

Baoming Qin

List of Publications by Year in descending order

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Version: 2024-02-01

40
papers

3,938
citations

331670

21
h-index

302126

39
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all docs

42
docs citations

42
times ranked

5928
citing authors

#	ARTICLE	IF	CITATIONS
1	A Mesenchymal-to-Epithelial Transition Initiates and Is Required for the Nuclear Reprogramming of Mouse Fibroblasts. <i>Cell Stem Cell</i> , 2010, 7, 51-63.	11.1	1,038
2	Vitamin C Enhances the Generation of Mouse and Human Induced Pluripotent Stem Cells. <i>Cell Stem Cell</i> , 2010, 6, 71-79.	11.1	878
3	The Histone Demethylases Jhdm1a/1b Enhance Somatic Cell Reprogramming in a Vitamin-C-Dependent Manner. <i>Cell Stem Cell</i> , 2011, 9, 575-587.	11.1	407
4	MicroRNA Cluster 302a-367 Enhances Somatic Cell Reprogramming by Accelerating a Mesenchymal-to-Epithelial Transition. <i>Journal of Biological Chemistry</i> , 2011, 286, 17359-17364.	3.4	231
5	Capturing the interactome of newly transcribed RNA. <i>Nature Methods</i> , 2018, 15, 213-220.	19.0	170
6	The p53-induced lincRNA-p21 derails somatic cell reprogramming by sustaining H3K9me3 and CpG methylation at pluripotency gene promoters. <i>Cell Research</i> , 2015, 25, 80-92.	12.0	160
7	Nuclear localization of the phosphatidylserine receptor protein via multiple nuclear localization signals. <i>Experimental Cell Research</i> , 2004, 293, 154-163.	2.6	103
8	Rolling back human pluripotent stem cells to an eight-cell embryo-like stage. <i>Nature</i> , 2022, 605, 315-324.	27.8	87
9	Autophagy and mTORC1 regulate the stochastic phase of somatic cell reprogramming. <i>Nature Cell Biology</i> , 2015, 17, 715-725.	10.3	81
10	Cell transcriptomic atlas of the non-human primate <i>Macaca fascicularis</i> . <i>Nature</i> , 2022, 604, 723-731.	27.8	81
11	NCoR/SMRT co-repressors cooperate with c-MYC to create an epigenetic barrier to somatic cell reprogramming. <i>Nature Cell Biology</i> , 2018, 20, 400-412.	10.3	64
12	Sorting Nexin 10 Induces Giant Vacuoles in Mammalian Cells. <i>Journal of Biological Chemistry</i> , 2006, 281, 36891-36896.	3.4	62
13	The mesenchymal-to-epithelial transition in somatic cell reprogramming. <i>Current Opinion in Genetics and Development</i> , 2012, 22, 423-428.	3.3	62
14	Transcriptional Pause Release Is a Rate-Limiting Step for Somatic Cell Reprogramming. <i>Cell Stem Cell</i> , 2014, 15, 574-588.	11.1	60
15	Identification of EGFR kinase domain mutations among lung cancer patients in China: implication for targeted cancer therapy. <i>Cell Research</i> , 2005, 15, 212-217.	12.0	58
16	Boosting ATM activity alleviates aging and extends lifespan in a mouse model of progeria. <i>ELife</i> , 2018, 7, .	6.0	54
17	The Hypoxia-inducible Factor Renders Cancer Cells More Sensitive to Vitamin C-induced Toxicity. <i>Journal of Biological Chemistry</i> , 2014, 289, 3339-3351.	3.4	45
18	MicroRNAs in somatic cell reprogramming. <i>Current Opinion in Cell Biology</i> , 2013, 25, 208-214.	5.4	43

