Li-Ying Yan

List of Publications by Year in descending order

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Ι Ι-ΥΙΝΟ ΥΛΝ

#	Article	IF	CITATIONS
1	Single-cell RNA-Seq profiling of human preimplantation embryos and embryonic stem cells. Nature Structural and Molecular Biology, 2013, 20, 1131-1139.	8.2	1,416
2	The DNA methylation landscape of human early embryos. Nature, 2014, 511, 606-610.	27.8	787
3	A single-cell RNA-seq survey of the developmental landscape of the human prefrontal cortex. Nature, 2018, 555, 524-528.	27.8	551
4	The Transcriptome and DNA Methylome Landscapes of Human Primordial Germ Cells. Cell, 2015, 161, 1437-1452.	28.9	500
5	Single-Cell RNA-Seq Analysis Maps Development of Human Germline Cells and Gonadal Niche Interactions. Cell Stem Cell, 2017, 20, 858-873.e4.	11.1	376
6	Single-Cell Transcriptome Analysis Maps the Developmental Track of the Human Heart. Cell Reports, 2019, 26, 1934-1950.e5.	6.4	355
7	Single-Cell RNA Sequencing Analysis Reveals Sequential Cell Fate Transition during Human Spermatogenesis. Cell Stem Cell, 2018, 23, 599-614.e4.	11.1	309
8	Genome Analyses of Single Human Oocytes. Cell, 2013, 155, 1492-1506.	28.9	279
9	Transcriptome Landscape of Human Folliculogenesis Reveals Oocyte and Granulosa Cell Interactions. Molecular Cell, 2018, 72, 1021-1034.e4.	9.7	262
10	Single-cell multiomics sequencing and analyses of human colorectal cancer. Science, 2018, 362, 1060-1063.	12.6	256
11	Single-cell DNA methylome sequencing of human preimplantation embryos. Nature Genetics, 2018, 50, 12-19.	21.4	248
12	Reconstituting the transcriptome and DNA methylome landscapes of human implantation. Nature, 2019, 572, 660-664.	27.8	207
13	Spatial transcriptomic survey of human embryonic cerebral cortex by single-cell RNA-seq analysis. Cell Research, 2018, 28, 730-745.	12.0	179
14	Single-cell multi-omics sequencing of human early embryos. Nature Cell Biology, 2018, 20, 847-858.	10.3	142
15	Tracing the expression of circular RNAs in human pre-implantation embryos. Genome Biology, 2016, 17, 130.	8.8	140
16	Tracing the temporal-spatial transcriptome landscapes of the human fetal digestive tract using single-cell RNA-sequencing. Nature Cell Biology, 2018, 20, 721-734.	10.3	125
17	Live births after simultaneous avoidance of monogenic diseases and chromosome abnormality by next-generation sequencing with linkage analyses. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 15964-15969.	7.1	115
18	Oocyte-expressed yes-associated protein is a key activator of the early zygotic genome in mouse. Cell Research, 2016, 26, 275-287.	12.0	108

