

Danika L Bannasch

List of Publications by Year in descending order

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Version: 2024-02-01

98
papers

4,751
citations

136740

32
h-index

106150

65
g-index

105
all docs

105
docs citations

105
times ranked

5092
citing authors

#	ARTICLE	IF	CITATIONS
1	The Effects of FGF4 Retrogenes on Canine Morphology. <i>Genes</i> , 2022, 13, 325.	1.0	7
2	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
3	SLC25A12 Missense Variant in Nova Scotia Duck Tolling Retrievers Affected by Cerebellar Degenerationâ€”Myositis Complex (CDMC). <i>Genes</i> , 2022, 13, 1223.	1.0	2
4	Special Issue â€œMolecular Basis of Inherited Diseases in Companion Animalsâ€• <i>Genes</i> , 2021, 12, 68.	1.0	0
5	Canine DVL2 variant contributes to brachycephalic phenotype and caudal vertebral anomalies. <i>Human Genetics</i> , 2021, 140, 1535-1545.	1.8	9
6	Variants at the ASIP locus contribute to coat color darkening in Nellore cattle. <i>Genetics Selection Evolution</i> , 2021, 53, 40.	1.2	25
7	Improving the resolution of canine genome-wide association studies using genotype imputation: A study of two breeds. <i>Animal Genetics</i> , 2021, 52, 703-713.	0.6	5
8	Dog colour patterns explained by modular promoters of ancient canid origin. <i>Nature Ecology and Evolution</i> , 2021, 5, 1415-1423.	3.4	24
9	The effect of inbreeding, body size and morphology on health in dog breeds. <i>Canine Medicine and Genetics</i> , 2021, 8, 12.	1.4	26
10	Current Understanding of the Genetics of Intervertebral Disc Degeneration. <i>Frontiers in Veterinary Science</i> , 2020, 7, 431.	0.9	28
11	Multiple FGF4 Retrocopies Recently Derived within Canids. <i>Genes</i> , 2020, 11, 839.	1.0	12
12	A Missense Variant in ALDH5A1 Associated with Canine Succinic Semialdehyde Dehydrogenase Deficiency (SSADHD) in the Saluki Dog. <i>Genes</i> , 2020, 11, 1033.	1.0	3
13	ATP2A2 SINE Insertion in an Irish Terrier with Darier Disease and Associated Infundibular Cyst Formation. <i>Genes</i> , 2020, 11, 481.	1.0	5
14	Genetic Variants Affecting Skeletal Morphology in Domestic Dogs. <i>Trends in Genetics</i> , 2020, 36, 598-609.	2.9	15
15	Novel Brown Coat Color (Cocoa) in French Bulldogs Results from a Nonsense Variant in HPS3. <i>Genes</i> , 2020, 11, 636.	1.0	6
16	Quality of DNA extracted from formalin-fixed, paraffin-embedded canine tissues. <i>Journal of Veterinary Diagnostic Investigation</i> , 2020, 32, 556-559.	0.5	2
17	Quantitative Translation of Dog-to-Human Aging by Conserved Remodeling of the DNA Methylome. <i>Cell Systems</i> , 2020, 11, 176-185.e6.	2.9	67
18	A Missense Variant Affecting the C-Terminal Tail of UNC93B1 in Dogs with Exfoliative Cutaneous Lupus Erythematosus (ECLE). <i>Genes</i> , 2020, 11, 159.	1.0	13

