

Jakub Cieslak

List of Publications by Year in descending order

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Version: 2024-02-01

26
papers

335
citations

840119

11
h-index

839053

18
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27
all docs

27
docs citations

27
times ranked

507
citing authors

#	ARTICLE	IF	CITATIONS
1	Genetics of fat tissue accumulation in pigs: a comparative approach. <i>Journal of Applied Genetics</i> , 2010, 51, 153-168.	1.0	88
2	Genetic diversity in Hucul and Polish primitive horse breeds. <i>Archives Animal Breeding</i> , 2015, 58, 23-31.	0.5	25
3	SNPs in the porcine PPARC1a gene: Interbreed differences and their phenotypic effects. <i>Cellular and Molecular Biology Letters</i> , 2007, 12, 231-9.	2.7	21
4	Screening for the Most Suitable Reference Genes for Gene Expression Studies in Equine Milk Somatic Cells. <i>PLoS ONE</i> , 2015, 10, e0139688.	1.1	16
5	Association studies on the porcine RETN, UCP1, UCP3 and ADRB3 genes polymorphism with fatness traits. <i>Meat Science</i> , 2009, 83, 551-554.	2.7	15
6	Polymorphisms in 5' flanking regions of genes encoding adiponectin, leptin, and resistin are not associated with obesity of Polish children and adolescents. <i>Molecular Biology Reports</i> , 2011, 38, 1793-1798.	1.0	15
7	Missense mutations and polymorphisms of the MC4R gene in Polish obese children and adolescents in relation to the relative body mass index. <i>Journal of Applied Genetics</i> , 2011, 52, 319-323.	1.0	14
8	Association of MC3R gene polymorphisms with body weight in the red fox and comparative gene organization in four canids. <i>Animal Genetics</i> , 2011, 42, 104-107.	0.6	13
9	Effect of three common SNPs in 5' flanking region of LEP and ADIPOQ genes on their expression in Polish obese children and adolescents. <i>Molecular Biology Reports</i> , 2012, 39, 3951-3955.	1.0	13
10	The pig <i>CART</i> (cocaine- and amphetamine-regulated transcript) gene and association of its microsatellite polymorphism with production traits. <i>Journal of Animal Breeding and Genetics</i> , 2009, 126, 37-42.	0.8	11
11	Three-Dimensional Arrangement of Genes Involved in Lipid Metabolism in Nuclei of Porcine Adipocytes and Fibroblasts in Relation to Their Transcription Level. <i>Cytogenetic and Genome Research</i> , 2012, 136, 295-302.	0.6	11
12	Common polymorphism (81Val>Ile) and rare mutations (257Arg>Ser and 335Ile>Ser) of the MC3R gene in obese Polish children and adolescents. <i>Molecular Biology Reports</i> , 2013, 40, 6893-6898.	1.0	11
13	Variability of lysozyme and lactoferrin bioactive protein concentrations in equine milk in relation to <i>LYZ</i> and <i>LTF</i> gene polymorphisms and expression. <i>Journal of the Science of Food and Agriculture</i> , 2017, 97, 2174-2181.	1.7	11
14	5' flanking variants of equine casein genes (CSN1S1, CSN1S2, CSN2, CSN3) and their relationship with gene expression and milk composition. <i>Journal of Applied Genetics</i> , 2019, 60, 71-78.	1.0	11
15	Genes encoding equine β -lactoglobulin (LGB1 and LGB2): Polymorphism, expression, and impact on milk composition. <i>PLoS ONE</i> , 2020, 15, e0232066.	1.1	10
16	Polymorphisms in the promoter region of the adiponectin (<i>ADIPOQ</i>) gene are presumably associated with transcription level and carcass traits in pigs. <i>Animal Genetics</i> , 2013, 44, 340-343.	0.6	8
17	Characterization of the Polish Primitive Horse (Konik) maternal lines using mitochondrial D-loop sequence variation. <i>PeerJ</i> , 2017, 5, e3714.	0.9	7
18	TBX3 and ASIP genotypes reveal discrepancies in officially recorded coat colors of Hucul horses. <i>Animal</i> , 2019, 13, 1811-1816.	1.3	5

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19	The equine graying with age mutation of the <i>STX17</i> gene: A copy number study using droplet digital PCR reveals a new pattern. <i>Animal Genetics</i> , 2021, 52, 223-227.	0.6	5
20	Identification of target sequences for association studies - analysis of the pig <i>FABP3</i> and <i>FABP4</i> loci using comparative genomics methods. <i>Journal of Animal and Feed Sciences</i> , 2008, 17, 191-201.	0.4	5
21	5'-flanking variants of the equine β -lactalbumin (<i>LALBA</i>) gene relationship with gene expression and mare's milk composition. <i>Journal of Animal and Feed Sciences</i> , 2018, 27, 317-326.	0.4	5
22	Characterization of equine <i>CSN1S2</i> variants considering genetics, transcriptomics, and proteomics. <i>Journal of Dairy Science</i> , 2016, 99, 1277-1285.	1.4	4
23	Between-breed variation in frequency of five novel missense SNPs in porcine <i>casein beta</i> (<i>CSN2</i>) and <i>casein kappa</i> (<i>CSN3</i>) genes. <i>Animal Genetics</i> , 2012, 43, 363-364.	0.6	3
24	Interbreed Distribution of the Myostatin (MSTN) Gene 5'-Flanking Variants and Their Relationship With Horse Biometric Traits. <i>Journal of Equine Veterinary Science</i> , 2018, 60, 83-89.e1.	0.4	3
25	Genetic Background of the Polish Primitive Horse (Konik) Coat Color Variation – New Insight into Dun Dilution Phenotypic Effect. <i>Journal of Heredity</i> , 2021, 112, 436-442.	1.0	2
26	Equine <i>STX17</i> intronic triplication confirmed by droplet digital PCR analysis of its breakpoints. <i>Animal Genetics</i> , 2021, 52, 567-568.	0.6	0