

# Resource Summary Report

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

## pCAG-GST-ATG101-W110D/F112D

RRID:Addgene\_188106

Type: Plasmid

### Proper Citation

RRID:Addgene\_188106

### Plasmid Information

**URL:** <http://www.addgene.org/188106>

**Proper Citation:** RRID:Addgene\_188106

**Insert Name:** ATG101

**Organism:** Homo sapiens

**Bacterial Resistance:** Ampicillin

**Defining Citation:** [PMID:36791199](#)

**Vector Backbone Description:** Vector Backbone:pCAG; Vector Types:Mammalian Expression; Bacterial Resistance:Ampicillin

**Plasmid Name:** pCAG-GST-ATG101-W110D/F112D

**Relevant Mutation:** codon-optimized ATG101-W110D/F112D

**Record Creation Time:** 20220819T080230+0000

**Record Last Update:** 20260829T081416+0000

### Ratings and Alerts

No rating or validation information has been found for pCAG-GST-ATG101-W110D/F112D.

No alerts have been found for pCAG-GST-ATG101-W110D/F112D.

### 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 [RRID](#) .

Ren X, et al. (2023) Structural basis for ATG9A recruitment to the ULK1 complex in mitophagy initiation. Science advances, 9(7), eadg2997.