

Doreen Becker

List of Publications by Citations

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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

478

citations

11

h-index

21

g-index

34

ext. papers

606

ext. citations

3.7

avg, IF

2.92

L-index

#	Paper	IF	Citations
26	A missense mutation in the SERPINH1 gene in Dachshunds with osteogenesis imperfecta. <i>PLoS Genetics</i> , 2009 , 5, e1000579	6	102
25	A comprehensive biomedical variant catalogue based on whole genome sequences of 582 dogs and eight wolves. <i>Animal Genetics</i> , 2019 , 50, 695-704	2.5	64
24	Microphthalmia in Texel sheep is associated with a missense mutation in the paired-like homeodomain 3 (PITX3) gene. <i>PLoS ONE</i> , 2010 , 5, e8689	3.7	43
23	A missense change in the ATG4D gene links aberrant autophagy to a neurodegenerative vacuolar storage disease. <i>PLoS Genetics</i> , 2015 , 11, e1005169	6	37
22	A mutation in the FAM83G gene in dogs with hereditary footpad hyperkeratosis (HFH). <i>PLoS Genetics</i> , 2014 , 10, e1004370	6	37
21	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
20	A deletion in the N-myc downstream regulated gene 1 (NDRG1) gene in Greyhounds with polyneuropathy. <i>PLoS ONE</i> , 2010 , 5, e11258	3.7	28
19	The brown coat colour of Coppernecked goats is associated with a non-synonymous variant at the TYRP1 locus on chromosome 8. <i>Animal Genetics</i> , 2015 , 46, 50-4	2.5	27
18	An ARHGEF10 deletion is highly associated with a juvenile-onset inherited polyneuropathy in Leonberger and Saint Bernard dogs. <i>PLoS Genetics</i> , 2014 , 10, e1004635	6	20
17	Wattles in goats are associated with the FMN1/GREM1 region on chromosome 10. <i>Animal Genetics</i> , 2015 , 46, 316-20	2.5	18
16	A GJA9 frameshift variant is associated with polyneuropathy in Leonberger dogs. <i>BMC Genomics</i> , 2017 , 18, 662	4.5	17
15	Complex Structural Variant Associated with Non-syndromic Canine Retinal Degeneration. <i>G3: Genes, Genomes, Genetics</i> , 2019 , 9, 425-437	3.2	10
14	Genome-wide association study and whole-genome sequencing identify a deletion in LRIT3 associated with canine congenital stationary night blindness. <i>Scientific Reports</i> , 2019 , 9, 14166	4.9	9
13	Canine NAPEPLD-associated models of human myelin disorders. <i>Scientific Reports</i> , 2018 , 8, 5818	4.9	8
12	Hepatic Transcriptome Analysis Identifies Divergent Pathogen-Specific Targeting-Strategies to Modulate the Innate Immune System in Response to Intramammary Infection. <i>Frontiers in Immunology</i> , 2020 , 11, 715	8.4	7
11	Molecular studies of phenotype variation in canine RPGR-XLPRA1. <i>Molecular Vision</i> , 2016 , 22, 319-31	2.3	7
10	Cows selected for divergent mastitis susceptibility display a differential liver transcriptome profile after experimental <i>Staphylococcus aureus</i> mammary gland inoculation. <i>Journal of Dairy Science</i> , 2020 , 103, 6364-6373	4	2

9	Candidate Genetic Modifiers for RPGR Retinal Degeneration 2020 , 61, 20		2
8	Coat Color Roan Shows Association with Variants and No Evidence of Lethality in Icelandic Horses. <i>Genes</i> , 2020 , 11,	4.2	1
7	A genome-wide significant association on chromosome 15 for congenital entropion in Swiss White Alpine sheep. <i>Animal Genetics</i> , 2020 , 51, 278-283	2.5	1
6	A variant in MYO10 is associated with hind limb conformation in Swiss Large White boars. <i>Animal Genetics</i> , 2014 , 45, 308	2.5	1
5	Single-cell RNA sequencing of freshly isolated bovine milk cells and cultured primary mammary epithelial cells. <i>Scientific Data</i> , 2021 , 8, 177	8.2	1
4	A 50-kb deletion disrupting the RSPO2 gene is associated with tetradysmelia in Holstein Friesian cattle. <i>Genetics Selection Evolution</i> , 2020 , 52, 68	4.9	0
3	CCDC66 frameshift variant associated with a new form of early-onset progressive retinal atrophy in Portuguese Water Dogs. <i>Scientific Reports</i> , 2020 , 10, 21162	4.9	0
2	Optic nerve hypoplasia in miniature poodle dogs: A preliminary genetic and candidate gene association study. <i>Veterinary Ophthalmology</i> , 2020 , 23, 67-76	1.4	0
1	Exploring the Origin and Relatedness of Maternal Lineages Through Analysis of Mitochondrial DNA in the Holstein Horse. <i>Frontiers in Genetics</i> , 2021 , 12, 632500	4.5	0