## Yongjie Wan

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/883466/publications.pdf

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29	505	759055	677027
papers	citations	h-index	22 g-index
30	30	30	496
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Expression pattern of alkB homolog 5 in goat testis and its role in spermatogonial stem cells. Cell and Tissue Research, 2022, 387, 131-142.	1.5	3
2	The function of the m6A methyltransferase METTL3 in goat early embryo development under hypoxic and normoxic conditions. Theriogenology, 2022, 177, 140-150.	0.9	5
3	YBX1 mediates alternative splicing and maternal mRNA decay during pre-implantation development. Cell and Bioscience, 2022, 12, 12.	2.1	11
4	Comprehensive Transcriptome Analysis of mRNA Expression Patterns of Early Embryo Development in Goat under Hypoxic and Normoxic Conditions. Biology, 2021, 10, 381.	1.3	5
5	Long non-coding RNA lnc_3712 impedes nuclear reprogramming via repressing Kdm5b. Molecular Therapy - Nucleic Acids, 2021, 24, 54-66.	2.3	9
6	Characterization of transcriptional activity during ZGA in mammalian SCNT embryo. Biology of Reproduction, 2021, 105, 905-917.	1,2	8
7	Melatonin alleviated oxidative stress induced by energy restriction on sheep Leydig cells through Sirt1/Sod2 pathway. Theriogenology, 2021, 173, 83-92.	0.9	7
8	Inhibition of lysine-specific histone demethylase 1A results in meiotic aberration during oocyte maturation inÂvitro in goats. Theriogenology, 2020, 143, 168-178.	0.9	16
9	Exchanges of histone methylation and variants during mouse zygotic genome activation. Zygote, 2020, 28, 51-58.	0.5	7
10	YTHDF2 Regulates Maternal Transcriptome Degradation and Embryo Development in Goat. Frontiers in Cell and Developmental Biology, 2020, 8, 580367.	1.8	16
11	DNA methylation dynamics during zygotic genome activation in goat. Theriogenology, 2020, 156, 144-154.	0.9	14
12	Locus-Specific Regulation of <i>Xist</i> Expression Using the CRISPR-Cas9-Based System. DNA and Cell Biology, 2020, 39, 572-578.	0.9	2
13	Aberrant DNA and histone methylation during zygotic genome activation in goat cloned embryos. Theriogenology, 2020, 148, 27-36.	0.9	33
14	The Expression Pattern of p32 in Sheep Muscle and Its Role in Differentiation, Cell Proliferation, and Apoptosis of Myoblasts. International Journal of Molecular Sciences, 2019, 20, 5161.	1.8	5
15	Effect of Curcumin Supplement in Summer Diet on Blood Metabolites, Antioxidant Status, Immune Response, and Testicular Gene Expression in Hu Sheep. Animals, 2019, 9, 720.	1.0	31
16	Highly methylated Xist in SCNT embryos was retained in deceased cloned female goats. Reproduction, Fertility and Development, 2019, 31, 855.	0.1	12
17	Efficient generation of CLPG1 â€edited rabbits using the CRISPR/Cas9 system. Reproduction in Domestic Animals, 2019, 54, 538-544.	0.6	6
18	Comprehensive analysis of long noncoding RNA and mRNA expression patterns in sheep testicular maturationâ€. Biology of Reproduction, 2018, 99, 650-661.	1.2	47

#	Article	IF	CITATIONS
19	Analysis of <i>H19/lgf2</i> Methylation Status in the Sperm of Cloned Goats and Their Offspring. Cellular Reprogramming, 2018, 20, 66-75.	0.5	3
20	Long noncoding RNAs exchange during zygotic genome activation in goatâ€. Biology of Reproduction, 2018, 99, 707-717.	1.2	48
21	Epigenetic Status of <i>H19</i> - <i>Igf2</i> Imprinted Genes and Loss of 5-Hydroxymethylcytosine in the Brain of Cloned Goats. Cellular Reprogramming, 2017, 19, 199-207.	0.5	4
22	Vitamin D receptor expression and potential role of vitamin D on cell proliferation and steroidogenesis in goat ovarian granulosa cells. Theriogenology, 2017, 102, 162-173.	0.9	53
23	MBD1 and MeCP2 expression in embryos and placentas from transgenic cloned goats. Zygote, 2017, 25, 462-471.	0.5	1
24	Generation of beta-lactoglobulin knock-out goats using CRISPR/Cas9. PLoS ONE, 2017, 12, e0186056.	1.1	47
25	Using cysteine/cystine to overcome oxidative stress in goat oocytes and embryos cultured in vitro. Molecular Medicine Reports, 2016, 14, 1219-1226.	1.1	20
26	Generation and evaluation of Myostatin knock-out rabbits and goats using CRISPR/Cas9 system. Scientific Reports, 2016, 6, 29855.	1.6	71
27	Abnormal expression of DNA methyltransferases and genomic imprinting in cloned goat fibroblasts. Cell Biology International, 2016, 40, 74-82.	1.4	15
28	Transgenesis of humanized fat1 promotes nâ^'3 polyunsaturated fatty acid synthesis and expression of genes involved in lipid metabolism in goat cells. Gene, 2016, 576, 249-255.	1.0	2
29	A novel fluorescence reporter system for the characterization of dairy goat mammary epithelial cells. Biochemical and Biophysical Research Communications, 2015, 458, 783-789.	1.0	2