

		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 ♣	
//		• >			<u> </u>

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

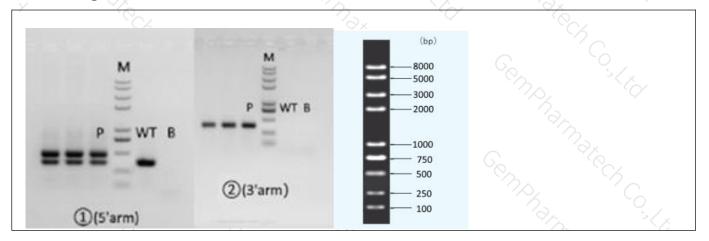
Wild type: ①PCR reaction obtains a single WT band; ②PCR reaction obtains a single WT band. Heterozygote: ①PCR reaction obtains a WT band and a Targeted band; ②PCR reaction obtains a WT band and a Targeted band. Homozygote: ①PCR reaction obtains a single Targeted band; ②PCR reaction obtains a single Targeted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

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 ⁽⁴⁾	72℃	45s*			
G C	98℃	30s	15×		
; ⁵ 75.	55℃*	30s	3. [°] 30		
	72℃	45s*			
3	72℃	5min	3. 4		
jC _C	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 ⁷ 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.



