Huaming Chen

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/138610/publications.pdf

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		759055	1125617	
12	3,332	12	13	
papers	citations	h-index	g-index	
10	10	10	7020	
19	19	19	7039	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	Epigenomic Analysis of Multilineage Differentiation of Human Embryonic Stem Cells. Cell, 2013, 153, 1134-1148.	13.5	689
2	Human body epigenome maps reveal noncanonical DNA methylation variation. Nature, 2015, 523, 212-216.	13.7	605
3	Epigenomic Diversity in a Global Collection of Arabidopsis thaliana Accessions. Cell, 2016, 166, 492-505.	13.5	594
4	A transcription factor hierarchy defines an environmental stress response network. Science, 2016, 354, .	6.0	394
5	Temporal transcriptional response to ethylene gas drives growth hormone cross-regulation in Arabidopsis. ELife, 2013, 2, e00675.	2.8	379
6	Integrated multi-omics framework of the plant response to jasmonic acid. Nature Plants, 2020, 6, 290-302.	4.7	145
7	DNA methylation atlas of the mouse brain at single-cell resolution. Nature, 2021, 598, 120-128.	13.7	135
8	The complex architecture and epigenomic impact of plant T-DNA insertions. PLoS Genetics, 2019, 15, e1007819.	1.5	109
9	Spatiotemporal DNA methylome dynamics of the developing mouse fetus. Nature, 2020, 583, 752-759.	13.7	84
10	Mapping transcription factor interactome networks using HaloTag protein arrays. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E4238-47.	3.3	67
11	PHYTOCHROME-INTERACTING FACTORs trigger environmentally responsive chromatin dynamics in plants. Nature Genetics, 2021, 53, 955-961.	9.4	54
12	Epigenetic silencing of a multifunctional plant stress regulator. ELife, 2019, 8, .	2.8	28