Resource Summary Report

Generated by FDI Lab - SciCrunch.org on May 13, 2025

AffyPipe

RRID:SCR_002032
Type: Tool

Proper Citation

AffyPipe (RRID:SCR_002032)

Resource Information

URL: https://github.com/nicolazzie/AffyPipe

Proper Citation: AffyPipe (RRID:SCR_002032)

Description: An open-source software pipeline for Affymetrix Axiom genotyping workflow.

Synonyms: AffyPipe: an open-source pipeline for Affymetrix Axiom genotyping workflow

Resource Type: software resource

Defining Citation: PMID:25028724

Keywords: affymetrix, bio.tools

Funding: Italian Ministry of Education University and Research 505/Ric;
project GenHome ;
European Union FP7 project Gene2Farm 289592

Availability: GNU General Public License, Acknowledgement requested

Resource Name: AffyPipe

Resource ID: SCR_002032

Alternate IDs: biotools:affypipe, OMICS_05203

Alternate URLs: https://bio.tools/affypipe

Record Creation Time: 20220129T080211+0000

Record Last Update: 20250420T014049+0000

Ratings and Alerts

No rating or validation information has been found for AffyPipe.

No alerts have been found for AffyPipe.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 5 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch.org.

Díaz-de Usera A, et al. (2022) Developing CIRdb as a catalog of natural genetic variation in the Canary Islanders. Scientific reports, 12(1), 16132.

Hernández-Beeftink T, et al. (2021) Whole-Blood Mitochondrial DNA Copies Are Associated With the Prognosis of Acute Respiratory Distress Syndrome After Sepsis. Frontiers in immunology, 12, 737369.

Ghoreishifar SM, et al. (2020) Genomic measures of inbreeding coefficients and genome-wide scan for runs of homozygosity islands in Iranian river buffalo, Bubalus bubalis. BMC genetics, 21(1), 16.

Mokhber M, et al. (2019) Study of whole genome linkage disequilibrium patterns of Iranian water buffalo breeds using the Axiom Buffalo Genotyping 90K Array. PloS one, 14(5), e0217687.

Mokhber M, et al. (2018) A genome-wide scan for signatures of selection in Azeri and Khuzestani buffalo breeds. BMC genomics, 19(1), 449.