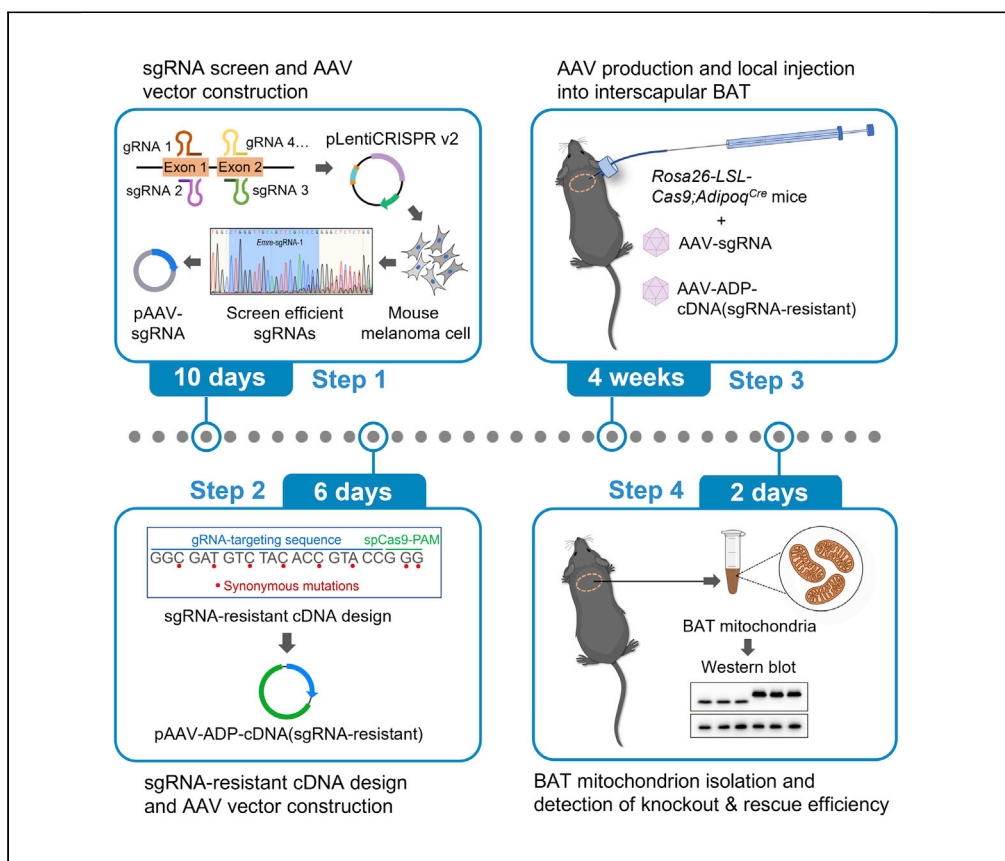


## Protocol

# Specific and efficient gene knockout and overexpression in mouse interscapular brown adipocytes *in vivo*



The classical Cre-LoxP system is time consuming. Here we detail a protocol that leverages Rosa26-LSL-Cas9;Adiponectin-Cre mice to restrict Cas9 expression in adipocytes. This enables specific deletion of target genes in brown adipocytes within 6 weeks by local injection of AAV-sgRNA into interscapular brown adipose tissue. We also describe an adiponectin-promoter-driven AAV vector to express sgRNA-resistant cDNA-encoded protein for subsequent rescue. This protocol thus provides an efficient means to specifically knockout and overexpress genes in brown adipocytes *in vivo*.

Publisher's note: Undertaking any experimental protocol requires adherence to local institutional guidelines for laboratory safety and ethics.

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### Highlights

Generation of adipocyte-specific Cas9 transgenic mice for gene editing

Brown-adipocyte-specific knockout by local injection of AAV-sgRNA within 6 weeks

Adiponectin-promoter-driven expression of sgRNA-resistant protein for rescue

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## Protocol

Specific and efficient gene knockout and overexpression in mouse interscapular brown adipocytes *in vivo*Kaili Xue,<sup>1,3,\*</sup> Dongmei Wu,<sup>1,2</sup> and Yifu Qiu<sup>1,2,4,\*</sup><sup>1</sup>Institute of Molecular Medicine, Beijing Key Laboratory of Cardiometabolic Molecular Medicine, College of Future Technology, Peking University, Beijing 100871, China<sup>2</sup>Peking-Tsinghua Center for Life Sciences, Peking University, Beijing 100871, China<sup>3</sup>Technical contact: [kaili.xue@pku.edu.cn](mailto:kaili.xue@pku.edu.cn)<sup>4</sup>Lead contact\*Correspondence: [kaili.xue@pku.edu.cn](mailto:kaili.xue@pku.edu.cn) (K.X.), [yifu.qiu@pku.edu.cn](mailto:yifu.qiu@pku.edu.cn) (Y.Q.)  
<https://doi.org/10.1016/j.xpro.2022.101895>

## SUMMARY

The classical Cre-LoxP system is time consuming. Here we detail a protocol that leverages Rosa26-LSL-Cas9;Adiponectin-Cre mice to restrict Cas9 expression in adipocytes. This enables specific deletion of target genes in brown adipocytes within 6 weeks by local injection of AAV-sgRNA into interscapular brown adipose tissue. We also describe an adiponectin-promoter-driven AAV vector to express sgRNA-resistant cDNA-encoded protein for subsequent rescue. This protocol thus provides an efficient means to specifically knockout and overexpress genes in brown adipocytes *in vivo*.

For complete details on the use and execution of this protocol, please refer to Xue et al. (2022).<sup>1</sup>

## BEFORE YOU BEGIN

This protocol describes how to specifically and efficiently knockout genes of interest and perform their rescues in brown adipocytes *in vivo*. This protocol leverages key technique advances towards the uses of AAVs to target adipose tissues and the methods to inject interscapular brown adipose tissue (iBAT).<sup>2–8</sup>

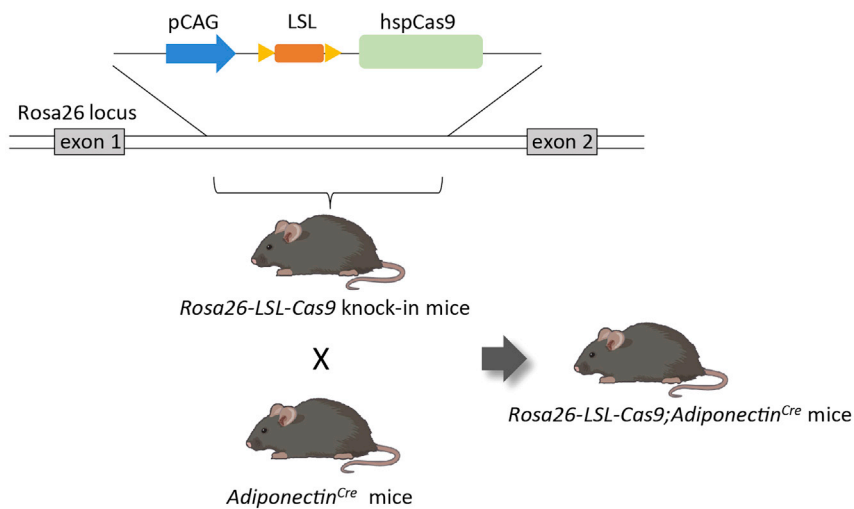
*Ucp1*<sup>Cre</sup> transgenic mice are utilized in the Cre-LoxP system to allow “BAT-specific” gene knockout. However, Kristin E. Clafin et al. recently reported that the *Ucp1*<sup>Cre</sup> expression is not restricted to BAT but throughout the brain.<sup>9</sup> Thus, the interpretation of many findings obtained by using *Ucp1*<sup>Cre</sup>-mediated knockout mouse models should consider their potential non-specific effects through the central nervous system. In this protocol, we employ Rosa26-LSL-Cas9;Adiponectin<sup>Cre</sup> mice to restrict Cas9 expression in adipocytes and further combine with iBAT local injection of AAV-sgRNA, which provides a dual guarantee of the specificity of gene knockout in brown adipocytes.

