

Resource Summary Report

Generated by [RRID](#) on Aug 12, 2026

Sargasso

RRID:SCR_023489

Type: Tool

Proper Citation

Sargasso (RRID:SCR_023489)

Resource Information

URL: <http://statbio.github.io/Sargasso/>

Proper Citation: Sargasso (RRID:SCR_023489)

Description: Software Python tool to disambiguate mixed species RNA-seq reads according to their species of origin.

Resource Type: software application, software resource

Keywords: disambiguate mixed species RNA-seq reads, mixed species RNA-seq reads, mixed species, RNA-seq reads

Availability: Free, Available for download, Freely available

Resource Name: Sargasso

Resource ID: SCR_023489

Alternate URLs: <https://zenodo.org/record/260123#.ZEG6ynbMK3B>

License: MIT license

Record Creation Time: 20230421T050214+0000

Record Last Update: 20260810T040839+0000

Ratings and Alerts

No rating or validation information has been found for Sargasso.

No alerts have been found for Sargasso.

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 [RRID](#) .

Martinez-Lozada Z, et al. (2023) Cooperative and competitive regulation of the astrocytic transcriptome by neurons and endothelial cells: Impact on astrocyte maturation. Journal of neurochemistry, 167(1), 52.