

Resource Summary Report

Generated by SPARC SAWG on Jul 28, 2026

Narcisse

RRID:SCR_007813

Type: Tool

Proper Citation

Narcisse (RRID:SCR_007813)

Resource Information

URL: <http://narcisse.toulouse.inra.fr/multi/cgi-bin/narcisse.cgi>

Proper Citation: Narcisse (RRID:SCR_007813)

Description: Narcisse is a comparative genome browser. The aim is to provide a simple and intuitive access to the results of comparison of genomes completely or partially sequenced. The principle of construction of conserved segments based on the idea that the concept of conservation depends on the level of resolution. The number of levels of conservation is fixed, more or less arbitrary for animals 5 and 2 for the other kingdoms. Narcisse software allows users to move between these levels of conservation.

Synonyms: Narcisse

Resource Type: database, data or information resource

Resource Name: Narcisse

Resource ID: SCR_007813

Alternate IDs: nif-0000-03175

Alternate URLs: <http://narcisse.toulouse.inra.fr>

Record Creation Time: 20220129T080243+0000

Record Last Update: 20260725T191151+0000

Ratings and Alerts

No rating or validation information has been found for Narcisse.

No alerts have been found for Narcisse.

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 [SPARC SAWG](#) .

Bourneuf E, et al. (2018) New susceptibility loci for cutaneous melanoma risk and progression revealed using a porcine model. Oncotarget, 9(45), 27682.