To overcome potential off-target mutations by CRISPR/Cas9-mediated gene knockout, we suggest using at least two different sgRNA to independently verify the phenotypes. In addition, we develop an adiponectin promoter-driven AAV vector to express sgRNA-resistant cDNA encoded wild-type protein as a rescue, which serves as an on-target control for the sgRNA. Moreover, one also can express sgRNA-resistant cDNA encoded functional mutant protein such as kinase dead one.

## Institutional permissions

The experiments should be approved by the Institutional Animal Care and Use Committee (IACUC) of laboratory animals to protect the welfare of animals. All mouse experiments were performed according to PKU IACUC guidelines.





**Figure 1. Breeding strategy for generating *Rosa26-LSL-Cas9;Adipoq<sup>Cre</sup>* mice**

### Preparation of *Rosa26-LSL-Cas9;Adipoq<sup>Cre</sup>* mice

⌚ Timing: n/a

The *Rosa26-LSL(loxP-stop-loxP)-Cas9* knock-in mice<sup>10</sup> were crossed with *Adiponectin<sup>Cre</sup>* mice to generate *Rosa26-LSL-Cas9;Adipoq<sup>Cre</sup>* mice that restrict Cas9 expression in adipocytes. Local injection of AAV-sgRNA into iBAT of *Rosa26-LSL-Cas9;Adipoq<sup>Cre</sup>* mice allows a specific deletion of the target gene in brown adipocytes (Figure 1). In this protocol, AAV injections were performed with sex-matched mice at the age of 6–8 weeks.

### sgRNA design and synthesis

⌚ Timing: 1 day

sgRNAs can be designed using online CRISPR design tools such as Benchling, Broad Institute GPP, CRISPOR, etc. In this protocol, we use CRISPOR to design sgRNAs targeting mouse gene *Smdt1* (encoding essential MCU regulator, EMRE).<sup>11</sup>

1. Obtain the genomic sequence of the target gene from the NCBI-Genome Data Viewer by searching the gene name.
2. Select the first exon containing the coding sequence preferentially, and input the sequence (starting from ATG) on CRISPOR. Select *Mus musculus* genome and spCas9 PAM motif.
3. Choose higher-scoring sgRNAs from the predicted guide sequences, which may have higher editing efficiency and lower off-target effects. Design at least three sgRNAs and screen for efficient ones (as described later in the protocol).
4. Synthesize two sgRNA oligonucleotides as follows:

sgRNA-F: 5'-CACCGNNNNNNNNNNNNNNNNNNNN-3'.

sgRNA-R: 5'-AAACNNNNNNNNNNNNNNNNNNNNNC-3'.

**Note:** The 20 bp sgRNA sequence is shown as polyN. The sgRNA-F (forward) contains the sgRNA sequence upstream of the PAM motif.



<b>Continued</b>		
REAGENT or RESOURCE	SOURCE	IDENTIFIER
sgRNA targeting sequence against <i>Emre</i> #2: GGCGATGTCTACACCGTACC	This paper	N/A
sgRNA targeting sequence against <i>Emre</i> #3: GTCTCAGCCAGGTACCGTCC	This paper	N/A
sgRNA targeting sequence against <i>Emre</i> #4: TGGCGATGTCTACACCGTACC	This paper	N/A
sgRNA targeting sequence against <i>Emre</i> #5: GCCTGGGTTGCAGTTGACCC	This paper	N/A
sgRNA targeting sequence against <i>LacZ</i> : CCCGAATCTCTATCGTGCGG	Addgene	Cat. #74179
Seq-F: CTTCCCCTGTGTGTACAGAG	This paper	N/A
Seq-R: TGAGTCCTATGTCCGGTCCC	This paper	N/A
SLIC-EMRE-F: GGTTGGGGCAACCGGTA TGGCGTCCACGGCGGCTCG	This paper	N/A
SLIC-EMRE-R: AGGCCCGGGCGTCCGACC ATCGTCGTCGTCATCCT	This paper	N/A
sgRNA-resis-mutation-F: GGGGAGGAGAGGTGGAGACGTG TATACGGTTCCTCCAGCTCAGG	This paper	N/A
sgRNA-resis-mutation-R: CTGAGACCTGAGCTGGAGGGAACC GTATACACGCTCCACCTCTC	This paper	N/A
<b>Recombinant DNA</b>		
lentiCRISPRv2	Addgene	Cat. #52961
pAAV-U6-gRNA-CBh-mCherry	Addgene	Cat. #91947
pAAV-ADP-MCS-FLAG	Addgene	Cat. #192360
pRC2/8	Addgene	Cat. #112864
pHelper	Addgene	Cat. #112867
<b>Software and algorithms</b>		
CRISPOR	TEFOR Infrastructure	<a href="#">Version 5.01</a>
ICE	Synthego	<a href="#">Version 3.0</a>
<b>Other</b>		
Surgical swabs	N/A	N/A
Animal hair clipper	N/A	N/A
Heating pad	N/A	N/A
Surgical scissors	Fine Science Tools	Cat. #14002-12
Curved forceps	Fine Science Tools	Cat. #11052-10
Surgical suture clips	Fine Science Tools	Cat. #12022-09
Clip applicator	Fine Science Tools	Cat. #12018-12
Microsyringe (50 $\mu$ L)	Hamilton	Cat. #80500
Polyethylene tubing	Smiths Medical	Cat. #10793527
Nanopass33 (33G needle for pen injectors)	Terumo Corp.	N/A

## MATERIALS AND EQUIPMENT

<b>LB (Lysogeny Broth) Medium</b>		
Reagent	Final concentration	Amount
NaCl	1%	10 g
Yeast extract	0.5%	5 g
Trypton	1%	10 g
H <sub>2</sub> O	N/A	Bring up to 1 L
<b>Total</b>	<b>N/A</b>	<b>1 L</b>

The LB medium should be sterilized by autoclaving.

### 2,2,2-Tribromoethanol stock solution

Reagent	Final concentration	Amount
2,2,2-Tribromoethanol	N/A	5 g
2-Methyl-2-butanol	N/A	10 mL
<b>Total</b>	<b>N/A</b>	<b>N/A</b>

The solution should be freshly made and fully dissolved with vortex mixer.

