Maqview

RRID:SCR_005632
Type: Tool

Proper Citation

Maqview (RRID:SCR_005632)

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: software resource, data visualization software, software application, data processing software

Defining Citation: DOI:10.1101/gr.078212.108

Keywords: bio.tools

Funding:

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

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Ratings and Alerts

No rating or validation information has been found for Maqview.

No alerts have been found for Maqview.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.