Resource Summary Report

Generated by NIF on May 3, 2025

Australian Sheep Gene Mapping

RRID:SCR_008152
Type: Tool

Proper Citation

Australian Sheep Gene Mapping (RRID:SCR_008152)

Resource Information

URL: http://rubens.its.unimelb.edu.au/~jillm/jill.htm

Proper Citation: Australian Sheep Gene Mapping (RRID:SCR_008152)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented August 23, 2016. This page contains a list of websites relating to sheep gene mapping. Users may chose from the list to view any desired webpage. The website also contains information on sheep maps, sheep markers, map comparisons, experiments, DNA tests, and other genome sites.

Synonyms: ASGM

Resource Type: topical portal, portal, data or information resource

Funding:

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Australian Sheep Gene Mapping

Resource ID: SCR_008152

Alternate IDs: nif-0000-20994

Record Creation Time: 20220129T080245+0000

Record Last Update: 20250503T060003+0000

Ratings and Alerts

No rating or validation information has been found for Australian Sheep Gene Mapping.

No alerts have been found for Australian Sheep Gene Mapping.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 7 mentions in open access literature.

Listed below are recent publications. The full list is available at NIF.

Popa A, et al. (2012) The sex-specific impact of meiotic recombination on nucleotide composition. Genome biology and evolution, 4(3), 412.

Lühken G, et al. (2012) Familiar hypopigmentation syndrome in sheep associated with homozygous deletion of the entire endothelin type-B receptor gene. PloS one, 7(12), e53020.

Raadsma HW, et al. (2009) Mapping quantitative trait loci (QTL) in sheep. I. A new male framework linkage map and QTL for growth rate and body weight. Genetics, selection, evolution : GSE, 41(1), 34.

Gutiérrez-Gil B, et al. (2009) Quantitative trait loci for resistance to trichostrongylid infection in Spanish Churra sheep. Genetics, selection, evolution : GSE, 41(1), 46.

Raadsma HW, et al. (2009) Mapping quantitative trait loci (QTL) in sheep. II. Meta-assembly and identification of novel QTL for milk production traits in sheep. Genetics, selection, evolution : GSE, 41(1), 45.

Liu H, et al. (2006) A BAC clone-based physical map of ovine major histocompatibility complex. Genomics, 88(1), 88.

Fadiel A, et al. (2005) Farm animal genomics and informatics: an update. Nucleic acids research, 33(19), 6308.