

Resource Summary Report

Generated by [dkNET](#) on Aug 18, 2026

'+/mT1 [umnIs52] II; famh-136(hd7024[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/mT1 [dpy-10(e128)] III.

RRID:WB-STRAIN:WBStrain00055534

Type: Organism

Proper Citation

RRID:WB-STRAIN:WBStrain00055534

Organism Information

URL: <http://www.wormbase.org/db/get?name=WBStrain00055534>

Proper Citation: RRID:WB-STRAIN:WBStrain00055534

Description: Caenorhabditis elegans with name '+/mT1 [umnIs52] II; famh-136(hd7024[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/mT1 [dpy-10(e128)] III. from WB.

Species: Caenorhabditis elegans

Synonyms: '+/mT1 [umnIs52] II; famh-136(hd7024[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/mT1 [dpy-10(e128)] III.

Notes: umnIs52 [myo-2p::mKate2 + NeoR, III: 8856215 (intergenic)] II. Pick wild-type GFP+ and mKate2+ animals to maintain. Apparent homozygous lethal deletion balanced over mT1. Heterozygotes are wild-type GFP+ mKate2+, and segregate wild-type GFP+ mKate2+, GFP+ non-mKate2 (hd7024 homozygotes), sterile Dpy non-GFP mKate2+ mT1 homozygotes, and large numbers of arrested aneuploid embryos. Derived from parental strains VH7027 and CGC66. hd7024 is a 542 bp deletion with Calarco/Colaiacovo selection cassette conferring myo-2 GFP and G418 resistance inserted at break. Left flanking sequence: TTTTGAGCTTTAGCTATCTTCCTTTATCCT; Right flanking sequence: ATCACAGGTGTGAAGATTTATTAAATTTTA. sgRNA #1: ATGAATGACGCAAGAATTAA; sgRNA #2: TCTGAATACTTATTGACATC. Please reference Au et al., G3 9(1): 135-144 2019 in any work resulting from use of this mutation.

Affected Gene: WBGene00001072(dpy-10)|WBGene00003514(myo-2)|WBGene00004496(rps-27)|WBGene00006789(unc-54)|WBGene00014022(famh-136)

Genomic Alteration: WBGene00001072(dpy-10), WBGene00003514(myo-2), WBGene00004496(rps-27), WBGene00006789(unc-54), WBGene00014022(famh-136)

Catalog Number: WB-STRAIN:WBStrain00055534

Database: WormBase (WB)

Database Abbreviation: WB

Availability: unknown

Organism Name: '+/mT1 [umnIs52] II; famh-136(hd7024[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/mT1 [dpy-10(e128)] III.

Record Creation Time: 20240716T202611+0000

Record Last Update: 20260815T163847+0000

Ratings and Alerts

No rating or validation information has been found for '+/mT1 [umnIs52] II; famh-136(hd7024[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/mT1 [dpy-10(e128)] III..

No alerts have been found for '+/mT1 [umnIs52] II; famh-136(hd7024[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/mT1 [dpy-10(e128)] III..

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.