Armand SÃ nchez Anchez SÃ;nchez

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Advantages of real-time PCR assay for diagnosis and monitoring of canine leishmaniosis. Veterinary Parasitology, 2006, 137, 214-221.	1.8	303
2	Genetic diversity measures of local European beef cattle breeds for conservation purposes. Genetics Selection Evolution, 2001, 33, 311-32.	3.0	146
3	Vector-borne infections in cats: Molecular study in Barcelona area (Spain). Veterinary Parasitology, 2008, 151, 332-336.	1.8	141
4	First birth of an animal from an extinct subspecies (Capra pyrenaica pyrenaica) by cloning. Theriogenology, 2009, 71, 1026-1034.	2.1	136
5	Real-time quantitative PCR-based system for determining transgene copy number in transgenic animals. BioTechniques, 2004, 37, 610-613.	1.8	120
6	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.	3.5	118
7	Integrating Y-Chromosome, Mitochondrial, and Autosomal Data to Analyze the Origin of Pig Breeds. Molecular Biology and Evolution, 2009, 26, 2061-2072.	8.9	103
8	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.	2.2	96
9	Genetic Characterization of Southwestern European Bovine Breeds: A Historical and Biogeographical Reassessment With a Set of 16 Microsatellites. , 2003, 94, 243-250.		78
10	Detection of QTL affecting fatty acid composition in the pig. Mammalian Genome, 2003, 14, 650-656.	2.2	74
11	Magnetic Bead/Gold Nanoparticle Double-Labeled Primers for Electrochemical Detection of Isothermal Amplified <i>Leishmania</i> DNA. Small, 2016, 12, 205-213.	10.0	70
12	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.	0.9	68
13	Microbiota profiling with long amplicons using Nanopore sequencing: full-length 16S rRNA gene and whole rrn operon. F1000Research, 2018, 7, 1755.	1.6	68
14	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.	1.6	67
15	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.	10.4	66
16	Drosophila koepferae: a New Member of the Drosophila serido (Diptera: Drosophilidae) Superspecies Taxon1. Annals of the Entomological Society of America, 1988, 81, 380-385.	2.5	63
17	Isolation of Genomic DNA from Feathers. Journal of Veterinary Diagnostic Investigation, 2001, 13, 162-164.	1.1	63
18	A genome-wide perspective about the diversity and demographic history of seven Spanish goat breeds. Genetics Selection Evolution, 2016, 48, 52.	3.0	63

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19	Canine leishmaniasis: the key points for qPCR result interpretation. Parasites and Vectors, 2011, 4, 57.	2.5	59
20	Mapping and Sequencing of the Canine NRAMP1 Gene and Identification of Mutations in Leishmaniasis-Susceptible Dogs. Infection and Immunity, 2002, 70, 2763-2771.	2.2	56
21	A RNA-Seq Analysis to Describe the Boar Sperm Transcriptome and Its Seasonal Changes. Frontiers in Genetics, 2019, 10, 299.	2.3	53
22	Gene frequencies of caprine αs1-casein polymorphism in Spanish goat breeds. Small Ruminant Research, 1996, 20, 215-221.	1.2	52
23	Population structure of eleven Spanish ovine breeds and detection of selective sweeps with BayeScan and hapFLK. Scientific Reports, 2016, 6, 27296.	3.3	52
24	Strong phylogeographic relationships among three goat breeds from the Canary Islands. Journal of Dairy Research, 2004, 71, 257-262.	1.4	51
25	A selective sweep of >8 Mb on chromosome 26 in the Boxer genome. BMC Genomics, 2011, 12, 339.	2.8	50
26	Determination of Reference microRNAs for Relative Quantification in Porcine Tissues. PLoS ONE, 2012, 7, e44413.	2.5	49
27	Small <i>Demodex</i> populations colonize most parts of the skin of healthy dogs. Veterinary Dermatology, 2013, 24, 168.	1.2	49
28	Signatures of demographic bottlenecks in European wolf populations. Conservation Genetics, 2011, 12, 701-712.	1.5	48
29	Slc11a1 (formerly Nramp1) and susceptibility to canine visceral leishmaniasis. Veterinary Research, 2008, 39, 36.	3.0	48
30	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.	2.0	46
31	Polymorphism of Slc11a1 (Nramp1) Gene and Canine Leishmaniasis in a Case-Control Study. Journal of Heredity, 2005, 96, 755-758.	2.4	46
32	Mitochondrial DNA diversity and origins of South and Central American goats. Animal Genetics, 2009, 40, 315-322.	1.7	46
33	A technical assessment of the porcine ejaculated spermatozoa for a sperm-specific RNA-seq analysis. Systems Biology in Reproductive Medicine, 2018, 64, 291-303.	2.1	45
34	Isolation of genomic DNA from milk samples by using Chelex resin. Journal of Dairy Research, 1997, 64, 231-238.	1.4	42
35	Genetic assessment of the Iberian wolf Canis lupus signatus captive breeding program. Conservation Genetics, 2006, 7, 861-878.	1.5	42
36	Phylogenetic relationships in three species of canine <i>Demodex</i> mite based on partial sequences of mitochondrial 16S rDNA. Veterinary Dermatology, 2012, 23, 509.	1.2	42

