

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 19, 2026

Gossamer

RRID:SCR_010612

Type: Tool

Proper Citation

Gossamer (RRID:SCR_010612)

Resource Information

URL: <http://bioinformatics.research.nicta.com.au/software/gossamer/>

Proper Citation: Gossamer (RRID:SCR_010612)

Description: A software application for the de novo assembly of genomes from fragments of DNA that specifically attacks the question of scalability.

Abbreviations: Gossamer

Synonyms: Gossamer: A Space-Efficient Genome Assembler

Resource Type: software resource

Resource Name: Gossamer

Resource ID: SCR_010612

Alternate IDs: OMICS_00017

Record Creation Time: 20220129T080259+0000

Record Last Update: 20260919T195157+0000

Ratings and Alerts

No rating or validation information has been found for Gossamer.

No alerts have been found for Gossamer.

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 [PRECISE-TBI](#) .

Gyan E, et al. (2025) Maintenance therapy with the FMS-like tyrosine kinase 3 inhibitor gilteritinib in patients with FMS-like tyrosine kinase 3-internal tandem duplication acute myeloid leukemia: A phase 2 study. *Cancer*, 131(4), e35746.