

# Resource Summary Report

Generated by [RRID](#) on Sep 3, 2026

## AICSDP-35:AAVS1-mEGFP

RRID:Addgene\_91565

Type: Plasmid

### Proper Citation

RRID:Addgene\_91565

### Plasmid Information

**URL:** <http://www.addgene.org/91565>

**Proper Citation:** RRID:Addgene\_91565

**Insert Name:** AAVS1 Homology Arms with CAGGS driven mEGFP

**Organism:** Homo sapiens

**Bacterial Resistance:** Ampicillin

**Vector Backbone Description:** Vector Backbone:unknown; Vector Types:Mammalian Expression, CRISPR, Donor Template; Bacterial Resistance:Ampicillin

**Comments:** This plasmid has been used with locus-specific CRISPR/Cas9 to insert a CAGGS-mEGFP at the AAVS1 safe harbor locus in WTC human induced pluripotent stem cells by the Allen Institute for Cell Science. After insertion of the reporter using this donor template plasmid and CRISPR/Cas9 reagents, transfected cells may exhibit varying intensity levels of fluorescence, likely due to editing precision. To obtain cells of uniform intensity levels, see our protocol for fluorescence-assisted cell sorting and subcloning of transfected cells (<https://www.allencell.org/instructional-videos-and-tutorials-for-cell-methods.html>). Further, we recommend PCR-based assays for identifying precisely edited clones as previously described (<https://www.molbiolcell.org/doi/abs/10.1091/mbc.e17-03-0209>). For more information on the entire plasmid collection, please see <https://www.addgene.org/allen-institute-cell-science/>.

**Plasmid Name:** AICSDP-35:AAVS1-mEGFP

**Record Creation Time:** 20220422T222619+0000

**Record Last Update:** 20260902T050517+0000

### Ratings and Alerts

No rating or validation information has been found for AICSDP-35:AAVS1-mEGFP.

No alerts have been found for AICSDP-35:AAVS1-mEGFP.

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## Data and Source Information

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

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## Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Hayes BH, et al. (2023) Confinement plus myosin-II suppression maximizes heritable loss of chromosomes, as revealed by live-cell ChReporters. *Journal of cell science*, 136(11).

B??hler B, et al. (2023) Avidity-based bright and photostable light-up aptamers for single-molecule mRNA imaging. *Nature chemical biology*, 19(4), 478.

Sunbul M, et al. (2021) Super-resolution RNA imaging using a rhodamine-binding aptamer with fast exchange kinetics. *Nature biotechnology*, 39(6), 686.