

Resource Summary Report

Generated by [RRID](#) on Aug 5, 2026

Automatic Primer Design

RRID:SCR_007155

Type: Tool

Proper Citation

Automatic Primer Design (RRID:SCR_007155)

Resource Information

URL: <http://primers.niob.knaw.nl/>

Proper Citation: Automatic Primer Design (RRID:SCR_007155)

Description: This automatic primer design service uses the Primer3 Software (through EMBOSS primer3 interface) to design two sets of primers for doing a nested PCR. The input can either be in Fasta format or in GENOTRACE output format. You will have to indicate with square brackets which region should be included in the amplicon. Primers for multiple amplicons (for example all exons) can be designed in a single run.

Synonyms: Primer Design Form

Resource Type: data processing software, software resource, software application, production service resource, service resource, data acquisition software

Keywords: primer, design, service, software, pcr

Resource Name: Automatic Primer Design

Resource ID: SCR_007155

Alternate IDs: nif-0000-30136

Record Creation Time: 20220129T080240+0000

Record Last Update: 20260805T044146+0000

Ratings and Alerts

No rating or validation information has been found for Automatic Primer Design.

No alerts have been found for Automatic Primer Design.

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](#) .

Wansleebe C, et al. (2011) An ENU-mutagenesis screen in the mouse: identification of novel developmental gene functions. PloS one, 6(4), e19357.