### 2,2,2-Tribromoethanol working solution

Reagent	Final concentration	Amount
2,2,2-Tribromoethanol stock solution	N/A	1.25 mL
0.9% NaCl	N/A	Bring up to 50 mL
<b>Total</b>	<b>N/A</b>	<b>50 mL</b>

The working solution should be sterilized using a 0.22  $\mu$ m filter and stored at 4°C away from light for a couple of weeks.

### BAT mitochondrion isolation buffer

Reagent	Final concentration	Amount
Sucrose	250 mM	8.56 g
1 M Hepes	10 mM	1 mL
EGTA	1 mM	0.038 g
BSA (fatty-acid free)	0.3%	0.3 g
3 M KOH	N/A	Adjust pH to 7.0
ddH <sub>2</sub> O	N/A	Bring up to 1 L
<b>Total</b>	<b>N/A</b>	<b>100 mL</b>

The buffer should be sterilized using a 0.22  $\mu$ m filter and stored at 4°C for a couple of weeks.

## STEP-BY-STEP METHOD DETAILS

### Screen efficient sgRNAs in mouse melanoma cells using pAAV-sgRNA vector

⌚ Timing: 10 days

The sgRNAs targeting mouse genes with editing efficiency are screened using lentiCRISPRv2 vector transfected in mouse melanoma cells.

1. Prepare lentiCRISPRv2-sgRNA vector.
  - a. Anneal the sgRNA oligonucleotides as inserts.
    - i. Anneal each pair of sgRNA oligonucleotides as the following reaction.

Reagent	Amount
Oligo 1 (100 $\mu$ M)	1 $\mu$ L
Oligo 2 (100 $\mu$ M)	1 $\mu$ L
10 $\times$ T4 ligation buffer (NEB)	1 $\mu$ L
T4 PNK (NEB)	0.5 $\mu$ L
ddH <sub>2</sub> O	6.5 $\mu$ L

- ii. Anneal in a thermocycler at 37°C for 30 min, then 95°C for 5 min and ramp down to 25°C (5°C/min).
- b. Digest 2  $\mu$ g of lentiCRISPRv2 plasmid at 55°C for 40 min and purify the digested plasmids from the 0.8% agarose gel using Thermo Scientific GeneJET Gel Extraction Kit (K0691) following the manufacturer's instructions.

([https://www.thermofisher.cn/document-connect/document-connect.html?url=https://assets.thermofisher.cn/TFS-Assets%2FMSG%2Fmanuals%2FMAN0012661\\_GeneJET\\_Gel\\_Extraction\\_UG.pdf](https://www.thermofisher.cn/document-connect/document-connect.html?url=https://assets.thermofisher.cn/TFS-Assets%2FMSG%2Fmanuals%2FMAN0012661_GeneJET_Gel_Extraction_UG.pdf)).

Reagent	Amount
lentiCRISPRv2	2 µg
BsmBI	1 µL
NEbuffer3.1 (10×)	2 µL
ddH <sub>2</sub> O	15 µL

- c. Clone the annealed oligos into the digested lentiCRISPRv2 vector.
  - i. Dilute the annealed oligos at 1:10.
  - ii. Set up the ligation reaction as follows and incubate the reaction at 25°C for 2 h.

Reagent	Amount
Digested and purified lentiCRISPRv2	50 ng
Diluted annealed oligos	1 µL
10× T4 ligation buffer (NEB)	1 µL
T4 ligase (NEB)	1 µL
ddH <sub>2</sub> O	Bring up to 10 µL

- iii. Transform the ligation mixture into Stbl3 competent cells. Transform 2 µL of each ligation mixture into 50 µL of Stbl3 competent cells and incubate the mixture for 20 min on ice. After heat shock at 42°C for 90 s, the transformation mixture is immediately incubated on ice for 2 min. Then, add 200 µL of LB medium to the mixture and allow the transformed cells to recover at 37°C for 1 h with 220 rpm shaking.
  - iv. Add the transformed cells onto the LB plate with 100 mg/mL ampicillin and incubate the LB plate at 37°C for 14 h to allowing colonies to grow.
  - v. Pick 2 colonies from each lentiCRISPRv2-sgRNA plate into 5 mL LB medium with 100 mg/mL ampicillin to be incubated at 37°C with 220 rpm shaking for 8–10 h and perform plasmid miniprep using Thermo Scientific GeneJET Plasmid Miniprep Kit (K0502) following the manufacturer's instructions ([https://www.thermofisher.cn/document-connect/document-connect.html?url=https://assets.thermofisher.cn/TFS-Assets%2FMSG%2Fmanuals%2FMAN0012655\\_GeneJET\\_Plasmid\\_Miniprep\\_UG.pdf](https://www.thermofisher.cn/document-connect/document-connect.html?url=https://assets.thermofisher.cn/TFS-Assets%2FMSG%2Fmanuals%2FMAN0012655_GeneJET_Plasmid_Miniprep_UG.pdf)).
  - vi. Sequence each colony to confirm the sgRNA insertion into lentiCRISPRv2 vector.
2. Screen sgRNAs with editing efficiency in mouse melanoma cells ([troubleshooting 1](#)) ([troubleshooting 2](#)).
    - a. Mouse melanoma cells (B16F10) are cultured in DMEM supplemented with 10% FBS at 37°C in a 5% CO<sub>2</sub> incubator.
    - b. Plate B16F10 cells onto 12-well plate in DMEM supplemented with 10% FBS.
    - c. When cells reach 80%–90% confluent, transfect 1 µg of lentiCRISPRv2-sgRNA plasmid into each well using PEI as follows. Preheat PEI solution and DMEM at 37°C and prepare plasmid/DMEM mixture for each well of cells. Add 3 µL of PEI into the plasmid/DMEM mixture (the ratio of PEI (µL): plasmid (µg) is 3:1), mix well, and place at room temperature for 10 min. Add the transfection mixture into each well of B16F10 cells.

Reagent	Amount
lentiCRISPRv2-sgRNA	1 µg
DMEM	N/A
PEI	3 µL
Total	100 µL

**Note:** Each lentiCRISPRv2-sgRNA plasmid is transfected in 2 wells, one for genomic DNA sequencing and another one for western blot analysis.

**△ CRITICAL:** The control sgRNA targets a gene not existing in the mouse genome (LacZ here).

- d. 24 h after transfection, pass the B16F10 cells from the 12-well plate onto the 6-well plate. Culture one well of cells in 2 mL of DMEM supplemented with 10% FBS and 10  $\mu$ M puromycin for 48–60 h to remove non-infected dead cells.
- e. Wash cells with PBS buffer to remove the dead cells and collect the remaining cells for genomic DNA extraction.
- f. Genomic DNA is extracted using Invitrogen Purelink Genomic DNA Kit (K182001) following the manufacturer's instructions ([https://www.thermofisher.cn/document-connect/document-connect.html?url=https://assets.thermofisher.cn/TFS-Assets%2FManuals%2FPurelink\\_genomic\\_man.pdf](https://www.thermofisher.cn/document-connect/document-connect.html?url=https://assets.thermofisher.cn/TFS-Assets%2FManuals%2FPurelink_genomic_man.pdf)).
- g. Amplify the sgRNA target region on genomic DNA via PCR using a pair of PCR primers (Seq-F/R primers) to produce 400–600 bp amplicons with the sgRNA target sequence in the center. Set up the PCR reaction as follows.

