

Resource Summary Report

Generated by [RRID](#) on Aug 4, 2026

Diploidocus

RRID:SCR_021231

Type: Tool

Proper Citation

Diploidocus (RRID:SCR_021231)

Resource Information

URL: <https://github.com/slimsuite/diploidocus>

Proper Citation: Diploidocus (RRID:SCR_021231)

Description: Software package for diploid genome assembly analysis. Sequence analysis toolkit for number of different analyses related to diploid genome assembly.

Resource Type: sequence analysis software, software toolkit, software application, data processing software, data analysis software, software resource

Keywords: Diploid genome assembly analysis, genome, genome assembly, sequence analysis

Availability: Free, Available for download, Freely available

Resource Name: Diploidocus

Resource ID: SCR_021231

License: GNU GPL v3.0

Record Creation Time: 20220129T080354+0000

Record Last Update: 20260804T043723+0000

Ratings and Alerts

No rating or validation information has been found for Diploidocus.

No alerts have been found for Diploidocus.

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 [RRID](#) .

Silver LW, et al. (2024) A reference genome for the eastern bettong (*Bettongia gaimardi*). *F1000Research*, 13, 1544.

Chen SH, et al. (2022) A high-quality pseudo-phased genome for *Melaleuca quinquenervia* shows allelic diversity of NLR-type resistance genes. *GigaScience*, 12.

Stuart KC, et al. (2022) Transcript- and annotation-guided genome assembly of the European starling. *Molecular ecology resources*, 22(8), 3141.