

Resource Summary Report

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plc-3 & srh-135(ve884[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/mIn1[dpy-10(e128) umnIs43] II.

RRID:WB-STRAIN:WBStrain00062990

Type: Organism

Proper Citation

RRID:WB-STRAIN:WBStrain00062990

Organism Information

URL: <http://www.wormbase.org/db/get?name=WBStrain00062990>

Proper Citation: RRID:WB-STRAIN:WBStrain00062990

Description: Caenorhabditis elegans with name plc-3 & srh-135(ve884[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/mIn1[dpy-10(e128) umnIs43] II. from WB.

Species: Caenorhabditis elegans

Synonyms: plc-3 & srh-135(ve884[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/mIn1[dpy-10(e128) umnIs43] II.

Notes: Made_by: RG KO Group|"umnIs43 [myo-2p::mKate2 + NeoR, II: 11755713 (intergenic)] II. Homozygous Ste. Deletion of 6,190 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 (ve884 homozygotes), and Dpy non-GFP mKate2+ (mIn1 homozygotes). Maintain by picking wild-type GFP+ mKate2+ and check for correct segregation of progeny. Left flanking Sequence: CTCACCTTGGCCCATCATCGTCGTCGCGAAA; Right flanking sequence: TTTTGGAGCATT TTTTATAGTTTAAGTTTA. plc-3_srh-135 sgRNA A: GACTACAGTAACCAGCACAG; plc-3_srh-135 sgRNA B: TTTTGTGGCAAAAAACAGAA. 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)|WBGene00004038(plc-3)|WBGene00004496(rps-27)|WBGene00005352(srh-135)|WBGene00006789(unc-54)

Genomic Alteration: WBGene00001072(dpy-10), WBGene00003514(myo-2), WBGene00004038(plc-3), WBGene00004496(rps-27), WBGene00005352(srh-135), WBGene00006789(unc-54)

Catalog Number: WB-STRAIN:WBStrain00062990

Database: WormBase (WB)

Database Abbreviation: WB

Availability: unknown

Organism Name: plc-3 & srh-135(ve884[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/mIn1[dpy-10(e128) umnIs43] II.

Record Creation Time: 20250827T213352+0000

Record Last Update: 20260829T163111+0000

Ratings and Alerts

No rating or validation information has been found for plc-3 & srh-135(ve884[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/mIn1[dpy-10(e128) umnIs43] II..

No alerts have been found for plc-3 & srh-135(ve884[LoxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + LoxP])/mIn1[dpy-10(e128) umnIs43] II..

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.