

# Resource Summary Report

Generated by [dkNET](#) on Aug 7, 2026

## GUSHR

RRID:SCR\_021166

Type: Tool

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### Proper Citation

GUSHR (RRID:SCR\_021166)

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### Resource Information

**URL:** <https://github.com/Gaius-Augustus/GUSHR>

**Proper Citation:** GUSHR (RRID:SCR\_021166)

**Description:** Software tool for assembly free construction of UTRs from short read RNA-Seq data on basis of coding sequence annotation.

**Resource Type:** data analysis software, software application, software resource, data processing software

**Keywords:** UTRs assembly free construction, short read RNA-Seq data, coding sequence annotation

**Availability:** Free, Available for download, Freely available

**Resource Name:** GUSHR

**Resource ID:** SCR\_021166

**License:** GNU General Public License v3.0

**Record Creation Time:** 20220129T080354+0000

**Record Last Update:** 20260807T042936+0000

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### Ratings and Alerts

No rating or validation information has been found for GUSHR .

No alerts have been found for GUSHR .

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### Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Cristofari R, et al. (2026) Lifestyle change accelerates epigenetic ageing in King penguins. Nature communications, 17(1).

Fujiwara K, et al. (2025) A Near Complete Genome Assembly of the Oshima Cherry *Cerasus speciosa*. Scientific data, 12(1), 162.

Yi L, et al. (2025) Chromosome-level genome assemblies of sunflower oilseed and confectionery cultivars. Scientific data, 12(1), 24.

Çiftçi O, et al. (2024) Long-Read-Based Hybrid Genome Assembly and Annotation of Snow Algal Strain CCCryo 101-99 (cf. *Sphaerocystis* sp., Chlamydomonadales). Genome biology and evolution, 16(7).

Legeai F, et al. (2024) Chromosome-Level Assembly and Annotation of the Pearly Heath *Coenonympha arcania* Butterfly Genome. Genome biology and evolution, 16(3).

Cicconardi F, et al. (2023) Evolutionary dynamics of genome size and content during the adaptive radiation of Heliconiini butterflies. Nature communications, 14(1), 5620.