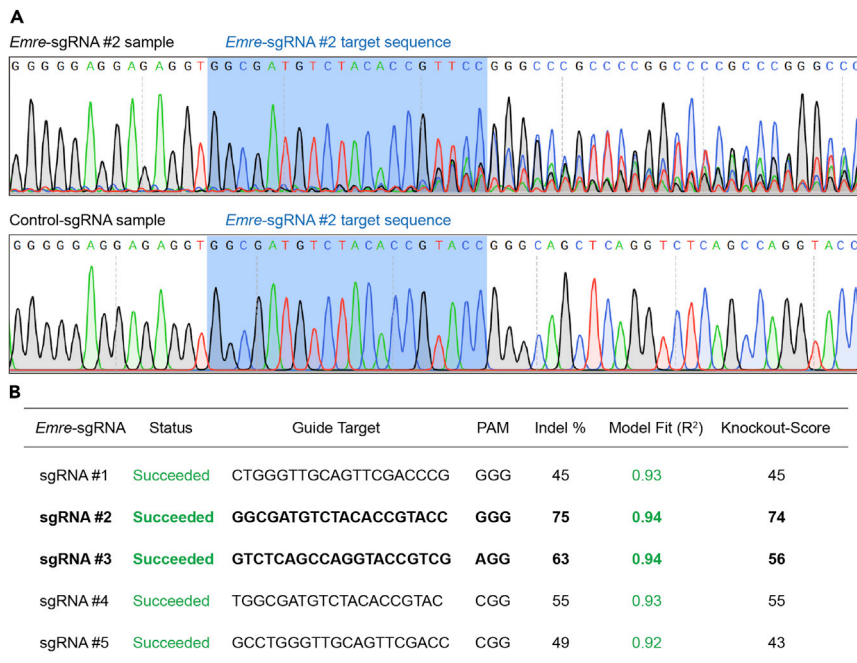
Reagent	Amount
2x Phanta Max Buffer	25 $\mu$ L
dNTP Mix (10 mM each)	1 $\mu$ L
500 ng/ $\mu$ L Genomic DNA	3 $\mu$ L
10 $\mu$ M Seq-F primer	1 $\mu$ L
10 $\mu$ M Seq-R primer	1 $\mu$ L
Phanta Max Super-Fidelity DNA Polymerase (Vazyme)	1 $\mu$ L
ddH <sub>2</sub> O	Up to 50 $\mu$ L

Then run PCR using the following cycling condition.

Steps	Temperature	Time	Cycles
Initial Denaturation	95°C	3 min	1
Denaturation	95°C	15 s	29 cycles
Annealing	58°C	15 s	
Extension	72°C	30 s/kb	
Final extension	72°C	5 min	1
Hold	4°C	forever	

- h. Purify the PCR products from the 2% agarose gel using Thermo Scientific GeneJET Gel Extraction Kit (K0691).
  - i. Sequence the amplicons of control sgRNA and target gene sgRNA samples using Seq-F primer.
  - j. Analyze the editing efficiency through uploading sgRNA sequence and sequencing results of control and experimental amplicons onto the software ICE (<https://ice.synthego.com/#/>).
  - k. Select at least two sgRNAs with the highest editing activity based on the Indel %, Model Fit and Knockout-Score got from ICE analyses (Figure 2).
  - l. Another well of transfected cells is used for protein extraction to confirm the knock-out efficiency of the target gene on protein level by western blot.
3. pAAV-sgRNA-mCherry vector construction.





**Figure 2. Analysis of the editing efficiency of the sgRNAs by sequencing the amplicons of sgRNA-targeted region on genomic DNA**

(A) The disorderly sequencing peaks suggest potential editing activity of *Emre*-sgRNAs compared to control-sgRNA. (B) The editing efficiency analyzed by the ICE software.

- a. sgRNA synthesis. Select two sgRNAs with highest editing efficiency verified in step 2 and synthesize the oligonucleotides as follows.  
 sgRNA-F: 5'-ACCGNNNNNNNNNNNNNNNNNNNNNN-3'.  
 sgRNA-R: 5'-AACANNNNNNNNNNNNNNNNNNNNNNC-3'.
- b. Anneal the sgRNA oligos as inserts.
- c. Digest 2 µg of pAAV-U6-gRNA-CBh-mCherry plasmid with BspQI at 50°C for 40 min.

Reagent	Amount
pAAV-U6-gRNA-CBh-mCherry	2 µg
BsmBI	1 µL
NEbuffer3.1 (10×)	2 µL
ddH <sub>2</sub> O	15 µL

- d. Purify the digested plasmids from the 0.8% agarose gel using Thermo Scientific GeneJET Gel Extraction Kit (K0691).
- e. Ligate the annealed oligos into the digested pAAV-U6-gRNA-CBh-mCherry plasmid.
  - i. Dilute the annealed oligos at 1:10.
  - ii. Set up the ligation reaction as follows and incubate the reaction at 25°C for 2 h.

Reagent	Amount
Digested and purified pAAV-U6-gRNA-CBh-mCherry	50 ng
Diluted annealed oligos	1 µL
10× T4 ligation buffer (NEB)	1 µL
T4 ligase (NEB)	1 µL
ddH <sub>2</sub> O	Up to 10 µL

- iii. Transform the ligation mixture into T1 competent cells. Transform 2  $\mu\text{L}$  of the ligation mixture into 50  $\mu\text{L}$  of T1 competent cells and incubate the mixture for 20 min on ice. After heat shock at 42°C for 90 s, the transformation mixture is immediately incubated on ice for 2 min.
- iv. Plate transformed cells onto the LB plate with 100 mg/mL ampicillin and incubate the LB plate at 37°C for 14 h allowing colonies to grow.
- v. Pick 2 colonies from each pAAV-U6-gRNA-CBh-mCherry plate into 5 mL LB medium with 100 mg/mL ampicillin to be incubated at 37°C with 220 rpm shaking for 8–10 h and perform plasmid miniprep.
- vi. Sequence each colony to confirm the sgRNA insertions into the pAAV-U6-gRNA-mCherry vector.

### Design sgRNA-resistant cDNA of target gene and construct AAV vector

⌚ Timing: 6 days

To avoid recognition by sgRNA-CRISPR, sgRNA-resistant cDNA of the target gene is designed by the introduction of synonymous mutations on the critical “GG” in the PAM motif (NGG) and on the sgRNA-targeted sequence region. Using sgRNA-resistant cDNA, we can rescue with tagged wild-type or mutant protein according to experimental purpose in the knockout context of the endogenous gene.

4. Construct pAAV-ADP (*adiponectin* promoter)-EMRE-FLAG ([troubleshooting 4](#)).
  - a. Use SLIC (sequence and ligation-independent cloning) to insert EMRE-cDNA into the MCS (multiclonal site) of pAAV-ADP-MCS-FLAG. Design SLIC PCR primers according to the selected dual enzyme cleavage sites in the MCS of pAAV-ADP-MCS-FLAG.

