

# Resource Summary Report

Generated by [dkNET](#) on Sep 1, 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 [dkNET](#) .

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.