

		Genoty	oing Report		Colored Colored
Strain ID	T020587	Strain Type	CKO(Cas9)	Genetic Background	C57BL/6JGpt
Designer	Ya'nan Xu	Gene Name	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~	Capn6	20
Strategy of	Genotyping	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~	Array .	Stern S	
	F1 → ①5'arr	n 🕇	F2	2)3'arm ♣	
_//_		• <b>&gt;</b>			<u> </u>

Loxp - Primer (1) (2) PCR Reaction Number - Sequencing Region

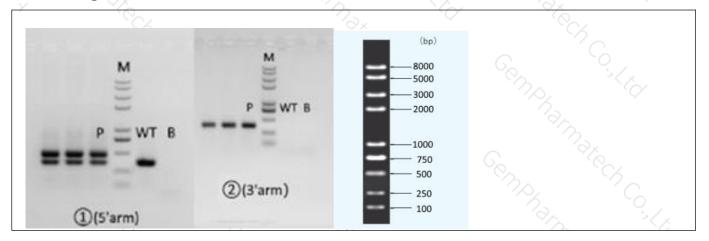
Wild type: ①PCR reaction obtains a single WT band; ②PCR reaction obtains a single WT band. Heterozygote: ①PCR reaction obtains a WT band and a Targeted band; ②PCR reaction obtains a WT band and a Targeted band. Homozygote: ①PCR reaction obtains a single Targeted band; ②PCR reaction obtains a single Targeted band.

Note: The sizes of WT and Targeted band are shown below.

## 2. Primer Information

PCR No.	Primer No.	Sequence		Band Size	
1)(5'arm)	T020587-F1	CACCAGATGAACTCTG	GGGATAC	WT:341bp Targeted:445bp	
	T020587-R1	CCATAACAATCAGCTTT	ГСССААСС		
@(3'arm)	T020587-F2	TCTGAGGCGGAAAGAA	CCAG	WT:0bp	
	T020587-R2 CACTCAGCATACTTATTCTAGCCTGC		CTAGCCTGC	Targeted:346bp	

## 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

4. PCR Condi	tion		Party istra		
PCR Reaction Co	mponent	3. 0	24		
Seg.	reaction co	reaction component			
ı M	2 × Rapid Taq Master Mix (Vazym	2 × Rapid Taq Master Mix (Vazyme P222)			
2 3.	ddH2O	0	9.5		
3	Primer A(10pmol/µl)	Primer A(10pmol/µl)			
ļ	Primer B(10pmol/µl)	Primer B(10pmol/µl)			
; 6,	Template(20~80ng/μl)	$^{\prime}$ C (			
CR program I	priority selection	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~	Mr. Co		
Seg.	Temp.	Time	Cycle		
L C	95°C	5min	The second		
	98°C	30s	20× 20×		
s Tr	65℃*(-0.5℃/cycle)	30s	K. G.		
1 <sup>(4)</sup>	72℃	45s*			
G C	98℃	30s	15×		
; <sup>5</sup> 75.	55℃*	30s	3. <sup>°</sup> 30		
	72℃	45s*			
3	72℃	5min	3. 4		
jC <sub>C</sub>	10°C	hold			
CR program II	the second choice				
Seg.	Temp.	Time	Cycle		
r ?	95°C	5min	1737 × 14		
2	98°C	30s	35×		
6	58°C*	30s	G C		
1 <sup>7</sup> 0	72°C	45s*			
;	∂72℃	5min	The second se		
;	10°C	hold	- Bar		

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