**Note:** In this case we select AgeI and Sall as dual enzyme cleavage sites, and synthesize the SLIC PCR primers as shown in [Figure 3](#).

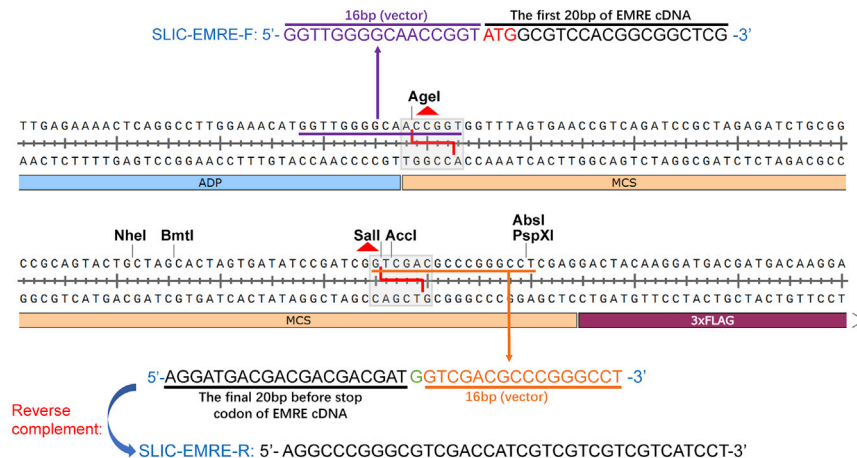
- b. Use the above SLIC PCR primers to amplify the EMRE insert. Perform PCR using Vazyme Phanta Max Super-Fidelity DNA Polymerase kit (P505d1) and set up PCR reaction as below.

Reagent	Amount
2× Phanta Max Buffer	25 $\mu\text{L}$
dNTP Mix (10 mM each)	1 $\mu\text{L}$
100 ng/ $\mu\text{L}$ template plasmid containing EMRE cDNA	1 $\mu\text{L}$
10 $\mu\text{M}$ SLIC-EMRE-F	1 $\mu\text{L}$
10 $\mu\text{M}$ SLIC-EMRE-R	1 $\mu\text{L}$
Phanta Max Super-Fidelity DNA Polymerase (Vazyme)	1 $\mu\text{L}$
ddH <sub>2</sub> O	Up to 50 $\mu\text{L}$

- c. Run PCR using the following cycling condition.

Steps	Temperature	Time	Cycles
Initial Denaturation	95°C	3 min	1
Denaturation	95°C	15 s	29 cycles
Annealing	58°C	15 s	
Extension	72°C	30 s/kb	
Final extension	72°C	5 min	1
Hold	4°C	forever	

- d. Purify the PCR amplicons as SLIC inserts using Thermo Scientific GeneJET Gel Extraction Kit (K0691).



**Figure 3. Design SLIC PCR primers to clone EMRE-cDNA into pAAV-ADP-MCS-3×FLAG**

e. Digest 2 μg of pAAV-ADP-MCS-FLAG plasmid with Agel and Sall at 37°C for 60 min as below.

Reagent	Amount
pAAV-ADP-MCS-3×FLAG	2 μg
Agel-HF (NEB)	1 μL
Sall-HF (NEB)	1 μL
10× Cutsmart buffer (NEB)	2 μL
ddH <sub>2</sub> O	14 μL

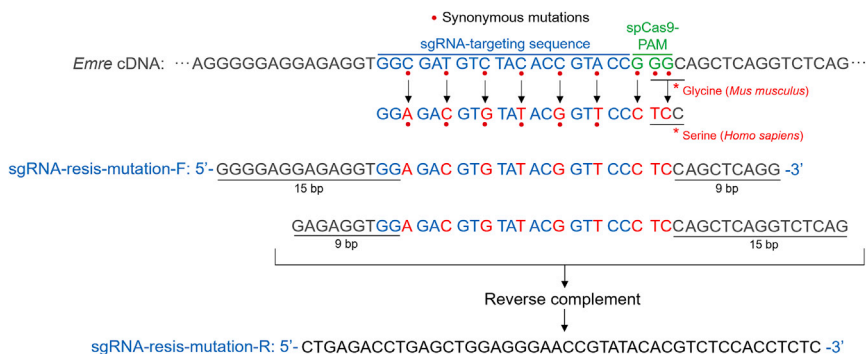
f. Purify the digested plasmid as the SLIC vector using Thermo Scientific GeneJET Gel Extraction Kit (K0691).

g. Perform SLIC reaction to construct pAAV-ADP-EMRE-3×FLAG.

i. Set up SLIC reaction as below.

Reagent	Amount
SLIC insert	50 ng
SLIC vector	25 ng
10× NEbuffer 1	0.5 μL
ddH <sub>2</sub> O	Up to 5 μL

- ii. Place the reaction tube on ice for 5 min and the following steps should be performed on ice.
- iii. Add 0.25 μL Exonuclease III (40 units) to the tube and mix well by pipetting for several times, and then place on ice for 30 min.
- iv. Add 0.5 μL 0.5 M EDTA (pH 8.0) into the tube and mix by pipetting for several times, and then incubate the mixture at 65°C for 5 min to stop the reaction.
- v. Centrifuge the tube to concentrate the mixture.
- vi. Replace the reaction tube on ice to cool down for 5 min.
- vii. Transform the total reaction mixture into 50 μL T1 competent cells and incubate the mixture for 20 min on ice. After heat shock at 42°C for 90 s, the mixture is immediately incubated on ice for 2 min.
- viii. Plate transformed cells onto the LB plate with 100 mg/mL ampicillin and incubate the LB plate at 37°C for 14 h allowing colonies to grow.



**Figure 4. Design synonymously mutagenic primers for sgRNA-resistant cDNA of target gene**

- ix. Pick 2 colonies from the plate into 5 mL LB medium with 100 mg/mL ampicillin to be incubated at 37°C with 220 rpm shaking for 8–10 h and perform plasmid miniprep.
- x. Sequence each colony to confirm the correct construction of pAAV-ADP-EMRE-3×FLAG.
5. Design synonymously mutagenic primers for sgRNA-resistant cDNA of target gene ([troubleshooting 2](#)).
  - a. Synonymously mutate the “GG” in the PAM motif (NGG) which is critical for the recognition by sgRNA-spCas9.

**Note:** In this case, since synonymous mutations cannot be performed on the “GG”, we mutate the glycine containing “GG” to serine in *smdt1* of *homo sapiens*, as illustrated in [Figure 4](#).

- b. Synonymously mutate the sgRNA-targeting sequence as illustrated in [Figure 4](#).
- c. Design and synthesize the synonymously mutagenic primers as follows.
 

sgRNA-resis-mutation-F:  
5'-GGGGAGGAGAGGTGGA GAC GTG TAT ACG GTT CCC TCCAGCTCAGG-3'.

sgRNA-resis-mutation-R:  
5'-CTGAGACCTGAGCTGGAGGGAACCGTATACACGTCTCCACCTCTC-3'.
6. Take pAAV-ADP-EMRE-3×FLAG plasmid as PCR template and use synonymously mutagenic primers to produce sgRNA-resistant pAAV-ADP-EMRE-resis-3×FLAG.
  - a. Set up PCR reaction as follows to perform site-directed mutagenesis using Vazyme Phanta Max Super-Fidelity DNA Polymerase kit.

