

Resource Summary Report

Generated by [RRID](#) on Aug 12, 2026

HomoeologPrimer

RRID:SCR_017559

Type: Tool

Proper Citation

HomoeologPrimer (RRID:SCR_017559)

Resource Information

URL: <https://github.com/lufuhao/HomoeologPrimer>

Proper Citation: HomoeologPrimer (RRID:SCR_017559)

Description: Software tool as script used to design primers for homoeologs, to amplify either homoeolog specific or all homoeologs.

Resource Type: software application, software resource

Keywords: Design, primer, homoeolog, amplify

Availability: Free, Available for download, Freely available

Resource Name: HomoeologPrimer

Resource ID: SCR_017559

License: GNU General Public License v3.0

Record Creation Time: 20220129T080335+0000

Record Last Update: 20260810T040722+0000

Ratings and Alerts

No rating or validation information has been found for HomoeologPrimer.

No alerts have been found for HomoeologPrimer.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Lu FH, et al. (2020) Reduced chromatin accessibility underlies gene expression differences in homologous chromosome arms of diploid *Aegilops tauschii* and hexaploid wheat. *GigaScience*, 9(6).