

# Resource Summary Report

Generated by [NIF](#) on Apr 27, 2025

## GPU-BLAST

RRID:SCR\_011820

Type: Tool

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### Proper Citation

GPU-BLAST (RRID:SCR\_011820)

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### Resource Information

**URL:** <http://archimedes.cheme.cmu.edu/?q=gpublast>

**Proper Citation:** GPU-BLAST (RRID:SCR\_011820)

**Description:** Software for an accelerated version of the popular NCBI-BLAST using a general-purpose graphics processing unit (GPU). It is nearly four times faster, while producing identical results. GPU-BLAST supports: protein alignment according to blastp (it does not support psiblast), multiple CPU threads working in parallel with a single GPU, and input files with multiple protein queries.

**Abbreviations:** GPU-BLAST

**Resource Type:** software resource

**Defining Citation:** [PMID:21088027](#)

**Keywords:** c++, gpu/cuda

**Funding:**

**Availability:** Free, Public, Acknowledgement requested

**Resource Name:** GPU-BLAST

**Resource ID:** SCR\_011820

**Alternate IDs:** OMICS\_00995

**Old URLs:** <http://eudoxus.cheme.cmu.edu/gpublast/gpublast.html>

**Record Creation Time:** 20220129T080306+0000

**Record Last Update:** 20250420T014600+0000

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## Ratings and Alerts

No rating or validation information has been found for GPU-BLAST.

No alerts have been found for GPU-BLAST.

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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.