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#	Article	IF	CITATIONS
19	Dissecting the transcriptome landscape of the human fetal neural retina and retinal pigment epithelium by single-cell RNA-seq analysis. PLoS Biology, 2019, 17, e3000365.	5.6	108
20	DNA methylation and chromatin accessibility profiling of mouse and human fetal germ cells. Cell Research, 2017, 27, 165-183.	12.0	102
21	Single-cell transcriptome analysis of the novel coronavirus (SARS-CoV-2) associated gene ACE2 expression in normal and non-obstructive azoospermia (NOA) human male testes. Science China Life Sciences, 2020, 63, 1006-1015.	4.9	96
22	Single-cell transcriptomics identifies divergent developmental lineage trajectories during human pituitary development. Nature Communications, 2020, 11, 5275.	12.8	79
23	Single-cell multiomics sequencing reveals the functional regulatory landscape of early embryos. Nature Communications, 2021, 12, 1247.	12.8	79
24	Identification of a human subcortical maternal complex. Molecular Human Reproduction, 2015, 21, 320-329.	2.8	75
25	Bioinspired <scp>l</scp> -Proline Oligomers for the Cryopreservation of Oocytes <i>via</i> Controlling Ice Growth. ACS Applied Materials & Interfaces, 2020, 12, 18352-18362.	8.0	52
26	Effects of oocyte vitrification on histone modifications. Reproduction, Fertility and Development, 2010, 22, 920.	0.4	51
27	Human single follicle growth <i>in vitro</i> from cryopreserved ovarian tissue after slow freezing or vitrification. Human Reproduction, 2016, 31, 763-773.	0.9	50
28	Mesenchymal stem cell-derived angiogenin promotes primodial follicle survival and angiogenesis in transplanted human ovarian tissue. Reproductive Biology and Endocrinology, 2017, 15, 18.	3.3	50
29	Epigenomic Landscape of Human Fetal Brain, Heart, and Liver. Journal of Biological Chemistry, 2016, 291, 4386-4398.	3.4	45
30	Validation of a next-generation sequencing–based protocol for 24-chromosome aneuploidy screening of blastocysts. Fertility and Sterility, 2016, 105, 1532-1536.	1.0	41
31	Testosterone Represses Estrogen Signaling by Upregulating miR-22. Hypertension, 2017, 69, 721-730.	2.7	41
32	Integrated multi-omics reveal epigenomic disturbance of assisted reproductive technologies in human offspring. EBioMedicine, 2020, 61, 103076.	6.1	41
33	High-fat diets exaggerate endocrine and metabolic phenotypes in a rat model of DHEA-induced PCOS. Reproduction, 2016, 151, 431-441.	2.6	37
34	Re-analysis of aneuploidy blastocysts with an inner cell mass and different regional trophectoderm cells. Journal of Assisted Reproduction and Genetics, 2017, 34, 487-493.	2.5	37
35	L-proline: a highly effective cryoprotectant for mouse oocyte vitrification. Scientific Reports, 2016, 6, 26326.	3.3	36
36	Validation of multiple annealing and looping-based amplification cycle sequencing for 24-chromosome aneuploidy screening of cleavage-stage embryos. Fertility and Sterility, 2014, 102, 1685-1691.	1.0	31

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37	Epigenetic Regulation and Risk Factors During the Development of Human Gametes and Early Embryos. Annual Review of Genomics and Human Genetics, 2019, 20, 21-40.	6.2	29
38	Endometrial miR-543 Is Downregulated During the Implantation Window in Women With Endometriosis-Related Infertility. Reproductive Sciences, 2019, 26, 900-908.	2.5	29
39	Mesenchymal Stem Cells Facilitate In Vitro Development of Human Preantral Follicle. Reproductive Sciences, 2015, 22, 1367-1376.	2.5	28
40	Dissecting the epigenomic dynamics of human fetal germ cell development at single-cell resolution. Cell Research, 2021, 31, 463-477.	12.0	28
41	Clinical applications of MARSALA for preimplantation genetic diagnosis of spinal muscular atrophy. Journal of Genetics and Genomics, 2016, 43, 541-547.	3.9	27
42	SARS-CoV-2 Entry Factors: ACE2 and TMPRSS2 Are Expressed in Peri-Implantation Embryos and the Maternal–Fetal Interface. Engineering, 2020, 6, 1162-1169.	6.7	27
43	A novel homozygous mutation of phospholipase C zeta leading to defective human oocyte activation and fertilization failure. Human Reproduction, 2020, 35, 977-985.	0.9	27
44	Trioâ€wholeâ€exome sequencing and preimplantation genetic diagnosis for unexplained recurrent fetal malformations. Human Mutation, 2020, 41, 432-448.	2.5	26
45	Activation of hedgehog signaling and its association with cisplatin resistance in ovarian epithelial tumors. Oncology Letters, 2018, 15, 5569-5576.	1.8	24
46	Effects of vitrification and cryostorage duration on single-cell RNA-Seq profiling of vitrified-thawed human metaphase II oocytes. Frontiers of Medicine, 2021, 15, 144-154.	3.4	23
47	NAT10-mediated <i>N</i> 4-acetylcytidine modification is required for meiosis entry and progression in male germ cells. Nucleic Acids Research, 2022, 50, 10896-10913.	14.5	20
48	Age-related changes in human conventional semen parameters and sperm chromatin structure assay-defined sperm DNA/chromatin integrity. Reproductive BioMedicine Online, 2021, 42, 973-982.	2.4	18
49	DevOmics: an integrated multi-omics database of human and mouse early embryo. Briefings in Bioinformatics, 2021, 22, .	6.5	16
50	Bone mesenchymal stem cells improve pregnancy outcome by inducing maternal tolerance to the allogeneic fetus in abortion-prone matings in mouse. Placenta, 2016, 47, 29-36.	1.5	15
51	Advances in preimplantation genetic diagnosis/screening. Science China Life Sciences, 2014, 57, 665-671.	4.9	13
52	Cryobiological Characteristics of L-proline in Mammalian Oocyte Cryopreservation. Chinese Medical Journal, 2016, 129, 1963-1968.	2.3	13
53	Poor intracytoplasmic sperm injection outcome in infertile males with azoospermia factor c microdeletions. Fertility and Sterility, 2021, 116, 96-104.	1.0	13
54	The â€~normal' range of FMR1 triple CGG repeats may be associated with primary ovarian insufficiency in China. Reproductive BioMedicine Online, 2017, 34, 175-180.	2.4	12