Reagent	Amount
2× Phanta Max Buffer	5 μL
dNTP Mix (10 mM each)	0.2 μL
Template plasmid	20 ng
10 μM sgRNA-resis-mutation-F	0.25 μL
Phanta Max Super-Fidelity DNA Polymerase (Vazyme)	0.2 μL
ddH <sub>2</sub> O	Up to 9.5 μL

- b. Run PCR using the following cycling condition.

Steps	Temperature	Time	Cycles
Initial Denaturation	95°C	3 min	1
Denaturation	95°C	15 s	9 cycles
Annealing	58°C	15 s	
Extension	72°C	30 s/kb	

(Continued on next page)

**Continued**

Steps	Temperature	Time	Cycles
Final extension	72°C	5 min	1
Hold	4°C	forever	

- c. Add 0.5  $\mu\text{L}$  of 10  $\mu\text{M}$  sgRNA-resis-mutant-R to the PCR reaction mixture and run PCR using the following cycling condition.

Steps	Temperature	Time	Cycles
Initial Denaturation	95°C	3 min	1
Denaturation	95°C	15 s	19 cycles
Annealing	58°C	15 s	
Extension	72°C	30 s/kb	
Final extension	72°C	5 min	1
Hold	4°C	forever	

- d. Digest the PCR amplicons with DpnI at 37°C for 2 h to remove the template plasmid as follows.

Reagent	Amount
PCR amplicons	10 $\mu\text{L}$
DpnI (NEB)	0.2 $\mu\text{L}$
10 $\times$ Cutsmart buffer (NEB)	1 $\mu\text{L}$

- e. Add 5  $\mu\text{L}$  of the digestion mixture into 50  $\mu\text{L}$  T1 competent cells and incubate the mixture on ice for 20 min. After heat shock at 42°C for 90 s, the mixture is immediately incubated on ice for 2 min.
- f. Plate the transformed cells onto LB plate with 100 mg/mL ampicillin and incubate the LB plate at 37°C for 14 h allowing colonies to grow.
- g. Pick 2 colonies from the plate into 5 mL LB medium with 100 mg/mL ampicillin to be incubated at 37°C with 220 rpm shaking for 8–10 h and perform plasmid miniprep.
- h. Sequence each colony to confirm the correct mutation of pAAV-ADP-EMRE-resis-3 $\times$ FLAG.
- i. Use SLIC to insert EMRE-resis-cDNA into the MCS of pAAV-ADP-MCS-3 $\times$ FLAG via SLIC-EMRE-F/R primers to avoid vector mutation, following step 4.

### AAV production

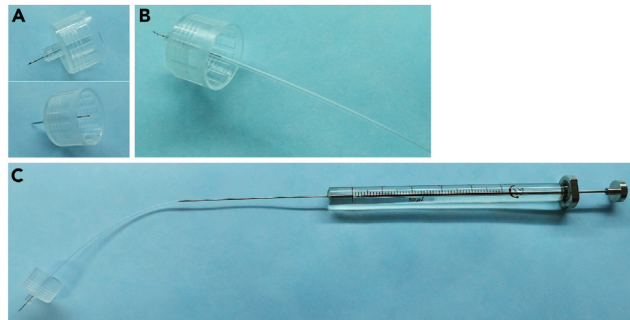
⌚ Timing: 1 week

In this protocol, we use AAV8 serotype that can effectively infect BAT.<sup>12</sup> AAV-U6-*Emre*-sgRNA-mCherry, AAV-U6-*Ctrl*-sgRNA-mCherry and AAV-ADP-EMRE-resis-3 $\times$ FLAG are produced following the previous step-by-step instructions.<sup>13</sup> The AAV titer for the following AAV injection experiments should be above  $1 \times 10^{13}$  vg/mL (vector genomes per mL).

### Local injection of AAV into intrascapular BAT (iBAT)

⌚ Timing: 3 weeks

7. Preparation for local injection of AAV.
- Prepare surgical equipment as in [key resources table](#).



**Figure 5. Set up injection device**

(A) 33G needle for pen injectors.

(B) Connect the tail end of a pen needle to a polyethylene tubing.

(C) The injection device assembled by connecting a needle of microsyringe (50  $\mu$ L) to the other side of the polyethylene tubing.

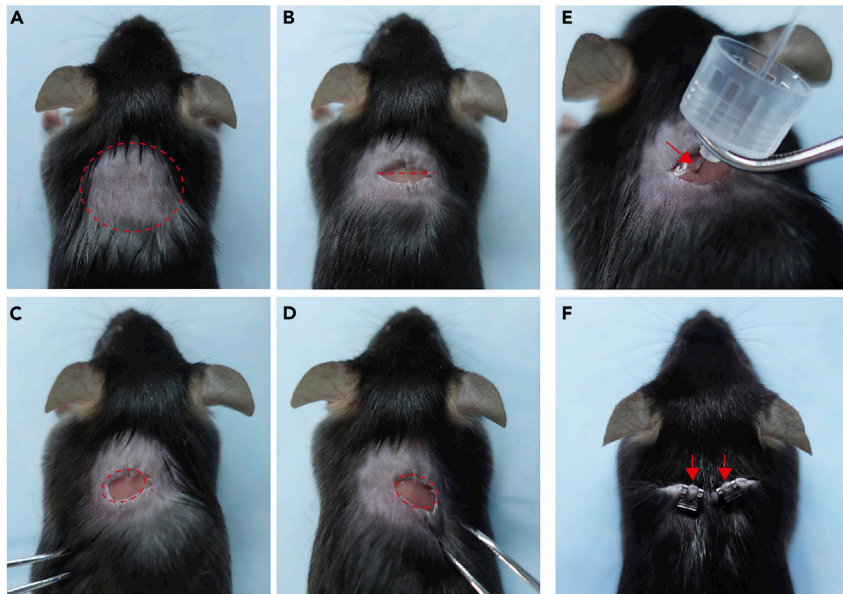
**Note:** all the surgical equipment should be sterilized with 75% ethanol before surgical operation.

- b. Set up injection device.
  - i. Connect the tail end of a pen needle to a polyethylene tubing (5 cm in length), as illustrated in [Figures 5A](#) and [5B](#).
  - ii. Connect a needle of microsyringe (50  $\mu$ L) to the other side of the polyethylene tubing, as illustrated in [Figure 5C](#).
  - iii. Test the assembled injection device and make sure all the connections are sealed well without leakage.
- c. Prepare virus solution ([troubleshooting 5](#)).
  - i. Dilute AAVs with PBS as follows to achieve the volume for injection (50  $\mu$ L for each lobe of iBAT, 100  $\mu$ L for each mouse).

