Resource Summary Report

Generated by NIF on May 14, 2025

Exon Array Browser

RRID:SCR_008712
Type: Tool

Proper Citation

Exon Array Browser (RRID:SCR_008712)

Resource Information

URL: http://www.stanford.edu/group/exonarray/cgi-bin/plot_selector.pl

Proper Citation: Exon Array Browser (RRID:SCR_008712)

Description: Transcriptome database of acutely isolated purified astrocytes, neurons, and oligodendrocytes. Provides improved cell-type-specific markers for better understanding of neural development, function, and disease.

Abbreviations: Exon Array Browser

Resource Type: service resource, data or information resource, database

Defining Citation: PMID:18171944

Keywords: mature mouse, forebrain, transcriptome, astrocyte, neuron, oligodendrocyte, brain development, brain function, molecular neuroanatomy resource, visualization

Funding: NINDS R01NS045621;
NEI R01EY10257;
NEI EY07033;
Medical Scientist Training Program Grant MSTP GM07365;
Australian National Health and Medical Research Council CJ Martin Fellowship 400438;
NIDDK DK54388;
NCI CA095030

Availability: Free, Freely available

Resource Name: Exon Array Browser

Resource ID: SCR_008712

Alternate IDs: nlx_143565

Record Creation Time: 20220129T080248+0000

Record Last Update: 20250514T061455+0000

Ratings and Alerts

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

No alerts have been found for Exon Array Browser .

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 NIF.

Peng L, et al. (2013) Methodological limitations in determining astrocytic gene expression. Frontiers in endocrinology, 4, 176.