

Resource Summary Report

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

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 application, software resource

Keywords: gene, genetic, genomic

Resource Name: CHROMOSEG

Resource ID: SCR_009152

Alternate IDs: nlx_154271

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260829T183106+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 2 mentions in open access literature.

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

Horta P, et al. (2022) Counteracting forces of introgressive hybridization and interspecific competition shape the morphological traits of cryptic Iberian *Eptesicus* bats. *Scientific reports*, 12(1), 11695.

Gouskov 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.