Armand SÃ nchez Anchez SÃ;nchez

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Characterisation of sperm piRNAs and their correlation with semen quality traits in swine. Animal Genetics, 2021, 52, 114-120.	0.6	15
2	Characterization of the Impact of Density Gradient Centrifugation on the Profile of the Pig Sperm Transcriptome by RNA-Seq. Frontiers in Veterinary Science, 2021, 8, 668158.	0.9	4
3	Assessing the levels of intraspecific admixture and interspecific hybridization in Iberian wild goats (<i>Capra pyrenaica</i>). Evolutionary Applications, 2021, 14, 2618-2634.	1.5	6
4	Expression analysis of porcine miR-33a/b in liver, adipose tissue and muscle and its potential role in fatty acid metabolism. PLoS ONE, 2021, 16, e0245858.	1.1	3
5	Characterization of skin surface and dermal microbiota in dogs with mast cell tumor. Scientific Reports, 2020, 10, 12634.	1.6	3
6	A pilot RNA-seq study in 40 pietrain ejaculates to characterize the porcine sperm microbiome. Theriogenology, 2020, 157, 525-533.	0.9	19
7	Whole genome sequencing identifies allelic ratio distortion in sperm involving genes related to spermatogenesis in a swine model. DNA Research, 2020, 27, .	1.5	6
8	A systems biology framework integrating GWAS and RNA-seq to shed light on the molecular basis of sperm quality in swine. Genetics Selection Evolution, 2020, 52, 72.	1.2	25
9	Identification of circular RNAs in porcine sperm and evaluation of their relation to sperm motility. Scientific Reports, 2020, 10, 7985.	1.6	27
10	Short communication: Milk microbiota profiling on water buffalo with full-length 16S rRNA using nanopore sequencing. Journal of Dairy Science, 2020, 103, 2693-2700.	1.4	12
11	A RNA-Seq Analysis to Describe the Boar Sperm Transcriptome and Its Seasonal Changes. Frontiers in Genetics, 2019, 10, 299.	1.1	53
12	Skin mites in mice (Mus musculus): high prevalence of Myobia sp. (Acari, Arachnida) in Robertsonian mice. Parasitology Research, 2018, 117, 2139-2148.	0.6	5
13	A technical assessment of the porcine ejaculated spermatozoa for a sperm-specific RNA-seq analysis. Systems Biology in Reproductive Medicine, 2018, 64, 291-303.	1.0	45
14	African swine fever virus does not express viral microRNAs in experimentally infected pigs. BMC Veterinary Research, 2018, 14, 268.	0.7	10
15	Microbiota profiling with long amplicons using Nanopore sequencing: full-length 16S rRNA gene and whole rrn operon. F1000Research, 2018, 7, 1755.	0.8	68
16	Microbiota profiling with long amplicons using Nanopore sequencing: full-length 16S rRNA gene and the 16S-ITS-23S of theÂrrn operon. F1000Research, 2018, 7, 1755.	0.8	67
17	Individual Signatures Define Canine Skin Microbiota Composition and Variability. Frontiers in Veterinary Science, 2017, 4, 6.	0.9	26
18	Differential expression of porcine microRNAs in African swine fever virus infected pigs: a proof-of-concept study. Virology Journal, 2017, 14, 198.	1.4	22

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19	Individual signatures and environmental factors shape skin microbiota in healthy dogs. Microbiome, 2017, 5, 139.	4.9	38
20	Identification of protein-damaging mutations in 10 swine taste receptors and 191 appetite-reward genes. BMC Genomics, 2016, 17, 685.	1.2	5
21	Detection, Prevalence and Phylogenetic Relationships of Demodex spp and further Skin Prostigmata Mites (Acari, Arachnida) in Wild and Domestic Mammals. PLoS ONE, 2016, 11, e0165765.	1.1	27