Reagent	Amount
AAV-U6- <i>Emre</i> /Ctrl-sgRNA-mCherry	$1 \times 10^{12}$ vg
AAV-ADP-EMRE-resis-3 $\times$ FLAG	$5 \times 10^{11}$ vg
PBS	Up to 100 $\mu$ L

- ii. Draw 50  $\mu$ L of virus solution with microsyringe and connect the injection device.
8. Surgical operation for local injection of AAV into iBAT ([troubleshooting 3](#)) ([Methods video S1](#)).
  - a. Anaesthetize mice with 2,2,2-Tribromoethanol.  
Inject mice with 2,2,2-Tribromoethanol working solution (20  $\mu$ L/g body weight) through intraperitoneal injection. Two min after injection, the mice will be anaesthetized deeply enough to perform surgical operation.
 

**Note:** The anaesthetized mice should be on a heating pad during all the surgical process.
  - b. Remove the hair above iBAT using an animal hair clipper and wipe the exposed skin with ethanol ([Figure 6A](#)).
  - c. Cut a 0.5–0.8 cm incision in the skin above iBAT with a surgical scissor to expose the two lobes of iBAT ([Figures 6B–6D](#)).



**Figure 6. Surgical operation for local injection of AAV into iBAT**

- (A) Remove the hair above iBAT to exposure the skin.  
 (B) Cut a 0.5–0.8 cm incision in the skin above iBAT as indicated by the red dotted line.  
 (C) The left lobe of iBAT illustrated as the red dotted circle.  
 (D) The right lobe of iBAT illustrated as the red dotted circle.  
 (E) Insert the pen needle into the fat pad of iBAT, and the injection point is indicated by the red arrow.  
 (F) Close the incision with two surgical suture clips (9 mm) as indicated by the red arrows.

- d. Hold the injection device and carefully insert the pen needle into the fat pad of one iBAT lobe, as illustrated in [Figure 6E](#). When the needle is inserted well, inject 5  $\mu$ L virus for one point and inject 10 points (50  $\mu$ L virus) to distribute evenly throughout one lobe of iBAT.

**⚠ CRITICAL:** The depth of needle insertion should be 1–2 mm avoiding too deep or too superficial.

- e. Draw another 50  $\mu$ L of virus solution rapidly for another lobe of iBAT.  
 f. After injection, close the incision with two surgical suture clips (9 mm) ([Figure 6F](#)).

**Note:** The suture clips should be clamped tightly using a clip applicator avoiding shedding.

- g. Inject mice with analgesic (Tolfedine, 2 mg/kg body weight) by a single intramuscular injection.  
 h. Keep the mice on a heating pad until revivication.  
 i. Place one mouse per cage to avoid fighting. Monitor the health status of mice daily and inject the mice with analgesic (Tolfedine, 2 mg/kg body weight) at 24 h and 48 h after the surgery.

**Note:** Recovery time after surgery is about 7 days. It needs 3 weeks after AAV injection to allow sufficient CRISPR editing *in vivo* and enough overexpression of protein driven by *adiponectin* promoter.

### Knockout and rescue efficiency detection

⌚ Timing: 2 days

#### 9. Isolation of BAT mitochondria.



- a. Isolate the iBAT from mice injected with AAV for 3 weeks and immediately transfer the iBAT into 2 mL tube with 0.5 mL BAT mitochondrion isolation buffer.

**Note:** The whole isolation process should be performed on ice. The buffer and equipment used for mitochondrion isolation should be precooled on ice.

- b. Mince the iBAT into small pieces (1 mm<sup>3</sup> each) with a scissor in 2-mL tube and transfer the content into a Teflon-glass Dounce homogenizer (Wheaton, 2 mL). Wash the tube with 0.5 mL isolation buffer and collect it into the homogenizer (total 1 mL content in the homogenizer).
- c. Homogenize the iBAT pieces with a teflon pestle for 10 strokes. Transfer the homogenate into a new 2 mL tube and wash the homogenizer with 0.5 mL isolation buffer and collect it into the 2 mL tube (total 1.5 mL homogenate in the tube).
- d. Centrifuge at 8,500 g for 10 min to separate the lipids from the cell lysate in brown adipocytes.
- e. After centrifugation, discard the upper lipid layer and the supernatant, and keep the pellet.
- f. Resuspend the pellet with 0.5 mL isolation buffer and transfer it into homogenizer. Wash the tube with 0.5 mL isolation buffer and collect it into the homogenizer (total 1 mL content in the homogenizer).
- g. Homogenize the resuspended pellet with a teflon pestle for 5–8 strokes and transfer the homogenate into a new 2 mL tube. Wash the homogenizer with 0.5 mL isolation buffer and collect it into the 2 mL tube (total 1.5 mL homogenate in the tube).
- h. Centrifuge at 700 g for 10 min to separate mitochondria from other cell components.
- i. Transfer the supernatant containing mitochondria to a new 2 mL tube and centrifuge at 8,500 g for 10 min to precipitate the mitochondria.
- j. Discard the supernatant and dissolve the mitochondrion pellet with 1% digitonin buffer containing 150 mM NaCl, 50 mM Tris-HCl (pH 7.5), 1 mM EDTA and 1× complete protease inhibitor.
- k. After dissolution on ice for 30 min, centrifuge at 20,000 g for 10 min at 4°C, and the supernatant (mitochondrial protein lysate) can be used to detect the knockout and rescue efficiency by western blot.

## EXPECTED OUTCOMES

Through local injection of AAV-sgRNA into the iBAT of *Rosa26-LSL-Cas9;Adipoq<sup>Cre</sup>* mice with adipocyte-specific expression of Cas9, we can efficiently and specifically knock out *Emre* in brown adipocytes of iBAT (Figure 7). Meanwhile, we can specifically rescue with tagged sgRNA-resistant protein in brown adipocytes of iBAT using AAV-ADP (Figure 7).

## LIMITATIONS

One limitation of this protocol is that the CRISPR-knockout system requires *Rosa26-LSL-Cas9;Adipoq<sup>Cre</sup>* mice generated by crossing *Rosa26-LSL (loxP-stop-loxP)-Cas9* with *Adiponectin<sup>Cre</sup>* mice. Another limitation is that the pAAV-ADP vector can only express proteins encoded cDNA no longer than 1.9 kb due to AAV package limitation.

## TROUBLESHOOTING

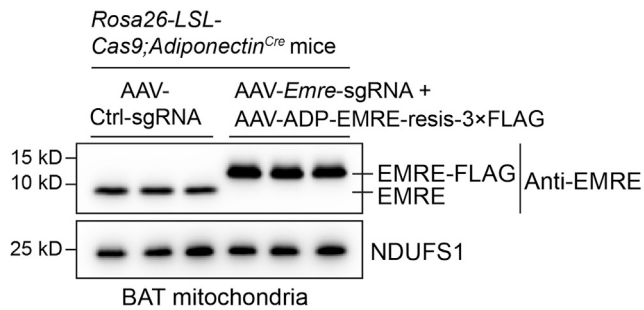
### Problem 1

Low knockout efficiency (in step 2 “Screen sgRNAs with editing efficiency in mouse melanoma cells”).

### Potential solution

The knockout efficiency in this protocol mainly relies on the editing efficiency of sgRNA for the target gene. Thus, more than three sgRNAs should be designed and screened in mouse melanoma cells so that at least two sgRNAs have relatively high editing activity.





