FACS
RRID:SCR_000055
Type: Tool

Proper Citation

FACS (RRID:SCR_000055)

Resource Information

**URL:** [https://github.com/SciLifeLab/facs](https://github.com/SciLifeLab/facs)

**Proper Citation:** FACS (RRID:SCR_000055)

**Description:** Software for classification of Sequences using Bloom filters that can accurately and rapidly align sequences to a reference sequence.

**Abbreviations:** FACS

**Synonyms:** Fast and Accurate Classification of Sequences

**Resource Type:** software resource

**Defining Citation:** PMID:20472541

**Keywords:** unix/linux, sequence, bio.tools

**Availability:** MIT License

**Resource Name:** FACS

**Resource ID:** SCR_000055

**Alternate IDs:** OMICS_02147, biotools:facs

**Alternate URLs:** [https://bio.tools/facs](https://bio.tools/facs)

Ratings and Alerts
No rating or validation information has been found for FACS.
No alerts have been found for FACS.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID.