22	Magnetic Bead/Gold Nanoparticle Double-Labeled Primers for Electrochemical Detection of Isothermal Amplified <i>Leishmania</i> DNA. Small, 2016, 12, 205-213.	5.2	70
23	Population structure of eleven Spanish ovine breeds and detection of selective sweeps with BayeScan and hapFLK. Scientific Reports, 2016, 6, 27296.	1.6	52
24	A genome-wide perspective about the diversity and demographic history of seven Spanish goat breeds. Genetics Selection Evolution, 2016, 48, 52.	1.2	63
25	Romanian wild boars and Mangalitza pigs have a European ancestry and harbour genetic signatures compatible with past population bottlenecks. Scientific Reports, 2016, 6, 29913.	1.6	16
26	Electrocatalytic Detection: Magnetic Bead/Gold Nanoparticle Double-Labeled Primers for Electrochemical Detection of Isothermal Amplified <i>Leishmania</i> DNA (Small 2/2016). Small, 2016, 12, 204-204.	5.2	2
27	Rapid genome resequencing of an atoxigenic strain of Aspergillus carbonarius. Scientific Reports, 2015, 5, 9086.	1.6	15
28	Endometrial gene expression profile of pregnant sows with extreme phenotypes for reproductive efficiency. Scientific Reports, 2015, 5, 14416.	1.6	16
29	East African pigs have a complex Indian, Far Eastern and Western ancestry. Animal Genetics, 2015, 46, 433-436.	0.6	15
30	A Genetic Predictive Model for Canine Hip Dysplasia: Integration of Genome Wide Association Study (GWAS) and Candidate Gene Approaches. PLoS ONE, 2015, 10, e0122558.	1.1	28
31	Evaluation of the capability of the PCV2 genome to encode miRNAs: lack of viral miRNA expression in an experimental infection. Veterinary Research, 2015, 46, 48.	1.1	6
32	An iridium oxide nanoparticle and polythionine thin film based platform for sensitive Leishmania DNA detection. Journal of Materials Chemistry B, 2015, 3, 5166-5171.	2.9	29
33	Identification of microRNAs in PCV2 subclinically infected pigs by high throughput sequencing. Veterinary Research, 2015, 46, 18.	1.1	15
34	Triple lines gold nanoparticle-based lateral flow assay for enhanced and simultaneous detection of Leishmania DNA and endogenous control. Nano Research, 2015, 8, 3704-3714.	5.8	66
35	The Role of Viral and Host MicroRNAs in the Aujeszky's Disease Virus during the Infection Process. PLoS ONE, 2014, 9, e86965.	1.1	21
36	Non-synonymous genetic variation in exonic regions of canine Toll-like receptors. Canine Genetics and Epidemiology, 2014, 1, 11.	2.9	9

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37	Development of 23 individual TaqMan® real-time PCR assays for identifying common foodborne pathogens using a single set of amplification conditions. Food Microbiology, 2014, 43, 35-40.	2.1	40
38	Development of a PCR technique specific for Demodex injai in biological specimens. Parasitology Research, 2013, 112, 3369-3372.	0.6	12
39	Small <i>Demodex</i> populations colonize most parts of the skin of healthy dogs. Veterinary Dermatology, 2013, 24, 168.	0.4	49
40	miRNA Expression Profile Analysis in Kidney of Different Porcine Breeds. PLoS ONE, 2013, 8, e55402.	1.1	23
41	Microarray analysis of mediastinal lymph node of pigs naturally affected by postweaning multisystemic wasting syndrome. Virus Research, 2012, 165, 134-142.	1.1	9
42	Phylogenetic relationships in three species of canine <i>Demodex</i> mite based on partial sequences of mitochondrial 16S rDNA. Veterinary Dermatology, 2012, 23, 509.	0.4	42
43	Determination of Reference microRNAs for Relative Quantification in Porcine Tissues. PLoS ONE, 2012, 7, e44413.	1.1	49
