Chuanying Pan

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Survey of the relationship between polymorphisms within the <i>BMPR1B</i> gene and sheep reproductive traits. Animal Biotechnology, 2023, 34, 718-727.	0.7	10
2	Screen of small fragment mutations within the sheep thyroid stimulating hormone receptor gene associated with litter size. Animal Biotechnology, 2023, 34, 658-663.	0.7	4
3	Investigation on mRNA expression and genetic variation within goat <i>SMAD2</i> gene and its association with litter size. Animal Biotechnology, 2023, 34, 2111-2119.	0.7	3
4	Relationships between novel nucleotide variants within the colony-stimulating factor 1 receptor (<i>CSF1R</i>) gene and mastitis indicators in sheep. Animal Biotechnology, 2022, 33, 731-738.	0.7	3
5	ldentification of three new microsatellites and their effects on body measurement traits in pigs using time of flight-mass spectrometry (TOF-MS). Animal Biotechnology, 2022, 33, 1035-1044.	0.7	1
6	A 7-nt nucleotide sequence variant within the sheep <i>KDM3B</i> gene affects female reproduction traits. Animal Biotechnology, 2022, 33, 1661-1667.	0.7	3
7	Early-life lead exposure induces long-term toxicity in the central nervous system: From zebrafish larvae to juveniles and adults. Science of the Total Environment, 2022, 804, 150185.	3.9	41
8	Developmental exposure to environmental levels of cadmium induces neurotoxicity and activates microglia in zebrafish larvae: From the perspectives of neurobehavior and neuroimaging. Chemosphere, 2022, 291, 132802.	4.2	24
9	Three novel simple sequence repeats (SSRs) identified by MALDI-TOF-MS method were associated with backfat in pig. Animal Biotechnology, 2022, , 1-8.	0.7	0
10	Investigation of Copy Number Variations (CNVs) of the Goat PPP3CA Gene and Their Effect on Litter Size and Semen Quality. Animals, 2022, 12, 445.	1.0	5
11	Whole-genome sequencing to identify candidate genes for litter size and to uncover the variant function in goats (Capra hircus). Genomics, 2021, 113, 142-150.	1.3	28
12	An insertion/deletion within the CREB1 gene identified using the RNA-sequencing is associated with sheep body morphometric traits. Gene, 2021, 775, 145444.	1.0	10
13	The mRNA expression profile of the goat prion protein testis-specific (PRNT) gene and its associations with litter size. Theriogenology, 2021, 165, 69-75.	0.9	13
14	A 24-bp indel within the sheep AHR gene is associated with litter size. Animal Biotechnology, 2021, , 1-6.	0.7	2
15	Detection of 15-bp Deletion Mutation within PLAG1 Gene and Its Effects on Growth Traits in Goats. Animals, 2021, 11, 2064.	1.0	8
16	Ronin governs the metabolic capacity of the embryonic lineage for postâ€implantation development. EMBO Reports, 2021, 22, e53048.	2.0	4
17	Insertion/deletion variants within the IGF2BP2 gene identified in reported genome-wide selective sweep analysis reveal a correlation with goat litter size. Journal of Zhejiang University: Science B, 2021, 22, 757-766.	1.3	8
18	Palliative effects of metformin on testicular damage induced by triptolide in male rats. Ecotoxicology and Environmental Safety, 2021, 222, 112536.	2.9	6

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19	Detection of mRNA Expression and Copy Number Variations Within the Goat FecB Gene Associated With Litter Size. Frontiers in Veterinary Science, 2021, 8, 758705.	0.9	13
20	A deletion mutation within the <i>ATBF1</i> gene is strongly associated with goat litter size. Animal Biotechnology, 2020, 31, 174-180.	0.7	11
21	Indel variants within the <i>PRL</i> and <i>GHR</i> genes associated with sheep litter size. Reproduction in Domestic Animals, 2020, 55, 1470-1478.	0.6	20
