

Resource Summary Report

Generated by [ASWG](#) on Aug 17, 2026

[vha-1\(luc161\) III/hT2\[bli-4\(e937\) let-?\(q782\) qIs48\] \(I;III\).](#)

RRID:WB-STRAIN:WBStrain00051769

Type: Organism

Proper Citation

RRID:WB-STRAIN:WBStrain00051769

Organism Information

URL: <http://www.wormbase.org/db/get?name=WBStrain00051769>

Proper Citation: RRID:WB-STRAIN:WBStrain00051769

Description: Caenorhabditis elegans with name vha-1(luc161) III/hT2[bli-4(e937) let-?(q782) qIs48] (I;III). from WB.

Species: Caenorhabditis elegans

Synonyms: vha-1(luc161) III/hT2[bli-4(e937) let-?(q782) qIs48] (I;III).

Notes: Heterozygotes are wild-type with pharyngeal GFP signal, and segregate wild-type GFP, arrested hT2 aneuploids, and non-GFP luc161 homozygotes (embryonic lethal). Homozygous hT2[bli-4 let-? qIs48] inviable. Pick wild-type GFP and check for correct segregation of progeny to maintain. vha-1(luc161) is a 454 bp deletion removing most of the coding sequence. Reference: Gutierrez-Prez, P. et al. A deeply conserved miR-1 dependent regulon supports muscle cell physiology. bioRxiv, 2020, doi.org/10.1101/2020.08.31.275644. | "Made_by: Paula Gutierrez-Prez"

Affected Gene: WBGene00000254(bli-4)|WBGene00006910(vha-1)

Genomic Alteration: WBGene00000254(bli-4), WBGene00006910(vha-1)

Catalog Number: WB-STRAIN:WBStrain00051769

Database: WormBase (WB)

Database Abbreviation: WB

Availability: unknown

Organism Name: vha-1(luc161) III/hT2[bli-4(e937) let-?(q782) qIs48] (I;III).

Record Creation Time: 20230227T013849+0000

Record Last Update: 20260815T163749+0000

Ratings and Alerts

No rating or validation information has been found for vha-1(luc161) III/hT2[bli-4(e937) let-?(q782) qIs48] (I;III)..

No alerts have been found for vha-1(luc161) III/hT2[bli-4(e937) let-?(q782) qIs48] (I;III)..

Data and Source Information

Source: [Integrated Animals](#)

Source Database: WormBase (WB)

Usage and Citation Metrics

We have not found any literature mentions for this resource.