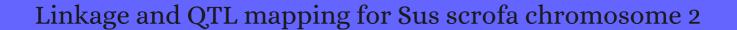
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#	Paper	IF	Citations
47	Polymorphism detection of porcine PSMC3, PSMC6 and PSMD3 genes and their association with partial growth, carcass traits, meat quality and immune traits. <i>Canadian Journal of Animal Science</i> , 2005 , 85, 475-480	0.9	5
46	Evolutionary breakpoints through a high-resolution comparative map between porcine chromosomes 2 and 16 and human chromosomes. <i>Genomics</i> , 2006 , 88, 504-12	4.3	12
45	Detection of quantitative trait loci associated with several internal organ traits and teat number trait in a pig population. <i>Journal of Genetics and Genomics</i> , 2007 , 34, 307-14	4	12
44	QTL for the heritable inverted teat defect in pigs. <i>Mammalian Genome</i> , 2008 , 19, 127-38	3.2	24
43	Quantitative trait loci for chemical body composition traits in pigs and their positional associations with body tissues, growth and feed intake. <i>Animal Genetics</i> , 2008 , 39, 130-40	2.5	34
42	Detection of quantitative trait loci for reproduction and production traits in Large White and French Landrace pig populations. <i>Genetics Selection Evolution</i> , 2008 , 40, 61-78	4.9	14
41	Porcine TEF1 and RTEF1: molecular characterization and association analyses with growth traits. <i>Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology</i> , 2008 , 150, 447-53	2.3	14
40	Single nucleotide polymorphisms in several porcine cathepsin genes are associated with growth, carcass, and production traits in Italian Large White pigs. <i>Journal of Animal Science</i> , 2008 , 86, 3300-14	0.7	46
39	A quantitative trait locus genome scan for porcine muscle fiber traits reveals overdominance and epistasis. <i>Journal of Animal Science</i> , 2008 , 86, 3290-9	0.7	31
38	Using microarrays to identify positional candidate genes for QTL: the case study of ACTH response in pigs. <i>BMC Proceedings</i> , 2009 , 3 Suppl 4, S14	2.3	8
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36	Genome-wide QTL mapping for three traits related to teat number in a White Duroc x Erhualian pig resource population. <i>BMC Genetics</i> , 2009 , 10, 6	2.6	31
35	Investigation of LDHA and COPB1 as candidate genes for muscle development in the MYOD1 region of pig chromosome 2. <i>Molecular Biology Reports</i> , 2010 , 37, 629-36	2.8	19
34	Epistatic QTL pairs associated with meat quality and carcass composition traits in a porcine Duroc Deletrain population. <i>Genetics Selection Evolution</i> , 2010 , 42, 39	4.9	15
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32	Polymorphism screening and mapping of nine meat performance-related genes in the pig. <i>Animal Genetics</i> , 2010 , 41, 334-5	2.5	2
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30	Quantitative trait loci analysis of swine meat quality traits. <i>Journal of Animal Science</i> , 2010 , 88, 2904-12	0.7	21
29	Epistatic analysis of carcass characteristics in pigs reveals genomic interactions between quantitative trait loci attributable to additive and dominance genetic effects. <i>Journal of Animal Science</i> , 2010 , 88, 2219-34	0.7	10
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27	Transcriptome analysis to identify differential gene expression affecting meat quality in heavy Italian pigs. <i>Animal Genetics</i> , 2011 , 42, 161-71	2.5	3
26	Epistatic quantitative trait loci affecting chemical body composition and deposition as well as feed intake and feed efficiency throughout the entire growth period of pigs. <i>Livestock Science</i> , 2011 , 138, 34-48	1.7	3
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11	Polymorphisms and expression analysis of SOX-6 in relation to porcine growth, carcass, and meat quality traits. <i>Meat Science</i> , 2015 , 107, 26-32	6.4	5
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2	A genome-wide association study to detect QTL for commercially important traits in Swiss Large White boars. <i>PLoS ONE</i> , 2013 , 8, e55951	3.7	29
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