8	5
2	Optimizing DNA hypomethylating therapy in acute myeloid leukemia and myelodysplastic syndromes. <i>BioEssays</i> , 2021, 43, 2100125.	2.5	4
3	Targeting Control of Cell Cycle Enhances the Activity of Conventional Chemotherapy in Chemotherapy-Resistant Acute Myeloid Leukemia. <i>Blood</i> , 2021, 138, 2241-2241.	1.4	0
4	Distinct effects of ruxolitinib and interferon-alpha on murine JAK2V617F myeloproliferative neoplasm hematopoietic stem cell populations. <i>Leukemia</i> , 2020, 34, 1075-1089.	7.2	29
5	Transcriptome dynamics of CD4+ T cells during malaria maps gradual transit from effector to memory. <i>Nature Immunology</i> , 2020, 21, 1597-1610.	14.5	43
6	MPN: The Molecular Drivers of Disease Initiation, Progression and Transformation and their Effect on Treatment. <i>Cells</i> , 2020, 9, 1901.	4.1	27
7	Hematopoietic stem and progenitor cell-restricted Cdx2 expression induces transformation to myelodysplasia and acute leukemia. <i>Nature Communications</i> , 2020, 11, 3021.	12.8	15
8	Q-Cell Glioblastoma Resource: Proteomics Analysis Reveals Unique Cell-States Are Maintained in 3D Culture. <i>Cells</i> , 2020, 9, 267.	4.1	12
9	Ex-vivo drug testing predicts chemosensitivity in acute myeloid leukemia. <i>Journal of Leukocyte Biology</i> , 2020, 107, 859-870.	3.3	15
10	Reactivation of Myc transcription in the mouse heart unlocks its proliferative capacity. <i>Nature Communications</i> , 2020, 11, 1827.	12.8	38
11	Proteomic Analysis of the Breast Cancer Brain Metastasis Microenvironment. <i>International Journal of Molecular Sciences</i> , 2019, 20, 2524.	4.1	22
12	The impact of age, NPM1mut, and FLT3ITD allelic ratio in patients with acute myeloid leukemia. <i>Blood</i> , 2018, 131, 1148-1153.	1.4	53
13	Jak2V617F and Dnmt3a loss cooperate to induce myelofibrosis through activated enhancer-driven inflammation. <i>Blood</i> , 2018, 132, 2707-2721.	1.4	56
14	CDX2 Expression in Hematopoietic Stem Cells Represents a Novel Model of De Novo Leukemia. <i>Experimental Hematology</i> , 2018, 64, S50-S51.	0.4	0
15	Recipient mucosal-associated invariant T cells control GVHD within the colon. <i>Journal of Clinical Investigation</i> , 2018, 128, 1919-1936.	8.2	78
16	Identification of Genetic Pathways Controlling Resistance to Standard Combination Chemotherapy in Acute Myeloid Leukemia. <i>Blood</i> , 2018, 132, 2771-2771.	1.4	0
17	Oncogenic-Drivers Dictate Immune Responses to Control Disease Progression in Acute Myeloid Leukaemia. <i>Blood</i> , 2018, 132, 904-904.	1.4	0
18	Ssb1 and Ssb2 cooperate to regulate mouse hematopoietic stem and progenitor cells by resolving replicative stress. <i>Blood</i> , 2017, 129, 2479-2492.	1.4	18

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19	DynOmics to identify delays and co-expression patterns across time course experiments. Scientific Reports, 2017, 7, 40131.	3.3	15
20	Interleukin-12 from CD103+ Batf3-Dependent Dendritic Cells Required for NK-Cell Suppression of Metastasis. Cancer Immunology Research, 2017, 5, 1098-1108.	3.4	98
21	In vivo CRISPR editing of DNMT3A in JAK2V617F hematopoietic stem cells induces myelofibrosis. Experimental Hematology, 2017, 53, S95.	0.4	0
22	Tumor immunoevasion by the conversion of effector NK cells into type 1 innate lymphoid cells. Nature Immunology, 2017, 18, 1004-1015.	14.5	504
23	Integrated Molecular Analysis Identifies Replicative Stress As Sensitizer to Imetelstat Therapy in AML. Blood, 2017, 130, 798-798.	1.4	2
24	A Linear Mixed Model Spline Framework for Analysing Time Course "Omics" Data. PLoS ONE, 2015, 10, e0134540.	2.5	46
25	ReadXplorer" visualization and analysis of mapped sequences. Bioinformatics, 2014, 30, 2247-2254.	4.1	127
26	New insights into <i>C. jejuni</i> hydrogen production processes by combined microarray/RNA-seq transcriptomics. Plant Biotechnology Journal, 2013, 11, 717-733.	8.3	47