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55	Loss of CEP70 function affects acrosome biogenesis and flagella formation during spermiogenesis. Cell Death and Disease, 2021, 12, 478.	6.3	12
56	Lipid Metabolism Was Associated With Oocyte in vitro Maturation in Women With Polycystic Ovarian Syndrome Undergoing Unstimulated Natural Cycle. Frontiers in Cell and Developmental Biology, 2021, 9, 719173.	3.7	12
57	Fractalkine restores the decreased expression of StAR and progesterone in granulosa cells from patients with polycystic ovary syndrome. Scientific Reports, 2016, 6, 26205.	3.3	10
58	TRAIP is involved in chromosome alignment and SAC regulation in mouse oocyte meiosis. Scientific Reports, 2016, 6, 29735.	3.3	10
59	The present and future of whole-exome sequencing in studying and treating human reproductive disorders. Journal of Genetics and Genomics, 2018, 45, 517-525.	3.9	10
60	Maternal and neonatal outcomes following blastocyst biopsy for PGT in single vitrified–warmed embryo transfer cycles. Reproductive BioMedicine Online, 2022, 44, 151-162.	2.4	10
61	A strategy using <scp>SNP</scp> linkage analysis for monogenic diseases <scp>PGD</scp> combined with <scp>HLA</scp> typing. Clinical Genetics, 2020, 98, 138-146.	2.0	9
62	Clinical application of an NGS-based method in the preimplantation genetic testing for Duchenne muscular dystrophy. Journal of Assisted Reproduction and Genetics, 2021, 38, 1979-1986.	2.5	9
63	Exploring the role of autophagy during early human embryonic development through single-cell transcriptome and methylome analyses. Science China Life Sciences, 2022, 65, 940-952.	4.9	9
64	scHaplotyper: haplotype construction and visualization for genetic diagnosis using single cell DNA sequencing data. BMC Bioinformatics, 2020, 21, 41.	2.6	9
65	Dissecting Human Gonadal Cell Lineage Specification and Sex Determination Using A Single-cell RNA-seq Approach. Genomics, Proteomics and Bioinformatics, 2022, 20, 223-245.	6.9	9
66	5-Formylcytosine landscapes of human preimplantation embryos at single-cell resolution. PLoS Biology, 2020, 18, e3000799.	5.6	8
67	The methylome of a human polar body reflects that of its sibling oocyte and its aberrance may indicate poor embryo development. Human Reproduction, 2021, 36, 318-330.	0.9	8
68	Selective impairment in glycogen synthase kinase-3 and mitogen-activated protein kinase phosphorylation: comparisons with the hyperandrogenic and the hyperinsulinemic rats. Fertility and Sterility, 2009, 92, 1447-1455.	1.0	7
69	ART do not increase the risk of Y-chromosome microdeletion in 19 candidate genes at AZF regions. Reproduction, Fertility and Development, 2014, 26, 778.	0.4	7
70	Identifying normal embryos from reciprocal translocation carriers by whole chromosome haplotyping. Journal of Genetics and Genomics, 2018, 45, 505-508.	3.9	7
71	Genetic testing and PGD for unexplained recurrent fetal malformations with MAGEL2 gene mutation. Science China Life Sciences, 2019, 62, 886-894.	4.9	6
72	Novel PGD strategy based on single sperm linkage analysis for carriers of single gene pathogenic variant and chromosome reciprocal translocation. Journal of Assisted Reproduction and Genetics, 2020, 37, 1239-1250.	2.5	6

