

Resource Summary Report

Generated by [NIF](#) on Sep 22, 2026

pSLIK 3xFLAG-MAVS-Ala148,Ala271 hygro

RRID:Addgene_158639

Type: Plasmid

Proper Citation

RRID:Addgene_158639

Plasmid Information

URL: <http://www.addgene.org/158639>

Proper Citation: RRID:Addgene_158639

Insert Name: MAVS

Organism: Homo sapiens

Bacterial Resistance: Ampicillin

Defining Citation: [PMID:33740397](#)

Vector Backbone Description: Backbone Marker:Addgene #25737; Vector Backbone:pSLIK hygro; Vector Types:Lentiviral; Bacterial Resistance:Ampicillin

Comments: The plasmid differs from the MAVS reference sequence at 3 position (Q93E, Q198K, and S409F). This plasmid has an additional Q148A and Q271A mutation. For pSLIK vectors, it is recommended to use zeocin in addition to ampicillin for bacterial selection.

Plasmid Name: pSLIK 3xFLAG-MAVS-Ala148,Ala271 hygro

Relevant Mutation: Q148A/Q271A

Record Creation Time: 20220422T221859+0000

Record Last Update: 20260919T091156+0000

Ratings and Alerts

No rating or validation information has been found for pSLIK 3xFLAG-MAVS-Ala148,Ala271 hygro.

No alerts have been found for pSLIK 3xFLAG-MAVS-Ala148,Ala271 hygro.

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at [NIF](#) .

Lopacinski AB, et al. (2021) Modeling the complete kinetics of coxsackievirus B3 reveals human determinants of host-cell feedback. Cell systems, 12(4), 304.