44	Genetic Control of Canine Leishmaniasis: Genome-Wide Association Study and Genomic Selection Analysis. PLoS ONE, 2012, 7, e35349.	1.1	31
45	Sequencing and gene expression of the porcine ITIH SSC13 cluster and its effect on litter size in an Iberian×Meishan F2 population. Animal Reproduction Science, 2011, 128, 85-92.	0.5	4
46	Effects of α _{s1} -casein (<i>CSN1S1</i>) and κ-casein (<i>CSN3</i>) genotypes on milk coagulation properties in Murciano-Granadina goats. Journal of Dairy Research, 2011, 78, 32-37.	0.7	24
47	A Novel Unstable Duplication Upstream of HAS2 Predisposes to a Breed-Defining Skin Phenotype and a Periodic Fever Syndrome in Chinese Shar-Pei Dogs. PLoS Genetics, 2011, 7, e1001332.	1.5	118
48	Quantitative trait loci for fatness at growing and reproductive stages in Iberianâ€f×â€fMeishan F ₂ sows. Animal Genetics, 2011, 42, 548-551.	0.6	5
49	Canine leishmaniasis: the key points for qPCR result interpretation. Parasites and Vectors, 2011, 4, 57.	1.0	59
50	Genetic assessment, illegal trafficking and management of the Mediterranean spur-thighed tortoise in Southern Spain and Northern Africa. Conservation Genetics, 2011, 12, 1-13.	0.8	13
51	Signatures of demographic bottlenecks in European wolf populations. Conservation Genetics, 2011, 12, 701-712.	0.8	48
52	Development of a real-time PCR to detect Demodex canis DNA in different tissue samples. Parasitology Research, 2011, 108, 305-308.	0.6	39
53	Analysis of porcine MUC4 gene as a candidate gene for prolificacy QTL on SSC13 in an Iberian × Meishan F2 population. BMC Genetics, 2011, 12, 93.	2.7	8
54	A selective sweep of >8 Mb on chromosome 26 in the Boxer genome. BMC Genomics, 2011, 12, 339.	1.2	50

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55	Mycoplasma and Herpesvirus PCR Detection in Tortoises with Rhinitis-stomatitis Complex in Spain. Journal of Wildlife Diseases, 2011, 47, 195-200.	0.3	20
56	Genetic variability, structure and assignment of Spanish and French pig populations based on a large sampling. Animal Genetics, 2010, 41, 608-618.	0.6	23
57	Time course differential gene expression in response to porcine circovirus typeÂ2 subclinical infection. Veterinary Research, 2010, 41, 12.	1.1	30
58	Presence of opportunistic bacteria (Rhizobium spp.) with potential for molecular misdiagnosis among canine and feline clinical samples. Canadian Veterinary Journal, 2010, 51, 895-7.	0.0	2
59	Exploratory Study on the Transcriptional Profile of Pigs Subclinically Infected with Porcine Circovirus Type 2. Animal Biotechnology, 2009, 20, 96-109.	0.7	12
60	Integrating Y-Chromosome, Mitochondrial, and Autosomal Data to Analyze the Origin of Pig Breeds. Molecular Biology and Evolution, 2009, 26, 2061-2072.	3.5	103
61	A bi-dimensional genome scan for prolificacy traits in pigs shows the existence of multiple epistatic QTL. BMC Genomics, 2009, 10, 636.	1.2	40
62	Sex identification of wolf (Canis lupus) using non-invasive samples. Conservation Genetics, 2009, 10, 555-558.	0.8	31
63	Mitochondrial DNA diversity and origins of South and Central American goats. Animal Genetics, 2009, 40, 315-322.	0.6	46
64	A nonâ€synonymous mutation in a conserved site of the <i>MTTP</i> gene is strongly associated with protein activity and fatty acid profile in pigs. Animal Genetics, 2009, 40, 813-820.	0.6	28
65	Evaluation of <i>FABP2</i> as candidate gene for a fatty acid composition QTL in porcine chromosome 8. Journal of Animal Breeding and Genetics, 2009, 126, 52-58.	0.8	10
