

Olga V Mitrofanova

List of Publications by Year in descending order

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Version: 2024-02-01

11
papers

66
citations

1937685

4
h-index

1588992

8
g-index

11
all docs

11
docs citations

11
times ranked

56
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide association studies targeting the yield of extraembryonic fluid and production traits in Russian White chickens. BMC Genomics, 2019, 20, 270.	2.8	22
2	Evolutionary Subdivision of Domestic Chickens: Implications for Local Breeds as Assessed by Phenotype and Genotype in Comparison to Commercial and Fancy Breeds. Agriculture (Switzerland), 2021, 11, 914.	3.1	14
3	Genetic Variability in Local and Imported Germplasm Chicken Populations as Revealed by Analyzing Runs of Homozygosity. Animals, 2020, 10, 1887.	2.3	11
4	Molecular-genetic bases of plumage coloring in chicken. Vavilovskii Zhurnal Genetiki I Seleksii, 2019, 23, 343-354.	1.1	8
5	Genetic Variability of Indels in the Prolactin and Dopamine Receptor D2 Genes and Their Association with the Yield of Allanto-Amniotic Fluid in Russian White Laying Hens. Tarim Bilimleri Dergisi, 0, , 373-379.	0.4	3
6	Assessment of variability of egg production traits based on analysis of SNP markers and search for traces of selection in the genome of Russian white chickens. Ecological Genetics, 2020, 18, 423-432.	0.5	3
7	Monitoring and significance of the recessive genetic defect AH1 of Ayrshire cattle. Czech Journal of Animal Science, 2020, 65, 323-329.	1.3	2
8	Efficiency of using SNP markers in the <i>MSTN</i> gene in the selection of the Pushkin breed chickens. Vavilovskii Zhurnal Genetiki I Seleksii, 2020, 23, 993-998.	1.1	2
9	Analysis of the genetic diversity of Ayrshire cattle in Russia (part 1). Ecological Genetics, 2022, 20, 5-12.	0.5	1
10	ASSOCIATION OF POLYMORPHIC TYPES OF STEAROYL-CoA DESATURASE GENE (SCD1) WITH ECONOMICALLY VALUABLE TRAITS IN RUSSIAN POPULATION OF AYRSHIRE COWS. Sel'skokhozyaistvennaya Biologiya, 2017, 52, 1139-1147.	0.3	0
11	Analysis of the accumulation of homozygosity regions in chickens of the Pushkin breed using data from whole genome genotyping. Ecological Genetics, 2022, 20, 31-39.	0.5	0