Resource Summary Report

Generated by RRID on Aug 17, 2026

RHCglo

RRID:Addgene_80169
Type: Plasmid

Proper Citation

RRID:Addgene_80169

Plasmid Information

URL: http://www.addgene.org/80169

Proper Citation: RRID:Addgene_80169

Insert Name: segments of chicken cardiac troponin T (Tnnt2) introns 4 and 5 flanking an artifical exon

Organism: Gallus gallus

Bacterial Resistance: Ampicillin

Defining Citation: PMID:16925019

Vector Backbone Description: Vector Backbone:Bluescript KS+; Vector Types:Mammalian Expression; Bacterial Resistance:Ampicillin

Comments: Please note that there are some discrepancies between Addgene's quality control sequences and the depositor's sequence. The depositor noted that these discrepancies do NOT affect plasmid function.

Plasmid Name: RHCglo

Record Creation Time: 20220422T222524+0000

Record Last Update: 20260815T082037+0000

Ratings and Alerts

No rating or validation information has been found for RHCglo.

No alerts have been found for RHCglo.

Data and Source Information

Source: Addgene

Usage and Citation Metrics

We found 7 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID .

Gentile GM, et al. (2025) Alternative splicing of the Snap23 microexon is regulated by MBNL, QKI, and RBFOX2 in a tissue-specific manner and is altered in striated muscle diseases. RNA biology, 22(1), 1.

Guan B, et al. (2023) The qMini assay identifies an overlooked class of splice variants. medRxiv : the preprint server for health sciences.

Pagnamenta AT, et al. (2023) Structural and non-coding variants increase the diagnostic yield of clinical whole genome sequencing for rare diseases. Genome medicine, 15(1), 94.

Liu J, et al. (2023) Neuropathy target esterase activity predicts retinopathy among PNPLA6 disorders. bioRxiv : the preprint server for biology.

Vo T, et al. (2022) HNRNPH1 destabilizes the G-quadruplex structures formed by G-rich RNA sequences that regulate the alternative splicing of an oncogenic fusion transcript. Nucleic acids research, 50(11), 6474.

Bender C, et al. (2022) Predominant Founder Effect among Recurrent Pathogenic Variants for an X-Linked Disorder. Genes, 13(4).

Parry DA, et al. (2020) PRIM1 deficiency causes a distinctive primordial dwarfism syndrome. Genes & development, 34(21-22), 1520.