Yoshiyuki Seki

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/356786/publications.pdf

Version: 2024-02-01

		687363	888059	
18	2,156	13	17	
papers	citations	h-index	g-index	
18	18	18	2232	
all docs	docs citations	times ranked	citing authors	

#	Article	lF	Citations
1	PRDM14-CtBP1/2-PRC2 complex regulates transcriptional repression during transition from primed to na \tilde{A} -ve pluripotency. Journal of Cell Science, 2020, 133, .	2.0	10
2	Co-option of the PRDM14–CBFA2T complex from motor neurons to pluripotent cells during vertebrate evolution. Development (Cambridge), 2019, 146, .	2.5	9
3	PRDM14 Is a Unique Epigenetic Regulator Stabilizing Transcriptional Networks for Pluripotency. Frontiers in Cell and Developmental Biology, 2018, 6, 12.	3.7	30
4	PRDM14 Drives OCT3/4 Recruitment via Active Demethylation in the Transition from Primed to Naive Pluripotency. Stem Cell Reports, 2016, 7, 1072-1086.	4.8	31
5	PRDM14 maintains pluripotency of embryonic stem cells through TET-mediated active DNA demethylation. Biochemical and Biophysical Research Communications, 2015, 466, 138-145.	2.1	18
6	PRDM14 promotes active DNA demethylation through the Ten-eleven translocation (TET)-mediated base excision repair pathway in embryonic stem cells. Development (Cambridge), 2014, 141, 269-280.	2.5	113
7	Serum-mediated transgenerational effects on sperm: Evidence for lamarckian inheritance?. Hepatology, 2013, 57, 1663-1665.	7.3	3
8	Epigenetic Reprogramming in Primordial Germ Cells in Mice. Journal of Mammalian Ova Research, 2013, 30, 95-100.	0.1	0
9	A replication-dependent passive mechanism modulates DNA demethylation in mouse primordial germ cells. Development (Cambridge), 2013, 140, 2892-2903.	2.5	71
10	Locus- and domain-dependent control of DNA methylation at mouse B1 retrotransposons during male germ cell development. Genome Research, 2011, 21, 2058-2066.	5.5	50
11	Critical function of Prdm14 for the establishment of the germ cell lineage in mice. Nature Genetics, 2008, 40, 1016-1022.	21.4	516
12	Specification of the germ cell lineage in mice: A process orchestrated by the PR-domain proteins, Blimp1 and Prdm14. Cell Cycle, 2008, 7, 3514-3518.	2.6	84
13	Cellular dynamics associated with the genome-wide epigenetic reprogramming in migrating primordial germ cells in mice. Development (Cambridge), 2007, 134, 2627-2638.	2.5	388
14	Gene Expression Dynamics During Germline Specification in Mice Identified by Quantitative Single-Cell Gene Expression Profiling 1. Biology of Reproduction, 2006, 75, 705-716.	2.7	256
15	Extensive and orderly reprogramming of genome-wide chromatin modifications associated with specification and early development of germ cells in mice. Developmental Biology, 2005, 278, 440-458.	2.0	484
16	Myeloid Elf-1-like Factor, an ETS Transcription Factor, Up-regulates Lysozyme Transcription in Epithelial Cells through Interaction with Promyelocytic Leukemia Protein. Journal of Biological Chemistry, 2004, 279, 19091-19098.	3.4	24
17	ETS2 is involved in protein kinase C-activated expression of granulocyte–macrophage colony-stimulating factor in human non-small lung carcinoma cell line, A549. Biochemical and Biophysical Research Communications, 2003, 303, 190-195.	2.1	7
18	The ETS transcription factor MEF is a candidate tumor suppressor gene on the X chromosome. Cancer Research, 2002, 62, 6579-86.	0.9	62