Resource Summary Report

Generated by RRID on Aug 29, 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: 20260829T075307+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.