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37	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.	2.3	40
38	A bi-dimensional genome scan for prolificacy traits in pigs shows the existence of multiple epistatic QTL. BMC Genomics, 2009, 10, 636.	2.8	40
39	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.	4.2	40
40	Estrogen receptor polymorphism in Landrace pigs and its association with litter size performance. Livestock Science, 2003, 82, 53-59.	1.2	39
41	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.	3.4	39
42	Development of a real-time PCR to detect Demodex canis DNA in different tissue samples. Parasitology Research, 2011, 108, 305-308.	1.6	39
43	Individual signatures and environmental factors shape skin microbiota in healthy dogs. Microbiome, 2017, 5, 139.	11.1	38
44	<i>Malic enzyme 1</i> genotype is associated with backfat thickness and meat quality traits in pigs. Animal Genetics, 2006, 37, 28-32.	1.7	37
45	Genetic polymorphism of the caprine kappa casein gene. Journal of Dairy Research, 2001, 68, 209-216.	1.4	36
46	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.	0.5	36
47	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.	1.2	35
48	Estimating the frequency of Asian cytochrome B haplotypes in standard European and local Spanish pig breeds. Genetics Selection Evolution, 2004, 36, 97-104.	3.0	35
49	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.	1.7	35
50	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.	3.9	34
51	Nucleotide sequence of the goat κ-casein cDNA. Journal of Animal Science, 1993, 71, 2833-2833.	0.5	33
52	On growth, fatness, and form: A further look at porcine Chromosome 4 in an Iberian × Landrace cross. Mammalian Genome, 2005, 16, 374-382.	2.2	33
53	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.	1.7	33
54	Low diversity in the major histocompatibility complex class II DRB1 gene of the Spanish ibex, Capra pyrenaica. Heredity, 2004, 93, 266-272.	2.6	32

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55	Sex identification of wolf (Canis lupus) using non-invasive samples. Conservation Genetics, 2009, 10, 555-558.	1.5	31
56	Genetic Control of Canine Leishmaniasis: Genome-Wide Association Study and Genomic Selection Analysis. PLoS ONE, 2012, 7, e35349.	2.5	31
57	Time course differential gene expression in response to porcine circovirus typeÂ2 subclinical infection. Veterinary Research, 2010, 41, 12.	3.0	30
58	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.	1.1	29
59	An iridium oxide nanoparticle and polythionine thin film based platform for sensitive Leishmania DNA detection. Journal of Materials Chemistry B, 2015, 3, 5166-5171.	5.8	29
60	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.	1.7	28
61	A Genetic Predictive Model for Canine Hip Dysplasia: Integration of Genome Wide Association Study (GWAS) and Candidate Gene Approaches. PLoS ONE, 2015, 10, e0122558.	2.5	28
62	Characterization of genetic polymorphism in the goat β-lactoglobulin gene. Journal of Dairy Research, 2000, 67, 217-224.	1.4	27
63	Assignment of the fatty acid synthase (FASN) gene to pig chromosome 12 by physical and linkage mapping. Animal Genetics, 2003, 34, 234-235.	1.7	27
64	Detection of Leishmania infantum in captive wolves from Southwestern Europe. Veterinary Parasitology, 2008, 158, 117-120.	1.8	27
65	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.	2.5	27
66	Identification of circular RNAs in porcine sperm and evaluation of their relation to sperm motility. Scientific Reports, 2020, 10, 7985.	3.3	27
67	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.	2.2	26
68	Adipocyte fatty-acid binding protein is closely associated to the porcine FAT1 locus on chromosome 41. Journal of Animal Science, 2006, 84, 2907-2913.	0.5	26
69	Individual Signatures Define Canine Skin Microbiota Composition and Variability. Frontiers in Veterinary Science, 2017, 4, 6.	2.2	26
70	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.	2.0	25
71	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.	3.0	25
72	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.	1.4	24

