Eric M Mendenhall

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5934633/publications.pdf

Version: 2024-02-01

21 papers 7,598 citations

471371 17 h-index 713332 21 g-index

25 all docs

25 docs citations

25 times ranked 11959 citing authors

#	Article	IF	CITATIONS
1	Genome-wide maps of chromatin state in pluripotent and lineage-committed cells. Nature, 2007, 448, 553-560.	13.7	3,733
2	Expanded encyclopaedias of DNA elements in the human and mouse genomes. Nature, 2020, 583, 699-710.	13.7	1,252
3	Genomewide Analysis of PRC1 and PRC2 Occupancy Identifies Two Classes of Bivalent Domains. PLoS Genetics, 2008, 4, e1000242.	1.5	878
4	GC-Rich Sequence Elements Recruit PRC2 in Mammalian ES Cells. PLoS Genetics, 2010, 6, e1001244.	1.5	368
5	Locus-specific editing of histone modifications at endogenous enhancers. Nature Biotechnology, 2013, 31, 1133-1136.	9.4	339
6	The molecular repertoire of the 'almighty' stem cell. Nature Reviews Molecular Cell Biology, 2005, 6, 726-737.	16.1	183
7	Perspectives on ENCODE. Nature, 2020, 583, 693-698.	13.7	123
8	CETCh-seq: CRISPR epitope tagging ChIP-seq of DNA-binding proteins. Genome Research, 2015, 25, 1581-1589.	2.4	122
9	Functional Analysis of Human Hematopoietic Stem Cell Gene Expression Using Zebrafish. PLoS Biology, 2005, 3, e254.	2.6	96
10	Occupancy maps of 208 chromatin-associated proteins in one human cell type. Nature, 2020, 583, 720-728.	13.7	90
11	Chromatin state maps: new technologies, new insights. Current Opinion in Genetics and Development, 2008, 18, 109-115.	1.5	77
12	Genome-Wide Reverse Genetics Framework to Identify Novel Functions of the Vertebrate Secretome. PLoS ONE, 2006, 1 , e104.	1.1	67
13	Tissue-specific SMARCA4 binding at active and repressed regulatory elements during embryogenesis. Genome Research, 2014, 24, 920-929.	2.4	63
14	A small molecule that induces translational readthrough of CFTR nonsense mutations by eRF1 depletion. Nature Communications, 2021, 12, 4358.	5.8	59
15	Htt CAG repeat expansion confers pleiotropic gains of mutant huntingtin function in chromatin regulation. Human Molecular Genetics, 2015, 24, 2442-2457.	1.4	53
16	Characterization of expanded intermediate cell mass in zebrafish chordin morphant embryos. Developmental Biology, 2005, 277, 235-254.	0.9	43
17	Identification and function of enhancers in the human genome. Human Molecular Genetics, 2016, 25, R190-R197.	1.4	26
18	Every transcription factor deserves its map: Scaling up epitope tagging of proteins to bypass antibody problems. BioEssays, 2016, 38, 801-811.	1.2	13

#	Article	IF	CITATIONS
19	DNA-protein interactions in high definition. Genome Biology, 2012, 13, 139.	13.9	2
20	Functional Analysis of the Differential Gene Expression Profile of Human HSC Using a Functional Genomics Screen in the Zebrafish Blood, 2004, 104, 136-136.	0.6	1
21	SPRY1 Is a Negative Regulator of Long-Term In Vivo Engraftment and Ex Vivo Expansion of Primitive Human Umbilical Cord Blood Cells Blood, 2005, 106, 1715-1715.	0.6	O