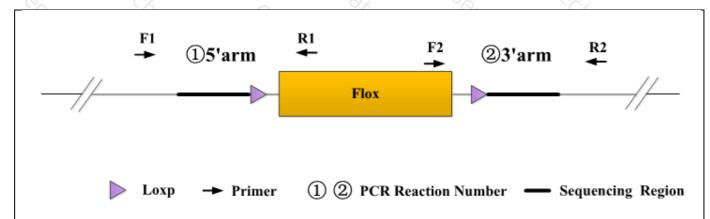


Genotyping Report

| Strain ID | T064960 | Strain Type | CKO(Cas9) | Genetic Background | C57BL/6JGpt |
|-----------|-------------|-------------|-----------|--------------------|-------------|
| Designer | Binjie Jiao | Gene Name | 3/2/ | Smarca4 | 6 |

1. Strategy of Genotyping



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

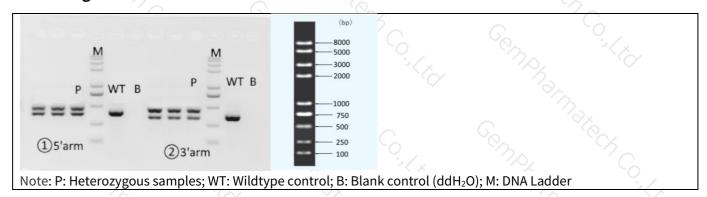
Heterozygote: ①PCR reaction obtains a WT band and a Targeted band; ②PCR reaction obtains a WT band and a Targeted band.

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 | T064960(P1)-F1 | AGTGACACCTCTTACAACAGACCA | WT: 380bp Targeted: 485bp |
| | R1 | T064960(P1)-R1 | GACACTCACAAAGATACGATACTGG | |
| ②(3'arm) | F2 | T064960(P1)-F2 | ATGGCCTTGCCTCATA | WT: 326bp |
| | R2 T064960(P1 | | TCTGCTAGTGACCCCAAGGACAA | Targeted: 432bp |

3. Gel Image & Conclusion





- 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 | ```` | 120 | |
|--------------|----------------------|--|---|--|
| Seg. | r | reaction component | | |
| 1 | 9% | 2 × Rapid Taq Master Mix(Vazyme P222) or 2 × Phanta Max Master Mix (Vazyme P515) | | |
| 2 | 16 | ddH2O | | |
| 3 % | 34 ₆ , f | Primer A(10pmol/μl) | | |
| 1 | 3. F | Primer B(10pmol/μl) | | |
| 5 | Y/2 To | Template(20~80ng/μl) | | |
| PCR program | I priority selection | , CX | 0 3/2 | |
| Seg. | Temp. | Time | Cycle | |
| 1 7 | 95℃ | 5min / | 1/2/2 5./ | |
| 2 | 98°C | 30s | 20× | |
| 3 66 | 65°C* (-0.5°C/cycle) | 65°C* (-0.5°C/cycle) 30s | | |
| 1 70/ | 72°C | 45s* | 7/s) 7/C | |
| 5 9 | 98℃ | 30s | 15× | |
| 5 | 55°C* | 30s | , J ⁹ × | |
| Co | 72°C | 45s* | | |
| 3 % | 72°C | 5min | 7% (C) | |
| 9/2 | 10°C | hold | 3/2 | |
| PCR program | II the second choice | 3/2 | 7/2 · · · · · · · · · · · · · · · · · · · | |
| Seg. | Temp. | Time | Cycle | |
| | 95℃ | 5mín | | |
| <u> </u> | 98℃ | 30s | 35× | |
| 3 | 58°C* | 30s | 79/2 | |
| 1 | 72°C | 45s* | 73x | |
| 5 | 72°C ∕ | 5min | 600 | |
| 5 % | 10°C | hold | 702 (C) | |



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