

Resource Summary Report

Generated by [NIF](#) on Aug 28, 2026

VC30194

RRID:WB-STRAIN:WBStrain00038917

Type: Organism

Proper Citation

RRID:WB-STRAIN:WBStrain00038917

Organism Information

URL: <http://www.wormbase.org/db/get?name=WBStrain00038917>

Proper Citation: RRID:WB-STRAIN:WBStrain00038917

Description: Caenorhabditis elegans with name Whole-genome sequenced strain. from WB.

Species: Caenorhabditis elegans

Synonyms: Whole-genome sequenced strain.

Notes: Made_by: Vancouver KO Group|"Million Mutation Project strain. This strain was isolated after ENU mutagenesis of VC2010, propagated clonally through F10 to drive mutations to homozygosity, and subjected to whole-genome sequencing. It is homozygous for a large number of mutations determined from sequence data. It may also carry large copy number variations that are not homozygous. Alleles numbered between gk100000 and gk962522 are homozygous; those numbered from gk962523 up should be assumed to be non-homozygous. A graphical representation of these large copy number differences can be seen in the Plot section for each strain on the MMP web site (URL: genome.sfu.ca/mmp/).|"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."

Catalog Number: WB-STRAIN:WBStrain00038917

Database: WormBase (WB)

Database Abbreviation: WB

Availability: available

Alternate IDs: WB-STRAIN:VC30194, CGC_VC30194

Organism Name: VC30194

Record Creation Time: 20230227T013712+0000

Record Last Update: 20260815T163351+0000

Ratings and Alerts

No rating or validation information has been found for VC30194.

No alerts have been found for VC30194.

Data and Source Information

Source: [Integrated Animals](#)

Source Database: WormBase (WB)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Hodgkin J, et al. (2025) Mutations of nhr-49 affect C. elegans susceptibility to Yersinia biofilms. microPublication biology, 2025.