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List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Ketogenic diet and chemotherapy combine to disrupt pancreatic cancer metabolism and growth. Med, 2022, 3, 119-136.e8.	4.4	31
2	Targeting LIF-mediated paracrine interaction for pancreatic cancer therapy and monitoring. Nature, 2019, 569, 131-135.	27.8	287
3	Cell cycle and p53 gate the direct conversion of human fibroblasts to dopaminergic neurons. Nature Communications, 2015, 6, 10100.	12.8	108
4	Genetic and Epigenetic Variations in iPSCs: Potential Causes and Implications for Application. Cell Stem Cell, 2013, 13, 149-159.	11.1	326
5	Embryonic stem cell and induced pluripotent stem cell: an epigenetic perspective. Cell Research, 2013, 23, 49-69.	12.0	152
6	Kdm2b promotes induced pluripotent stem cell generation by facilitating gene activation early inÂreprogramming. Nature Cell Biology, 2012, 14, 457-466.	10.3	166
7	Butyrate Promotes Induced Pluripotent Stem Cell Generation. Journal of Biological Chemistry, 2010, 285, 25516-25521.	3.4	123
8	A Conserved Central Region of Yeast Ada2 Regulates the Histone Acetyltransferase Activity of Gcn5 and Interacts with Phospholipids. Journal of Molecular Biology, 2008, 384, 743-755.	4.2	14
9	Generation of Insulin-secreting Islet-like Clusters from Human Skin Fibroblasts. Journal of Biological Chemistry, 2008, 283, 31601-31607.	3.4	313
10	Demethylation of Histone H3K36 and H3K9 by Rph1: a Vestige of an H3K9 Methylation System in Saccharomyces cerevisiae ?. Molecular and Cellular Biology, 2007, 27, 3951-3961.	2.3	79
11	The Saccharomyces cerevisiae Histone Demethylase Jhd1 Fine-Tunes the Distribution of H3K36me2. Molecular and Cellular Biology, 2007, 27, 5055-5065.	2.3	32
12	C-terminal processing of yeast Spt7 occurs in the absence of functional SAGA complex. BMC Biochemistry, 2007, 8, 16.	4.4	10
13	Yeast Jhd2p is a histone H3 Lys4 trimethyl demethylase. Nature Structural and Molecular Biology, 2007, 14, 243-245.	8.2	111
14	The role of histone ubiquitylation and deubiquitylation in gene expression as determined by the analysis of an HTB1 K123R Saccharomyces cerevisiae strain. Molecular Genetics and Genomics, 2007, 277, 491-506.	2.1	32