

Resource Summary Report

Generated by [nidm-terms](#) on Sep 21, 2026

COMBIN

RRID:SCR_009050

Type: Tool

Proper Citation

COMBIN (RRID:SCR_009050)

Resource Information

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

Proper Citation: COMBIN (RRID:SCR_009050)

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 application, software resource

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: 20260919T195757+0000

Ratings and Alerts

No rating or validation information has been found for COMBIN.

No alerts have been found for COMBIN.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.