Resource Summary Report

Generated by RRID on Aug 2, 2026

Gene Array Analyzer

RRID:SCR_008323
Type: Tool

Proper Citation

Gene Array Analyzer (RRID:SCR_008323)

Resource Information

URL: http://gaa.mpi-bn.mpg.de/

Proper Citation: Gene Array Analyzer (RRID:SCR_008323)

Description: Data analysis service that allows to process CEL files from Affymetrix, Inc. GeneChip Gene 1.0 ST Arrays to identify alternative splicing.

Abbreviations: GAA

Resource Type: data analysis service, service resource, production service resource, analysis service resource

Defining Citation: PMID:22123740

Keywords: bio.tools

Availability: Acknowledgement requested

Resource Name: Gene Array Analyzer

Resource ID: SCR_008323

Alternate IDs: OMICS_00759, biotools:gene_array_analyzer

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

Record Creation Time: 20220129T080246+0000

Record Last Update: 20260801T191003+0000

Ratings and Alerts

No rating or validation information has been found for Gene Array Analyzer.

No alerts have been found for Gene Array Analyzer.

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 RRID .

Marques-da-Silva C, et al. (2025) Type I interferons induce guanylate-binding proteins and lysosomal defense in hepatocytes to control malaria. Cell host & microbe, 33(4), 529.

Bosteels C, et al. (2022) Loss of GM-CSF-dependent instruction of alveolar macrophages in COVID-19 provides a rationale for inhaled GM-CSF treatment. Cell reports. Medicine, 3(12), 100833.

Vassen L, et al. (2014) Growth factor independence 1b (gfi1b) is important for the maturation of erythroid cells and the regulation of embryonic globin expression. PloS one, 9(5), e96636.

Buck D, et al. (2014) Removal of immunoglobulin-like domains from titin's spring segment alters titin splicing in mouse skeletal muscle and causes myopathy. The Journal of general physiology, 143(2), 215.

Dayeh TA, et al. (2013) Identification of CpG-SNPs associated with type 2 diabetes and differential DNA methylation in human pancreatic islets. Diabetologia, 56(5), 1036.