

Jakub Cieslak

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

26

papers

266

citations

10

h-index

15

g-index

27

ext. papers

296

ext. citations

2.9

avg, IF

2.77

L-index

#	Paper	IF	Citations
26	Equine STX17 intronic triplication confirmed by droplet digital PCR analysis of its breakpoints. <i>Animal Genetics</i> , 2021 , 52, 567-568	2.5	
25	The equine graying with age mutation of the STX17 gene: A copy number study using droplet digital PCR reveals a new pattern. <i>Animal Genetics</i> , 2021 , 52, 223-227	2.5	1
24	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	2.4	1
23	Genes encoding equine β -lactoglobulin (LGB1 and LGB2): Polymorphism, expression, and impact on milk composition. <i>PLoS ONE</i> , 2020 , 15, e0232066	3.7	5
22	and genotypes reveal discrepancies in officially recorded coat colors of Hucul horses. <i>Animal</i> , 2019 , 13, 1811-1816	3.1	3
21	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	2.5	5
20	5'-flanking variants of the equine β -lactalbumin (LALBA) gene [relationship with gene expression and mare's milk composition. <i>Journal of Animal and Feed Sciences</i> , 2018 , 27, 317-326	1.5	4
19	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	1.2	1
18	Variability of lysozyme and lactoferrin bioactive protein concentrations in equine milk in relation to LYZ and LTF gene polymorphisms and expression. <i>Journal of the Science of Food and Agriculture</i> , 2017 , 97, 2174-2181	4.3	7
17	Characterization of the Polish Primitive Horse (Konik) maternal lines using mitochondrial D-loop sequence variation. <i>PeerJ</i> , 2017 , 5, e3714	3.1	5
16	Characterization of equine CSN1S2 variants considering genetics, transcriptomics, and proteomics. <i>Journal of Dairy Science</i> , 2016 , 99, 1277-1285	4	4
15	Screening for the Most Suitable Reference Genes for Gene Expression Studies in Equine Milk Somatic Cells. <i>PLoS ONE</i> , 2015 , 10, e0139688	3.7	13
14	Genetic diversity in Hucul and Polish primitive horse breeds. <i>Archives Animal Breeding</i> , 2015 , 58, 23-31	1.6	18
13	Polymorphisms in the promoter region of the adiponectin (ADIPOQ) gene are presumably associated with transcription level and carcass traits in pigs. <i>Animal Genetics</i> , 2013 , 44, 340-3	2.5	7
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, 6893-8	2.8	10
11	Between-breed variation in frequency of five novel missense SNPs in porcine casein beta (CSN2) and casein kappa (CSN3) genes. <i>Animal Genetics</i> , 2012 , 43, 363-4	2.5	1
10	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, 3951-5	2.8	11

9	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	1.9	11
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, 104-7	2.5	12
7	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, 1793-8	2.8	14
6	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, 319-23	2.5	12
5	Genetics of fat tissue accumulation in pigs: a comparative approach. <i>Journal of Applied Genetics</i> , 2010 , 51, 153-68	2.5	74
4	The pig CART (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	2.9	10
3	Association studies on the porcine RETN, UCP1, UCP3 and ADRB3 genes polymorphism with fatness traits. <i>Meat Science</i> , 2009 , 83, 551-4	6.4	13
2	Identification of target sequences for association studies - analysis of the pig FABP3 and FABP4 loci using comparative genomics methods. <i>Journal of Animal and Feed Sciences</i> , 2008 , 17, 191-201	1.5	4
1	SNPs in the porcine PPARGC1a gene: interbreed differences and their phenotypic effects. <i>Cellular and Molecular Biology Letters</i> , 2007 , 12, 231-9	8.1	20