

# Resource Summary Report

Generated by T1D on Aug 2, 2026

## COMBIN

RRID:SCR\_009050

Type: Tool

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### Proper Citation

COMBIN (RRID:SCR\_009050)

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### Resource Information

**URL:** <https://github.com/gaow/genetic-analysis-software/blob/master/pages/COMBIN.md>

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**Description:** Software application designed for the construction of highly saturated linkage maps, based on BC1, DH, Radiation Hybrid or CP (CrossPollinators) data sets. F2 is not supported. (entry from Genetic Analysis Software)

**Abbreviations:** COMBIN

**Resource Type:** software resource, software application

**Keywords:** gene, genetic, genomic, visual basic 5, ms-windows, (95/98/nt)

**Resource Name:** COMBIN

**Resource ID:** SCR\_009050

**Alternate IDs:** nlx\_154025

**Old URLs:** <http://www.dpw.wau.nl/pv/pub/combin/>

**Record Creation Time:** 20220129T080250+0000

**Record Last Update:** 20260801T191054+0000

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### Ratings and Alerts

No rating or validation information has been found for COMBIN.

No alerts have been found for COMBIN.

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### Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.