

Resource Summary Report

Generated by [dkNET](#) on Aug 16, 2026

VC2293

RRID:WB-STRAIN:WBStrain00037234

Type: Organism

Proper Citation

RRID:WB-STRAIN:WBStrain00037234

Organism Information

URL: <http://www.wormbase.org/db/get?name=WBStrain00037234>

Proper Citation: RRID:WB-STRAIN:WBStrain00037234

Description: *Caenorhabditis elegans* with name B0511.6(ok2948) I/hT2 [bli-4(e937) let-?(q782) qIs48] (I;III). from WB.

Species: *Caenorhabditis elegans*

Synonyms: B0511.6(ok2948) I/hT2 [bli-4(e937) let-?(q782) qIs48] (I;III).

Notes: B0511.6. Homozygous lethal deletion chromosome balanced by bli-4- and GFP-marked translocation. Heterozygotes are WT with pharyngeal GFP signal, and segregate WT GFP, arrested hT2 aneuploids, and non-GFP ok1948 homozygotes (early larval arrest). Homozygous hT2[bli-4 let-? qIs48] inviable. Pick WT GFP and check for correct segregation of progeny to maintain. External left primer: TCGTGTCTTTTCCCTTCTGG. External right primer: CGTTCATTTCCGGTCTTTGT. Internal left primer: TCATAAAATTTGTTAATTTTGCAGG. Internal right primer: CCAATGAACGAAACAACGTG. Internal WT amplicon: 1363 bp. Deletion size: 577 bp. Deletion left flank: TGGACGACTTTTGGATCATCTTCAGAATAC. Deletion right flank: ACAACTCGCGGAAATGTTTTTATTTAATT."|B0511.6. Homozygous lethal deletion chromosome balanced by bli-4- and GFP-marked translocation. Heterozygotes are WT with pharyngeal GFP signal, and segregate WT GFP, arrested hT2 aneuploids, and non-GFP ok2948 homozygotes (early larval arrest). Homozygous hT2[bli-4 let-? qIs48] inviable. Pick WT GFP and check for correct segregation of progeny to maintain. External left primer: TCGTGTCTTTTCCCTTCTGG. External right primer: CGTTCATTTCCGGTCTTTGT. Internal left primer: TCATAAAATTTGTTAATTTTGCAGG. Internal right primer: CCAATGAACGAAACAACGTG. Internal WT amplicon: 1363 bp. Deletion size: 577 bp. Deletion left flank: TGGACGACTTTTGGATCATCTTCAGAATAC. Deletion right flank: ACAACTCGCGGAAATGTTTTTATTTAATT."|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."

Affected Gene: WBGene00000254(bli-4)|WBGene00015232(B0511.6)

Genomic Alteration: WBGene00000254(bli-4), WBGene00015232(B0511.6)

Catalog Number: WB-STRAIN:WBStrain00037234

Database: WormBase (WB)

Database Abbreviation: WB

Availability: available

Alternate IDs: WB-STRAIN:VC2293, CGC_VC2293

Organism Name: VC2293

Record Creation Time: 20230227T013659+0000

Record Last Update: 20260815T163324+0000

Ratings and Alerts

No rating or validation information has been found for VC2293.

No alerts have been found for VC2293.

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.