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73	A PCR-RFLP typing method for the caprine Mhc class II DRB gene. Veterinary Immunology and Immunopathology, 1996, 55, 255-260.	1.2	23
74	Genetic variability, structure and assignment of Spanish and French pig populations based on a large sampling. Animal Genetics, 2010, 41, 608-618.	1.7	23
75	miRNA Expression Profile Analysis in Kidney of Different Porcine Breeds. PLoS ONE, 2013, 8, e55402.	2.5	23
76	Differential expression of porcine microRNAs in African swine fever virus infected pigs: a proof-of-concept study. Virology Journal, 2017, 14, 198.	3.4	22
77	Characterization of a caprine \hat{l}^2 -lactoglobulin pseudogene, identification and chromosomal localization by in situ hybridization in goat, sheep and cow. Gene, 1996, 177, 87-91.	2.2	21
78	Polymorphisms in the goat β-lactoglobulin gene. Journal of Dairy Research, 2005, 72, 379-384.	1.4	21
79	The Role of Viral and Host MicroRNAs in the Aujeszky's Disease Virus during the Infection Process. PLoS ONE, 2014, 9, e86965.	2.5	21
80	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.	3.4	20
81	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.	1.2	20
82	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.	1.4	20
83	Mycoplasma and Herpesvirus PCR Detection in Tortoises with Rhinitis-stomatitis Complex in Spain. Journal of Wildlife Diseases, 2011, 47, 195-200.	0.8	20
84	A pilot RNA-seq study in 40 pietrain ejaculates to characterize the porcine sperm microbiome. Theriogenology, 2020, 157, 525-533.	2.1	19
85	Epithelioid Malignant Peripheral Nerve Sheath Tumour in a Dog. Journal of Comparative Pathology, 2004, 131, 87-91.	0.4	16
86	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.	2.0	16
87	Endometrial gene expression profile of pregnant sows with extreme phenotypes for reproductive efficiency. Scientific Reports, 2015, 5, 14416.	3.3	16
88	Romanian wild boars and Mangalitza pigs have a European ancestry and harbour genetic signatures compatible with past population bottlenecks. Scientific Reports, 2016, 6, 29913.	3.3	16
89	Analysis of genetic relationships in horse breeds. Journal of Equine Veterinary Science, 1995, 15, 320-328.	0.9	15
90	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.	1.3	15

