

# Resource Summary Report

Generated by [NIF](#) on Sep 2, 2026

## dCAS

RRID:SCR\_016612

Type: Tool

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### Proper Citation

dCAS (RRID:SCR\_016612)

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### Resource Information

**URL:** <https://niaid.github.io/dcas/>

**Proper Citation:** dCAS (RRID:SCR\_016612)

**Description:** Web tool to import raw cDNA sequences, clean sequences, build sequence contigs, perform SignalP analysis, BLAST contigs against numerous BLAST databases, and view the results. Automates large scale cDNA sequence analysis., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

**Abbreviations:** dCAS

**Synonyms:** Desktop cDNA Annotation System

**Resource Type:** software resource, web application

**Keywords:** automate, large, scale, cDNA, sequence, analysis, BLAST, database, gene

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** dCAS

**Resource ID:** SCR\_016612

**Record Creation Time:** 20220129T080331+0000

**Record Last Update:** 20260829T182550+0000

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### Ratings and Alerts

No rating or validation information has been found for dCAS.

No alerts have been found for dCAS.

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### Data and Source Information

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

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.