

Resource Summary Report

Generated by SPARC SAWG on Sep 14, 2026

ArrayQuest

RRID:SCR_010935

Type: Tool

Proper Citation

ArrayQuest (RRID:SCR_010935)

Resource Information

URL: <http://proteogenomics.musc.edu/ma/arrayQuest.php?page=home&act=manage>

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Description: A web-accessible program for the analysis of DNA microarray data. ArrayQuest is designed to apply any type of DNA microarray analysis program executable on a Linux system (i.e., Bioconductor statistical and graphical methods written in R as well as BioPerl and C++ based scripts) to DNA microarray data stored in the MUSC DNA Microarray Database, the Gene Expression Omnibus (GEO) or in a password protected private database uploaded to the center point server. ArrayQuest analyses are performed on a computer cluster.

Abbreviations: ArrayQuest

Synonyms: ArrayQuest - An On-line DNA Microarray Analysis System

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

Defining Citation: [PMID:16321157](#)

Funding: University Research Resource Foundation ;
NCI R24CA095841;
NCRR P20RR016434

Availability: Free, Public

Resource Name: ArrayQuest

Resource ID: SCR_010935

Alternate IDs: OMICS_00746

Record Creation Time: 20220129T080301+0000

Record Last Update: 20260912T200206+0000

Ratings and Alerts

No rating or validation information has been found for ArrayQuest.

No alerts have been found for ArrayQuest.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Barth JL, et al. (2010) Jarid2 is among a set of genes differentially regulated by Nkx2.5 during outflow tract morphogenesis. Developmental dynamics : an official publication of the American Association of Anatomists, 239(7), 2024.