66	First birth of an animal from an extinct subspecies (Capra pyrenaica pyrenaica) by cloning. Theriogenology, 2009, 71, 1026-1034.	0.9	136
67	An age-dependent association between a leptin C3469T single nucleotide polymorphism and intramuscular fat content in pigs. Livestock Science, 2009, 121, 335-338.	0.6	6
68	Short communication: Effect of αS1-casein (CSN1S1) and κ-casein (CSN3) genotypes on milk composition in Murciano-Granadina goats. Journal of Dairy Science, 2009, 92, 2960-2964.	1.4	39
69	Vector-borne infections in cats: Molecular study in Barcelona area (Spain). Veterinary Parasitology, 2008, 151, 332-336.	0.7	141
70	Detection of Leishmania infantum in captive wolves from Southwestern Europe. Veterinary Parasitology, 2008, 158, 117-120.	0.7	27
71	Using haplotype probabilities in categorical survival analysis: a case study with three candidate genes in an Iberianâ€f×â€fMeishan F ₂ population of newborn piglets. Journal of Animal Breeding and Genetics, 2008, 125, 5-12.	0.8	2
72	Plasma leptin levels in pigs with different leptin and leptin receptor genotypes. Journal of Animal Breeding and Genetics, 2008, 125, 228-233.	0.8	13

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73	Longitudinal analysis of cytokine gene expression and parasite load in PBMC in Leishmania infantum experimentally infected dogs. Veterinary Immunology and Immunopathology, 2008, 125, 168-175.	0.5	20
74	Sequence Analysis of Goat Major Histocompatibility Complex Class I Genes. Journal of Dairy Science, 2008, 91, 814-817.	1.4	8
75	Effect of α _{s1} -casein (<i>CSN1S1</i>) genotype on milk CSN1S1 content in Malagueña and Murciano-Granadina goats. Journal of Dairy Research, 2008, 75, 481-484.	0.7	20
76	Empirical Bayes factor analyses of quantitative trait loci for gestation length in Iberian × Meishan F2 sows. Animal, 2008, 2, 177-183.	1.3	6
77	Slc11a1 (formerly Nramp1) and susceptibility to canine visceral leishmaniasis. Veterinary Research, 2008, 39, 36.	1.1	48
78	Association of CA repeat polymorphism at intron 1 of insulin-like growth factor (IGF-I) gene with circulating IGF-I concentration, growth, and fatness in swine. Physiological Genomics, 2007, 31, 236-243.	1.0	40
79	Nucleotide Sequence and Polymorphism of the Pig Acyl Coenzyme A Synthetase Long-Chain 1 (ACSL1) Gene. Animal Biotechnology, 2007, 18, 117-122.	0.7	2
80	Lack ofIn VitroandIn VivoEffects of Lipopolysaccharide on Porcine Circovirus Type 2 Infection. Viral Immunology, 2007, 20, 541-552.	0.6	15
81	Characterization and Physical Mapping of the PorcineCDS1andCDS2Genes. Animal Biotechnology, 2007, 18, 23-35.	0.7	8
82	Short Communication: Identification of Two Polymorphisms in the Goat Lipoprotein Lipase Gene and Their Association with Milk Production Traits. Journal of Dairy Science, 2007, 90, 3012-3017.	1.4	20
83	Individual identification and genetic traceability in the pig using the SNPlex TM genotyping system. Animal Genetics, 2007, 38, 663-665.	0.6	8
84	Genetic polymorphism of αs1- and αs2-caseins in Hungarian Milking Goats. Small Ruminant Research, 2007, 68, 329-332.	0.6	15
85	Adipocyte fatty-acid binding protein is closely associated to the porcine FAT1 locus on chromosome 41. Journal of Animal Science, 2006, 84, 2907-2913.	0.2	26
86	High amino acid variation in the intracellular domain of the pig prolactin receptor (PRLR) and its relation to ovulation rate and piglet survival traits1. Journal of Animal Science, 2006, 84, 1991-1998.	0.2	36
