

# Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 7, 2026

## QSRA

RRID:SCR\_010733

Type: Tool

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### Proper Citation

QSRA (RRID:SCR\_010733)

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### Resource Information

**URL:** <http://mocklerlab.org/tools/2>

**Proper Citation:** QSRA (RRID:SCR\_010733)

**Description:** A quality-value guided de novo short read assembler.

**Abbreviations:** QSRA

**Resource Type:** software resource

**Keywords:** bio.tools

**Resource Name:** QSRA

**Resource ID:** SCR\_010733

**Alternate IDs:** OMICS\_00026, biotools:qsra

**Alternate URLs:** <https://bio.tools/qsra>

**Record Creation Time:** 20220129T080300+0000

**Record Last Update:** 20260905T132645+0000

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### Ratings and Alerts

No rating or validation information has been found for QSRA.

No alerts have been found for QSRA.

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### 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 [PRECISE-TBI](#) .

El-Metwally S, et al. (2013) Next-generation sequence assembly: four stages of data processing and computational challenges. PLoS computational biology, 9(12), e1003345.