## **Resource Summary Report**

Generated by RRID on Apr 28, 2025

# **QSRA**

RRID:SCR\_010733 Type: Tool

**Proper Citation** 

QSRA (RRID:SCR\_010733)

#### **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

Funding:

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: 20250420T014507+0000

#### **Ratings and Alerts**

No rating or validation information has been found for QSRA.

No alerts have been found for QSRA.

#### Data and Source Information

Source: <u>SciCrunch Registry</u>

### **Usage and Citation Metrics**

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>RRID</u>.

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