87	Malic enzyme 1 genotype is associated with backfat thickness and meat quality traits in pigs. Animal Genetics, 2006, 37, 28-32.	0.6	37
88	Characterization of the porcine acyl-CoA synthetase long-chain 4 gene and its association with growth and meat quality traits. Animal Genetics, 2006, 37, 219-224.	0.6	35
89	An association study between polymorphisms of the porcine bone morphogenetic protein receptor type1beta(BMPR1B) and reproductive performance of Iberian x Meishan F2 sows. Animal Genetics, 2006, 37, 297-298.	0.6	8
90	Polymorphisms of the porcine dopaminebeta-hydroxylase gene and their relation to reproduction and piglet survivability in an Iberian x Meishan F2 intercross. Animal Genetics, 2006, 37, 279-282.	0.6	11

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91	Characterization of the porcine FABP5 gene and its association with the FAT1 QTL in an Iberian by Landrace cross. Animal Genetics, 2006, 37, 589-591.	0.6	33
92	Advantages of real-time PCR assay for diagnosis and monitoring of canine leishmaniosis. Veterinary Parasitology, 2006, 137, 214-221.	0.7	303
93	Expression of Recombinant Human Follicle-Stimulating Hormone in the Mammary Gland of Transgenic Mice. Molecular Biotechnology, 2006, 34, 37-44.	1.3	0
94	Genetic assessment of the Iberian wolf Canis lupus signatus captive breeding program. Conservation Genetics, 2006, 7, 861-878.	0.8	42
95	Potential benefit from using the alphas1-casein genotype information in a selection scheme for dairy goats. Journal of Animal Breeding and Genetics, 2005, 122, 21-29.	0.8	25
96	Exclusion of the acyl CoA:diacylglycerol acyltransferase 1 gene (DGAT1) as a candidate for a fatty acid composition QTL on porcine chromosome 4. Journal of Animal Breeding and Genetics, 2005, 122, 161-164.	0.8	10
97	Assignment of theacyl-CoA synthetase long-chain family member 4 (ACSL4)gene to porcine chromosome X. Animal Genetics, 2005, 36, 76-76.	0.6	3
98	Assignment of the microsomal triglyceride transfer protein large subunit (MTP) gene to porcine chromosome 8. Animal Genetics, 2005, 36, 354-355.	0.6	5
99	Assignment of the beta-lactoglobulin (BLG) gene to porcine chromosome 1. Animal Genetics, 2005, 36, 356-358.	0.6	2
100	Assignment of the <i>phospholipase C</i> β <i>1</i> (<i>PLCB1</i>) gene to porcine chromosome 17. Animal Genetics, 2005, 36, 516-517.	0.6	1
101	On growth, fatness, and form: A further look at porcine Chromosome 4 in an Iberian × Landrace cross. Mammalian Genome, 2005, 16, 374-382.	1.0	33
102	Polymorphism of Slc11a1 (Nramp1) Gene and Canine Leishmaniasis in a Case-Control Study. Journal of Heredity, 2005, 96, 755-758.	1.0	46
103	Polymorphisms in the goat β-lactoglobulin gene. Journal of Dairy Research, 2005, 72, 379-384.	0.7	21
104	Nucleotide sequence and polymorphism of the caprine major histocompatibility complex class II (-) gene. Molecular Immunology, 2005, 42, 375-379.	1.0	14
105	Genetic polymorphism of the K-casein (CSN3) gene in goats reared in Southern Italy. Italian Journal of Animal Science, 2005, 4, 97-101.	0.8	8
106	Real-time quantitative PCR-based system for determining transgene copy number in transgenic animals. BioTechniques, 2004, 37, 610-613.	0.8	120
107	Genetic analysis of SLC11A1 polymorphisms in multiple sclerosis patients. Multiple Sclerosis Journal, 2004, 10, 618-620.	1.4	9
108	Low diversity in the major histocompatibility complex class II DRB1 gene of the Spanish ibex, Capra pyrenaica. Heredity, 2004, 93, 266-272.	1.2	32