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19	Pigment Intensity in Dogs is Associated with a Copy Number Variant Upstream of KITLG. <i>Genes</i> , 2020, 11, 75.	1.0	27
20	Whole genome sequencing for mutation discovery in a single case of lysosomal storage disease (MPS) Tj ETQq0 0 Q rgBT /Overlock 10 T	1.6	7
21	Genetic Tests for Large Animals. , 2020, , 1709-1716.e5.		0
22	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
23	NME5 frameshift variant in Alaskan Malamutes with primary ciliary dyskinesia. <i>PLoS Genetics</i> , 2019, 15, e1008378.	1.5	21
24	Pathologic Features of the Intervertebral Disc in Young Nova Scotia Duck Tolling Retrievers Confirms Chondrodystrophy Degenerative Phenotype Associated With Genotype. <i>Veterinary Pathology</i> , 2019, 56, 895-902.	0.8	13
25	Phenotypic Effects of FGF4 Retrogenes on Intervertebral Disc Disease in Dogs. <i>Genes</i> , 2019, 10, 435.	1.0	33
26	Evaluation of the major histocompatibility complex (MHC) class II as a candidate for sudden acquired retinal degeneration syndrome (SARDS) in Dachshunds. <i>Veterinary Ophthalmology</i> , 2019, 22, 751-759.	0.6	9
27	Comparative Cytogenetic Analysis of Dog and Human Choroid Plexus Tumors Defines Syntenic Regions of Genomic Loss. <i>Journal of Neuropathology and Experimental Neurology</i> , 2018, 77, 413-419.	0.9	4
28	Genetic analysis of optic nerve head coloboma in the Nova Scotia Duck Tolling Retriever identifies discordance with the <i><sc>NHEJ</sc>1</i> intronic deletion (collie eye anomaly mutation). <i>Veterinary Ophthalmology</i> , 2018, 21, 144-150.	0.6	7
29	Identification of a Candidate Mutation in the COL1A2 Gene of a Chow Chow With Osteogenesis Imperfecta. <i>Journal of Heredity</i> , 2018, 109, 308-314.	1.0	13
30	Whole genome variant association across 100 dogs identifies a frame shift mutation in DISHEVELLED 2 which contributes to Robinow-like syndrome in Bulldogs and related screw tail dog breeds. <i>PLoS Genetics</i> , 2018, 14, e1007850.	1.5	61
31	A Missense Mutation in the Vacuolar Protein Sorting 11 (<i>VPS11</i>) Gene Is Associated with Neuroaxonal Dystrophy in Rottweiler Dogs. <i>G3: Genes, Genomes, Genetics</i> , 2018, 8, 2773-2780.	0.8	19
32	Association of cancer-related mortality, age and gonadectomy in golden retriever dogs at a veterinary academic center (1989-2016). <i>PLoS ONE</i> , 2018, 13, e0192578.	1.1	45
33	Association of early onset myasthenia gravis in Newfoundland dogs with the canine major histocompatibility complex class I. <i>Neuromuscular Disorders</i> , 2017, 27, 409-416.	0.3	15
34	<i>FGF4</i> retrogene on CFA12 is responsible for chondrodystrophy and intervertebral disc disease in dogs. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 11476-11481.	3.3	92
35	Correlation of neuter status and expression of heritable disorders. <i>Canine Genetics and Epidemiology</i> , 2017, 4, 6.	2.9	29
36	Developing a 670k genotyping array to tag ~2M SNPs across 24 horse breeds. <i>BMC Genomics</i> , 2017, 18, 565.	1.2	116