**Figure 7. Knockout and rescue efficiency analysis of EMRE protein in BAT mitochondria via Western blot**

### Problem 2

Off-target effects of CRISPR-Cas9 system (in step 2 “Screen sgRNAs with editing efficiency in mouse melanoma cells” and step 5 “Design synonymously mutagenic primers for sgRNA-resistant cDNA of target gene”).

### Potential solution

Use at least two sgRNAs targeting different coding regions and use rescue experiments via expressing sgRNA-resistant cDNA of the target gene to validate the on-target phenotypes.

### Problem 3

Low efficiency of gene knockout or overexpression due to nonstandard operation of AAV local injection, i.e., without widespread infection of AAV in iBAT (in step 8 “Surgical operation for local injection of AAV into iBAT”).

### Potential solution

The virus solution should be injected into each lobe of iBAT by 10 evenly dispersed points to allow thorough infection. The injection depth of the needle should be in the middle of the fat pad. One can practice injection operation using trypan blue solution.

### Problem 4

Low success rate of vector construction by SLIC (in step 4 “Construct pAAV-ADP (adiponectin promoter)-EMRE-3xFLAG”).

### Potential solution

SLIC PCR primers for amplifying SLIC insert should be designed strictly according to the rules in [Figure 3](#). Insert and vector used for SLIC must be obtained from gel purification. When setting up SLIC reaction, amount of insert should be 2 folds more than that of vector.

### Problem 5

Poor control of the expression level of wild-type and mutant protein to a comparable level in the rescue experiments (in step 7 “Prepare virus solution”).

### Potential solution

The titers of AAVs expressing wild-type or mutant protein should be measured at the same time to avoid titration errors among batches. For proteins with the same molecular weight, an equal amount of AAVs should be injected. Usually, AAVs carrying cDNA encoding smaller protein will express more, such that one should inject mice with less amount of AAVs to achieve comparable protein expression.

## RESOURCE AVAILABILITY

### Lead contact

Further information and requests for resources and reagents should be directed to and will be fulfilled by the lead contact, Yifu Qiu ([yifu.qiu@pku.edu.cn](mailto:yifu.qiu@pku.edu.cn)).

### Materials availability

This study did not generate new unique reagents.

### Data and code availability

This study did not generate any unique datasets or code.

## SUPPLEMENTAL INFORMATION

Supplemental information can be found online at <https://doi.org/10.1016/j.xpro.2022.101895>.

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## AUTHOR CONTRIBUTIONS

K.X., D.W., and Y.Q. conceptualized the study and designed experiments. K.X. performed experiments and analyzed data. K.X., D.W., and Y.Q. wrote the paper.

## DECLARATION OF INTERESTS

A Chinese patent application (No. 202210961425.3) related to this work was filed (Y.Q., K.X., D.W., and H.S.).

## REFERENCES

- Xue, K., Wu, D., Wang, Y., Zhao, Y., Shen, H., Yao, J., Huang, X., Li, X., Zhou, Z., Wang, Z., and Qiu, Y. (2022). The mitochondrial calcium uniporter engages UCP1 to form a thermoprotector that promotes thermogenesis. *Cell Metab.* 34, 1325–1341.e6. <https://doi.org/10.1016/j.cmet.2022.07.011>.
- Romanelli, S.M., Lewis, K.T., Nishii, A., Rupp, A.C., Li, Z., Mori, H., Schill, R.L., Learman, B.S., Rhodes, C.J., and MacDougald, O.A. (2021). BAd-CRISPR: inducible gene knockout in interscapular brown adipose tissue of adult mice. *J. Biol. Chem.* 297, 101402. <https://doi.org/10.1016/j.jbc.2021.101402>.
- Jimenez, V., Muñoz, S., Casana, E., Mallol, C., Elias, I., Jambriña, C., Ribera, A., Ferre, T., Franckhauser, S., and Bosch, F. (2013). In vivo adeno-associated viral vector-mediated genetic engineering of white and brown adipose tissue in adult mice. *Diabetes* 62, 4012–4022. <https://doi.org/10.2337/db13-0311>.
- O'Neill, S.M., Hinkle, C., Chen, S.J., Sandhu, A., Hovhannisyanyan, R., Stephan, S., Lagor, W.R., Ahima, R.S., Johnston, J.C., and Reilly, M.P. (2014). Targeting adipose tissue via systemic gene therapy. *Gene Ther.* 21, 653–661. <https://doi.org/10.1038/gt.2014.38>.
- Huang, W., Queen, N.J., and Cao, L. (2019). rAAV-mediated gene delivery to adipose tissue. *Methods Mol. Biol.* 1950, 389–405. [https://doi.org/10.1007/978-1-4939-9139-6\\_23](https://doi.org/10.1007/978-1-4939-9139-6_23).
- Gómez-Banoy, N., and Lo, J.C. (2017). Genetic manipulation with viral vectors to assess metabolism and adipose tissue function. *Methods Mol. Biol.* 1566, 109–124. [https://doi.org/10.1007/978-1-4939-6820-6\\_11](https://doi.org/10.1007/978-1-4939-6820-6_11).
- Balkow, A., Hoffmann, L.S., Klepac, K., Glöde, A., Gnad, T., Zimmermann, K., and Pfeifer, A. (2016). Direct lentivirus injection for fast and efficient gene transfer into brown and beige adipose tissue. *J. Biol. Methods* 3, e48. <https://doi.org/10.14440/jbm.2016.123>.
- Bagchi, D.P., Forss, I., Mandrup, S., and MacDougald, O.A. (2018). SnapShot: niche determines adipocyte character I. *Cell Metab.* 27, 264–264.e1. <https://doi.org/10.1016/j.cmet.2017.11.013>.
- Clafkin, K.E., Flippo, K.H., Sullivan, A.I., Naber, M.C., Zhou, B., Neff, T.J., Jensen-Cody, S.O., and Potthoff, M.J. (2022). Conditional gene targeting using UCP1-Cre mice directly targets the central nervous system beyond thermogenic adipose tissues. *Mol. Metab.* 55, 101405. <https://doi.org/10.1016/j.molmet.2021.101405>.
- Platt, R.J., Chen, S., Zhou, Y., Yim, M.J., Swiech, L., Kempton, H.R., Dahlman, J.E., Parnas, O., Eisenhaure, T.M., Jovanovic, M., et al. (2014). CRISPR-Cas9 knockin mice for genome editing and cancer modeling. *Cell* 159, 440–455. <https://doi.org/10.1016/j.cell.2014.09.014>.
- Concordet, J.P., and Haeussler, M. (2018). CRISPOR: intuitive guide selection for CRISPR/Cas9 genome editing experiments and screens. *Nucleic Acids Res.* 46, W242–W245. <https://doi.org/10.1093/nar/gky354>.
- Chen, Y., Zeng, X., Huang, X., Serag, S., Woolf, C.J., and Spiegelman, B.M. (2017). Crosstalk between KCNK3-mediated ion current and adrenergic signaling regulates adipose thermogenesis and obesity. *Cell* 171, 836–848.e13. <https://doi.org/10.1016/j.cell.2017.09.015>.
- Fripont, S., Marneffe, C., Marino, M., Rincon, M.Y., and Holt, M.G. (2019). Production, purification, and quality control for adeno-associated virus-based vectors. *J. Vis. Exp.* <https://doi.org/10.3791/58960>.