

Christina Weimann

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

507
citations

1040056

9
h-index

996975

15
g-index

15
all docs

15
docs citations

15
times ranked

535
citing authors

#	ARTICLE	IF	CITATIONS
1	Whole-genome resequencing of wild and domestic sheep identifies genes associated with morphological and agronomic traits. <i>Nature Communications</i> , 2020, 11, 2815.	12.8	142
2	Combined analysis of data from two granddaughter designs: A simple strategy for QTL confirmation and increasing experimental power in dairy cattle. <i>Genetics Selection Evolution</i> , 2003, 35, 319-38.	3.0	71
3	Whole-Genome Resequencing of Worldwide Wild and Domestic Sheep Elucidates Genetic Diversity, Introgression, and Agronomically Important Loci. <i>Molecular Biology and Evolution</i> , 2022, 39, .	8.9	50
4	Genetic diversity, introgression and relationships among West/Central African cattle breeds. <i>Genetics Selection Evolution</i> , 2004, 36, 673-90.	3.0	49
5	Paternal Origins and Migratory Episodes of Domestic Sheep. <i>Current Biology</i> , 2020, 30, 4085-4095.e6.	3.9	49
6	Historical Introgression from Wild Relatives Enhanced Climatic Adaptation and Resistance to Pneumonia in Sheep. <i>Molecular Biology and Evolution</i> , 2021, 38, 838-855.	8.9	44
7	Multiple Quantitative Trait Loci Mapping With Cofactors and Application of Alternative Variants of the False Discovery Rate in an Enlarged Granddaughter Design. <i>Genetics</i> , 2004, 168, 1019-1027.	2.9	41
8	Alpha S1-casein polymorphisms in camel (<i>Camelus dromedarius</i>) and descriptions of biological active peptides and allergenic epitopes. <i>Tropical Animal Health and Production</i> , 2016, 48, 879-887.	1.4	24
9	The role of agro-ecological factors and transboundary transhumance in shaping the genetic diversity in four indigenous cattle populations of Benin. <i>Journal of Animal Breeding and Genetics</i> , 2020, 137, 622-640.	2.0	9
10	Genome-Wide Associations for Microscopic Differential Somatic Cell Count and Specific Mastitis Pathogens in Holstein Cows in Compost-Bedded Pack and Cubicle Farming Systems. <i>Animals</i> , 2021, 11, 1839.	2.3	9
11	Identification of Thermophilic Aerobic Sporeformers in Bedding Material of Compost-Bedded Dairy Cows Using Microbial and Molecular Methods. <i>Animals</i> , 2021, 11, 2890.	2.3	6
12	Allele substitution and dominance effects of CD166/ALCAM gene polymorphisms for endoparasite resistance and test-day traits in a small cattle population using logistic regression analyses. <i>Mammalian Genome</i> , 2019, 30, 301-317.	2.2	4
13	Microscopic differential cell count and specific mastitis pathogens in cow milk from compost-bedded pack barns and cubicle barns. <i>Journal of Dairy Research</i> , 2021, 88, 413-419.	1.4	4
14	Genetic diversity and population structure in divergent German cattle selection lines on the basis of milk protein polymorphisms. <i>Archives Animal Breeding</i> , 2021, 64, 91-102.	1.4	3
15	Associations between maternal milk protein genotypes with preweaning calf growth traits in beef cattle. <i>Journal of Animal Science</i> , 2020, 98, .	0.5	2