

Resource Summary Report

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Human Brain Transcriptome

RRID:SCR_013742

Type: Tool

Proper Citation

Human Brain Transcriptome (RRID:SCR_013742)

Resource Information

URL: <http://hbatlas.org>

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Description: A data repository containing transcriptome and associated metadata for the developing and adult human brain. It provides genome-wide, exon-level transcriptome data from both sexes and multiple ethnicities.

Abbreviations: HBT

Resource Type: data or information resource, database

Defining Citation: [PMID:19477152](#)

Keywords: human brain, transcriptome data, brain regions, FASEB list

Funding: NIMH U01MH081896

Resource Name: Human Brain Transcriptome

Resource ID: SCR_013742

Record Creation Time: 20220129T080317+0000

Record Last Update: 20260811T043454+0000

Ratings and Alerts

No rating or validation information has been found for Human Brain Transcriptome.

No alerts have been found for Human Brain Transcriptome.

Data and Source Information

Usage and Citation Metrics

We found 107 mentions in open access literature.

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

Saenz-Anto??anzas A, et al. (2026) Differential Gene Expression in Human Hippocampus With Aging. Aging cell, 25(4), e70459.

Lourinho A, et al. (2026) RNArchitects: how hnRNPs shape neuronal landscapes. Brain : a journal of neurology, 149(5), 1469.

Xie Y, et al. (2026) Shared genetic architecture of gray matter deficits in schizophrenia and bipolar disorder: evidence from structural neuroimaging-genetic analyses. Psychological medicine, 56, e149.

Liu J, et al. (2026) Recent developments in imaging transcriptomics for psychiatric disorders. Psychoradiology, 6, kag010.

Wei Y, et al. (2025) Key regions aberrantly connected within cerebello-thalamo-cortical circuit and their genetic mechanism in schizophrenia: an fMRI meta-analysis and transcriptome study. Schizophrenia (Heidelberg, Germany), 11(1), 8.

Mamat M, et al. (2025) Molecular architecture of the altered cortical complexity in autism. Molecular autism, 16(1), 1.

Liu J, et al. (2025) PSMB4: a potential biomarker and therapeutic target for depression, perspective from integration analysis of depression GWAS data and human plasma proteome. Translational psychiatry, 15(1), 62.

Guan M, et al. (2025) Brain connectivity and transcriptional changes induced by rTMS in first-episode major depressive disorder. Translational psychiatry, 15(1), 159.

Zhong YL, et al. (2025) Genetic mechanisms of dynamic functional connectivity density in diabetic retinopathy brains: a combined transcriptomic and resting-state functional magnetic resonance imaging study. Frontiers in cellular neuroscience, 19, 1476038.

Lee YJ, et al. (2025) Molecular and developmental deficits in Smith-Magenis syndrome human stem cell-derived cortical neural models. American journal of human genetics, 112(10), 2338.

Huang X, et al. (2025) Genetic mechanisms of hemispheric functional connectivity in diabetic retinopathy: a joint neuroimaging and transcriptomic study. Frontiers in cell and developmental biology, 13, 1590627.

Hu Y, et al. (2025) Differences in the amplitude of dynamic low-frequency fluctuations in primary angle-closure glaucoma are associated with gene-molecular multi-omics. Frontiers in medicine, 12, 1679910.

Gao S, et al. (2025) SARM regulates cell apoptosis and inflammation during *Toxoplasma gondii* infection through a multistep mechanism. *Parasites & vectors*, 18(1), 103.

Yao W, et al. (2024) Associations between the multitrajectory neuroplasticity of neuronavigated rTMS-mediated angular gyrus networks and brain gene expression in AD spectrum patients with sleep disorders. *Alzheimer's & dementia : the journal of the Alzheimer's Association*, 20(11), 7885.

Zhou C, et al. (2024) Convergent and divergent genes expression profiles associated with brain-wide functional connectome dysfunction in deficit and non-deficit schizophrenia. *Translational psychiatry*, 14(1), 124.

Choi H, et al. (2024) Case Report: Intellectual disability and borderline intellectual functioning in two sisters with a 12p11.22 loss. *Frontiers in genetics*, 15, 1355823.

Do QB, et al. (2024) Early deficits in an in vitro striatal microcircuit model carrying the Parkinson's GBA-N370S mutation. *NPJ Parkinson's disease*, 10(1), 82.

Zhao P, et al. (2024) Analysis of epilepsy-associated variants in HCN3 - Functional implications and clinical observations. *Epilepsia open*, 9(6), 2294.

Le Grand Q, et al. (2024) Diffusion imaging genomics provides novel insight into early mechanisms of cerebral small vessel disease. *Molecular psychiatry*, 29(11), 3567.

Lederbauer J, et al. (2024) The role of DEAD- and DExH-box RNA helicases in neurodevelopmental disorders. *Frontiers in molecular neuroscience*, 17, 1414949.