

Resource Summary Report

Generated by [nidm-terms](#) on Jul 30, 2026

ASC

RRID:SCR_001013

Type: Tool

Proper Citation

ASC (RRID:SCR_001013)

Resource Information

URL: <http://www.stat.brown.edu/ZWU/research.aspx>

Proper Citation: ASC (RRID:SCR_001013)

Description: Borrows information across sequences to establish prior distribution of sample variation, so that biological variation can be accounted for even when replicates are not available.

Abbreviations: ASC

Resource Type: software resource

Keywords: sample variation, rna, dna, biology, bio.tools

Resource Name: ASC

Resource ID: SCR_001013

Alternate IDs: biotools:sqn, OMICS_01298

Alternate URLs: <https://bio.tools/sqn>

Record Creation Time: 20220129T080205+0000

Record Last Update: 20260725T190455+0000

Ratings and Alerts

No rating or validation information has been found for ASC.

No alerts have been found for ASC.

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 [nidm-terms](#) .

Wu Z, et al. (2010) Empirical bayes analysis of sequencing-based transcriptional profiling without replicates. BMC bioinformatics, 11, 564.