

Resource Summary Report

Generated by T1D on Sep 5, 2026

seqwish

RRID:SCR_024332

Type: Tool

Proper Citation

seqwish (RRID:SCR_024332)

Resource Information

URL: <https://github.com/ekg/seqwish>

Proper Citation: seqwish (RRID:SCR_024332)

Description: Software tool for alignment to variation graph inducer.

Resource Type: software application, software resource

Keywords: alignment, variation graph inducer,

Availability: Free, Available for download, Freely available,

Resource Name: seqwish

Resource ID: SCR_024332

Old URLs: <https://sources.debian.org/src/seqwish/>

License: MIT license

Record Creation Time: 20230830T050217+0000

Record Last Update: 20260904T000546+0000

Ratings and Alerts

No rating or validation information has been found for seqwish.

No alerts have been found for seqwish.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#) .

Tennakoon C, et al. (2026) Panalyze: automated virus pangenome variation graph construction, analysis and annotation. Bioinformatics advances, 6(1), vbag071.

Bouhouch Y, et al. (2024) Early Detection of Both *Pyrenophora teres* f. *teres* and f. *maculata* in Asymptomatic Barley Leaves Using Digital Droplet PCR (ddPCR). International journal of molecular sciences, 25(22).