## **Resource Summary Report**

Generated by NIF on May 3, 2025

# **Mammalian Brain Methylomes**

RRID:SCR 001648

Type: Tool

### **Proper Citation**

Mammalian Brain Methylomes (RRID:SCR\_001648)

#### **Resource Information**

URL: http://neomorph.salk.edu/brain\_methylomes/

**Proper Citation:** Mammalian Brain Methylomes (RRID:SCR\_001648)

**Description:** Datasets described in the manuscript: "Global Epigenomic Reconfiguration During Mammalian Brain Development" (Science, 2013 - DOI: 10.1126/science.1237905. This study provides genome-wide composition, patterning, cell specificity, and dynamics of DNA methylation at single-base resolution in human and mouse frontal cortex throughout their lifespan. Widespread methylome reconfiguration occurs during fetal to young adult development, coincident with synaptogenesis.

**Abbreviations:** Mammalian Brain Methylomes

**Synonyms:** Mammalian Brain Methylomes

**Resource Type:** data or information resource, data set

**Defining Citation:** PMID:23828890

Keywords: epigenetics, methylation, frontal cortex, development, neuron, methylome,

maturation, learning, young adult, fetus

**Funding:** 

Resource Name: Mammalian Brain Methylomes

Resource ID: SCR\_001648

Alternate IDs: nlx 153926

**Record Creation Time:** 20220129T080208+0000

**Record Last Update:** 20250429T054653+0000

## **Ratings and Alerts**

No rating or validation information has been found for Mammalian Brain Methylomes.

No alerts have been found for Mammalian Brain Methylomes.

### **Data and Source Information**

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We have not found any literature mentions for this resource.