

# Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

## Maqview

RRID:SCR\_005632

Type: Tool

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### Proper Citation

Maqview (RRID:SCR\_005632)

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### Resource Information

**URL:** <http://maq.sourceforge.net/maqview.shtml>

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**Description:** A graphical read alignment viewer specifically designed for the Maq alignment file and allows you to see the mismatches, base qualities and mapping qualities. It is highly efficient in speed, memory and disk usage. Maqview is based on OpenGL and is known to work on both Mac OS X and Linux. Porting to Windows is in principle easy.

**Abbreviations:** MaqView

**Synonyms:** Mapping and Assembly with Qualities Viewer, M.A.Q Viewer

**Resource Type:** data processing software, data visualization software, software application, software resource

**Defining Citation:** [DOI:10.1101/gr.078212.108](https://doi.org/10.1101/gr.078212.108)

**Keywords:** bio.tools

**Availability:** GNU General Public License

**Resource Name:** Maqview

**Resource ID:** SCR\_005632

**Alternate IDs:** OMICS\_00889, biotools:maqview

**Alternate URLs:** <https://bio.tools/maqview>, <https://sources.debian.org/src/maqview/>

**Record Creation Time:** 20220129T080231+0000

**Record Last Update:** 20260905T132539+0000

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### Ratings and Alerts

No rating or validation information has been found for Maqview.

No alerts have been found for Maqview.

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## Data and Source Information

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

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.