

Resource Summary Report

Generated by [SPARC SAWG](#) on Aug 17, 2026

SaKKH-ABEmax

RRID:Addgene_119815

Type: Plasmid

Proper Citation

RRID:Addgene_119815

Plasmid Information

URL: <http://www.addgene.org/119815>

Proper Citation: RRID:Addgene_119815

Insert Name: ecTadA-ecTadA*(7.10)-nSaCas9 (KKH)

Organism: Other

Bacterial Resistance: Ampicillin

Defining Citation: [PMID:31110355](#)

Vector Backbone Description: Vector Backbone:pCMV with a BR322 origin; Vector Types:Mammalian Expression; Bacterial Resistance:Ampicillin

Plasmid Name: SaKKH-ABEmax

Record Creation Time: 20220422T221626+0000

Record Last Update: 20260815T080317+0000

Ratings and Alerts

No rating or validation information has been found for SaKKH-ABEmax.

No alerts have been found for SaKKH-ABEmax.

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 [SPARC SAWG](#) .

Tachida Y, et al. (2025) Systematic empirical evaluation of individual base editing targets: Validating therapeutic targets in USH2A and comparison of methods. Molecular therapy : the journal of the American Society of Gene Therapy, 33(4), 1466.

Geurts MH, et al. (2023) One-step generation of tumor models by base editor multiplexing in adult stem cell-derived organoids. Nature communications, 14(1), 4998.

Villiger L, et al. (2021) Replacing the SpCas9 HNH domain by deaminases generates compact base editors with an alternative targeting scope. Molecular therapy. Nucleic acids, 26, 502.