Amy E Young

List of Publications by Year in descending order

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623734 642732 22 950 14 23 citations g-index h-index papers 27 27 27 1478 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Feline Polycystic Kidney Disease Mutation Identified in PKD1. Journal of the American Society of Nephrology: JASN, 2004, 15, 2548-2555.	6.1	120
2	Identification of Gene Networks for Residual Feed Intake in Angus Cattle Using Genomic Prediction and RNA-seq. PLoS ONE, 2016, 11, e0152274.	2.5	106
3	Applied Animal Genomics: Results from the Field. Annual Review of Animal Biosciences, 2014, 2, 105-139.	7.4	102
4	Localization of Canine Brachycephaly Using an Across Breed Mapping Approach. PLoS ONE, 2010, 5, e9632.	2.5	101
5	Mutations in the SLC2A9 Gene Cause Hyperuricosuria and Hyperuricemia in the Dog. PLoS Genetics, 2008, 4, e1000246.	3.5	94
6	<i>FGF4</i> retrogene on CFA12 is responsible for chondrodystrophy and intervertebral disc disease in dogs. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 11476-11481.	7.1	92
7	Genomic and phenotypic analyses of six offspring of a genome-edited hornless bull. Nature Biotechnology, 2020, 38, 225-232.	17.5	69
8	A LINE-1 Insertion in DLX6 Is Responsible for Cleft Palate and Mandibular Abnormalities in a Canine Model of Pierre Robin Sequence. PLoS Genetics, 2014, 10, e1004257.	3.5	49
9	Activity, Expression and Genetic Variation of Canine \hat{I}^2 -Defensin 103: A Multifunctional Antimicrobial Peptide in the Skin of Domestic Dogs. Journal of Innate Immunity, 2012, 4, 248-259.	3.8	45
10	Feline polycystic kidney disease is linked to the PKD1 region. Mammalian Genome, 2005, 16, 59-65.	2.2	32
11	Validation of a urine test and characterization of the putative genetic mutation for hyperuricosuria in Bulldogs and Black Russian Terriers. American Journal of Veterinary Research, 2010, 71, 909-914.	0.6	21
12	SERPINB11 Frameshift Variant Associated with Novel Hoof Specific Phenotype in Connemara Ponies. PLoS Genetics, 2015, 11, e1005122.	3.5	21
13	Evaluation of FOXC2 as a candidate gene for chronic progressive lymphedema in draft horses. Veterinary Journal, 2007, 174, 397-399.	1.7	19
14	Whole-Genome Sequencing and Concordance Between Antimicrobial Susceptibility Genotypes and Phenotypes of Bacterial Isolates Associated with Bovine Respiratory Disease. G3: Genes, Genomes, Genetics, 2017, 7, 3059-3071.	1.8	19
15	One-step generation of a targeted knock-in calf using the CRISPR-Cas9 system in bovine zygotes. BMC Genomics, 2021, 22, 118.	2.8	14
16	SNPSin the Promoter Regions of the Canine RMRP and SHOX Genes are not Associated with Canine Chondrodysplasia. Animal Biotechnology, 2008, 19, 1-5.	1.5	8
17	Genome Report: Identification and Validation of Antigenic Proteins from <i>Pajaroellobacter abortibovis</i> Using <i>De Novo</i> Genome Sequence Assembly and Reverse Vaccinology. G3: Genes, Genomes, Genetics, 2017, 7, 321-331.	1.8	8
18	Deletions in the COL10A1 gene are not associated with skeletal changes in dogs. Mammalian Genome, 2006, 17, 761-768.	2.2	6

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#	Article	IF	CITATION
19	Canine fibroblast growth factor receptor 3 sequence is conserved across dogs of divergent skeletal size. BMC Genetics, 2008, 9, 67.	2.7	5
20	Animal agriculture and the importance of agnostic governance of biotechnology. Agriculture and Food Security, 2015, 4, .	4.2	5
21	Harnessing endogenous repair mechanisms for targeted gene knock-in of bovine embryos. Scientific Reports, 2020, 10, 16031.	3.3	5
22	Animal Health and Food Safety Analyses of Six Offspring of a Genome-Edited Hornless Bull. , 2022, 1, 192-206.		3