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19	Transcriptional Control of Somatic Cell Reprogramming. <i>Trends in Cell Biology</i> , 2016, 26, 272-288.	7.9	35
20	Generation of Human Liver Chimeric Mice with Hepatocytes from Familial Hypercholesterolemia Induced Pluripotent Stem Cells. <i>Stem Cell Reports</i> , 2017, 8, 605-618.	4.8	27
21	Profiling the long noncoding RNA interaction network in the regulatory elements of target genes by chromatin in situ reverse transcription sequencing. <i>Genome Research</i> , 2019, 29, 1521-1532.	5.5	27
22	JMJD3 acts in tandem with KLF4 to facilitate reprogramming to pluripotency. <i>Nature Communications</i> , 2020, 11, 5061.	12.8	24
23	Roles of small molecules in somatic cell reprogramming. <i>Acta Pharmacologica Sinica</i> , 2013, 34, 719-724.	6.1	18
24	Fumarylacetoacetate Hydrolase Knock-out Rabbit Model for Hereditary Tyrosinemia Type 1. <i>Journal of Biological Chemistry</i> , 2017, 292, 4755-4763.	3.4	15
25	Nuclear-cytoplasmic shuttling of class IIa histone deacetylases regulates somatic cell reprogramming. <i>Cell Regeneration</i> , 2019, 8, 21-29.	2.6	13
26	mTORC1 axis regulates mitochondrial remodeling during reprogramming. <i>FEBS Journal</i> , 2020, 287, 108-121.	4.7	13
27	Class IIa Histone Deacetylases and Myocyte Enhancer Factor 2 Proteins Regulate the Mesenchymal-to-Epithelial Transition of Somatic Cell Reprogramming. <i>Journal of Biological Chemistry</i> , 2013, 288, 12022-12031.	3.4	11
28	β-Catenin safeguards the ground state of mouse pluripotency by strengthening the robustness of the transcriptional apparatus. <i>Science Advances</i> , 2020, 6, eaba1593.	10.3	10
29	PML2-mediated threadlike nuclear bodies mark late senescence in Hutchinson Gilford progeria syndrome. <i>Aging Cell</i> , 2020, 19, e13147.	6.7	10
30	Role of Long Non-coding RNAs in Reprogramming to Induced Pluripotency. <i>Genomics, Proteomics and Bioinformatics</i> , 2020, 18, 16-25.	6.9	10
31	The mTORC1-eIF4F axis controls paused pluripotency. <i>EMBO Reports</i> , 2022, 23, e53081.	4.5	8
32	Exonuclease combinations reduce noises in 3D genomics technologies. <i>Nucleic Acids Research</i> , 2020, 48, e44-e44.	14.5	6
33	PX domain and CD domain play different roles in localization and vacuolation of Sorting Nexin 10. <i>Science Bulletin</i> , 2009, 54, 3965-3971.	1.7	5
34	Capture of the newly transcribed RNA interactome using click chemistry. <i>Nature Protocols</i> , 2021, 16, 5193-5219.	12.0	5
35	A Familial Hypercholesterolemia Human Liver Chimeric Mouse Model Using Induced Pluripotent Stem Cell-derived Hepatocytes. <i>Journal of Visualized Experiments</i> , 2018, , .	0.3	3
36	Generation of two LRRK2 homozygous knockout human induced pluripotent stem cell lines using CRISPR/Cas9. <i>Stem Cell Research</i> , 2020, 45, 101804.	0.7	3

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37	Generation of Rybp homozygous knockout murine ES cell line GIBHe001-A-1 by using CRISPR/Cas9 technology. Stem Cell Research, 2019, 41, 101638.	0.7	2
38	Generation of an induced pluripotent stem cell line (GIBHi003-A) from a Parkinson's disease patient with mutant PINK1 (p. I368N). Stem Cell Research, 2019, 41, 101607.	0.7	1
39	Reprogramming mouse embryonic fibroblasts using different systems. Protocol Exchange, 0, , .	0.3	1
40	Generation of a PARK2 homozygous knockout induced pluripotent stem cell line (GIBHi002-A-1) with two common isoforms abolished. Stem Cell Research, 2019, 41, 101602.	0.7	0