

Design and construct **retroviral** sgRNA plasmids for gene knockout

1. Vector backbone

pMSCV-U6sgRNA(BbsI)-PGKpuro2ABFP (Addgene, #102796)

2. Design sgRNA oligos

I. Access CRISPick website (Broad institute)

<https://portals.broadinstitute.org/gppx/crispick/public>

II. Select Reference Genome: human or mouse?

- CRISPRa/i...NCBI reference data set
- CRISPRko...Ensembl data set

III. Select Mechanism

- CRISPRko

IV. Select the type of Enzyme

- SpyCas9(NGG)

V. On Target Scorer

Use Hsu(2013) if the tracrRNA in your vector starts with GTTTT.

(pLKO5.sgRNA.EFS.tRFP657, pLKO5.sgRNA.EFS.GFP, lentiGuide-Puro start with GTTTT, and pMSCV-U6sgRNA(BbsI)-PGKpuro2ABFP)

VI. Input Transcript IDs: Put RefSeq transcript ID

Click **validate** and then click **submit**

Status	Input	Resolved target	Notes
Valid	TP53	7157 (TP53)	

Targets (QUOTA = 5)		Additional Guides		USER-PROVIDED GUIDE SEQUENCES		Estimated # Picked
Experimental	1	NO_SITE	0	Additional negative controls	0	5
Positive Control	0	ONE_INTERGENIC_SITE	0	Preselects	0	

Click **picking results**

VII. Target Sequences

原則として上位 1, 2 の “target sequences (sgRNA sequences)” を使う (Pick order)

VIII. Design oligos

Oligo 1: **CACCG**(Target Sequence)**GT**

Oligo 2: **TAAAC**(Reverse Complement of Target Sequence)**C**

e.g.

Target Sequence: AATCACGAAATGGCCAACGG

Reverse Complement: CCGTTGGCCATTTTCGTGATT

(http://www.bioinformatics.org/sms/rev_comp.html)

Oligo1: **CACCG**AATCACGAAATGGCCAACGG**GT**

Oligo 2: **TAAAC**CCGTTGGCCATTTTCGTGATT**C**

3. Cloning Oligos into pMSCV-U6sgRNA(BbsI)-PGKpuro2ABFP**(1) Digesting pMSCV-U6sgRNA(BbsI)-PGKpuro2ABFP vector**

Vector (2 µg)	2 µl
10x rCutSmart™ Buffer	2 µl
BbsI-HF (NEB)	1 µl
DDW	15 µl
Total	20 µl

Incubate at 37°C for 2 hours

Inactivate at 65°C for 20 min and run 1% agarose gel

(2) Annealing oligos

Forward oligo (100 µM)	1 µl
Reverse oligo (100 µM)	1 µl
10x NEBuffer™ r2.1 (NEB) or 10x T buffer (Takara)	5 µl
DDW	43 µl
Total	50 µl

Vortex, spin down

95°C 4 min

70°C 10 min

Take out the heat block and cool it down to room temperature (at least 1 hour)

*Do not directly put it on ice!

(3) Run 1% agarose gel (100V, 25 min) with step 3-1 digested vectors

Expected fragment size:

pMSCV-U6sgRNA(BbsI)-PGKpuro2ABFP: **7457 bp (cut vector)**, 22 bp

-> Cut 7-8 kb digested vector and perform gel extraction

-> Elute DNA with 25 µl DDW

(4) Ligation

*10x dilution for annealed oligos: 5 µl mixture (from step 3-2) + 45 µl DDW

10x diluted oligos	3 µl
Purified vector (step 3-3)	1 µl
2x mighty ligation mix (Takara, 6023)	4 µl
Total	8 µl

Incubate at 16°C for 1-3 hours

(5) Transformation (DH5α)

-> Take 2 µl of ligation product (step 3-4) into 25 µl of DH5α

-> on ice 30 min -> heat shock 42°C for 45 sec -> on ice 3 min -> add 80 µl SOC

-> all for plating with Amp plate

(6) Mini-culture

Pick up 6 colonies each group for mini-culture (2 ml LB+2 µl Amp)

(7) Mini-prep

Dissolve DNA with 30 µl TE

(8) Enzyme digestion check

Mini-prep DNA	1 µl
10x rCutSmart™ Buffer	2 µl
BbsI-HF (NEB)	0.4 µl
BsiWI-HF (NEB)	0.4 µl
DDW	16.2 µl
Total	20 µl

Correct expected fragment size: 7500 bp (single cut)

Run 1% agarose gel (100V, 20 min)

(9) Select a correct plasmid for sequencing

U6 primer: gagggcctatttcccatgatt