

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

Exon Array Analyzer

RRID:SCR_008684

Type: Tool

Proper Citation

Exon Array Analyzer (RRID:SCR_008684)

Resource Information

URL: <http://eaa.mpi-bn.mpg.de/>

Proper Citation: Exon Array Analyzer (RRID:SCR_008684)

Description: Service that allows you to process CEL files from Affymetrix, Inc. GeneChip Exon 1.0 ST Arrays to identify alternative splicing.

Abbreviations: Exon Array Analyzer

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

Resource Name: Exon Array Analyzer

Resource ID: SCR_008684

Alternate IDs: OMICS_00754

Record Creation Time: 20220129T080248+0000

Record Last Update: 20260728T043308+0000

Ratings and Alerts

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

No alerts have been found for Exon Array Analyzer.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#) .

Glaser SF, et al. (2020) The histone demethylase JMJD2B regulates endothelial-to-mesenchymal transition. *Proceedings of the National Academy of Sciences of the United States of America*, 117(8), 4180.

Perfetti A, et al. (2014) Genome wide identification of aberrant alternative splicing events in myotonic dystrophy type 2. *PloS one*, 9(4), e93983.

Weigand JE, et al. (2012) Hypoxia-induced alternative splicing in endothelial cells. *PloS one*, 7(8), e42697.

Ricci M, et al. (2012) Myocardial alternative RNA splicing and gene expression profiling in early stage hypoplastic left heart syndrome. *PloS one*, 7(1), e29784.