

Resource Summary Report

Generated by [NIF](#) on Aug 11, 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/>

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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: production service resource, data analysis service, analysis service resource, 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: 20260811T043327+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 [NIF](#) .

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.