

Resource Summary Report

Generated by [RRID](#) on Aug 6, 2026

Human Gene Expression Index

RRID:SCR_007726

Type: Tool

Proper Citation

Human Gene Expression Index (RRID:SCR_007726)

Resource Information

URL: <http://zlab.bu.edu/HugeSearch/nph-HugeSearch.cgi>

Proper Citation: Human Gene Expression Index (RRID:SCR_007726)

Description: The Human Gene Expression Index (HuGE Index) aims to provide a comprehensive database to further our understanding of the expression of human genes in normal human tissues. mRNA expression levels of thousands of genes are obtained using high-density oligonucleotide array technology and used to create a public database. The website also provides interactive tools for researchers to query and visualize data over the Internet. To facilitate data analysis, genes are also cross-referenced with their annotation in the LocusLink database at NCBI.

Synonyms: HUGE Index

Resource Type: data or information resource, database

Keywords: human, human genome

Resource Name: Human Gene Expression Index

Resource ID: SCR_007726

Alternate IDs: nif-0000-02991

Old URLs: <http://www.hugeindex.org>

Record Creation Time: 20220129T080243+0000

Record Last Update: 20260806T054447+0000

Ratings and Alerts

No rating or validation information has been found for Human Gene Expression Index.

No alerts have been found for Human Gene Expression Index.

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 [RRID](#) .

Lin WH, et al. (2009) Topological and organizational properties of the products of house-keeping and tissue-specific genes in protein-protein interaction networks. BMC systems biology, 3, 32.

Sandberg R, et al. (2005) The molecular portrait of in vitro growth by meta-analysis of gene-expression profiles. Genome biology, 6(8), R65.

Maurer-Stroh S, et al. (2003) Protein prenyltransferases. Genome biology, 4(4), 212.

Lee SI, et al. (2003) Application of independent component analysis to microarrays. Genome biology, 4(11), R76.