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91	Lack ofIn VitroandIn VivoEffects of Lipopolysaccharide on Porcine Circovirus Type 2 Infection. Viral Immunology, 2007, 20, 541-552.	1.3	15
92	Genetic polymorphism of αs1- and αs2-caseins in Hungarian Milking Goats. Small Ruminant Research, 2007, 68, 329-332.	1.2	15
93	Rapid genome resequencing of an atoxigenic strain of Aspergillus carbonarius. Scientific Reports, 2015, 5, 9086.	3.3	15
94	East African pigs have a complex Indian, Far Eastern and Western ancestry. Animal Genetics, 2015, 46, 433-436.	1.7	15
95	Identification of microRNAs in PCV2 subclinically infected pigs by high throughput sequencing. Veterinary Research, 2015, 46, 18.	3.0	15
96	Characterisation of sperm piRNAs and their correlation with semen quality traits in swine. Animal Genetics, 2021, 52, 114-120.	1.7	15
97	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.	2.0	14
98	Nucleotide sequence and polymorphism of the caprine major histocompatibility complex class II (-) gene. Molecular Immunology, 2005, 42, 375-379.	2.2	14
99	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.	2.5	13
100	Genetic variation (protein markers and microsatellites) in endangered Catalonian donkeys. Biochemical Systematics and Ecology, 1999, 27, 791-798.	1.3	13
101	Plasma leptin levels in pigs with different leptin and leptin receptor genotypes. Journal of Animal Breeding and Genetics, 2008, 125, 228-233.	2.0	13
102	Genetic assessment, illegal trafficking and management of the Mediterranean spur-thighed tortoise in Southern Spain and Northern Africa. Conservation Genetics, 2011, 12, 1-13.	1.5	13
103	Cloning and sequencing of the cDNA encoding goat Î ² -lactoglobulin. Journal of Animal Science, 1993, 71, 2832.	0.5	12
104	Exploratory Study on the Transcriptional Profile of Pigs Subclinically Infected with Porcine Circovirus Type 2. Animal Biotechnology, 2009, 20, 96-109.	1.5	12
105	Development of a PCR technique specific for Demodex injai in biological specimens. Parasitology Research, 2013, 112, 3369-3372.	1.6	12
106	Short communication: Milk microbiota profiling on water buffalo with full-length 16S rRNA using nanopore sequencing. Journal of Dairy Science, 2020, 103, 2693-2700.	3.4	12
107	Gene transfer technology in aquaculture. Hydrobiologia, 2000, 420, 91-94.	2.0	11
108	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.	1.7	11

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109	Adh expression in species of themulleri subgroup ofDrosophila. Biochemical Genetics, 1987, 25, 729-738.	1.7	10
110	Isolation, sequencing and relative quantitation by fluorescent-ratio PCR of feline β-lactoglobulin I, II, and III cDNAs. Mammalian Genome, 1999, 10, 560-564.	2.2	10
111	Assignment of the 2,4-dienoyl-CoA reductase (DECR) gene to porcine chromosome 4. Animal Genetics, 2002, 33, 164-165.	1.7	10
112	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.	2.0	10
113	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.	2.0	10
114	African swine fever virus does not express viral microRNAs in experimentally infected pigs. BMC Veterinary Research, 2018, 14, 268.	1.9	10
115	Chromatin Structures of Goat and Sheep β-Lactoglobulin Gene Differ. Biochemical and Biophysical Research Communications, 1998, 252, 649-653.	2.1	9
116	Genetic analysis of SLC11A1 polymorphisms in multiple sclerosis patients. Multiple Sclerosis Journal, 2004, 10, 618-620.	3.0	9
117	Microarray analysis of mediastinal lymph node of pigs naturally affected by postweaning multisystemic wasting syndrome. Virus Research, 2012, 165, 134-142.	2.2	9
118	Non-synonymous genetic variation in exonic regions of canine Toll-like receptors. Canine Genetics and Epidemiology, 2014, 1, 11.	2.8	9
119	Expression of caprine beta-lactoglobulin in the milk of transgenic mice. Transgenic Research, 1997, 6, 69-74.	2.4	8
120	DogMap: an international collaboration toward a low-resolution canine genetic marker map. , 1999, 90, 3-6.		8
121	Structural characterization of the caprine major histocompatibility complex class II DQB1 (Cahi-DQB1) gene. Molecular Immunology, 2004, 41, 843-846.	2.2	8
122	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.	1.7	8
123	Characterization and Physical Mapping of the PorcineCDS1andCDS2Genes. Animal Biotechnology, 2007, 18, 23-35.	1.5	8
124	Individual identification and genetic traceability in the pig using the SNPlex TM genotyping system. Animal Genetics, 2007, 38, 663-665.	1.7	8
125	Sequence Analysis of Goat Major Histocompatibility Complex Class I Genes. Journal of Dairy Science, 2008, 91, 814-817.	3.4	8
126	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