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37	Gonadectomy effects on the risk of immune disorders in the dog: a retrospective study. <i>BMC Veterinary Research</i> , 2016, 12, 278.	0.7	45
38	Transcriptome profiling of equine vitamin E deficient neuroaxonal dystrophy identifies upregulation of liver X receptor target genes. <i>Free Radical Biology and Medicine</i> , 2016, 101, 261-271.	1.3	33
39	Serum levels of innate immunity cytokines are elevated in dogs with metaphyseal osteopathy (hypertrophic osteodystrophy) during active disease and remission. <i>Veterinary Immunology and Immunopathology</i> , 2016, 179, 32-35.	0.5	16
40	Chromosomal Aberrations in Canine Gliomas Define Candidate Genes and Common Pathways in Dogs and Humans. <i>Journal of Neuropathology and Experimental Neurology</i> , 2016, 75, 700-710.	0.9	31
41	Utilizing the Dog Genome in the Search for Novel Candidate Genes Involved in Glioma Developmentâ€”Genome Wide Association Mapping followed by Targeted Massive Parallel Sequencing Identifies a Strongly Associated Locus. <i>PLoS Genetics</i> , 2016, 12, e1006000.	1.5	54
42	Blood and Cerebrospinal Fluid Î±â€”Tocopherol and Selenium Concentrations in Neonatal Foals with Neuroaxonal Dystrophy. <i>Journal of Veterinary Internal Medicine</i> , 2015, 29, 1667-1675.	0.6	26
43	DNA Sequence Variants in the Five Prime Untranslated Region of the Cyclooxygenase-2 Gene Are Commonly Found in Healthy Dogs and Gray Wolves. <i>PLoS ONE</i> , 2015, 10, e0133127.	1.1	2
44	Genome-Wide Association Studies in Dogs and Humans Identify ADAMTS20 as a Risk Variant for Cleft Lip and Palate. <i>PLoS Genetics</i> , 2015, 11, e1005059.	1.5	82
45	Wolf Subspecies: Reply to Weckworth et al. and Fredrickson et al.: Table 1.. <i>Journal of Heredity</i> , 2015, 106, 417-419.	1.0	16
46	SERPINB11 Frameshift Variant Associated with Novel Hoof Specific Phenotype in Connemara Ponies. <i>PLoS Genetics</i> , 2015, 11, e1005122.	1.5	21
47	Thiamine Deficiencyâ€”Mediated Brain Mitochondrial Pathology in <i>A</i> askan Huskies with Mutation in <i>SLC19A3.1</i> . <i>Brain Pathology</i> , 2015, 25, 441-453.	2.1	27
48	Single Nucleotide Polymorphism (SNP) Variation of Wolves (<i>Canis lupus</i>) in Southeast Alaska and Comparison with Wolves, Dogs, and Coyotes in North America. <i>Journal of Heredity</i> , 2015, 106, 26-36.	1.0	35
49	Early Developmental and Evolutionary Origins of Gene Body DNA Methylation Patterns in Mammalian Placentas. <i>PLoS Genetics</i> , 2015, 11, e1005442.	1.5	93
50	A LINE-1 Insertion in DLX6 Is Responsible for Cleft Palate and Mandibular Abnormalities in a Canine Model of Pierre Robin Sequence. <i>PLoS Genetics</i> , 2014, 10, e1004257.	1.5	49
51	Risk of false positive genetic associations in complex traits with underlying population structure: A case study. <i>Veterinary Journal</i> , 2014, 202, 543-549.	0.6	27
52	A mutation in the canine gene encoding folliculinâ€”interacting protein 2 (FNIP2) associated with a unique disruption in spinal cord myelination. <i>Glia</i> , 2014, 62, 39-51.	2.5	10
53	Urate urolithiasis and hyperuricosuria in a Weimaraner, secondary to the SLC2A9 transporter defect. <i>Veterinary Record Case Reports</i> , 2014, 2, e000016.	0.1	1
54	Applied equine genetics. <i>Equine Veterinary Journal</i> , 2014, 46, 538-544.	0.9	27

