

Resource Summary Report

Generated by [Kravitz](#) on Sep 18, 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, data analysis service, production service resource, service resource

Resource Name: Exon Array Analyzer

Resource ID: SCR_008684

Alternate IDs: OMICS_00754

Record Creation Time: 20220129T080248+0000

Record Last Update: 20260912T200201+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 [Kravitz](#) .

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.

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.

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