22	Goat sperm associated antigen 17 protein gene (SPAG17): Small and large fragment genetic variation detection, association analysis, and mRNA expression in gonads. Genomics, 2020, 112, 5115-5121.	1.3	16
23	C2C12 Mouse Myoblasts Damage Induced by Oxidative Stress Is Alleviated by the Antioxidant Capacity of the Active Substance Phloretin. Frontiers in Cell and Developmental Biology, 2020, 8, 541260.	1.8	14
24	Identification and characterization of male reproduction-related genes in pig (Sus scrofa) using transcriptome analysis. BMC Genomics, 2020, 21, 381.	1.2	7
25	Early cleavage of preimplantation embryos is regulated by tRNAGIn-TTG–derived small RNAs present in mature spermatozoa. Journal of Biological Chemistry, 2020, 295, 10885-10900.	1.6	33
26	Two Novel Rare Strongly Linked Missense SNPs (P27R and A85G) Within the GDF9 Gene Were Significantly Associated With Litter Size in Shaanbei White Cashmere (SBWC) Goats. Frontiers in Veterinary Science, 2020, 7, 406.	0.9	6
27	A 5-bp mutation withinMSTN/GDF8gene was significantly associated with growth traits in Inner Mongolia White Cashmere goats. Animal Biotechnology, 2020, 32, 1-6.	0.7	8
28	miR-205 Expression Elevated With EDS Treatment and Induced Leydig Cell Apoptosis by Targeting RAP2B via the PI3K/AKT Signaling Pathway. Frontiers in Cell and Developmental Biology, 2020, 8, 448.	1.8	4
29	Chlorpyrifos inhibits sperm maturation and induces a decrease in mouse male fertility. Environmental Research, 2020, 188, 109785.	3.7	20
30	Myostatin (MSTN) Gene Indel Variation and Its Associations with Body Traits in Shaanbei White Cashmere Goat. Animals, 2020, 10, 168.	1.0	19
31	Goat CMTM2: mRNA expression profiles of different alternative spliced variants and associations analyses with growth traits. 3 Biotech, 2020, 10, 131.	1.1	6
32	Genetic effects of DSCAML1 identified in genome-wide association study revealing strong associations with litter size and semen quality in goat (Capra hircus). Theriogenology, 2020, 146, 20-25.	0.9	52
33	Multiple morphological abnormalities of the sperm flagella (MMAF)-associated genes: The relationships between genetic variation and litter size in goats. Gene, 2020, 753, 144778.	1.0	12
34	Pig SOX9: Expression profiles of Sertoli cell (SCs) and a functional 18 bp indel affecting testis weight. Theriogenology, 2019, 138, 94-101.	0.9	29
35	A 14-bp functional deletion within the CMTM2 gene is significantly associated with litter size in goat. Theriogenology, 2019, 139, 49-57.	0.9	46
36	Genetic Effects of Single Nucleotide Polymorphisms in the Goat GDF9 Gene on Prolificacy: True or False Positive?. Animals, 2019, 9, 886.	1.0	27

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37	Pig Hsd17b3: Alternative splice variants expression, insertion/deletion (indel) in promoter region and their associations with male reproductive traits. Journal of Steroid Biochemistry and Molecular Biology, 2019, 195, 105483.	1.2	13
38	Goat SPEF2: Expression profile, indel variants identification and association analysis with litter size. Theriogenology, 2019, 139, 147-155.	0.9	33
39	A novel missense mutation (L280V) within POU1F1 gene strongly affects litter size and growth traits in goat. Theriogenology, 2019, 135, 198-203.	0.9	20
40	<scp>d</scp> - <i>chiro</i> -Inositol Ameliorates High Fat Diet-Induced Hepatic Steatosis and Insulin Resistance via PKCÎμ-PI3K/AKT Pathway. Journal of Agricultural and Food Chemistry, 2019, 67, 5957-5967.	2.4	38
41	Relationship between SNPs of POU1F1 Gene and Litter Size and Growth Traits in Shaanbei White Cashmere Goats. Animals, 2019, 9, 114.	1.0	37
42	Polymorphisms within the Boule Gene Detected by Tetra-Primer Amplification Refractory Mutation System PCR (T-ARMS-PCR) are Significantly Associated with Goat Litter Size. Animals, 2019, 9, 910.	1.0	3
