

Resource Summary Report

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F10G8.9(ve769[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/hT2 [umnIs73] I; +/-hT2 [bli-4(e937) let-?(h661)] III.

RRID:WB-STRAIN:WBStrain00055419

Type: Organism

Proper Citation

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Organism Information

URL: <http://www.wormbase.org/db/get?name=WBStrain00055419>

Proper Citation: RRID:WB-STRAIN:WBStrain00055419

Description: Caenorhabditis elegans with name F10G8.9(ve769[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/hT2 [umnIs73] I; +/-hT2 [bli-4(e937) let-?(h661)] III. from WB.

Species: Caenorhabditis elegans

Synonyms: F10G8.9(ve769[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/hT2 [umnIs73] I; +/-hT2 [bli-4(e937) let-?(h661)] III.

Notes: Made_by: RG KO Group| "umnIs73 [myo-2p::mKate2 + NeoR, III: 9421936 (intergenic)] I. Homozygous sterile. Deletion of 2540 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-mKate2 sterile adults (ve769 homozygotes), non-GFP mKate2+ arrested animals (arrest stage unknown)(hT2 homozygotes) and dead eggs (aneuploids). Pick wild-type GFP+ mKate2+ and check for correct segregation of progeny to maintain. [NOTE: Apparently the lethal mutation is closely linked but not within the balanced region of hT2. It can occasionally recombine away so that the strain will segregate Bli-4 hT2 homozygotes. (Mark Edgley)] Left flanking Sequence: atgaaattaaaaataaaaaattttgag ; Right flanking sequence: ATTTGTAATTCATTTGGATTCGGTGCCACA. sgRNA #1: ATGCACCGTGTTGTTATAAC; sgRNA #2: TGAGATTTCGCGATTTATTTC. Please reference Au et al., G3 9(1): 135-144 2019 in any work resulting from use of this mutation."

Affected Gene: WBGene00000254(bli-4)|WBGene00003514(myo-2)|WBGene00004496(rps-27)|WBGene00006789(unc-54)

Genomic Alteration: WBGene00000254(bli-4), WBGene00003514(myo-2), WBGene00004496(rps-27), WBGene00006789(unc-54)

Catalog Number: WB-STRAIN:WBStrain00055419

Database: WormBase (WB)

Database Abbreviation: WB

Availability: unknown

Organism Name: F10G8.9(ve769[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/hT2 [umnIs73] I; +/-hT2 [bli-4(e937) let-?(h661)] III..

Record Creation Time: 20240716T202610+0000

Record Last Update: 20260829T162855+0000

Ratings and Alerts

No rating or validation information has been found for F10G8.9(ve769[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/hT2 [umnIs73] I; +/-hT2 [bli-4(e937) let-?(h661)] III..

No alerts have been found for F10G8.9(ve769[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/hT2 [umnIs73] I; +/-hT2 [bli-4(e937) let-?(h661)] 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.