

Resource Summary Report

Generated by [RRID](#) on Aug 26, 2026

pcDNA3-sACE2v1-Fc(IgG1)

RRID:Addgene_145164

Type: Plasmid

Proper Citation

RRID:Addgene_145164

Plasmid Information

URL: <http://www.addgene.org/145164>

Proper Citation: RRID:Addgene_145164

Insert Name: Angiotensin-converting enzyme 2

Organism: Homo sapiens

Bacterial Resistance: Ampicillin

Defining Citation: [PMID:32753553](#)

Vector Backbone Description: Backbone Marker:Invitrogen; Backbone Size:5332; Vector Backbone:pcDNA3.1(+); Vector Types:Mammalian Expression; Bacterial Resistance:Ampicillin

Comments: Please visit <https://www.biorxiv.org/content/10.1101/2020.03.16.994236v1> for BioRxiv preprint.

Plasmid Name: pcDNA3-sACE2v1-Fc(IgG1)

Relevant Mutation: Extracellular, soluble protease domain (a.a. M1-D615) with mutations H34A, T92Q, Q325P, A386L

Record Creation Time: 20220422T221818+0000

Record Last Update: 20260815T080653+0000

Ratings and Alerts

No rating or validation information has been found for pcDNA3-sACE2v1-Fc(IgG1).

No alerts have been found for pcDNA3-sACE2v1-Fc(IgG1).

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](#) .

Hollis JA, et al. (2025) Molecular exaptation by the integrin α 5 β 1 domain. Science advances, 11(37), eadx9567.

Hollis JA, et al. (2023) Evolutionary origin and structural ligand mimicry by the inserted domain of alpha-integrin proteins. bioRxiv : the preprint server for biology.

Procko E, et al. (2020) The sequence of human ACE2 is suboptimal for binding the S spike protein of SARS coronavirus 2. bioRxiv : the preprint server for biology.