

Resource Summary Report

Generated by [RRID](#) on Sep 1, 2026

RRL.sin.cPPT.CMV/Flag-NCOA7 variant 6.IRES-puro.WPRE (CG494)

RRID:Addgene_139447

Type: Plasmid

Proper Citation

RRID:Addgene_139447

Plasmid Information

URL: <http://www.addgene.org/139447>

Proper Citation: RRID:Addgene_139447

Insert Name: FLAG-tag NCOA7 variant 6 (coding for isoform 4, the short, interferon-inducible isoform)

Organism: Homo sapiens

Bacterial Resistance: Ampicillin

Defining Citation: [PMID:30478388](#)

Vector Backbone Description: Backbone Marker:PMID: 24048477; Vector Backbone:RRL.sin.cPPT.CMV/IRES-puro.WPRE; Vector Types:Mammalian Expression, Lentiviral; Bacterial Resistance:Ampicillin

Plasmid Name: RRL.sin.cPPT.CMV/Flag-NCOA7 variant 6.IRES-puro.WPRE (CG494)

Record Creation Time: 20220422T221806+0000

Record Last Update: 20260902T041953+0000

Ratings and Alerts

No rating or validation information has been found for RRL.sin.cPPT.CMV/Flag-NCOA7 variant 6.IRES-puro.WPRE (CG494).

No alerts have been found for RRL.sin.cPPT.CMV/Flag-NCOA7 variant 6.IRES-puro.WPRE (CG494).

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Ruggiero-Ruff RE, et al. (2026) Discovery proteomics identification of factors contributing to gonadotropin ?? expression. Endocrinology, 167(8).

Oot RA, et al. (2024) Human V-ATPase function is positively and negatively regulated by TLDC proteins. Structure (London, England : 1993), 32(7), 989.

Le Pen J, et al. (2024) A genome-wide arrayed CRISPR screen identifies PLSCR1 as an intrinsic barrier to SARS-CoV-2 entry that recent virus variants have evolved to resist. PLoS biology, 22(9), e3002767.