

Resource Summary Report

Generated by [NIF](#) on Sep 2, 2026

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>

Proper Citation: CHROMOSCAN (RRID:SCR_009151)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, 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 application, software resource

Keywords: gene, genetic, genomic, java

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: 20260829T183054+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 [NIF](#) .

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.