

# CITATION REPORT

List of articles citing

The second generation of the International Equine Gene Mapping Workshop half-sibling linkage map

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Animal Genetics, 2003, 34, 161-8.

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#	Paper	IF	Citations
39	Equine genomics: galloping to new frontiers. <i>Cytogenetic and Genome Research</i> , <b>2003</b> , 102, 184-8	1.9	16
38	Genetic variation in Przewalski's horses, with special focus on the last wild caught mare, 231 Orlitza III. <i>Cytogenetic and Genome Research</i> , <b>2003</b> , 102, 226-34	1.9	29
37	The first-generation whole-genome radiation hybrid map in the horse identifies conserved segments in human and mouse genomes. <i>Genome Research</i> , <b>2003</b> , 13, 742-51	9.7	121
36	First-generation linkage map of the gray, short-tailed opossum, <i>Monodelphis domestica</i> , reveals genome-wide reduction in female recombination rates. <i>Genetics</i> , <b>2004</b> , 166, 307-29	4	47
35	Exceptional conservation of horse-human gene order on X chromosome revealed by high-resolution radiation hybrid mapping. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2004</b> , 101, 2386-91	11.5	64
34	Genetic mapping of dominant white (W), a homozygous lethal condition in the horse ( <i>Equus caballus</i> ). <i>Journal of Animal Breeding and Genetics</i> , <b>2004</b> , 121, 374-383	2.9	12
33	Identification of putative homology between horse microsatellite flanking sequences and cross-species ESTs, mRNAs and genomic sequences. <i>Animal Genetics</i> , <b>2004</b> , 35, 28-33	2.5	14
32	Assignment of the appaloosa coat colour gene (LP) to equine chromosome 1. <i>Animal Genetics</i> , <b>2004</b> , 35, 134-7	2.5	29
31	Isolation, characterization and chromosome assignment of 341 newly isolated equine TKY microsatellite markers. <i>Animal Genetics</i> , <b>2004</b> , 35, 487-96	2.5	7
30	Variation in human meiotic recombination. <i>Annual Review of Genomics and Human Genetics</i> , <b>2004</b> , 5, 317-49	9.7	129
29	A 1.4-Mb interval RH map of horse chromosome 17 provides detailed comparison with human and mouse homologues. <i>Genomics</i> , <b>2004</b> , 83, 203-15	4.3	16
28	Phylogenetic analysis and species allocation of individual equids using microsatellite data. <i>Journal of Animal Breeding and Genetics</i> , <b>2005</b> , 122 Suppl 1, 78-86	2.9	26
27	Sequence, detection of polymorphisms and radiation hybrid mapping of the equine catechol-o-methyltransferase gene. <i>Animal Genetics</i> , <b>2005</b> , 36, 190	2.5	6
26	A high-resolution physical map of equine homologs of HSA19 shows divergent evolution compared with other mammals. <i>Mammalian Genome</i> , <b>2005</b> , 16, 631-49	3.2	21
25	Equine FISH mapping of 36 genes known to locate on human chromosome ends. <i>Cytogenetic and Genome Research</i> , <b>2005</b> , 111, 46-50	1.9	6
24	Advances in livestock genomics: opening the barn door. <i>Genome Research</i> , <b>2005</b> , 15, 1699-705	9.7	59
23	Prospects for whole genome linkage disequilibrium mapping in thoroughbreds. <i>Gene</i> , <b>2005</b> , 346, 127-32	3.8	15

22	High-resolution RH map of horse chromosome 22 reveals a putative ancestral vertebrate chromosome. <i>Genomics</i> , <b>2005</b> , 85, 188-200	4.3	14
21	International Equine Gene Mapping Workshop Report: a comprehensive linkage map constructed with data from new markers and by merging four mapping resources. <i>Cytogenetic and Genome Research</i> , <b>2005</b> , 111, 5-15	1.9	54
20	Single linkage group per chromosome genetic linkage map for the horse, based on two three-generation, full-sibling, crossbred horse reference families. <i>Genomics</i> , <b>2006</b> , 87, 1-29	4.3	59
19	Construction of a medium-density horse gene map. <i>Animal Genetics</i> , <b>2006</b> , 37, 145-55	2.5	29
18	High-resolution gene maps of horse chromosomes 14 and 21: additional insights into evolution and rearrangements of HSA5 homologs in mammals. <i>Genomics</i> , <b>2007</b> , 89, 89-112	4.3	9
17	Multidirectional cross-species painting illuminates the history of karyotypic evolution in Perissodactyla. <i>Chromosome Research</i> , <b>2008</b> , 16, 89-107	4.4	57
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15	A BAC contig map over the proximal approximately 3.3 Mb region of horse chromosome 21. <i>Cytogenetic and Genome Research</i> , <b>2008</b> , 120, 164-72	1.9	2
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13	Inherited Ocular Disorders. <b>2011</b> , 434-442		3
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