

Julia Franzen

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3573189/publications.pdf>

Version: 2024-02-01

9
papers

480
citations

1162889

8
h-index

1474057

9
g-index

10
all docs

10
docs citations

10
times ranked

904
citing authors

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