

# Resource Summary Report

Generated by SPARC SAWG on Aug 9, 2026

## JAX Cre Repository

RRID:SCR\_005566

Type: Tool

### Proper Citation

JAX Cre Repository (RRID:SCR\_005566)

### Resource Information

**URL:** <http://cre.jax.org/index.html>

**Proper Citation:** JAX Cre Repository (RRID:SCR\_005566)

**Description:** Repository of Cre Driver lines and related information resources. Their services include analysis of Cre line excision function in both target and non-target tissues using Cre reporter lines and presenting the annotated data in the expression data portion of this website, <http://cre.jax.org/data.html>.

**Abbreviations:** Cre Repository

**Synonyms:** Jackson Laboratory Cre Repository, Cre Driver Strain Resources, The Jackson Laboratory Cre Repository

**Resource Type:** biomaterial supply resource, material resource, organism supplier

**Keywords:** cre, cre reporter, organism, cre driver line, cryopreserved, live, cre, cre expression, cre reporter strain, cre strain with floxed allele, image, strain, expression data, neurobiology

**Related Condition:** Cre, Cre expression, Cre reporter strain, Cre strain with floxed allele

**Funding:** NIH Blueprint for Neuroscience Research ;  
NIDCR ;  
NCRR RR03 2656;  
NCRR RR026117;  
NCRR RR001183;  
NIH Office of the Director OD011190;  
NIH Office of the Director OD010972;  
NIH Office of the Director DE020052

**Availability:** Public

**Resource Name:** JAX Cre Repository

**Resource ID:** SCR\_005566

**Alternate IDs:** nlx\_144662

**Record Creation Time:** 20220129T080231+0000

**Record Last Update:** 20260808T185829+0000

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## Ratings and Alerts

No rating or validation information has been found for JAX Cre Repository.

No alerts have been found for JAX Cre Repository.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 5 mentions in open access literature.

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

LaSarge CL, et al. (2016) Disrupted hippocampal network physiology following PTEN deletion from newborn dentate granule cells. *Neurobiology of disease*, 96, 105.

Nelms BD, et al. (2016) CellMapper: rapid and accurate inference of gene expression in difficult-to-isolate cell types. *Genome biology*, 17(1), 201.

Bersten DC, et al. (2015) Inducible and reversible lentiviral and Recombination Mediated Cassette Exchange (RMCE) systems for controlling gene expression. *PloS one*, 10(3), e0116373.

Guo J, et al. (2012) Deletion of FoxN1 in the thymic medullary epithelium reduces peripheral T cell responses to infection and mimics changes of aging. *PloS one*, 7(4), e34681.

Smedley D, et al. (2011) Cre recombinase resources for conditional mouse mutagenesis. *Methods (San Diego, Calif.)*, 53(4), 411.