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109	Epithelioid Malignant Peripheral Nerve Sheath Tumour in a Dog. Journal of Comparative Pathology, 2004, 131, 87-91.	0.1	16
110	Estimating the frequency of Asian cytochrome B haplotypes in standard European and local Spanish pig breeds. Genetics Selection Evolution, 2004, 36, 97-104.	1.2	35
111	Genetic diversity measures of the bovine Alberes breed using microsatellites: variability among herds and types of coat colour*. Journal of Animal Breeding and Genetics, 2004, 121, 101-110.	0.8	16
112	Disruption of the mouse phospholipase C-β1 gene in a β-lactoglobulin transgenic line affects viability, growth, and fertility in mice. Gene, 2004, 341, 279-289.	1.0	26
113	Structural characterization of the caprine major histocompatibility complex class II DQB1 (Cahi-DQB1) gene. Molecular Immunology, 2004, 41, 843-846.	1.0	8
114	Strong phylogeographic relationships among three goat breeds from the Canary Islands. Journal of Dairy Research, 2004, 71, 257-262.	0.7	51
115	Detection of QTL affecting fatty acid composition in the pig. Mammalian Genome, 2003, 14, 650-656.	1.0	74
116	Estrogen receptor polymorphism in Landrace pigs and its association with litter size performance. Livestock Science, 2003, 82, 53-59.	1.2	39
117	Genetic structure of eighteen local south European beef cattle breeds by comparative F-statistics analysis. Journal of Animal Breeding and Genetics, 2003, 120, 73-87.	0.8	46
118	Structural characterization of the porcine pyruvate carboxylase (PC) gene. Journal of Animal Breeding and Genetics, 2003, 120, 338-345.	0.8	1
119	Assignment of the fatty acid synthase (FASN) gene to pig chromosome 12 by physical and linkage mapping. Animal Genetics, 2003, 34, 234-235.	0.6	27
120	Linkage mapping of the porcine hairless gene (HR  ) to chromosome 14. Animal Genetics, 2003, 34, 317-318.	0.6	5
121	Mapping of the goat stearoyl coenzyme A desaturase gene to chromosome 26. Animal Genetics, 2003, 34, 474-475.	0.6	5
122	Genetic Characterization of Southwestern European Bovine Breeds: A Historical and Biogeographical Reassessment With a Set of 16 Microsatellites. , 2003, 94, 243-250.		78
123	A growth hormone-based phylogenetic analysis of euteleostean fishes including a representative species of the Atheriniformes Order, Odontesthes argentinensis. Genetics and Molecular Biology, 2003, 26, 295-300.	0.6	15
124	Mapping and Sequencing of the Canine NRAMP1 Gene and Identification of Mutations in Leishmaniasis-Susceptible Dogs. Infection and Immunity, 2002, 70, 2763-2771.	1.0	56
125	QTL mapping for growth and carcass traits in an Iberian by Landrace pig intercross: additive, dominant and epistatic effects. Genetical Research, 2002, 80, 145-154.	0.3	68
126	Physical and linkage mapping of the porcinepink-eye dilutiongene (OCA2). Animal Genetics, 2002, 33, 392-394.	0.6	3

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127	Assignment of the 2,4-dienoyl-CoA reductase (DECR) gene to porcine chromosome 4. Animal Genetics, 2002, 33, 164-165.	0.6	10
128	Isolation of Genomic DNA from Feathers. Journal of Veterinary Diagnostic Investigation, 2001, 13, 162-164.	0.5	63
129	Genetic polymorphism of the caprine kappa casein gene. Journal of Dairy Research, 2001, 68, 209-216.	0.7	36
130	A within-breed comparison of RYR1 pig genotypes for performance, feeding behaviour, and carcass, meat and fat quality traits. Journal of Animal Breeding and Genetics, 2001, 118, 417-427.	0.8	14
131	Genetic diversity measures of local European beef cattle breeds for conservation purposes. Genetics Selection Evolution, 2001, 33, 311-32.	1.2	146
