## Genotyping Report

| Strain ID | T019286 | Strain Type | CKO（Cas9） | Genetic Background | C57BL／6JGpt |
| :--- | :---: | :---: | :---: | :---: | :---: |
| Designer | Yin Chen | Gene Name |  | Mcam |  |

## 1．Strategy of Genotyping



Wild type：（1）PCR reaction obtains a single WT band；（2）PCR reaction obtains none band．
Heterozygote：（1）PCR reaction obtains a WT band and a Targeted band；（2）PCR reaction obtains a Targeted band．
Homozygote：（1）PCR reaction obtains a single Targeted band；（2）PCR reaction obtains a 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 | $\begin{aligned} & \text { JS09580-Mcam- } \\ & \text { 5wt-tF1 } \end{aligned}$ | TGTAAAGAGAGCACAGCTCCTTCTG | WT：296bp Targeted：398bp |
|  | R1 | JS09580－Mcam－ 5wt－tR1 | CTAGTGTTAGGGGGTTGAAAGTGTG |  |
| （2）（3＇arm） | F2 | ZMK2F4 | CATCGCATTGTCTGAGTAGGTG | WT：0bp Targeted：336bp |
|  | R2 | $\begin{aligned} & \text { JS09580-Mcam- } \\ & \text { 3wt-tR1 } \end{aligned}$ | GCTTGGTATTTACCCTAGAAAGCACT |  |

## 3．Gel Image \＆Conclusion



Note：P：Heterozygous samples；WT：Wildtype control；B：Blank control（ $\mathrm{ddH}_{2} \mathrm{O}$ ）；M：DNA Ladder
（1）Control（WT）：It is an important reference mark for whether the PCR reaction is successful and whether the

## 4．PCR Condition

（Generally recommend to use Vazyme P222；If the sequences contain special structures such as GC\％$\geqslant 60 \%$ or $\mathrm{GC} \% \leqslant 40 \%$ ，recommend to use Vazyme P515．）

| PCR Reaction Component |  |  |
| :--- | :---: | :--- |
| Seg． | reaction component | Volume $(\boldsymbol{\mu l})$ |
| 1 | $2 \times$ Rapid Taq Master Mix（Vazyme P222） |  |
| or |  |  |
|  | $2 \times$ Phanta Max Master Mix（Vazyme P515） | 12.5 |
| 2 | ddH2O |  |
| 3 | Primer A（10pmol／$\mu \mathrm{l})$ | 9.5 |
| 4 | Primer $\mathrm{B}(10 \mathrm{pmol} / \mu \mathrm{l})$ | 1 |
| 5 | Template $\left.20^{\sim} 80 \mathrm{ng} / \mu \mathrm{l}\right)$ | 1 |

PCR program I priority selection

| Seg． | Temp． | Time | Cycle |
| :--- | :--- | :--- | :--- |
| 1 | $95^{\circ} \mathrm{C}$ | 5 min |  |
| 2 | $98^{\circ} \mathrm{C}$ | 30 s |  |
| 3 | $65^{\circ} \mathrm{C}^{*}\left(-0.5^{\circ} \mathrm{C} /\right.$ cycle $)$ | 30 s |  |
| 4 | $72^{\circ} \mathrm{C}$ | $45 \mathrm{~s}^{*}$ |  |
| 5 | $98^{\circ} \mathrm{C}$ | 30 s |  |
| 6 | $55^{\circ} \mathrm{C}^{*}$ | 30 s |  |
| 7 | $72^{\circ} \mathrm{C}$ | $45 s^{*}$ |  |
| 8 | $72^{\circ} \mathrm{C}$ | 5 min |  |
| 9 | $10^{\circ} \mathrm{C}$ | hold |  |
| PCR program II the second choice |  |  |  |


| Seg． | Temp． | Time | Cycle |
| :--- | :--- | :--- | :--- |
| 1 | $95^{\circ} \mathrm{C}$ | 5 min |  |
| 2 | $98^{\circ} \mathrm{C}$ | 30 s | $35 \times$ |
| 3 | $58^{\circ} \mathrm{C}^{*}$ | 30 s |  |
| 4 | $72^{\circ} \mathrm{C}$ | $45 \mathrm{~s}^{*}$ |  |
| 5 | $72^{\circ} \mathrm{C}$ | 5 min |  |
| 6 | $10^{\circ} \mathrm{C}$ | hold |  |

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

