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List of Publications by Year in descending order

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Version: 2024-02-01

21
papers

468
citations

932766

10
h-index

713013

21
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24
all docs

24
docs citations

24
times ranked

1306
citing authors

#	ARTICLE	IF	CITATIONS
1	CD9 expression in porcine blood CD4+ T cells delineates two subsets with phenotypic characteristics of central and effector memory cells. <i>Developmental and Comparative Immunology</i> , 2022, 133, 104431.	1.0	1
2	<i>Salmonella Typhimurium</i> Infection Along the Porcine Gastrointestinal Tract and Associated Lymphoid Tissues. <i>Veterinary Pathology</i> , 2019, 56, 681-690.	0.8	14
3	Identification and functional characterization of polymorphisms in promoter sequences of porcine NOD1 and NOD2 genes. <i>Research in Veterinary Science</i> , 2019, 124, 310-316.	0.9	1
4	Comparative proteomic analysis reveals different responses in porcine lymph nodes to virulent and attenuated homologous African swine fever virus strains. <i>Veterinary Research</i> , 2018, 49, 90.	1.1	14
5	Early <i>Salmonella Typhimurium</i> infection in pigs disrupts Microbiome composition and functionality principally at the ileum mucosa. <i>Scientific Reports</i> , 2018, 8, 7788.	1.6	61
6	Comparative Proteomics Reveals Differences in Host-Pathogen Interaction between Infectious and Commensal Relationship with <i>Campylobacter jejuni</i> . <i>Frontiers in Cellular and Infection Microbiology</i> , 2017, 7, 145.	1.8	11
7	Live attenuated African swine fever viruses as ideal tools to dissect the mechanisms involved in viral pathogenesis and immune protection. <i>Veterinary Research</i> , 2015, 46, 135.	1.1	74
8	Interaction between <i>Campylobacter</i> and intestinal epithelial cells leads to a different proinflammatory response in human and porcine host. <i>Veterinary Immunology and Immunopathology</i> , 2014, 162, 14-23.	0.5	12
9	Molecular cloning, characterization and gene expression of the full length cDNA encoding the porcine CD11b(±M) and chromosomal localization of the porcine CD11a(±L) and CD11b(±M) and CD11b(±D) gene clusters. <i>Veterinary Immunology and Immunopathology</i> , 2012, 145, 505-510.	0.5	2
10	Two cDNAs coding for the porcine CD51 (±v) integrin subunit: Cloning, expression analysis, adhesion assays and chromosomal localization. <i>Gene</i> , 2011, 481, 29-40.	1.0	6
11	Immunohistochemical distribution of the tetraspanin CD9 in normal porcine tissues. <i>Molecular Biology Reports</i> , 2011, 38, 1021-1028.	1.0	7
12	Gene expression pattern in swine neutrophils after lipopolysaccharide exposure: a time course comparison. <i>BMC Proceedings</i> , 2011, 5, S11.	1.8	3
13	Methods for interpreting lists of affected genes obtained in a DNA microarray experiment. <i>BMC Proceedings</i> , 2009, 3, S5.	1.8	29
14	Biological pathway analysis by ArrayUnlock and Ingenuity Pathway Analysis. <i>BMC Proceedings</i> , 2009, 3, S6.	1.8	135
15	Selection of housekeeping genes for normalization by real-time RT-PCR: Analysis of Or-MYB1 gene expression in <i>Orobanche ramosa</i> development. <i>Analytical Biochemistry</i> , 2008, 379, 176-181.	1.1	46
16	Molecular characterization and expression analysis of the gene coding for the porcine β 23 integrin subunit (CD61). <i>Gene</i> , 2008, 408, 9-17.	1.0	11
17	Molecular cloning, chromosomal location, and expression analysis of porcine CD14. <i>Developmental and Comparative Immunology</i> , 2007, 31, 738-747.	1.0	12
18	Analysis of a simulated microarray dataset: Comparison of methods for data normalisation and detection of differential expression (Open Access publication). <i>Genetics Selection Evolution</i> , 2007, 39, 669.	1.2	5

#	ARTICLE	IF	CITATIONS
19	Localization of porcine CD29 transcripts and protein in pig cells and tissues by RT-PCR and immunohistochemistry. <i>Veterinary Immunology and Immunopathology</i> , 2005, 104, 281-288.	0.5	6
20	Molecular cloning, expression pattern and chromosomal mapping of pig CD9 antigen. <i>Cytogenetic and Genome Research</i> , 2003, 101, 143-146.	0.6	7
21	A polymorphic microsatellite located on pig chromosome band 12p11-2/3p13, within the 3' UTR of the ITGB3 gene. <i>Animal Genetics</i> , 2002, 33, 239-240.	0.6	2