

Resource Summary Report

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CHROMOSEG

RRID:SCR_009152

Type: Tool

Proper Citation

CHROMOSEG (RRID:SCR_009152)

Resource Information

URL: <http://ib.berkeley.edu/labs/slatkin/eriq/software/software.htm>

Proper Citation: CHROMOSEG (RRID:SCR_009152)

Description: A simulation program that simulates and plots (in real time) ancestral recombination graphs. This is currently primarily a teaching/educational tool. (entry from Genetic Analysis Software)

Abbreviations: CHROMOSEG

Resource Type: software resource, software application

Keywords: gene, genetic, genomic

Funding:

Resource Name: CHROMOSEG

Resource ID: SCR_009152

Alternate IDs: nlx_154271

Record Creation Time: 20220129T080251+0000

Record Last Update: 20250411T055256+0000

Ratings and Alerts

No rating or validation information has been found for CHROMOSEG.

No alerts have been found for CHROMOSEG.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#).

Gousskov A, et al. (2016) Postglacial recolonizations, watershed crossings and human translocations shape the distribution of chub lineages around the Swiss Alps. BMC evolutionary biology, 16(1), 185.