43	An 11-bp Indel Polymorphism within the CSN1S1 Gene Is Associated with Milk Performance and Body Measurement Traits in Chinese Goats. Animals, 2019, 9, 1114.	1.0	25
44	One 16†bp insertion/deletion (indel) within the KDM6A gene revealing strong associations with growth traits in goat. Gene, 2019, 686, 16-20.	1.0	29
45	Two strongly linked single nucleotide polymorphisms (Q320P and V397I) in GDF9 gene are associated with litter size in cashmere goats. Theriogenology, 2019, 125, 115-121.	0.9	77
46	Development of a touch-down multiplex PCR method for simultaneously rapidly detecting three novel insertion/deletions (indels) within one gene: an example for goat GHR gene. Animal Biotechnology, 2019, 30, 366-371.	0.7	38
47	Activation of Nrf2 by Phloretin Attenuates Palmitic Acid-Induced Endothelial Cell Oxidative Stress via AMPK-Dependent Signaling. Journal of Agricultural and Food Chemistry, 2019, 67, 120-131.	2.4	55
48	A 20-bp insertion/deletion (indel) polymorphism within the <i>CDC25A</i> gene and its associations with growth traits in goat. Archives Animal Breeding, 2019, 62, 353-360.	0.5	18
49	Chlorpyrifos Induction of Testicular-Cell Apoptosis through Generation of Reactive Oxygen Species and Phosphorylation of AMPK. Journal of Agricultural and Food Chemistry, 2018, 66, 12455-12470.	2.4	50
50	Goat CTNNB1: mRNA expression profile of alternative splicing in testis and association analysis with litter size. Gene, 2018, 679, 297-304.	1.0	34
51	Goat Boule: Isoforms identification, mRNA expression in testis and functional study and promoter methylation profiles. Theriogenology, 2018, 116, 53-63.	0.9	5
52	Insertion/Deletion Within the KDM6A Gene Is Significantly Associated With Litter Size in Goat. Frontiers in Genetics, 2018, 9, 91.	1.1	112
53	A novel indel within goat casein alpha S1 gene is significantly associated with litter size. Gene, 2018, 671, 161-169.	1.0	48
54	Identification of a novel 12-bp insertion/deletion (indel) of iPS-related Oct4 gene and its association with reproductive traits in male piglets. Animal Reproduction Science, 2017, 178, 55-60.	0.5	32

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55	A novel 12â€bp indel polymorphism within the <i><scp>GDF</scp>9</i> gene is significantly associated with litter size and growth traits in goats. Animal Genetics, 2017, 48, 735-736.	0.6	75
56	Cysteine protects rabbit spermatozoa against reactive oxygen species-induced damages. PLoS ONE, 2017, 12, e0181110.	1.1	28
57	Associations of six SNPs of POU1F1-PROP1-PITX1-SIX3 pathway genes with growth traits in two Chinese indigenous goat breeds. Annals of Animal Science, 2017, 17, 399-411.	0.6	15
58	Identification of novel isoforms of dairy goat EEF1D and their mRNA expression characterization. Gene, 2016, 581, 14-20.	1.0	5
59	Identification of novel alternative splicing transcript and expression analysis of bovine TMEM95 gene. Gene, 2016, 575, 531-536.	1.0	16
60	Novel alternative splice variants of NFIX and their diverse mRNA expression patterns in dairy goat. Gene, 2015, 569, 250-258.	1.0	11
61	Expression of TAT recombinant Oct4, Sox2, Lin28, and Nanog proteins from baculovirus-infected Sf9 insect cells. Gene, 2015, 556, 245-248.	1.0	9
62	Reprogramming human fibroblasts using HIV-1 TAT recombinant proteins OCT4, SOX2, KLF4 and c-MYC. Molecular Biology Reports, 2010, 37, 2117-2124.	1.0	42
63	An economical single-sided antibody incubation method for Western blotting. Journal of Virological Methods, 2010, 169, 409-411.	1.0	5
64	SNL fibroblast feeder layers support derivation and maintenance of human induced pluripotent stem cells. Journal of Genetics and Genomics, 2010, 37, 241-248.	1.7	24
65	A Taql PCR-RFLP Detecting a Novel SNP in ExonÂ2 of the Bovine POU1F1 Gene. Biochemical Genetics, 2008, 46, 424-432.	0.8	9