

Resource Summary Report

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VC2125

RRID:WB-STRAIN:WBStrain00037094

Type: Organism

Proper Citation

RRID:WB-STRAIN:WBStrain00037094

Organism Information

URL: <http://www.wormbase.org/db/get?name=WBStrain00037094>

Proper Citation: RRID:WB-STRAIN:WBStrain00037094

Description: Caenorhabditis elegans with name tfg-1(ok2290)/hln1 [unc-101(sy241)] I. from WB.

Species: Caenorhabditis elegans

Synonyms: tfg-1(ok2290)/hln1 [unc-101(sy241)] I.

Notes: This strain was provided by the C. elegans Reverse Genetics Core Facility at the University of British Columbia, which is part of the international C. elegans Gene Knockout Consortium, which should be acknowledged in any publications resulting from its use. Y63D3A.5. Apparent homozygous lethal deletion chromosome balanced by unc-101-marked inversion. Heterozygotes are WT, and segregate WT, Unc-101 hln1 homozygotes, and ok2290 homozygotes (arrest stage/phenotype undetermined). Pick WT and check for correct segregation of progeny to maintain. External left primer: AGCTCCGGATAAACTTGGGT. External right primer: TATTCCCTTCCCACCAATCA. Internal left primer: TTCCAGATGGTGCATTCAA. Internal right primer: AGACAGGAGCCCGAGATTTT. Internal WT amplicon: 2440 bp. Deletion size: 1264 bp. Deletion left flank: AGCAGATTAAGGTAAGGAGGATTTTGAGCG. Deletion right flank: CCACCACCGCAGGGAGCTCCCCAGCAAGGA."

Affected Gene: WBGene00006565(tfg-1)|WBGene00006829(unc-101)

Genomic Alteration: WBGene00006565(tfg-1), WBGene00006829(unc-101)

Catalog Number: WB-STRAIN:WBStrain00037094

Database: WormBase (WB)

Database Abbreviation: WB

Availability: available

Alternate IDs: WB-STRAIN:VC2125, CGC_VC2125

Organism Name: VC2125

Record Creation Time: 20230227T013658+0000

Record Last Update: 20260808T171528+0000

Ratings and Alerts

No rating or validation information has been found for VC2125.

No alerts have been found for VC2125.

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.