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73	A comprehensive PGT-M strategy for ADPKD patients with de novo PKD1 mutations using affected embryo or gametes as proband. Journal of Assisted Reproduction and Genetics, 2021, 38, 2425-2434.	2.5	6
74	Epigenetic consequences of hormonal interactions between oppositeâ€sex twin fetuses. Clinical and Translational Medicine, 2020, 10, e234.	4.0	5
75	Risk of miscarriage in women with endometriosis undergoing IVF fresh cycles: a retrospective cohort study. Reproductive Biology and Endocrinology, 2019, 17, 21.	3.3	4
76	Meiotic chromatid recombination and segregation assessed with human single cell genome sequencing data. Journal of Medical Genetics, 2019, 56, 156-163.	3.2	4
77	Effects of oocyte vitrification on the behaviors and physiological indexes of aged first filial generation mice. Cryobiology, 2020, 95, 20-28.	0.7	4
78	Transcriptome analysis of PCOS arrested 2-cell embryos. Cell Cycle, 2018, 17, 1007-1013.	2.6	3
79	Genetic analysis and preimplantation genetic diagnosis of Chinese Marfan syndrome patients. Journal of Genetics and Genomics, 2019, 46, 319-323.	3.9	3
80	Effect of serum 25â€hydroxyvitamin D levels on sperm quality and assisted reproductive technology outcomes for men of infertile Chinese couples. Andrology, 2020, 8, 1277-1286.	3.5	3
81	OUP accepted manuscript. Molecular Human Reproduction, 2021, , .	2.8	3
82	Effect of vitrification at the germinal vesicle stage on the global methylation status in mouse oocytes subsequently matured in vitro. Chinese Medical Journal, 2014, 127, 4019-24.	2.3	3
83	Application of three-dimensional fluorescence in situ hybridization to human preimplantation genetic diagnosis. Fertility and Sterility, 2009, 92, 1492-1495.	1.0	2
84	Gonadotropin-Mediated Dynamic Alterations During Bovine Oocyte Maturation In Vitro1. Biology of Reproduction, 2014, 91, 44.	2.7	2
85	The function of Nucleoporin 37 on mouse oocyte maturation and preimplantation embryo development. Journal of Assisted Reproduction and Genetics, 2022, 39, 107.	2.5	2
86	A novel pathogenic mutation in FBN2 associated with congenital contractural arachnodactyly for preimplantation genetic diagnosis. Journal of Genetics and Genomics, 2020, 47, 281-284.	3.9	1
87	OP-IVM: Combining In vitro Maturation after Oocyte Retrieval with Gynecological Surgery. Journal of Visualized Experiments, 2021, , .	0.3	1
88	Application of next-generation sequencing to preimplantation genetic testing for recurrent hydatidiform mole patients. Journal of Assisted Reproduction and Genetics, 2021, 38, 2881-2891.	2.5	0