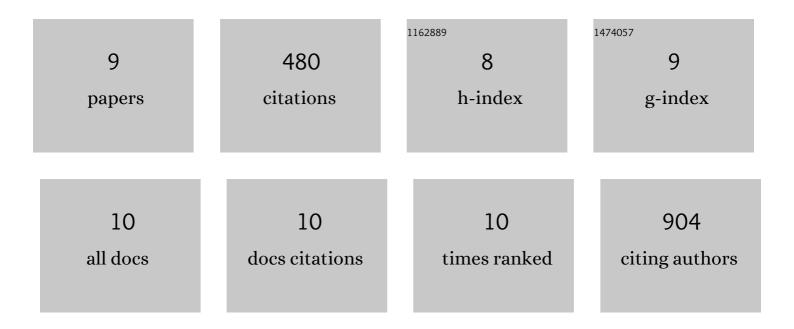
Julia Franzen

List of Publications by Year in descending order

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



IIIIIA FRANZEN

#	Article	IF	CITATIONS
1	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
2	Senescenceâ€associated <scp>DNA</scp> methylation is stochastically acquired in subpopulations of mesenchymal stem cells. Aging Cell, 2017, 16, 183-191.	3.0	70
3	Senescence-Associated Metabolomic Phenotype in Primary and iPSC-Derived Mesenchymal Stromal Cells. Stem Cell Reports, 2020, 14, 201-209.	2.3	62
4	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
5	New targeted approaches for epigenetic age predictions. BMC Biology, 2020, 18, 71.	1.7	55
6	DNA methylation changes during long-term in vitro cell culture are caused by epigenetic drift. Communications Biology, 2021, 4, 598.	2.0	27
7	Epigenetic Clocks Are Not Accelerated in COVID-19 Patients. International Journal of Molecular Sciences, 2021, 22, 9306.	1.8	21
8	Investigation of measurable residual disease in acute myeloid leukemia by DNA methylation patterns. Leukemia, 2022, 36, 80-89.	3.3	12
9	Hematopoietic differentiation persists in human iPSCs defective in de novo DNA methylation. BMC Biology, 2022, 20, .	1.7	3