

# Resource Summary Report

Generated by [RRID](#) on Aug 23, 2026

## GFP-HA-BACE1

RRID:Addgene\_165032

Type: Plasmid

### Proper Citation

RRID:Addgene\_165032

### Plasmid Information

**URL:** <http://www.addgene.org/165032>

**Proper Citation:** RRID:Addgene\_165032

**Insert Name:** BACE1

**Organism:** Homo sapiens

**Bacterial Resistance:** Kanamycin

**Defining Citation:** [PMID:22553349](#)

**Vector Backbone Description:** Backbone Marker:Clontech; Backbone Size:4733; Vector Backbone:pEGFP-N1; Vector Types:Mammalian Expression; Bacterial Resistance:Kanamycin

**Plasmid Name:** GFP-HA-BACE1

**Record Creation Time:** 20220422T221932+0000

**Record Last Update:** 20260815T080926+0000

### Ratings and Alerts

No rating or validation information has been found for GFP-HA-BACE1.

No alerts have been found for GFP-HA-BACE1.

### Data and Source Information

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

### Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Parutto P, et al. (2026) FidlTrack: high-fidelity structure-aware single particle tracking resolves intracellular molecular motion in organelles sensing APP processing. Nature communications, 17(1).

Rivera JF, et al. (2025) ATLAS: a rationally designed anterograde transsynaptic tracer. Nature methods, 22(5), 1101.

Rivera JF, et al. (2023) ATLAS: A rationally designed anterograde transsynaptic tracer. bioRxiv : the preprint server for biology.

Cunha C, et al. (2010) A simple role for BDNF in learning and memory? Frontiers in molecular neuroscience, 3, 1.