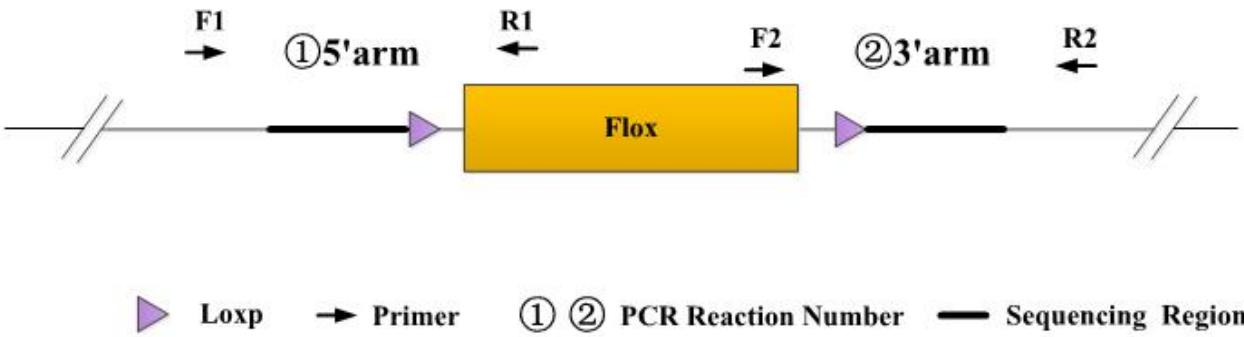




Genotyping Report

| | | | | | |
|-----------|--------------|-------------|-----------|--------------------|---------------|
| Strain ID | T006299 | Strain Type | CKO(Cas9) | Genetic Background | C57BL/6JGpt |
| Designer | Tiantian Sun | Gene Name | | | <i>Ptger4</i> |

1. Strategy of Genotyping



Wild type: ①PCR reaction obtains a single WT band; ②PCR reaction obtains a single WT band.

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

Homozygote: ①PCR reaction obtains a single Targeted band; ②PCR reaction obtains a single Targeted band.

Note: The sizes of WT and Targeted band are shown below.

2. Primer Information

| PCR No. | Primer No. | Primer Name | Sequence | Band Size |
|----------|------------|-------------|--------------------------|----------------------------|
| ①(5'arm) | F1 | T006299-F1 | CTAATCCAATAGAAACGATCTGCC | WT:331bp Targeted:411bp |
| | R1 | T006299-R1 | GAAGCTGGAATAAGATGGGCTTCT | |
| ②(3'arm) | F2 | T006299-F2 | ATGTAGGCTCTAGCCAGCATCTG | WT:384bp Targeted:462bp |
| | R2 | T006299-R2 | CTGTTCACTTACATCATGGTTCCT | |

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% $\geq 60\%$ or GC% $\leq 40\%$, recommend to use Vazyme P515.)

| PCR Reaction Component | | |
|------------------------|--|-------------------|
| Seg. | reaction component | Volume (μ l) |
| 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/ μ l) | 1 |
| 4 | Primer B(10pmol/ μ l) | 1 |
| 5 | Template(20~80ng/ μ l) | 1 |

PCR program I priority selection

| Seg. | Temp. | Time | Cycle |
|------|-------------------------|------|-------|
| 1 | 95 °C | 5min | |
| 2 | 98 °C | 30s | 20× |
| 3 | 65 °C * (-0.5 °C/cycle) | 30s | |
| 4 | 72 °C | 45s* | |
| 5 | 98 °C | 30s | 15× |
| 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 | |
| 2 | 98 °C | 30s | 35× |
| 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.