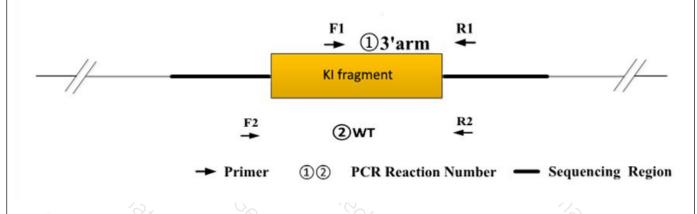


		Genoty	oing Report		~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
Strain ID	T004713	Strain Type	KI(Cas9)	Genetic Background	C57BL/6JGpt
Designer	Tiantian Sun	Gene Name	· · · / *	Myh6-iCre	°C -
Q <sub>1</sub> m	2 ( X	$\sim$	$\checkmark$	a contraction of the second se	3/4

## 1. Strategy of Genotyping

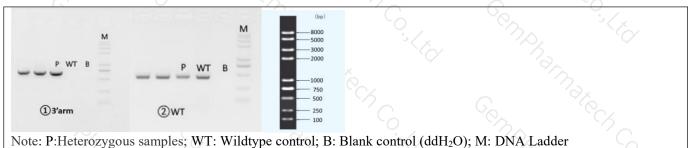


Wild type: ①PCR reaction obtains none band; ②PCR reaction obtains a WT band. Heterozygote: ①PCR reaction obtains a Targeted band; ②PCR reaction obtains a WT band. Homozygote: ①PCR reaction obtains a Targeted band; ②PCR reaction obtains none band. Note: The sizes of WT and Targeted band are shown below. 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	
(1)22	Fl	T004713-F1B	ATCAGAAAGGAGAATGTGGATGCTG	WT:0bp	
(1)3°arm	R1 (	T004713-R1B	ATGTTCACATTGGTCCAGCCACC	Targeted:602bp	
②WT	F2	T004713-F2	CAGCAAAACCTGGCTGTGGATC	WT:412bp Targeted:0bp	
	R2	T004713-R2	ATGAGCCACCATGTGGGTGTC	Targeted.00p	

### 3. Gel Image & Conclusion



① Control (WT) : It is an important reference mark for whether the PCR reaction is successful and whether the



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product band position and size meet the theoretical requirements.

<sup>(2)</sup> 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.)

1.2	1 N					
Seg.		reaction component	1	Volume (µl)		
1	Charne	2 × Rapid Taq Master Mix(Vazyme P222) or 2 × Phanta Max Master Mix (Vazyme P515)	- Chan	12.5		
2		ddH2O	9	9.5		
3		Primer A(10pmol/µl)		1 7 7		
4	Co.	Primer B(10pmol/µl)	S	1 0		
5	$\gamma_{\mathcal{A}_{\mathcal{A}}}$	Template(20~80ng/μl)	- Vy	1 <sub>0</sub> , <sup>1</sup> 0/		

#### PCR program I priority selection

PCR program	1 priority selection		the second s
Seg.	Temp.	Time	Cycle
1	95°C	5min	
2	98°C	30s 3	20×
3	65℃*(-0.5℃/cycle)	30s	
4 6	72℃	45s*	
5 <sup>3</sup> 2.	98°C	30s	15×
6	55°C*	30s	
7	72°C	45s*	The second
8	72°C	5min	
9 2	10°C	hold	ns, nc
PCR program	II the second choice	ns C	
Seg.	Temp.	Time	Cycle
1	95°C	5min	
2 6	98°C	30s	35×
3	58°C*	30s	
4	<b>72℃</b>	45s*	
5	72°C	5min	The second se
6	10°C	hold	

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