

Resource Summary Report

Generated by [RRID](#) on Aug 28, 2026

VRQR-ABEmax

RRID:Addgene_119811

Type: Plasmid

Proper Citation

RRID:Addgene_119811

Plasmid Information

URL: <http://www.addgene.org/119811>

Proper Citation: RRID:Addgene_119811

Insert Name: ecTadA-ecTadA*(7.10)-nSpCas9 (VRQR)

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: VRQR-ABEmax

Record Creation Time: 20220422T221626+0000

Record Last Update: 20260826T040533+0000

Ratings and Alerts

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

No alerts have been found for VRQR-ABEmax.

Data and Source Information

Source: [Addgene](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

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

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.

Valdez I, et al. (2025) A streamlined base editor engineering strategy to reduce bystander editing. *Nature communications*, 16(1), 8115.

Chai AC, et al. (2023) Base editing correction of hypertrophic cardiomyopathy in human cardiomyocytes and humanized mice. *Nature medicine*, 29(2), 401.

Su J, et al. (2023) In vivo adenine base editing corrects newborn murine model of Hurler syndrome. *Molecular biomedicine*, 4(1), 6.

Koblan LW, et al. (2021) In vivo base editing rescues Hutchinson-Gilford progeria syndrome in mice. *Nature*, 589(7843), 608.

Erwood S, et al. (2019) Modeling Niemann-Pick disease type C in a human haploid cell line allows for patient variant characterization and clinical interpretation. *Genome research*, 29(12), 2010.