Resource Summary Report

Generated by NIF on Apr 19, 2025

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

Funding:

Resource Name: Hawkeye

Resource ID: SCR_010982

Alternate IDs: OMICS_00884

Record Creation Time: 20220129T080301+0000

Record Last Update: 20250410T070035+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 NIF.

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 sulfuroxidizing 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.