

Yawei Gao

List of Publications by Year in descending order

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Version: 2024-02-01

32
papers

3,037
citations

331538

21
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395590

33
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38
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docs citations

38
times ranked

4308
citing authors

#	ARTICLE	IF	CITATIONS
1	Dynamic nucleosome organization after fertilization reveals regulatory factors for mouse zygotic genome activation. <i>Cell Research</i> , 2022, 32, 801-813.	5.7	14
2	Epigenetic regulation of cell fate transition: learning from early embryo development and somatic cell reprogramming. <i>Biology of Reproduction</i> , 2022, 107, 183-195.	1.2	7
3	FTO mediates LINE1 m ⁶ A demethylation and chromatin regulation in mESCs and mouse development. <i>Science</i> , 2022, 376, 968-973.	6.0	97
4	N6-methyladenosine regulates maternal RNA maintenance in oocytes and timely RNA decay during mouse maternal-to-zygotic transition. <i>Nature Cell Biology</i> , 2022, 24, 917-927.	4.6	28
5	Unique Patterns of H3K4me3 and H3K27me3 in 2-Cell-like Embryonic Stem Cells. <i>Stem Cell Reports</i> , 2021, 16, 458-469.	2.3	18
6	Nuclear m6A reader YTHDC1 regulates the scaffold function of LINE1 RNA in mouse ESCs and early embryos. <i>Protein and Cell</i> , 2021, 12, 455-474.	4.8	84
7	Dcaf11 activates Zscan4-mediated alternative telomere lengthening in early embryos and embryonic stem cells. <i>Cell Stem Cell</i> , 2021, 28, 732-747.e9.	5.2	30
8	Stabilization of ERK-Phosphorylated METTL3 by USP5 Increases m6A Methylation. <i>Molecular Cell</i> , 2020, 80, 633-647.e7.	4.5	83
9	Reprogramming competence of OCT factors is determined by transactivation domains. <i>Science Advances</i> , 2020, 6, .	4.7	25
10	N6-Deoxyadenosine Methylation in Mammalian Mitochondrial DNA. <i>Molecular Cell</i> , 2020, 78, 382-395.e8.	4.5	156
11	DNA 5-Methylcytosine-Specific Amplification and Sequencing. <i>Journal of the American Chemical Society</i> , 2020, 142, 4539-4543.	6.6	13
12	m ⁶ A-methyladenosine of chromosome-associated regulatory RNA regulates chromatin state and transcription. <i>Science</i> , 2020, 367, 580-586.	6.0	406
13	Distinct H3K9me3 and DNA methylation modifications during mouse spermatogenesis. <i>Journal of Biological Chemistry</i> , 2019, 294, 18714-18725.	1.6	38
14	Jump-seq: Genome-Wide Capture and Amplification of 5-Hydroxymethylcytosine Sites. <i>Journal of the American Chemical Society</i> , 2019, 141, 8694-8697.	6.6	26
15	Nuclear Exosome Targeting Complex Core Factor Zcchc8 Regulates the Degradation of LINE1 RNA in Early Embryos and Embryonic Stem Cells. <i>Cell Reports</i> , 2019, 29, 2461-2472.e6.	2.9	28
16	Esrrb plays important roles in maintaining self-renewal of trophoblast stem cells (TSCs) and reprogramming somatic cells to induced TSCs. <i>Journal of Molecular Cell Biology</i> , 2019, 11, 463-473.	1.5	19
17	Reprogramming of H3K9me3-dependent heterochromatin during mammalian embryo development. <i>Nature Cell Biology</i> , 2018, 20, 620-631.	4.6	292
18	Inhibition of Aberrant DNA Re-methylation Improves Post-implantation Development of Somatic Cell Nuclear Transfer Embryos. <i>Cell Stem Cell</i> , 2018, 23, 426-435.e5.	5.2	72

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19	Loss of YTHDF2-mediated m6A-dependent mRNA clearance facilitates hematopoietic stem cell regeneration. <i>Cell Research</i> , 2018, 28, 1035-1038.	5.7	95
20	<scp>DCAF</scp> 13 promotes pluripotency by negatively regulating <scp>SUV</scp> 39H1 stability during early embryonic development. <i>EMBO Journal</i> , 2018, 37, .	3.5	39
21	Baf60b-mediated ATM-p53 activation blocks cell identity conversion by sensing chromatin opening. <i>Cell Research</i> , 2017, 27, 642-656.	5.7	18
22	Maternal Sall4 Is Indispensable for Epigenetic Maturation of Mouse Oocytes. <i>Journal of Biological Chemistry</i> , 2017, 292, 1798-1807.	1.6	37
23	Direct induction of neural progenitor cells transiently passes through a partially reprogrammed state. <i>Biomaterials</i> , 2017, 119, 53-67.	5.7	10
24	Protein Expression Landscape of Mouse Embryos during Pre-implantation Development. <i>Cell Reports</i> , 2017, 21, 3957-3969.	2.9	135
25	High throughput sequencing identifies an imprinted gene, Grb10, associated with the pluripotency state in nuclear transfer embryonic stem cells. <i>Oncotarget</i> , 2017, 8, 47344-47355.	0.8	5
26	Identification of key factors conquering developmental arrest of somatic cell cloned embryos by combining embryo biopsy and single-cell sequencing. <i>Cell Discovery</i> , 2016, 2, 16010.	3.1	165
27	SIRT6 Controls Hematopoietic Stem Cell Homeostasis through Epigenetic Regulation of Wnt Signaling. <i>Cell Stem Cell</i> , 2016, 18, 495-507.	5.2	117
28	Distinct features of H3K4me3 and H3K27me3 chromatin domains in pre-implantation embryos. <i>Nature</i> , 2016, 537, 558-562.	13.7	538
29	Hierarchical Oct4 Binding in Concert with Primed Epigenetic Rearrangements during Somatic Cell Reprogramming. <i>Cell Reports</i> , 2016, 14, 1540-1554.	2.9	74
30	The Combination of Tet1 with Oct4 Generates High-Quality Mouse-Induced Pluripotent Stem Cells. <i>Stem Cells</i> , 2015, 33, 686-698.	1.4	39
31	Quality Control: H2A.X Links to Better iPSCs. <i>Cell Stem Cell</i> , 2014, 15, 259-260.	5.2	3
32	Replacement of Oct4 by Tet1 during iPSC Induction Reveals an Important Role of DNA Methylation and Hydroxymethylation in Reprogramming. <i>Cell Stem Cell</i> , 2013, 12, 453-469.	5.2	321