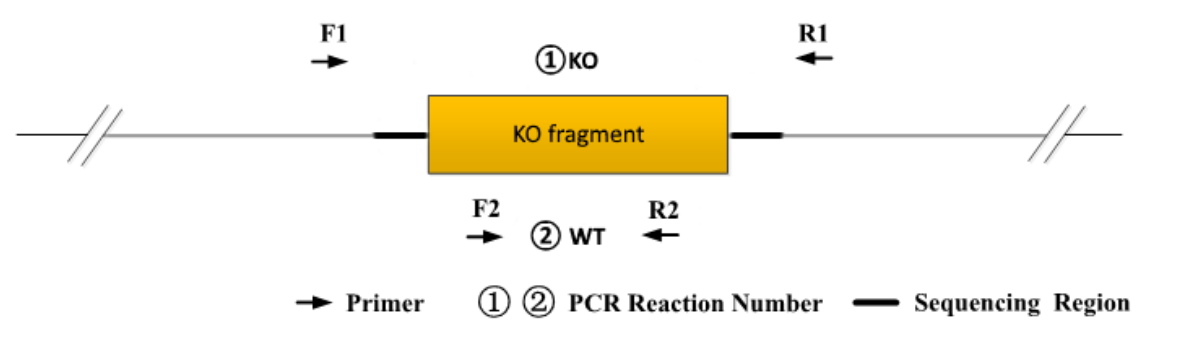


Genotyping Report

| | | | | | |
|-----------|-----------|-------------|---------------|--------------------|-------------|
| Strain ID | T005504 | Strain Type | KO(Cas9) | Genetic Background | C57BL/6JGpt |
| Designer | Zifan Lin | Gene Name | <i>Trim31</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 KO band; ②PCR reaction obtains a WT band.
 Homozygote: ①PCR reaction obtains a single KO band; ② PCR reaction without product.
 Note: 1)The sizes of WT and Targeted band are shown below.
 2)If the WT band is too large, it may not be possible to obtain a WT band.

2. Primer Information

| PCR No. | Primer No. | Primer Name | Sequence | Band Size |
|---------|------------|------------------------|-----------------------------|------------------------|
| PCR① | F1 | JS00271-Trim31-5wt-tF2 | TAAAGGTGTGTGCCAGCACTGC | WT:16108bp KO:184bp |
| | R1 | JS00271-Trim31-3wt-tR2 | CTTGTAAGTGAACAATGTTGCGGG | |
| PCR② | F2 | JS10271-Trim31-wt-F2 | GAGGAAGGGTTAAGAAGTGTGAGC | WT:334bp KO:0bp |
| | R2 | JS10271-Trim31-wt-R2 | GTGACAAAGTCCACTCAGGTGTAATTG | |

3. Gel Image

tacttttgctggttcacccaacgctttc-----15924bp-----catggtccaaccctcggggagttggcagg



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 | |
| 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.