

# Resource Summary Report

Generated by [dkNET](#) on Aug 21, 2026

## ninaE-stops m3-myc

RRID:Addgene\_26855

Type: Plasmid

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### Proper Citation

RRID:Addgene\_26855

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### Plasmid Information

**URL:** <http://www.addgene.org/26855>

**Proper Citation:** RRID:Addgene\_26855

**Insert Name:** Slow Termination of Phototransduction

**Organism:** Drosophila melanogaster

**Bacterial Resistance:** Ampicillin

**Defining Citation:** [PMID:18184564](#)

**Vector Backbone Description:** Backbone Size:0; Vector Backbone:pCNX; Vector Types:Insect Expression; Bacterial Resistance:Ampicillin

**Comments:** To express the stops cDNA under the control of the ninaE promoter, the cDNA (EST clone GH07253, DGRC) was subcloned between the NotI and XbaI sites of pCNX. To express Myc-tagged wild-type STOPS in flies, the sequence encoding a Myc tag was fused to the 3' end of the wild-type stops cDNA. The wild-type stops cDNAs was subsequently subcloned between the NotI and XbaI sites of pCNX. The pCNX vector was constructed as follows. First, the pNX vector (provided by Dr. J. O'Tousa, University of Notre Dame, Notre Dame, IN) was digested with KpnI and NotI, which liberated a 3 kb ninaE promoter sequence, and this fragment was subsequently introduced between the same sites of pCasper4. Second, to provide the signals for polyadenylation and 3' end cleavage of RNAs, a simian virus 40 3' region (0.5 kb) was excised from pCasper-heat shock protein 70 promoter (hs) and introduced between the XbaI and XhoI sites of the vector

**Plasmid Name:** ninaE-stops m3-myc

**Relevant Mutation:** Amino acids 296-299 (SLRH in wild-type) are mutated to GPGD

**Record Creation Time:** 20220422T222120+0000

**Record Last Update:** 20260815T081257+0000

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## Ratings and Alerts

No rating or validation information has been found for ninaE-stops m3-myc.

No alerts have been found for ninaE-stops m3-myc.

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## Data and Source Information

**Source:** [Addgene](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.