## **Resource Summary Report**

Generated by RRID on Apr 28, 2025

# **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 assembles 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

Funding:

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: 20250420T014348+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: <u>SciCrunch Registry</u>

### **Usage and Citation Metrics**

We found 2 mentions in open access literature.

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

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.