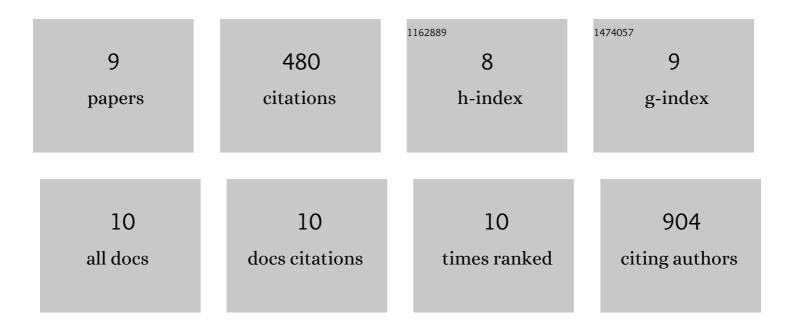
## Julia Franzen

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3573189/publications.pdf Version: 2024-02-01



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#	Article	IF	CITATIONS
1	Investigation of measurable residual disease in acute myeloid leukemia by DNA methylation patterns. Leukemia, 2022, 36, 80-89.	3.3	12
2	Hematopoietic differentiation persists in human iPSCs defective in de novo DNA methylation. BMC Biology, 2022, 20, .	1.7	3
3	DNA methylation changes during long-term in vitro cell culture are caused by epigenetic drift. Communications Biology, 2021, 4, 598.	2.0	27
4	Epigenetic Clocks Are Not Accelerated in COVID-19 Patients. International Journal of Molecular Sciences, 2021, 22, 9306.	1.8	21
5	New targeted approaches for epigenetic age predictions. BMC Biology, 2020, 18, 71.	1.7	55
6	Senescence-Associated Metabolomic Phenotype in Primary and iPSC-Derived Mesenchymal Stromal Cells. Stem Cell Reports, 2020, 14, 201-209.	2.3	62
7	HMGB2 Loss upon Senescence Entry Disrupts Genomic Organization and Induces CTCF Clustering across Cell Types. Molecular Cell, 2018, 70, 730-744.e6.	4.5	164
8	Human Platelet Lysate versus Fetal Calf Serum: These Supplements Do Not Select for Different Mesenchymal Stromal Cells. Scientific Reports, 2017, 7, 5132.	1.6	60
9	Senescenceâ€essociated <scp>DNA</scp> methylation is stochastically acquired in subpopulations of mesenchymal stem cells. Aging Cell, 2017, 16, 183-191.	3.0	70