

# Resource Summary Report

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[dna-2\(ve776\[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP\]\)/mnC1 \[dpy-10\(e128\) unc-52\(e444\) umnIs37\] II.](#)

RRID:WB-STRAIN:WBStrain00055425

Type: Organism

## Proper Citation

RRID:WB-STRAIN:WBStrain00055425

## Organism Information

**URL:** <http://www.wormbase.org/db/get?name=WBStrain00055425>

**Proper Citation:** RRID:WB-STRAIN:WBStrain00055425

**Description:** Caenorhabditis elegans with name dna-2(ve776[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/mnC1 [dpy-10(e128) unc-52(e444) umnIs37] II. from WB.

**Species:** Caenorhabditis elegans

**Synonyms:** dna-2(ve776[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/mnC1 [dpy-10(e128) unc-52(e444) umnIs37] II.

**Notes:** Made\_by: RG KO Group|"umnIs37 [myo-2p::mKate2 + NeoR, II: 11755713 (intergenic)] II. Maternal effect sterile. Deletion of 3982 bp with Calarco/Colaiacovo selection cassette conferring myo-2 GFP and G418 resistance inserted at break in parental strain N2. Heterozygotes are wild-type GFP+ mKate2+ and segregate wild-type GFP+ mKate2+ heterozygotes, GFP+ non-mKate2 adults that produce sterile progeny (ve776 homozygotes) and paralysed DpyUnc non-GFP mKate2+ (mnC1 homozygotes). Maintain by picking wild-type GFP+ mKate2+. Left flanking Sequence: aaatgtctgctgccccgcccgcgttgct ; Right flanking sequence: cctttttactcatttattagatttctcac. sgRNA #1: gattctggctgcgaaatacg; sgRNA #2: cggaattTTACAGTTGAGT. Please reference Au et al., G3 9(1): 135-144 2019 in any work resulting from use of this mutation."

**Affected Gene:** WBGene00001016(dna-2)|WBGene00001072(dpy-10)|WBGene00003514(myo-2)|WBGene00004496(rps-27)|WBGene00006787(unc-52)|WBGene00006789(unc-54)

**Genomic Alteration:** WBGene00001016(dna-2), WBGene00001072(dpy-10), WBGene00003514(myo-2), WBGene00004496(rps-27), WBGene00006787(unc-52), WBGene00006789(unc-54)

**Catalog Number:** WB-STRAIN:WBStrain00055425

**Database:** WormBase (WB)

**Database Abbreviation:** WB

**Availability:** unknown

**Organism Name:** dna-2(ve776[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/mnC1 [dpy-10(e128) unc-52(e444) umnIs37] II..

**Record Creation Time:** 20240716T202610+0000

**Record Last Update:** 20260725T172824+0000

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## Ratings and Alerts

No rating or validation information has been found for dna-2(ve776[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/mnC1 [dpy-10(e128) unc-52(e444) umnIs37] II..

No alerts have been found for dna-2(ve776[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/mnC1 [dpy-10(e128) unc-52(e444) umnIs37] II..

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## Data and Source Information

**Source:** [Integrated Animals](#)

**Source Database:** WormBase (WB)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.