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55	Clinical manifestations, response to treatment, and clinical outcome for Weimaraners with hypertrophic osteodystrophy: 53 cases (2009–2011). <i>Journal of the American Veterinary Medical Association</i> , 2013, 242, 1260-1266.	0.2	30
56	Pedigree Analysis and Exclusion of Alpha-Tocopherol Transfer Protein (<i>ATTPA</i>) as a Candidate Gene for Neuroaxonal Dystrophy in the American Quarter Horse. <i>Journal of Veterinary Internal Medicine</i> , 2013, 27, 177-185.	0.6	32
57	Prevalence of inherited disorders among mixed-breed and purebred dogs: 27,254 cases (1995–2010). <i>Journal of the American Veterinary Medical Association</i> , 2013, 242, 1549-1555.	0.2	119
58	Genetic Diversity in the Modern Horse Illustrated from Genome-Wide SNP Data. <i>PLoS ONE</i> , 2013, 8, e54997.	1.1	214
59	Genome-Wide Analysis Reveals Selection for Important Traits in Domestic Horse Breeds. <i>PLoS Genetics</i> , 2013, 9, e1003211.	1.5	240
60	Genome-Wide Association Mapping in Dogs Enables Identification of the Homeobox Gene, <i>NKX2-8</i> , as a Genetic Component of Neural Tube Defects in Humans. <i>PLoS Genetics</i> , 2013, 9, e1003646.	1.5	39
61	Genome-Wide Association Analysis Identifies a Mutation in the Thiamine Transporter 2 (<i>SLC19A3</i>) Gene Associated with Alaskan Husky Encephalopathy. <i>PLoS ONE</i> , 2013, 8, e57195.	1.1	40
62	A High Density SNP Array for the Domestic Horse and Extant Perissodactyla: Utility for Association Mapping, Genetic Diversity, and Phylogeny Studies. <i>PLoS Genetics</i> , 2012, 8, e1002451.	1.5	208
63	Activity, Expression and Genetic Variation of Canine β -Defensin 103: A Multifunctional Antimicrobial Peptide in the Skin of Domestic Dogs. <i>Journal of Innate Immunity</i> , 2012, 4, 248-259.	1.8	45
64	Electrophysiological studies in American Quarter horses with neuroaxonal dystrophy. <i>Veterinary Ophthalmology</i> , 2012, 15, 3-7.	0.6	13
65	Equine Degenerative Myeloencephalopathy in Lusitano Horses. <i>Journal of Veterinary Internal Medicine</i> , 2011, 25, 1439-1446.	0.6	34
66	Phylogenetic Distinctiveness of Middle Eastern and Southeast Asian Village Dog Y Chromosomes Illuminates Dog Origins. <i>PLoS ONE</i> , 2011, 6, e28496.	1.1	50
67	Examination of candidate genes for hypoadrenocorticism in Nova Scotia Duck Tolling Retrievers. <i>Veterinary Journal</i> , 2011, 187, 212-216.	0.6	17
68	Identification of Genomic Regions Associated with Phenotypic Variation between Dog Breeds using Selection Mapping. <i>PLoS Genetics</i> , 2011, 7, e1002316.	1.5	339
69	Genome-wide SNP and haplotype analyses reveal a rich history underlying dog domestication. <i>Nature</i> , 2010, 464, 898-902.	13.7	635
70	Genome-wide association mapping identifies multiple loci for a canine SLE-related disease complex. <i>Nature Genetics</i> , 2010, 42, 250-254.	9.4	99
71	Association of a dog leukocyte antigen class II haplotype with hypoadrenocorticism in Nova Scotia Duck Tolling Retrievers. <i>Tissue Antigens</i> , 2010, 75, 684-690.	1.0	37
72	Localization of Canine Brachycephaly Using an Across Breed Mapping Approach. <i>PLoS ONE</i> , 2010, 5, e9632.	1.1	101

