

Resource Summary Report

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Brain heatmap

RRID:SCR_023815

Type: Tool

Proper Citation

Brain heatmap (RRID:SCR_023815)

Resource Information

URL: <http://liorkirsch.github.io/brain-heatmap-web/>

Proper Citation: Brain heatmap (RRID:SCR_023815)

Description: Custom Brain Heatmaps on top of the Allen Brain Atlas. Platform to generate brain heatmaps for the human or mouse brain on top of the Allen Brain Atlas by editing google spreadsheet template with specific brain region and gene information.

Resource Type: data or information resource, portal, project portal

Keywords: Allen Mouse Brain Reference Atlas, Human Brain Reference Atlas,

Availability: Free, Freely available

Resource Name: Brain heatmap

Resource ID: SCR_023815

Alternate URLs: <https://github.com/LiorKirsch/brain-heatmap-web>,
<https://github.com/LiorKirsch/brain-heatmap>

Record Creation Time: 20230721T050220+0000

Record Last Update: 20260919T195616+0000

Ratings and Alerts

No rating or validation information has been found for Brain heatmap.

No alerts have been found for Brain heatmap.

Data and Source Information

Usage and Citation Metrics

We found 34 mentions in open access literature.

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

Gao C, et al. (2025) The Potential Role of Intestinal Microbiota on the Intestine-Protective and Lipid-Lowering Effects of Berberine in Zebrafish (*Danio rerio*) Under High-Lipid Stress. *Metabolites*, 15(2).

Chen Y, et al. (2025) Sirtuin1 mitigates hypoxia-induced cardiomyocyte apoptosis in myocardial infarction via PHD3/HIF-1 α ?. *Molecular medicine (Cambridge, Mass.)*, 31(1), 100.

Xie R, et al. (2024) *Helicobacter pylori*-Induced Angiopoietin-Like 4 Promotes Gastric Bacterial Colonization and Gastritis. *Research (Washington, D.C.)*, 7, 0409.

Ye X, et al. (2024) Identification of YUC genes associated with leaf wrinkling trait in Tacai variety of Chinese cabbage. *PeerJ*, 12, e17337.

Liu Z, et al. (2024) Expression profiling of ALOG family genes during inflorescence development and abiotic stress responses in rice (*Oryza sativa* L.). *Frontiers in genetics*, 15, 1381690.

Zhai T, et al. (2024) Genome-Wide Identification and Expression Analysis Reveal bZIP Transcription Factors Mediated Hormones That Functions during Early Somatic Embryogenesis in *Dimocarpus longan*. *Plants (Basel, Switzerland)*, 13(5).

Wang H, et al. (2024) A complete reference genome of broomcorn millet. *Scientific data*, 11(1), 657.

Wang T, et al. (2024) Flavonoid 4,4'-dimethoxychalcone selectively eliminates senescent cells via activating ferritinophagy. *Redox biology*, 69, 103017.

Xu Z, et al. (2023) Genome-Wide Identification and Expression Analysis of WRKY Gene Family in *Neolamarckia cadamba*. *International journal of molecular sciences*, 24(8).

Jiang D, et al. (2023) Exploring the Heat Shock Transcription Factor (HSF) Gene Family in Ginger: A Genome-Wide Investigation on Evolution, Expression Profiling, and Response to Developmental and Abiotic Stresses. *Plants (Basel, Switzerland)*, 12(16).

Teng T, et al. (2023) Characteristics of glucose and lipid metabolism and the interaction between gut microbiota and colonic mucosal immunity in pigs during cold exposure. *Journal of animal science and biotechnology*, 14(1), 84.

Li P, et al. (2023) Function and Characteristic Analysis of Candidate PEAR Proteins in *Populus yunnanensis*. *International journal of molecular sciences*, 24(17).

Yang L, et al. (2023) Genome-wide identification and expression analysis of xyloglucan endotransglucosylase/hydrolase genes family in Salicaceae during grafting. BMC genomics, 24(1), 676.

Guo M, et al. (2023) PiggyBac Transposon Mining in the Small Genomes of Animals. Biology, 13(1).

Yang JS, et al. (2022) Chromosome-level genome assembly of the aquatic plant *Nymphoides indica* reveals transposable element bursts and NBS-LRR gene family expansion shedding light on its invasiveness. DNA research : an international journal for rapid publication of reports on genes and genomes, 29(4).

Liu X, et al. (2022) Development and validation of a novel immune-related prognostic signature in lung squamous cell carcinoma patients. Scientific reports, 12(1), 20737.

Zhang Z, et al. (2022) Assessing the neurotoxicity of airborne nano-scale particulate matter in human iPSC-derived neurons using a transcriptomics benchmark dose model. Toxicology and applied pharmacology, 449, 116109.

Zhu JY, et al. (2021) Genomic and Transcriptomic Analysis Reveals Cuticular Protein Genes Responding to Different Insecticides in Fall Armyworm *Spodoptera frugiperda*. Insects, 12(11).

Zhang Z, et al. (2021) FDX1 can Impact the Prognosis and Mediate the Metabolism of Lung Adenocarcinoma. Frontiers in pharmacology, 12, 749134.

Dong X, et al. (2021) Single-cell analysis reveals the intra-tumor heterogeneity and identifies MLXIPL as a biomarker in the cellular trajectory of hepatocellular carcinoma. Cell death discovery, 7(1), 14.