

# Resource Summary Report

Generated by [kravitz2](#) on Sep 5, 2026

## PsyGeNET

RRID:SCR\_014406

Type: Tool

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### Proper Citation

PsyGeNET (RRID:SCR\_014406)

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### Resource Information

**URL:**

<http://www.psygenet.org/web/PsyGeNET/menu;jsessionid=y6kgy9lqlxymr0nwwkkfo84>

**Proper Citation:** PsyGeNET (RRID:SCR\_014406)

**Description:** Knowledge platform on psychiatric disorders and their genes. Resource for exploratory analysis of psychiatric diseases and their associated genes. PsyGeNET is composed of database and set of analysis tools and is the result of the integration of information from DisGeNET and data extracted from the literature by text mining, followed by curation by domain experts.

**Synonyms:** Psychiatric disorders Gene association NETwork, Psychiatric disorders Gene association Network

**Resource Type:** data analysis software, data or information resource, data processing software, database, software application, software resource

**Defining Citation:** [DOI:10.1093/bioinformatics/btv301](https://doi.org/10.1093/bioinformatics/btv301)

**Keywords:** psychiatric disease, associated gene, database, analysis tool, bio.tools

**Related Condition:** Psychiatric disorder

**Availability:** Available for the research community

**Resource Name:** PsyGeNET

**Resource ID:** SCR\_014406

**Alternate IDs:** biotools:psygenet2r

**Alternate URLs:** <https://bio.tools/psygenet2r>

**License:** Open Database License, Database Contents License

**License URLs:** <http://www.psygenet.org/ds/PsyGeNET/html/legal.html>

**Record Creation Time:** 20220129T080320+0000

**Record Last Update:** 20260903T235244+0000

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## Ratings and Alerts

No rating or validation information has been found for PsyGeNET.

No alerts have been found for PsyGeNET.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 11 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [kravitz2](#) .

Yue X, et al. (2025) Changes in RNA Splicing: A New Paradigm of Transcriptional Responses to Probiotic Action in the Mammalian Brain. *Microorganisms*, 13(1).

Francis A, et al. (2025) Deconvolution-Based Transcriptomic Analysis in the Hippocampus Reveals Cell Type-Specific Risk Genes and Pathways Associated With Depression and Suicide. *Biological psychiatry global open science*, 5(5), 100530.

Kaurani L, et al. (2023) Baseline levels of miR-223-3p correlate with the effectiveness of electroconvulsive therapy in patients with major depression. *Translational psychiatry*, 13(1), 294.

Zhao Y, et al. (2021) Learning interpretable cellular and gene signature embeddings from single-cell transcriptomic data. *Nature communications*, 12(1), 5261.

Takahashi Y, et al. (2020) Systems Level Analysis and Identification of Pathways and Key Genes Associated with Delirium. *Genes*, 11(10).

Ivanov R, et al. (2019) Reconstruction and Analysis of Gene Networks of Human Neurotransmitter Systems Reveal Genes with Contentious Manifestation for Anxiety, Depression, and Intellectual Disabilities. *Genes*, 10(9).

Almasi SM, et al. (2019) Measuring the importance of vertices in the weighted human disease network. *PLoS one*, 14(3), e0205936.

Jin C, et al. (2019) Shank3 regulates striatal synaptic abundance of Cyld, a deubiquitinase specific for Lys63-linked polyubiquitin chains. *Journal of neurochemistry*, 150(6), 776.

Lee Y, et al. (2017) Striatal Transcriptome and Interactome Analysis of Shank3-overexpressing Mice Reveals the Connectivity between Shank3 and mTORC1 Signaling. *Frontiers in molecular neuroscience*, 10, 201.

Gutiérrez-Sacristán A, et al. (2017) Text mining and expert curation to develop a database on psychiatric diseases and their genes. *Database : the journal of biological databases and curation*, 2017.

Lee SA, et al. (2016) Epigenetic profiling of human brain differential DNA methylation networks in schizophrenia. *BMC medical genomics*, 9(Suppl 3), 68.