

Resource Summary Report

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[npri-3\(ve778\[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP\]\)/nT1\[umnl49\]IV; +/-nT1 V.](#)

RRID:WB-STRAIN:WBStrain00055427

Type: Organism

Proper Citation

RRID:WB-STRAIN:WBStrain00055427

Organism Information

URL: <http://www.wormbase.org/db/get?name=WBStrain00055427>

Proper Citation: RRID:WB-STRAIN:WBStrain00055427

Description: *Caenorhabditis elegans* with name npri-3(ve778[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/nT1[umnl49]IV; +/-nT1 V. from WB.

Species: *Caenorhabditis elegans*

Synonyms: npri-3(ve778[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/nT1[umnl49]IV; +/-nT1 V.

Notes: Made_by: RG KO Group|"umnl49 [myo-2p::mKate2 + NeoR, V: 1005689 (intergenic)] IV. Possible maternal effect: descendants of homozygous animals may be sensitive to starvation (lethal). Deletion of 2127 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+, GFP+ non-mKate adults (ve778 homozygotes), Vul non-GFP mKate2+ (nT1 homozygotes) and dead eggs (aneuploids). Maintain by picking wild-type GFP+ mKate2+. Maintain by picking wild-type GFP+ mKate2+. Left flanking Sequence: AAATAATTTTGAATTTTCAGTGCTGTGATG ; Right flanking sequence: AGTAGCCAAGGATCTTTTATTCGGTTTTT. sgRNA #1: TGTTTTCTCACTGGACAAAG; sgRNA #2: CGGCAACAAAATCAGGACGG. Please reference Au et al., G3 9(1): 135-144 2019 in any work resulting from use of this mutation."|"umnl49 [myo-2p::mKate2 + NeoR, V: 1005689 (intergenic)] IV. Possible maternal effect: descendants of homozygous animals may be sensitive to starvation (lethal). Deletion of 2127 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+, GFP+ non-mKate adults (ve780 homozygotes), Vul

non-GFP mKate2+ (nT1 homozygotes) and dead eggs (aneuploids). Maintain by picking wild-type GFP+ mKate2+. Maintain by picking wild-type GFP+ mKate2+. Left flanking Sequence: AAATAATTTTGAATTTTCAGTGCTGTGATG ; Right flanking sequence: AGTAGCCAAGGATCTTTTATTCCGGTTTTT. sgRNA #1: TGTTTTCTCACTGGACAAAG; sgRNA #2: CGGCAACAAAATCAGGACGG. Please reference Au et al., G3 9(1): 135-144 2019 in any work resulting from use of this mutation."

Affected Gene: WBGene00003514(myo-2)|WBGene00004496(rps-27)|WBGene00006789(unc-54)|WBGene00018072(npri-3)

Genomic Alteration: WBGene00003514(myo-2), WBGene00004496(rps-27), WBGene00006789(unc-54), WBGene00018072(npri-3)

Catalog Number: WB-STRAIN:WBStrain00055427

Database: WormBase (WB)

Database Abbreviation: WB

Availability: unknown

Organism Name: npri-3(ve778[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/nT1[umnl49]IV; +/-nT1 V.

Record Creation Time: 20240716T202610+0000

Record Last Update: 20260829T162857+0000

Ratings and Alerts

No rating or validation information has been found for npri-3(ve778[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/nT1[umnl49]IV; +/-nT1 V..

No alerts have been found for npri-3(ve778[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/nT1[umnl49]IV; +/-nT1 V..

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.