

# Doreen Becker

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7384437/publications.pdf>

Version: 2024-02-01

31  
papers

703  
citations

686830

13  
h-index

552369

26  
g-index

34  
all docs

34  
docs citations

34  
times ranked

1091  
citing authors

#	ARTICLE	IF	CITATIONS
1	Mitochondrial DNA Variation Contributes to the Aptitude for Dressage and Show Jumping Ability in the Holstein Horse Breed. <i>Animals</i> , 2022, 12, 704.	1.0	1
2	An Overview of the Importance and Value of Porcine Species in Sialic Acid Research. <i>Biology</i> , 2022, 11, 903.	1.3	4
3	Canine reference genome accuracy impacts variant calling: Lessons learned from investigating embryonic lethal variants. <i>Animal Genetics</i> , 2022, 53, 706-708.	0.6	1
4	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.	1.1	2
5	Single-cell RNA sequencing of freshly isolated bovine milk cells and cultured primary mammary epithelial cells. <i>Scientific Data</i> , 2021, 8, 177.	2.4	7
6	Optic nerve hypoplasia in miniature poodle dogs: A preliminary genetic and candidate gene association study. <i>Veterinary Ophthalmology</i> , 2020, 23, 67-76.	0.6	1
7	A 50-kb deletion disrupting the <i>RSPO2</i> gene is associated with tetradysmelia in Holstein Friesian cattle. <i>Genetics Selection Evolution</i> , 2020, 52, 68.	1.2	4
8	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.	1.6	7
9	Candidate Genetic Modifiers for RPGR Retinal Degeneration. , 2020, 61, 20.		7
10	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.	2.2	15
11	Coat Color Roan Shows Association with <i>KIT</i> Variants and No Evidence of Lethality in Icelandic Horses. <i>Genes</i> , 2020, 11, 680.	1.0	4
12	A genome-wide significant association on chromosome 15 for congenital entropion in Swiss White Alpine sheep. <i>Animal Genetics</i> , 2020, 51, 278-283.	0.6	2
13	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.	1.4	5
14	A comprehensive biomedical variant catalogue based on whole genome sequences of 582 dogs and eight wolves. <i>Animal Genetics</i> , 2019, 50, 695-704.	0.6	138
15	Genome-wide association study and whole-genome sequencing identify a deletion in <i>LRR13</i> associated with canine congenital stationary night blindness. <i>Scientific Reports</i> , 2019, 9, 14166.	1.6	15
16	Complex Structural <i>PPT1</i> Variant Associated with Non-syndromic Canine Retinal Degeneration. <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 425-437.	0.8	13
17	Canine <i>NAPEPLD</i> -associated models of human myelin disorders. <i>Scientific Reports</i> , 2018, 8, 5818.	1.6	14
18	A <i>GJA9</i> frameshift variant is associated with polyneuropathy in Leonberger dogs. <i>BMC Genomics</i> , 2017, 18, 662.	1.2	20

#	ARTICLE	IF	CITATIONS
19	Molecular studies of phenotype variation in canine RPGR-XLPRA1. <i>Molecular Vision</i> , 2016, 22, 319-31.	1.1	8
20	Wattles in goats are associated with the <i>FMN1</i> / <i>GREM1</i> region on chromosome 10. <i>Animal Genetics</i> , 2015, 46, 316-320.	0.6	27
21	The brown coat colour of Coppernecked goats is associated with a non-synonymous variant at the <i>TYRP1</i> locus on chromosome 8. <i>Animal Genetics</i> , 2015, 46, 50-54.	0.6	42
22	A Missense Change in the <i>ATG4D</i> Gene Links Aberrant Autophagy to a Neurodegenerative Vacuolar Storage Disease. <i>PLoS Genetics</i> , 2015, 11, e1005169.	1.5	48
23	A Mutation in the <i>FAM83G</i> Gene in Dogs with Hereditary Footpad Hyperkeratosis (HFH). <i>PLoS Genetics</i> , 2014, 10, e1004370.	1.5	43
24	An <i>ARHGEF10</i> Deletion Is Highly Associated with a Juvenile-Onset Inherited Polyneuropathy in Leonberger and Saint Bernard Dogs. <i>PLoS Genetics</i> , 2014, 10, e1004635.	1.5	28
25	A variant in <i>MYO10</i> is associated with hind limb conformation in Swiss Large White boars. <i>Animal Genetics</i> , 2014, 45, 308-308.	0.6	2
26	A Genome-Wide Association Study to Detect QTL for Commercially Important Traits in Swiss Large White Boars. <i>PLoS ONE</i> , 2013, 8, e55951.	1.1	35
27	P3.24 A deletion in <i>ARHGEF10</i> is highly associated with early onset inherited polyneuropathy in Leonberger and St. Bernard dogs. <i>Neuromuscular Disorders</i> , 2011, 21, 689.	0.3	0
28	Microphthalmia in Texel Sheep Is Associated with a Missense Mutation in the Paired-Like Homeodomain 3 ( <i>PITX3</i> ) Gene. <i>PLoS ONE</i> , 2010, 5, e8689.	1.1	52
29	A Deletion in the N-Myc Downstream Regulated Gene 1 ( <i>NDRG1</i> ) Gene in Greyhounds with Polyneuropathy. <i>PLoS ONE</i> , 2010, 5, e11258.	1.1	33
30	A Missense Mutation in the <i>SERPINH1</i> Gene in Dachshunds with Osteogenesis Imperfecta. <i>PLoS Genetics</i> , 2009, 5, e1000579.	1.5	115
31	Roan coat color in livestock. <i>Animal Genetics</i> , 0, , .	0.6	2