132	Characterization of genetic polymorphism in the goat β-lactoglobulin gene. Journal of Dairy Research, 2000, 67, 217-224.	0.7	27
133	Gene transfer technology in aquaculture. Hydrobiologia, 2000, 420, 91-94.	1.0	11
134	Detection of <i>Leishmania</i> Infection in Paraffin-Embedded Skin Biopsies of Dogs Using Polymerase Chain Reaction. Journal of Veterinary Diagnostic Investigation, 1999, 11, 385-387.	0.5	29
135	DogMap: an international collaboration toward a low-resolution canine genetic marker map. , 1999, 90, 3-6.		8
136	The identification of a sex-specific DNA marker in the ostrich using a random amplified polymorphic DNA (RAPD) assay. Molecular Ecology, 1999, 8, 667-669.	2.0	34
137	Isolation, sequencing and relative quantitation by fluorescent-ratio PCR of feline β-lactoglobulin I, II, and III cDNAs. Mammalian Genome, 1999, 10, 560-564.	1.0	10
138	Genetic variation (protein markers and microsatellites) in endangered Catalonian donkeys. Biochemical Systematics and Ecology, 1999, 27, 791-798.	0.6	13
139	Chromatin Structures of Coat and Sheep β-Lactoglobulin Gene Differ. Biochemical and Biophysical Research Communications, 1998, 252, 649-653.	1.0	9
140	Isolation of genomic DNA from milk samples by using Chelex resin. Journal of Dairy Research, 1997, 64, 231-238.	0.7	42
141	Expression of caprine beta-lactoglobulin in the milk of transgenic mice. Transgenic Research, 1997, 6, 69-74.	1.3	8
142	Characterization of a caprine β-lactoglobulin pseudogene, identification and chromosomal localization by in situ hybridization in goat, sheep and cow. Gene, 1996, 177, 87-91.	1.0	21
143	A PCR-RFLP typing method for the caprine Mhc class II DRB gene. Veterinary Immunology and Immunopathology, 1996, 55, 255-260.	O.5	23
144	Gene frequencies of caprine αs1-casein polymorphism in Spanish goat breeds. Small Ruminant Research, 1996, 20, 215-221.	0.6	52

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145	Primer-directed synthesis of a molecular weight marker. Genetic Analysis, Techniques and Applications, 1996, 13, 147-149.	1.5	3
146	Northern analysis of highly folded goat $\hat{I}\pm s1$ Casein mRNA. Genetic Analysis, Techniques and Applications, 1996, 12, 143-145.	1.5	0
147	Analysis of genetic relationships in horse breeds. Journal of Equine Veterinary Science, 1995, 15, 320-328.	0.4	15
148	Nested PCR allows the characterization of TaqI and PstI RFLPs in the second exon of the caprine MHC class II DRB gene. Veterinary Immunology and Immunopathology, 1995, 48, 313-321.	0.5	35
149	Occurrence of a LINE sequence in the 3′ UTR of the goat αs1-casein E-encoding allele associated with reduced protein synthesis level. Gene, 1994, 147, 179-187.	1.0	96
150	Nucleotide sequence of the goat \hat{I}° -casein cDNA. Journal of Animal Science, 1993, 71, 2833-2833.	0.2	33
151	Cloning and sequencing of the cDNA encoding goat β-lactoglobulin. Journal of Animal Science, 1993, 71, 2832.	0.2	12
152	Description and Evolutionary Relationships of Two Species of the Drosophila mulleri Cluster (Diptera: Drosophilidae). Annals of the Entomological Society of America, 1990, 83, 444-452.	1.3	13
153	Drosophila koepferae: a New Member of the Drosophila serido (Diptera: Drosophilidae) Superspecies Taxon1. Annals of the Entomological Society of America, 1988, 81, 380-385.	1.3	63
154	Adh expression in species of themulleri subgroup ofDrosophila. Biochemical Genetics, 1987, 25, 729-738.	0.8	10