

# Resource Summary Report

Generated by [nidm-terms](#) on Sep 21, 2026

## NEXT-scASV

RRID:SCR\_027470

Type: Tool

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### Proper Citation

NEXT-scASV (RRID:SCR\_027470)

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### Resource Information

**URL:** <https://github.com/MedvedevaLab/NEXT-scASV.git>

**Proper Citation:** NEXT-scASV (RRID:SCR\_027470)

**Description:** Software Nextflow pipeline for calling allele-specific variants (ASVs) from 5? single-cell RNA sequencing data.

**Resource Type:** data analysis software, data processing software, software application, software resource, source code

**Keywords:** allele-specific variants, single-cell RNA-seq, Nextflow, computational pipeline, regulatory variants

**Availability:** Free, Available for download, Freely available

**Resource Name:** NEXT-scASV

**Resource ID:** SCR\_027470

**License:** MIT license

**Record Creation Time:** 20250927T062016+0000

**Record Last Update:** 20260919T200111+0000

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### Ratings and Alerts

No rating or validation information has been found for NEXT-scASV.

No alerts have been found for NEXT-scASV.

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### Data and Source Information

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

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [nidm-terms](#) .

Shevtsov A, et al. (2026) NEXT-scASV: a Nextflow pipeline for allele-specific variant calling from single-cell RNA-seq data. GigaScience, 15.