

Resource Summary Report

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

[NARWHAL](#)

RRID:SCR_011858

Type: Tool

Proper Citation

NARWHAL (RRID:SCR_011858)

Resource Information

URL: <https://trac.nbic.nl/narwhal/>

Proper Citation: NARWHAL (RRID:SCR_011858)

Description: Automates the primary analysis of massive parallel sequencing data.

Abbreviations: NARWHAL

Resource Type: software resource

Resource Name: NARWHAL

Resource ID: SCR_011858

Alternate IDs: OMICS_01130

Record Creation Time: 20220129T080307+0000

Record Last Update: 20260801T190421+0000

Ratings and Alerts

No rating or validation information has been found for NARWHAL.

No alerts have been found for NARWHAL.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 10 mentions in open access literature.

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

Silva CT, et al. (2018) A combined linkage, microarray and exome analysis suggests MAP3K11 as a candidate gene for left ventricular hypertrophy. BMC medical genomics, 11(1), 22.

van Nistelrooij AMJ, et al. (2018) Germline variant in MSX1 identified in a Dutch family with clustering of Barrett's esophagus and esophageal adenocarcinoma. Familial cancer, 17(3), 435.

Meuwissen ME, et al. (2016) Human USP18 deficiency underlies type 1 interferonopathy leading to severe pseudo-TORCH syndrome. The Journal of experimental medicine, 213(7), 1163.

Silva CT, et al. (2016) A Combined Linkage and Exome Sequencing Analysis for Electrocardiogram Parameters in the Erasmus Rucphen Family Study. Frontiers in genetics, 7, 190.

Demirkan A, et al. (2015) Insight in genome-wide association of metabolite quantitative traits by exome sequence analyses. PLoS genetics, 11(1), e1004835.

Wessel J, et al. (2015) Low-frequency and rare exome chip variants associate with fasting glucose and type 2 diabetes susceptibility. Nature communications, 6, 5897.

Derks KW, et al. (2015) Deciphering the RNA landscape by RNAome sequencing. RNA biology, 12(1), 30.

Derks KW, et al. (2015) RNAome sequencing delineates the complete RNA landscape. Genomics data, 5, 381.

Thangam M, et al. (2015) CRCDA--Comprehensive resources for cancer NGS data analysis. Database : the journal of biological databases and curation, 2015.

Stadhouders R, et al. (2012) Dynamic long-range chromatin interactions control Myb proto-oncogene transcription during erythroid development. The EMBO journal, 31(4), 986.