

# Resource Summary Report

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## eQTL Visualization Tool

RRID:SCR\_013413

Type: Tool

### Proper Citation

eQTL Visualization Tool (RRID:SCR\_013413)

### Resource Information

**URL:** <http://web.bioinformatics.ic.ac.uk/eqtlexplorer/>

**Proper Citation:** eQTL Visualization Tool (RRID:SCR\_013413)

**Description:** THIS RESOURCE IS NO LONGER IN SERVICE. Documented on June 1, 2023. eQTL Explorer was developed as a computational resource to visualize and explore data from combined genome-wide expression and linkage studies is essential for the development of testable hypotheses. This visualization tool stores expression profiles, linkage data and information from external sources in a relational database and enables simultaneous visualization and intuitive interpretation of the combined data via a Java graphical interface. eQTL Explorer also provides a new and powerful tool to interrogate these very large and complex datasets. eQTLExplorer allows users to mine and understand data from a repository of genetical genomics experiments. It will graphically display eQTL information based on a certain number of selection criteria, including: tissue type, p-value, cis/trans, probeset Affymetrix id and PQTL type. Sponsors: This work was funded by the MRC Clinical Sciences Centre and the Wellcome Trust programme for Cardiovascular Functional Genomics.

**Synonyms:** eQTLExplorer

**Resource Type:** data processing software, data visualization software, software application, software resource

**Keywords:** experiment, explore, expression, genome, genetic, genetical, cis, computational, data, database, genomic, grafical, interface, linkage, mine, pqtI type, p-value, repository, tissue, tissue type, trans, visualization, visualize

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** eQTL Visualization Tool

**Resource ID:** SCR\_013413

**Alternate IDs:** nif-0000-10222

**Record Creation Time:** 20220129T080316+0000

**Record Last Update:** 20260805T044312+0000

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## Ratings and Alerts

No rating or validation information has been found for eQTL Visualization Tool.

No alerts have been found for eQTL Visualization Tool.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Morrissey C, et al. (2011) Integrated genomic approaches to identification of candidate genes underlying metabolic and cardiovascular phenotypes in the spontaneously hypertensive rat. *Physiological genomics*, 43(21), 1207.