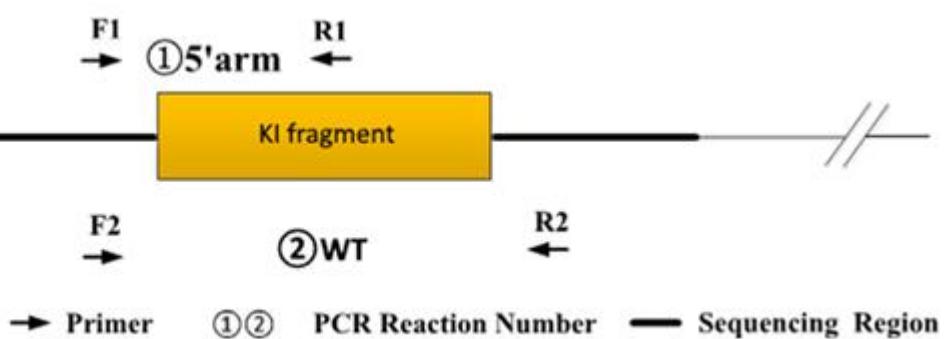




Genotyping Report

| | | | | | |
|-----------|-------------|-------------|----------|--------------------|--------------------------|
| Strain ID | T060079 | Strain Type | KI(Cas9) | Genetic Background | C57BL/6JGpt |
| Designer | Binjie Jiao | Gene Name | | | H11-Myh6-MerCreMer-ployA |

1. Strategy of Genotyping



Wild type: ①PCR reaction obtains none band; ②PCR reaction obtains a WT band.

Heterozygote: ①PCR reaction obtains a Targeted band; ②PCR reaction obtains a WT band.

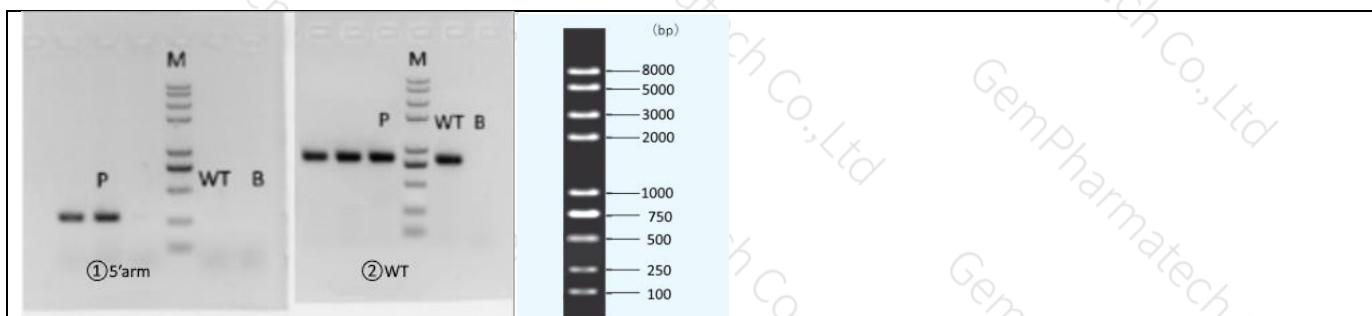
Homozygote: ①PCR reaction obtains a Targeted band; ②PCR reaction obtains none band.

Note: The sizes of WT and Targeted band are shown below. For ②PCR reaction, because the WT band is much smaller than the target band, it is likely to produce dominant amplification, the reaction is only used to judge whether there is a WT allele.

2. Primer Information

| PCR No. | Primer No. | Primer Name | Sequence | Band Size |
|---------|------------|-------------------------|-------------------------|-----------------------------|
| ①5'arm | F1 | H11-tF3 | GGGCAGTCTGGTACTTCCAAGCT | WT:0bp Targeted: 284bp |
| | R1 | GPT000491-01-CreER-5tR1 | TCTACTCCTCATTAGGCCCTTT | |
| ②WT | F2 | H11-wt-tF1a | AGTCTTCCTTGCCCTTGCT | WT:825bp Targeted:8963bp |
| | R2 | H11-wt-tR1a | GGGTCTTCCACCTTCTTCAG | |

3. Gel Image & Conclusion



Note: P:Heterozygous samples; WT: Wildtype control; B: Blank control (ddH₂O); M: DNA Ladder

① Control (WT) : It is an important reference mark for whether the PCR reaction is successful and whether the



product band position and size meet the theoretical requirements.

② Control (B) : PCR amplification was performed without template in the PCR reagent to monitor whether the reagent was contaminated.

4. PCR Condition

(Generally recommend to use Vazyme P222; If the sequences contain special structures such as GC% ≥ 60% or GC% ≤ 40%, recommend to use Vazyme P515.)

| PCR Reaction Component | | |
|------------------------|--|-------------|
| Seg. | reaction component | Volume (μl) |
| 1 | 2 × Rapid Taq Master Mix(Vazyme P222) or 2 × Phanta Max Master Mix (Vazyme P515) | 12.5 |
| 2 | ddH ₂ O | 9.5 |
| 3 | Primer A(10pmol/μl) | 1 |
| 4 | Primer B(10pmol/μl) | 1 |
| 5 | Template(20~80ng/μl) | 1 |

PCR program I priority selection

| Seg. | Temp. | Time | Cycle |
|------|-------------------------|------|-------|
| 1 | 95 °C | 5min | 20x |
| 2 | 98 °C | 30s | |
| 3 | 65 °C * (-0.5 °C/cycle) | 30s | |
| 4 | 72 °C | 45s* | |
| 5 | 98 °C | 30s | |
| 6 | 55 °C * | 30s | |
| 7 | 72 °C | 45s* | |
| 8 | 72 °C | 5min | |
| 9 | 10 °C | hold | |

PCR program II the second choice

| Seg. | Temp. | Time | Cycle |
|------|---------|------|-------|
| 1 | 95 °C | 5min | 35x |
| 2 | 98 °C | 30s | |
| 3 | 58 °C * | 30s | |
| 4 | 72 °C | 45s* | |
| 5 | 72 °C | 5min | |
| 6 | 10 °C | hold | |

Note*: Annealing temperature and extension time can be determined according to the actual amplification situation and amplification enzyme efficiency.



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