Resource Summary Report

Generated by ASWG on Aug 27, 2026

VC1662

RRID:WB-STRAIN:WBStrain00036775
Type: Organism

Proper Citation

RRID:WB-STRAIN:WBStrain00036775

Organism Information

URL: http://www.wormbase.org/db/get?name=WBStrain00036775

Proper Citation: RRID:WB-STRAIN:WBStrain00036775

Description: Caenorhabditis elegans with name lys-6(ok2075) IV. from WB.

Species: Caenorhabditis elegans

Synonyms: lys-6(ok2075) IV.

Notes: F58B3.3. External left primer: TTGTTTGATTGCACGTGGTT. External right primer: GATCGTGGTGTGGTTCACAG. Internal left primer: TTGGCTTCCAAACCATTTTC. Internal right primer: ATCAATGCCTCTGGATCGAC. Internal WT amplicon: 2135 bp. Deletion size: 1265 bp. Deletion left flank: GAGAACGCTTTCGTGAATCGGGATTTAAAA. Deletion right flank: TTTAGGCAAGGGAAGATGTATCCATCGACA. Insertion Sequence: GGCAAGGGAAGAAGAGA.|"Made_by: Vancouver KO Group"|"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: WBGene00003095(lys-6)

Genomic Alteration: WBGene00003095(lys-6)

Catalog Number: WB-STRAIN:WBStrain00036775

Database: WormBase (WB)

Database Abbreviation: WB

Availability: available

Alternate IDs: WB-STRAIN:VC1662, CGC_VC1662

Organism Name: VC1662

Record Creation Time: 20230227T013656+0000

Record Last Update: 20260815T163316+0000

Ratings and Alerts

No rating or validation information has been found for VC1662.

No alerts have been found for VC1662.

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 ASWG .

White CV, et al. (2018) Transcriptomic, Functional, and Network Analyses Reveal Novel Genes Involved in the Interaction Between Caenorhabditis elegans and Stenotrophomonas maltophilia. Frontiers in cellular and infection microbiology, 8, 266.