Simone Ecker

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/310083/publications.pdf

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18 papers

2,191 citations

15 h-index 18 g-index

24 all docs

24 docs citations

times ranked

24

5953 citing authors

#	Article	IF	CITATIONS
1	Genetic Drivers of Epigenetic and Transcriptional Variation in Human Immune Cells. Cell, 2016, 167, 1398-1414.e24.	13.5	573
2	Epigenomic analysis detects widespread gene-body DNA hypomethylation in chronic lymphocytic leukemia. Nature Genetics, 2012, 44, 1236-1242.	9.4	525
3	Whole-genome fingerprint of the DNA methylome during human B cell differentiation. Nature Genetics, 2015, 47, 746-756.	9.4	278
4	Transcriptome characterization by RNA sequencing identifies a major molecular and clinical subdivision in chronic lymphocytic leukemia. Genome Research, 2014, 24, 212-226.	2.4	175
5	Increased DNA methylation variability in type 1 diabetes across three immune effector cell types. Nature Communications, 2016, 7, 13555.	5 . 8	142
6	Genome-wide analysis of differential transcriptional and epigenetic variability across human immune cell types. Genome Biology, 2017, 18, 18.	3.8	97
7	Epigenetic and Transcriptional Variability Shape Phenotypic Plasticity. BioEssays, 2018, 40, 1700148.	1.2	71
8	Increased DNA methylation variability in rheumatoid arthritis-discordant monozygotic twins. Genome Medicine, 2018, 10, 64.	3 . 6	71
9	Higher gene expression variability in the more aggressive subtype of chronic lymphocytic leukemia. Genome Medicine, 2015, 7, 8.	3 . 6	57
10	Personal Genome Project UK (PGP-UK): a research and citizen science hybrid project in support of personalized medicine. BMC Medical Genomics, 2018, 11, 108.	0.7	34
11	Genetic perturbation of PU.1 binding and chromatin looping at neutrophil enhancers associates with autoimmune disease. Nature Communications, 2021, 12, 2298.	5 . 8	32
12	Characterization of Transcriptional Changes in ERG Rearrangement-Positive Prostate Cancer Identifies the Regulation of Metabolic Sensors Such as Neuropeptide Y. PLoS ONE, 2013, 8, e55207.	1.1	32
13	Exaggerated IL-17A activity in human in vivo recall responses discriminates active tuberculosis from latent infection and cured disease. Science Translational Medicine, 2021, 13, .	5.8	27
14	Chemotherapy induces canalization of cell state in childhood B-cell precursor acute lymphoblastic leukemia. Nature Cancer, 2021, 2, 835-852.	5.7	25
15	The synthetic glucocorticoids prednisolone and dexamethasone regulate the same genes in acute lymphoblastic leukemia cells. BMC Genomics, 2014, 15, 662.	1.2	20
16	The epigenetic clock: a molecular crystal ball for human aging?. Aging, 2019, 11, 833-835.	1.4	19
17	Epigenetic variation taking center stage in immunological research. Epigenomics, 2017, 9, 375-378.	1.0	4
18	GenomeChronicler: The Personal Genome Project UK Genomic Report Generator Pipeline. Frontiers in Genetics, 2020, 11, 518644.	1.1	4