

Resource Summary Report

Generated by [NIF](#) on Aug 30, 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: 20260829T182412+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 [NIF](#) .

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.

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.

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.