

Jeffrey J Schoenebeck

List of Publications by Year in descending order

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Version: 2024-02-01

37
papers

1,978
citations

535685

17
h-index

371746

37
g-index

41
all docs

41
docs citations

41
times ranked

3114
citing authors

#	ARTICLE	IF	CITATIONS
1	The impact of the COVID-19 pandemic on a cohort of Labrador retrievers in England. BMC Veterinary Research, 2022, 18, .	0.7	2
2	Canine reference genome accuracy impacts variant calling: Lessons learned from investigating embryonic lethal variants. Animal Genetics, 2022, 53, 706-708.	0.6	1
3	Exceptional Changes in Skeletal Anatomy under Domestication: The Case of Brachycephaly. Integrative Organismal Biology, 2021, 3, obab023.	0.9	15
4	Mining the 99 Lives Cat Genome Sequencing Consortium database implicates genes and variants for the <i>Ticked</i> locus in domestic cats (<i>Felis catus</i>). Animal Genetics, 2021, 52, 321-332.	0.6	9
5	A Highly Conserved Shh Enhancer Coordinates Hypothalamic and Craniofacial Development. Frontiers in Cell and Developmental Biology, 2021, 9, 595744.	1.8	3
6	Surveillance of a vomiting outbreak in dogs in the UK using owner-derived and internet search data. Veterinary Record, 2021, 189, e308.	0.2	1
7	Improving the resolution of canine genome-wide association studies using genotype imputation: A study of two breeds. Animal Genetics, 2021, 52, 703-713.	0.6	5
8	Dog colour patterns explained by modular promoters of ancient canid origin. Nature Ecology and Evolution, 2021, 5, 1415-1423.	3.4	24
9	From head to hind: Elucidating function through contrasting morphometrics of ancient and modern pedigree dogs. Anatomical Record, 2021, 304, 63-77.	0.8	4
10	More Than a Moggy; A Population Genetics Analysis of the United Kingdom's Non-Pedigree Cats. Genes, 2021, 12, 1619.	1.0	4
11	Characterisation of a second gain of function EDAR variant, encoding EDAR380R, in East Asia. European Journal of Human Genetics, 2020, 28, 1694-1702.	1.4	6
12	Is it time to stop sweeping data cleaning under the carpet? A novel algorithm for outlier management in growth data. PLoS ONE, 2020, 15, e0228154.	1.1	15
13	The ninth life of the cat reference genome, <i>Felis catus</i> . PLoS Genetics, 2020, 16, e1009045.	1.5	4
14	A comprehensive biomedical variant catalogue based on whole genome sequences of 582 dogs and eight wolves. Animal Genetics, 2019, 50, 695-704.	0.6	138
15	Network analysis of canine brain morphometry links tumour risk to oestrogen deficiency and accelerated brain ageing. Scientific Reports, 2019, 9, 12506.	1.6	11
16	Lactulose drives a reversible reduction and qualitative modulation of the faecal microbiota diversity in healthy dogs. Scientific Reports, 2019, 9, 13350.	1.6	11
17	Arginine to Glutamine Variant in Olfactomedin Like 3 (<i>OLFML3</i>) Is a Candidate for Severe Goniodysgenesis and Glaucoma in the Border Collie Dog Breed. G3: Genes, Genomes, Genetics, 2019, 9, 943-954.	0.8	11
18	Host-specialized fibrinogen-binding by a bacterial surface protein promotes biofilm formation and innate immune evasion. PLoS Pathogens, 2019, 15, e1007816.	2.1	34

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19	An ADAMTS3 missense variant is associated with Norwich Terrier upper airway syndrome. PLoS Genetics, 2019, 15, e1008102.	1.5	14
20	Morphological variation of the caudal fossa of domestic cat skulls assessed with CT and geometric morphometrics analysis. Journal of Feline Medicine and Surgery, 2018, 20, 752-758.	0.6	4
21	C7 vertebra homeotic transformation in domestic dogs “ are Pug dogs breaking mammalian evolutionary constraints?. Journal of Anatomy, 2018, 233, 255-265.	0.9	12
22	Canine Brachycephaly Is Associated with a Retrotransposon-Mediated Missplicing of SMOC2. Current Biology, 2017, 27, 1573-1584.e6.	1.8	80
23	Analysis of large versus small dogs reveals three genes on the canine X chromosome associated with body weight, muscling and back fat thickness. PLoS Genetics, 2017, 13, e1006661.	1.5	51
24	Developmental and Evolutionary Significance of the Zygomatic Bone. Anatomical Record, 2016, 299, 1616-1630.	0.8	11
25	The challenges of pedigree dog health: approaches to combating inherited disease. Canine Genetics and Epidemiology, 2015, 2, 3.	2.9	56
26	Insights into Morphology and Disease from the Dog Genome Project. Annual Review of Cell and Developmental Biology, 2014, 30, 535-560.	4.0	71
27	The Genetics of Canine Skull Shape Variation. Genetics, 2013, 193, 317-325.	1.2	82
28	Derived variants at six genes explain nearly half of size reduction in dog breeds. Genome Research, 2013, 23, 1985-1995.	2.4	131
29	Variation of BMP3 Contributes to Dog Breed Skull Diversity. PLoS Genetics, 2012, 8, e1002849.	1.5	159
30	A Simple Genetic Architecture Underlies Morphological Variation in Dogs. PLoS Biology, 2010, 8, e1000451.	2.6	429
31	Complex population structure in African village dogs and its implications for inferring dog domestication history. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 13903-13908.	3.3	141
32	Fine Mapping a Locus Controlling Leg Morphology in the Domestic Dog. Cold Spring Harbor Symposia on Quantitative Biology, 2009, 74, 327-333.	2.0	15
33	Modulation of β_3 receptor desensitization by the light chain of microtubule-associated protein 1B expressed in HEK 293 cells. Journal of Physiology, 2008, 586, 751-762.	1.3	32
34	Endocardium is necessary for cardiomyocyte movement during heart tube assembly. Development (Cambridge), 2007, 134, 2379-2386.	1.2	77
35	Early developmental specification of the thyroid gland depends on <i>hmx</i> -expressing surrounding tissue and on FGF signals. Development (Cambridge), 2007, 134, 2871-2879.	1.2	64
36	Illuminating cardiac development: Advances in imaging add new dimensions to the utility of zebrafish genetics. Seminars in Cell and Developmental Biology, 2007, 18, 27-35.	2.3	49

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37	Vessel and Blood Specification Override Cardiac Potential in Anterior Mesoderm. <i>Developmental Cell</i> , 2007, 13, 254-267.	3.1	201