

Rabbit genome analysis reveals a polygenic basis for ph domestication

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Advances and limits of using population genetics to understand local adaptation. <i>Trends in Ecology and Evolution</i> , 2014, 29, 673-680.	4.2	329
2	On the origin of Peter Rabbit. <i>Science</i> , 2014, 345, 1000-1001.	6.0	2
3	Candidate genes underlying heritable differences in reproductive seasonality between wild and domestic rabbits. <i>Animal Genetics</i> , 2015, 46, 418-425.	0.6	14
4	Survey of genetic diversity of IgG in wild and domestic rabbits. <i>International Journal of Immunogenetics</i> , 2015, 42, 364-367.	0.8	5
5	Levels and Patterns of Genetic Diversity and Population Structure in Domestic Rabbits. <i>PLoS ONE</i> , 2015, 10, e0144687.	1.1	38
6	Low persistence in nature of captive reared rabbits after restocking operations. <i>European Journal of Wildlife Research</i> , 2015, 61, 591-599.	0.7	9
7	Parallel Selection Revealed by Population Sequencing in Chicken. <i>Genome Biology and Evolution</i> , 2015, 7, 3299-3306.	1.1	25
8	Yak whole-genome resequencing reveals domestication signatures and prehistoric population expansions. <i>Nature Communications</i> , 2015, 6, 10283.	5.8	214
9	Selection for tameness, a key behavioral trait of domestication, increases adult hippocampal neurogenesis in foxes. <i>Hippocampus</i> , 2015, 25, 963-975.	0.9	46
10	Adaptation and possible ancient interspecies introgression in pigs identified by whole-genome sequencing. <i>Nature Genetics</i> , 2015, 47, 217-225.	9.4	288
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17	Atlantic salmon populations reveal adaptive divergence of immune related genes - a duplicated genome under selection. <i>BMC Genomics</i> , 2016, 17, 610.	1.2	44
18	The Immune System of Lagomorphs. , 2016, , 515-525.		4

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20	The genetic basis for ecological adaptation of the Atlantic herring revealed by genome sequencing. <i>ELife</i> , 2016, 5, .	2.8	143
21	Identification of a Bitter-Taste Receptor Gene Repertoire in Different Lagomorphs Species. <i>Frontiers in Genetics</i> , 2016, 7, 55.	1.1	0
22	Genome-wide patterns of copy number variation in the Chinese yak genome. <i>BMC Genomics</i> , 2016, 17, 379.	1.2	66
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24	Whole-genome sequencing of eight goat populations for the detection of selection signatures underlying production and adaptive traits. <i>Scientific Reports</i> , 2016, 6, 38932.	1.6	132
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34	Selective Sweeps. , 2016, , 23-32.		2
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56	Signatures of Selection and Interspecies Introgression in the Genome of Chinese Domestic Pigs. <i>Genome Biology and Evolution</i> , 2017, 9, 2592-2603.	1.1	43
57	An Evolutionary Genomic Perspective on the Breeding of Dwarf Chickens. <i>Molecular Biology and Evolution</i> , 2017, 34, 3081-3088.	3.5	42
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127	An evaluation of sequencing coverage and genotyping strategies to assess neutral and adaptive diversity. <i>Molecular Ecology Resources</i> , 2019, 19, 1497-1515.	2.2	31

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138	Animal models of arrhythmia: classic electrophysiology to genetically modified large animals. <i>Nature Reviews Cardiology</i> , 2019, 16, 457-475.	6.1	131
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159	Breve contexto legal y conceptual de la tenencia de animales domésticos, silvestres y exóticos en Colombia. <i>Forum Revista Departamento Ciencia Política</i> , 2020, , 72-93.	0.1	0
160	The Origin and Spread of Locally Adaptive Seasonal Camouflage in Snowshoe Hares. <i>American Naturalist</i> , 2020, 196, 316-332.	1.0	29
161	Breeding history and candidate genes responsible for black skin of Xichuan black-bone chicken. <i>BMC Genomics</i> , 2020, 21, 511.	1.2	32
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165	Brain Transcriptomics of Wild and Domestic Rabbits Suggests That Changes in Dopamine Signaling and Ciliary Function Contributed to Evolution of Tameless. <i>Genome Biology and Evolution</i> , 2020, 12, 1918-1928.	1.1	17
166	The Domestication Makeup: Evolution, Survival, and Challenges. <i>Frontiers in Ecology and Evolution</i> , 2020, 8, .	1.1	29
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