

# Resource Summary Report

Generated by SPARC SAWG on Jul 29, 2026

## Brainome portal

RRID:SCR\_018162

Type: Tool

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### Proper Citation

Brainome portal (RRID:SCR\_018162)

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### Resource Information

**URL:** [https://brainome.ucsd.edu/annoj/BICCN\\_MOp/](https://brainome.ucsd.edu/annoj/BICCN_MOp/)

**Proper Citation:** Brainome portal (RRID:SCR\_018162)

**Description:** Web based analytics environment for genomic data arising from BRAIN Initiative Cell Census Network, based on AnnoJ browser. AnnoJ browser displays epigenomic data, including base resolution DNA methylation, RNA-Seq, ChIP-Seq, as well as annotations.

**Synonyms:** Brainome

**Resource Type:** data or information resource, service resource, portal

**Keywords:** BICCN, BRAIN Initiative Cell Census Network, analytics environment, genomic data, AnnoJ, epigenomic, DNA methylation, RNAseq, ChIPseq

**Availability:** Free, Freely available

**Resource Name:** Brainome portal

**Resource ID:** SCR\_018162

**Record Creation Time:** 20220129T080339+0000

**Record Last Update:** 20260729T044450+0000

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### Ratings and Alerts

No rating or validation information has been found for Brainome portal.

No alerts have been found for Brainome portal.

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### Data and Source Information

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

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Hawrylycz M, et al. (2023) A guide to the BRAIN Initiative Cell Census Network data ecosystem. PLoS biology, 21(6), e3002133.

Guo C, et al. (2021) Graded heterogeneity of metabotropic signaling underlies a continuum of cell-intrinsic temporal responses in unipolar brush cells. Nature communications, 12(1), 5491.