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73	Validation of a urine test and characterization of the putative genetic mutation for hyperuricosuria in Bulldogs and Black Russian Terriers. <i>American Journal of Veterinary Research</i> , 2010, 71, 909-914.	0.3	21
74	Estimated Frequency of the Canine Hyperuricosuria Mutation in Different Dog Breeds. <i>Journal of Veterinary Internal Medicine</i> , 2010, 24, 1337-1342.	0.6	18
75	Changing Paradigms in Diagnosis of Inherited Defects Associated with Urolithiasis. <i>Veterinary Clinics of North America - Small Animal Practice</i> , 2009, 39, 111-125.	0.5	18
76	Evaluation of allele frequencies of inherited disease genes in subgroups of American Quarter Horses. <i>Journal of the American Veterinary Medical Association</i> , 2009, 234, 120-125.	0.2	100
77	Canine fibroblast growth factor receptor 3 sequence is conserved across dogs of divergent skeletal size. <i>BMC Genetics</i> , 2008, 9, 67.	2.7	5
78	Genetic Testing and the Future of Equine Genomics. <i>Journal of Equine Veterinary Science</i> , 2008, 28, 645-649.	0.4	6
79	SNPs in the Promoter Regions of the Canine RMRP and SHOX Genes are not Associated with Canine Chondrodysplasia. <i>Animal Biotechnology</i> , 2008, 19, 1-5.	0.7	8
80	Coyotes Demonstrate How Habitat Specialization by Individuals of a Generalist Species Can Diversify Populations in a Heterogeneous Ecoregion. <i>Molecular Biology and Evolution</i> , 2008, 25, 1384-1394.	3.5	65
81	Mutations in the SLC2A9 Gene Cause Hyperuricosuria and Hyperuricemia in the Dog. <i>PLoS Genetics</i> , 2008, 4, e1000246.	1.5	94
82	Clinical features and heritability of hypoadrenocorticism in Nova Scotia Duck Tolling Retrievers: 25 cases (1994-2006). <i>Journal of the American Veterinary Medical Association</i> , 2007, 231, 407-412.	0.2	41
83	Homozygosity mapping approach identifies a missense mutation in equine cyclophilin B (PPIB) associated with HERDA in the American Quarter Horse. <i>Genomics</i> , 2007, 90, 93-102.	1.3	99
84	Clinical and pathological findings in a HERDA-affected foal for 1.5 years of life. <i>Veterinary Dermatology</i> , 2007, 18, 36-40.	0.4	28
85	Evaluation of FOXC2 as a candidate gene for chronic progressive lymphedema in draft horses. <i>Veterinary Journal</i> , 2007, 174, 397-399.	0.6	19
86	Recent Advances in Small Animal Genetics. <i>Veterinary Clinics of North America - Small Animal Practice</i> , 2006, 36, 461-474.	0.5	3
87	Linkage analysis with an interbreed backcross maps Dalmatian hyperuricosuria to CFA03. <i>Mammalian Genome</i> , 2006, 17, 340-345.	1.0	16
88	Deletions in the COL10A1 gene are not associated with skeletal changes in dogs. <i>Mammalian Genome</i> , 2006, 17, 761-768.	1.0	6
89	Y chromosome haplotype analysis in purebred dogs. <i>Mammalian Genome</i> , 2005, 16, 273-280.	1.0	55
90	Exclusion of Urate Oxidase as a Candidate Gene for Hyperuricosuria in the Dalmatian Dog Using an Interbreed Backcross. <i>Journal of Heredity</i> , 2005, 96, 750-754.	1.0	19

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91	CanineCOL4A3andCOL4A4: Sequencing, mapping and genomic organization. DNA Sequence, 2005, 16, 241-251.	0.7	3
92	Evaluation of Canine COL4A3 and COL4A4 as Candidates for Familial Renal Disease in the Norwegian Elkhound. Journal of Heredity, 2005, 96, 739-744.	1.0	9
93	Inheritance of hereditary equine regional dermal asthenia in Quarter Horses. American Journal of Veterinary Research, 2005, 66, 437-442.	0.3	33
94	Inheritance of cricopharyngeal dysfunction in Golden Retrievers. American Journal of Veterinary Research, 2004, 65, 344-349.	0.3	16
95	Hereditary equine regional dermal asthenia ('hyperelastosis cutis') in 50 horses: clinical, histological, immunohistological and ultrastructural findings. Veterinary Dermatology, 2004, 15, 207-217.	0.4	56
96	Exclusion of galectin 9 as a candidate gene for hyperuricosuria in the Dalmatian dog. Animal Genetics, 2004, 35, 326-328.	0.6	7
97	Inheritance of urinary calculi in the Dalmatian. Journal of Veterinary Internal Medicine, 2004, 18, 483-7.	0.6	13
98	Melanocortin 1 receptor variation in the domestic dog. Mammalian Genome, 2000, 11, 24-30.	1.0	194