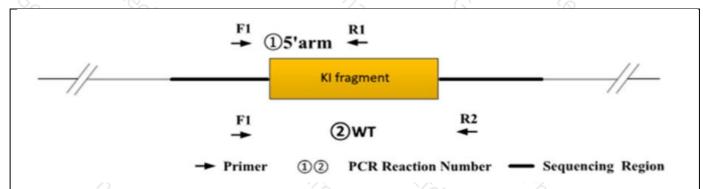


# **Genotyping Report**

Strain ID	T053087	Strain Type	KI(Cas9)	Genetic Background	C57BL/6JGpt
Designer	Dongdong Zhang	Gene Name	CAG-Rox-ZsGreen-Stop-Rox-tdTomato-ployA		ato-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.

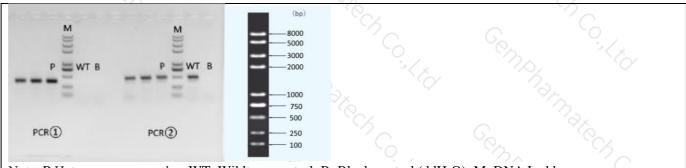
Homozygote: ①PCR reaction obtains a Targeted band; ②PCR reaction obtains none band.

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	T053087-F1	CCCAAAGTCGCTCTGAGTTGTTA	WT:0bp	
	R1	T053087-R1	TGGCGTTACTATGGGAACATACGTC	Targeted:375bp	
②WT	F1	T053087-F1	CCCAAAGTCGCTCTGAGTTGTTA	WT:479bp	
	R2	T053087-R2	TCGGGTGAGCATGTCTTTAATCT	Targeted:6427bp	

#### 3. Gel Image & Conclusion



Note: P:Heterozygous samples; WT: Wildtype control; B: Blank control (ddH<sub>2</sub>O); 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  $GC\% \ge 60\%$  or  $GC\% \le 40\%$ , recommend to use Vazyme P515.)

PCR Reaction Com	ponent ponent	25 6	9%	
Seg.	reaction comp	Volume (μl)		
1 73/7/73×	2 × Rapid Taq Master M or 2 × Phanta Max Master M	12.5		
2	ddH2O	ddH2O		
3	Primer A(10pr	Primer A(10pmol/µl)		
4	Primer B(10pr	1 3/x		
5 1/2/	Template(20~8	10,		
PCR program $\overset{\circ}{\mathrm{I}}$ p	riority selection	<u> </u>	9/2	
Seg.	Temp.	Time	Cycle	
	95℃	5min	200	
2 3	98℃	30s	20×	
3	65℃* (-0.5℃/cycle)	30s		
4 %	<b>72℃</b>	45s*	7/20/	
70/	98℃	30s	15×	
5 9/2	55℃*	30s	39/2 3/X	
7	<b>72℃</b>	45s*	J <sup>3×</sup> , 4	
3 %	<b>72</b> ℃	5min	200	
9 %	10°C	hold	5 6	
PCR program $ m II$ $$ t	he second choice		9/2 3/x	
Seg.	Temp.	Time	Cycle	
1	95℃	5min	, C/2	
2	98℃	30s	35×	
3 %	58℃*	30s	72/	
1	72℃	45s*	79/2	
5	<b>72℃</b>	5min	Dax.	
6	10°C	hold	, 000	

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