

Resource Summary Report

Generated by [NIF](#) on Aug 27, 2026

RB1787

RRID:WB-STRAIN:WBStrain00032478

Type: Organism

Proper Citation

RRID:WB-STRAIN:WBStrain00032478

Organism Information

URL: <http://www.wormbase.org/db/get?name=WBStrain00032478>

Proper Citation: RRID:WB-STRAIN:WBStrain00032478

Description: Caenorhabditis elegans with name F56F11.5(ok2305) III. from WB.

Species: Caenorhabditis elegans

Synonyms: F56F11.5(ok2305) III.

Notes: F56F11.5. Homozygous. Outer Left Sequence: ACAAGCTGAAAATGCCCAAC. Outer Right Sequence: ATTGCCAAAAGTTCGATTGC. Inner Left Sequence: CAGAAATGTCCATGAAATTAACAAA. Inner Right Sequence: CAGGCTCCAGTTCGAAGTTT. Inner Primer PCR Length: 3054 bp. Deletion Size: 2283 bp. Deletion left flank: TAACTCGGCTATAGCCTATAGCCGAGTTTC. Deletion right flank: TAATAGTGCCACACTGTGCGTAATTATTAT. Insertion Sequence: TGAGATTTTTCAACTTTTCAAAAAATCTTATAAAATCTAGAATTTTTTTGAATTTTTTAAGCATGATATTTTTGGTCTTTATGGCCCCATAGGCATGTTTTAAAGCAATTCCCACACATAGTGTAGTCCATCTTTAAGTTTCTATGTATAAAAGTAATTTTTACCATTGCTTTTGCTTGTAGGCAATCGCCATGATTTCAGACTTCGTTGAGACTTTTCAATTATATAATCACGGTAAGACTTCGAACTATCTTTCTTGATTGCGAAAACGAGGATGTGTAGTCAGCTTGCAATGCTTCTATTGCTTGACGTGGTCCCTGTTCCCCATTCAATTGAAGGTGAGTTATTACCTTACACGCAAGTTGTTCAAGTTCTCTGCAAATTTACAATTGAAAACCTCAC."F56F11.5. Homozygous. Outer Left Sequence: ACAAGCTGAAAATGCCCAAC. Outer Right Sequence: ATTGCCAAAAGTTCGATTGC. Inner Left Sequence: CAGAAATGTCCATGAAATTAACAAA. Inner Right Sequence: CAGGCTCCAGTTCGAAGTTT. Inner Primer PCR Length: 3054 bp. Deletion Size: 2283 bp. Deletion left flank: TAACTCGGCTATAGCCTATAGCCGAGTTTC. Deletion right flank: TAATAGTGCCACACTGTGCGTAATTATTAT. Insertion Sequence: TGAGATTTTTCAACTTTTCAAAAAATCTTATAAAATCTAGAATTTTTTTGAATTTTTTAAGCATGATATTTTTGGTCTTTATGGCCCCATAGGCATGTTTTAAAGCAATTCCCACACATAGTGTAGTCCATCTTTAAGTTTCTATGTATAAAAGTAATTTTTACCATTGCTTTTGCTTGTAGGCAATCGCCATGATTTCAGACTTCGTTGAGACTTTTCAATTATATAATCACGGTAAGACTTCGAACTATCTTTCTTGATTGCGAAAACGAGGATGTGTAGTCAGCTTGCAATGCTTCTATTGCTTGACGTGGTCCCTGTTCCCCATTCAATTGAAGGTGAGTTATTACCTTACACGCAAGTTGTTCAAGTTCTCTGCAAATTTACAATTGAAAACCTCAC."OMRF Knockout Group"]"This strain was provided by the C. elegans Gene Knockout Project at the Oklahoma Medical Research Foundation, which was part of the International

C. elegans Gene Knockout Consortium, which should be acknowledged in any publications resulting from its use."

Affected Gene: WBGene00018992(trpl-1)

Genomic Alteration: WBGene00018992(trpl-1)

Catalog Number: WB-STRAIN:WBStrain00032478

Database: WormBase (WB)

Database Abbreviation: WB

Availability: available

Alternate IDs: WB-STRAIN:RB1787, CGC_RB1787

Organism Name: RB1787

Record Creation Time: 20230227T013624+0000

Record Last Update: 20260815T163156+0000

Ratings and Alerts

No rating or validation information has been found for RB1787.

No alerts have been found for RB1787.

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](#) .

Flynn JM, et al. (2024) Dominant negative mutations in yeast Hsp90 reveal triage decision mechanism targeting client proteins for degradation. bioRxiv : the preprint server for biology.