

Resource Summary Report

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

EagleView

RRID:SCR_006859

Type: Tool

Proper Citation

EagleView (RRID:SCR_006859)

Resource Information

URL: <http://www.niehs.nih.gov/research/resources/software/biostatistics/eagleview/>

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Description: An information-rich viewer for next-generation genome assemblies with data integration capability. EagleView can display a dozen different types of information including base qualities, machine specific trace signals, and genome feature annotations. It provides an easy way for inspecting visually the quality of a genome assembly and validating polymorphism candidate sites (e.g., SNPs) reported by polymorphism discovery tools. It can also facilitate data interpretation and hypothesis generation. EagleView is a multi-platform application developed with C++ and is available for all three major platforms: Windows, Linux, and Mac OS.

Abbreviations: EagleView

Resource Type: software resource

Defining Citation: [PMID:18550804](#)

Keywords: bio.tools

Availability: Public, Free, Acknowledgement requested

Resource Name: EagleView

Resource ID: SCR_006859

Alternate IDs: biotools:eagleview, OMICS_00882

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

Record Creation Time: 20220129T080238+0000

Record Last Update: 20260905T132602+0000

Ratings and Alerts

No rating or validation information has been found for EagleView.

No alerts have been found for EagleView.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [Kravitz](#) .

Usher A, et al. (2025) Impact of sprint interval training on post-fatigue mitochondrial rate in professional boxers. European journal of applied physiology, 125(1), 261.

Pavlopoulos GA, et al. (2015) Visualizing genome and systems biology: technologies, tools, implementation techniques and trends, past, present and future. GigaScience, 4, 38.