

Resource Summary Report

Generated by [NIF](#) on Jul 29, 2026

pSLIK 3xFLAG-MAVS-Ala148 hygro

RRID:Addgene_158636

Type: Plasmid

Proper Citation

RRID:Addgene_158636

Plasmid Information

URL: <http://www.addgene.org/158636>

Proper Citation: RRID:Addgene_158636

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 positions: Q93E, Q198K, and S409F. This plasmid has an additional Q148A mutation. For pSLIK vectors, it is recommended to use zeocin in addition to ampicillin for bacterial selection.

Plasmid Name: pSLIK 3xFLAG-MAVS-Ala148 hygro

Relevant Mutation: Q148A

Record Creation Time: 20220422T221859+0000

Record Last Update: 20260725T074101+0000

Ratings and Alerts

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

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