## Cynthia L Fisher

List of Publications by Year in descending order

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759233 1058476 1,382 14 12 14 citations h-index g-index papers 15 15 15 2379 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Jarid2 is a PRC2 component in embryonic stem cells required for multi-lineage differentiation and recruitment of PRC1 and RNA Polymerase II to developmental regulators. Nature Cell Biology, 2010, 12, 618-624.	10.3	274
2	The interaction of PRC2 with RNA or chromatin is mutually antagonistic. Genome Research, 2016, 26, 896-907.	5 <b>.</b> 5	191
3	Chromatin states in pluripotent, differentiated, and reprogrammed cells. Current Opinion in Genetics and Development, 2011, 21, 140-146.	3.3	145
4	Auxetic nuclei in embryonic stem cells exiting pluripotency. Nature Materials, 2014, 13, 638-644.	27.5	145
5	Loss-of-function Additional sex combs like 1 mutations disrupt hematopoiesis but do not cause severe myelodysplasia or leukemia. Blood, 2010, 115, 38-46.	1.4	141
6	Maintenance of gene expression patterns. Developmental Dynamics, 2005, 232, 633-655.	1.8	102
7	Characterization of ABCB9, an ATP Binding Cassette Protein Associated with Lysosomes. Journal of Biological Chemistry, 2000, 275, 23287-23294.	3.4	91
8	A human homolog of Additional sex combs, ADDITIONAL SEX COMBS-LIKE 1, maps to chromosome 20q11. Gene, 2003, 306, 115-126.	2.2	88
9	Characterization of Asxl1, a murine homolog of Additional sex combs, and analysis of the Asx-like gene family. Gene, 2006, 369, 109-118.	2.2	87
10	M-ABC2, a new human mitochondrial ATP-binding cassette membrane protein. FEBS Letters, 2000, 478, 89-94.	2.8	39
11	A microfluidic device for characterizing nuclear deformations. Lab on A Chip, 2017, 17, 805-813.	6.0	33
12	Jmjd2c/Kdm4c facilitates the assembly of essential enhancer-protein complexes at the onset of embryonic stem cell differentiation. Development (Cambridge), 2017, 144, 567-579.	2.5	24
13	IQCELL: A platform for predicting the effect of gene perturbations on developmental trajectories using single-cell RNA-seq data. PLoS Computational Biology, 2022, 18, e1009907.	3.2	13
14	An efficient method for generation of bi-allelic null mutant mouse embryonic stem cells and its application for investigating epigenetic modifiers. Nucleic Acids Research, 2017, 45, e174-e174.	14.5	7