CHROMOSCAN

RRID:SCR_009151
Type: Tool

Proper Citation

CHROMOSCAN (RRID:SCR_009151)

Resource Information

URL: https://github.com/gaow/genetic-analysis-software/blob/master/pages/CHROMOSCAN.md

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Description: THIS RESOURCE IS NO LONGER IN SERVCE, documented September 22, 2016. Software application that is an implementation of a genome-based scan statistic that detects genomic regions.

Abbreviations: CHROMOSCAN

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, java

Funding:

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: CHROMOSCAN

Resource ID: SCR_009151

Alternate IDs: nlx_154270

Old URLs: http://www.epidkardia.sph.umich.edu/software/chromoscan/

Record Creation Time: 20220129T080251+0000

Record Last Update: 20250416T063538+0000

Ratings and Alerts

No rating or validation information has been found for CHROMOSCAN.

No alerts have been found for CHROMOSCAN.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Li Y, et al. (2017) A whole genome association study to detect additive and dominant single nucleotide polymorphisms for growth and carcass traits in Korean native cattle, Hanwoo. Asian-Australasian journal of animal sciences, 30(1), 8.

Schmuck E, et al. (2008) Deletion of Glu155 causes a deficiency of glutathione transferase Omega 1-1 but does not alter sensitivity to arsenic trioxide and other cytotoxic drugs. The international journal of biochemistry & cell biology, 40(11), 2553.