## **Resource Summary Report**

Generated by <u>NIF</u> on Apr 23, 2025

# **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 resource, software application

Keywords: gene, genetic, genomic, fortran77

Funding:

Resource Name: RHMAP

Resource ID: SCR\_009360

Alternate IDs: nlx\_154575

Record Creation Time: 20220129T080252+0000

Record Last Update: 20250421T053726+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: <u>SciCrunch Registry</u>

### **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.