

Resource Summary Report

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sss-1(gk5609[loxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + loxP]) IV.

RRID:WB-STRAIN:WBStrain00051226

Type: Organism

Proper Citation

RRID:WB-STRAIN:WBStrain00051226

Organism Information

URL: <http://www.wormbase.org/db/get?name=WBStrain00051226>

Proper Citation: RRID:WB-STRAIN:WBStrain00051226

Description: Caenorhabditis elegans with name sss-1(gk5609[loxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + loxP]) IV. from WB.

Species: Caenorhabditis elegans

Synonyms: sss-1(gk5609[loxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + loxP]) IV.

Notes: Homozygous viable. Deletion of 1256 bp with Calarco/Colaiacovo selection cassette conferring myo-2 GFP and G418 resistance inserted at break. Left flanking sequence: AGTTATAGAGAGAGGTTTAGAGAGAGAGCA. Right flanking sequence: AGGTTTAAGGTTTAAAAAATGCGACAAAGACATTTTTTGGACAATATTCT. Please reference Au et al., G3 9(1): 135-144 2019 in any work resulting from use of this mutation.|"Made_by: Vancouver KO Group"

Affected Gene: WBGene00003514(myo-2)|WBGene00004496(rps-27)|WBGene00006056(sss-1)|WBGene00006789(unc-54)

Genomic Alteration: WBGene00003514(myo-2), WBGene00004496(rps-27), WBGene00006056(sss-1), WBGene00006789(unc-54)

Catalog Number: WB-STRAIN:WBStrain00051226

Database: WormBase (WB)

Database Abbreviation: WB

Availability: unknown

Organism Name: sss-1(gk5609[loxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + loxP]) IV.

Record Creation Time: 20230227T013846+0000

Record Last Update: 20260808T171838+0000

Ratings and Alerts

No rating or validation information has been found for sss-1(gk5609[loxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + loxP]) IV..

No alerts have been found for sss-1(gk5609[loxP + myo-2p::GFP::unc-54 3' UTR + rps-27p::neoR::unc-54 3' UTR + loxP]) IV..

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.