

Resource Summary Report

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

Hawkeye

RRID:SCR_010982

Type: Tool

Proper Citation

Hawkeye (RRID:SCR_010982)

Resource Information

URL: <http://sourceforge.net/apps/mediawiki/amos/index.php?title=Hawkeye>

Proper Citation: Hawkeye (RRID:SCR_010982)

Description: A visual analytics tool for genome assembly analysis and validation, designed to aid in identifying and correcting assembly errors.

Abbreviations: Hawkeye

Resource Type: software resource

Resource Name: Hawkeye

Resource ID: SCR_010982

Alternate IDs: OMICS_00884

Record Creation Time: 20220129T080301+0000

Record Last Update: 20260905T132651+0000

Ratings and Alerts

No rating or validation information has been found for Hawkeye.

No alerts have been found for Hawkeye.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 9 mentions in open access literature.

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

Bresnahan PJ, et al. (2024) High-resolution ocean color imagery from the SeaHawk-HawkEye CubeSat mission. *Scientific data*, 11(1), 1246.

Akther S, et al. (2024) Natural selection and recombination at host-interacting lipoprotein loci drive genome diversification of Lyme disease and related bacteria. *mBio*, 15(9), e0174924.

Zhai Y, et al. (2021) Fabrication of Graphene Nanomesh FET Terahertz Detector. *Micromachines*, 12(6).

Brady SL, et al. (2019) Analysis of quantitative [I-123] mIBG SPECT/CT in a phantom and in patients with neuroblastoma. *EJNMMI physics*, 6(1), 31.

Fonseca A, et al. (2017) Genomic features of "Candidatus Venteria ishoeyi", a new sulfur-oxidizing macrobacterium from the Humboldt Sulfuretum off Chile. *PloS one*, 12(12), e0188371.

Sharaf A, et al. (2017) Functional and comparative genome analysis of novel virulent actinophages belonging to *Streptomyces flavovirens*. *BMC microbiology*, 17(1), 51.

Fioravanti V, et al. (2016) An Infrared Absorbance Sensor for the Detection of Melanoma in Skin Biopsies. *Sensors (Basel, Switzerland)*, 16(10).

Romano S, et al. (2016) Comparative Genomic Analysis Reveals a Diverse Repertoire of Genes Involved in Prokaryote-Eukaryote Interactions within the *Pseudovibrio* Genus. *Frontiers in microbiology*, 7, 387.

Antonuk LE, et al. (2009) An investigation of signal performance enhancements achieved through innovative pixel design across several generations of indirect detection, active matrix, flat-panel arrays. *Medical physics*, 36(7), 3322.