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127	Genetic polymorphism of the K-casein (CSN3) gene in goats reared in Southern Italy. Italian Journal of Animal Science, 2005, 4, 97-101.	1.9	8
128	Empirical Bayes factor analyses of quantitative trait loci for gestation length in Iberian × Meishan F2 sows. Animal, 2008, 2, 177-183.	3.3	6
129	An age-dependent association between a leptin C3469T single nucleotide polymorphism and intramuscular fat content in pigs. Livestock Science, 2009, 121, 335-338.	1.6	6
130	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.	3.0	6
131	Whole genome sequencing identifies allelic ratio distortion in sperm involving genes related to spermatogenesis in a swine model. DNA Research, 2020, 27, .	3.4	6
132	Assessing the levels of intraspecific admixture and interspecific hybridization in Iberian wild goats (<i>Capra pyrenaica</i>). Evolutionary Applications, 2021, 14, 2618-2634.	3.1	6
133	Linkage mapping of the porcine hairless gene (HR  ) to chromosome 14. Animal Genetics, 2003, 34, 317-318.	1.7	5
134	Mapping of the goat stearoyl coenzyme A desaturase gene to chromosome 26. Animal Genetics, 2003, 34, 474-475.	1.7	5
135	Assignment of the microsomal triglyceride transfer protein large subunit (MTP) gene to porcine chromosome 8. Animal Genetics, 2005, 36, 354-355.	1.7	5
136	Quantitative trait loci for fatness at growing and reproductive stages in Iberianâ€f×â€fMeishan F ₂ sows. Animal Genetics, 2011, 42, 548-551.	1.7	5
137	Identification of protein-damaging mutations in 10 swine taste receptors and 191 appetite-reward genes. BMC Genomics, 2016, 17, 685.	2.8	5
138	Skin mites in mice (Mus musculus): high prevalence of Myobia sp. (Acari, Arachnida) in Robertsonian mice. Parasitology Research, 2018, 117, 2139-2148.	1.6	5
139	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.	1.5	4
140	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.	2.2	4
141	Primer-directed synthesis of a molecular weight marker. Genetic Analysis, Techniques and Applications, 1996, 13, 147-149.	1.5	3
142	Physical and linkage mapping of the porcinepink-eye dilutiongene (OCA2). Animal Genetics, 2002, 33, 392-394.	1.7	3
143	Assignment of theacyl-CoA synthetase long-chain family member 4 (ACSL4)gene to porcine chromosome X. Animal Genetics, 2005, 36, 76-76.	1.7	3
144	Characterization of skin surface and dermal microbiota in dogs with mast cell tumor. Scientific Reports, 2020, 10, 12634.	3.3	3

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145	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.	2.5	3
146	Assignment of the beta-lactoglobulin (BLG) gene to porcine chromosome 1. Animal Genetics, 2005, 36, 356-358.	1.7	2
147	Nucleotide Sequence and Polymorphism of the Pig Acyl Coenzyme A Synthetase Long-Chain 1 (ACSL1) Gene. Animal Biotechnology, 2007, 18, 117-122.	1.5	2
148	Using haplotype probabilities in categorical survival analysis: a case study with three candidate genes in an Iberianâ€fĂ—â€fMeishan F ₂ population of newborn piglets. Journal of Animal Breeding and Genetics, 2008, 125, 5-12.	2.0	2
149	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.	10.0	2
150	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
151	Structural characterization of the porcine pyruvate carboxylase (PC) gene. Journal of Animal Breeding and Genetics, 2003, 120, 338-345.	2.0	1
152	Assignment of the <i>phospholipase C</i> l² <i>1</i> (<i>PLCB1</i>) gene to porcine chromosome 17. Animal Genetics, 2005, 36, 516-517.	1.7	1
153	Expression of Recombinant Human Follicle-Stimulating Hormone in the Mammary Gland of Transgenic Mice. Molecular Biotechnology, 2006, 34, 37-44.	2.4	0
154	Northern analysis of highly folded goat αs1 Casein mRNA. Genetic Analysis, Techniques and Applications, 1996, 12, 143-145.	1.5	0