

Resource Summary Report

Generated by [NIF](#) on Sep 1, 2026

[RHMAP](#)

RRID:SCR_009360

Type: Tool

Proper Citation

RHMAP (RRID:SCR_009360)

Resource Information

URL: <http://csg.sph.umich.edu/boehnke/rhmap.php>

Proper Citation: RHMAP (RRID:SCR_009360)

Description: Software application (entry from Genetic Analysis Software)

Abbreviations: RHMAP

Synonyms: Radiation Hybrid Map

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, fortran77

Resource Name: RHMAP

Resource ID: SCR_009360

Alternate IDs: nlx_154575

Record Creation Time: 20220129T080252+0000

Record Last Update: 20260829T183100+0000

Ratings and Alerts

No rating or validation information has been found for RHMAP.

No alerts have been found for RHMAP.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Li J, et al. (2021) A new duck genome reveals conserved and convergently evolved chromosome architectures of birds and mammals. *GigaScience*, 10(1).

Cui Z, et al. (2005) Molecular cloning, characterization, and chromosomal assignment of porcine cationic amino acid transporter-1. *Genomics*, 85(3), 352.

Housley DJ, et al. (2004) Comparative radiation hybrid map of canine chromosome 1 incorporating SNP and indel polymorphisms. *Genomics*, 84(2), 248.