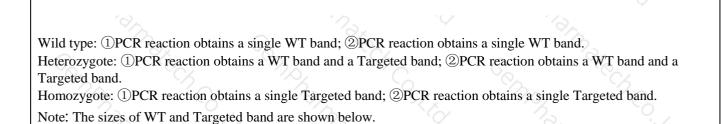


Strain ID	T054498	Strain Type	CKO(Cas9)	Genetic Background	C57BL/6JGp
Designer	Ya'nan Xu	Gene Name	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~	Foxk2	Co Co
Strategy of	Genotyping			annate	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~



1 2 PCR Reaction Number - Sequencing Region

Flox

Primer

Loxp

2. Primer Information

PCR No.	Primer No.	Sequence	Band Size	
(1)(5'arm)	T054498-F1	ACACGACATTACACGCACATGCT	WT: 300bp Targeted:405bp	$\langle \rangle$
	T054498-R1	GCCAAGCTTCTGCATATGAGGAC		< (
2)(3'arm)	T054498-F2	AGTGACTTCCAGTTTTGTAAACTCAAGA	WT: 259bp Targeted:365bp	
	T054498-R2	054498-R2 GTGAGGCTCAATGGACAGTACCA		

3. Gel Image & Conclusion





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Note: P: Positive control; 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

PCR Reaction Co	omponent	AX Con	20 ²	
Seg.	reactio	reaction component		
1 3/2	2 × Rapid Taq Master Mix (Va	2 × Rapid Taq Master Mix (Vazyme P222)		
2	ddH2O	ddH2O		
3	Primer A(10pmol/µl)	AX A	1 ~~~~	
4 🖓	Primer B(10pmol/µl)	Primer B(10pmol/µl)		
5	Template(≈100ng/µl)	Template(≈100ng/μl)		
PCR program (1)	priority selection	m the		
Seg.	Temp.	Time	Cycle	
1 6	95°C	5min		
2	98°C	30s	20×	
3	65℃*(-0.5℃/cycle)	30s	2/2 3/2	
4	72℃	45s*	12× 4	
5 %	98°C	30s	20×	
6	55℃*	30s		
7 7	72℃	45s*	Ph Str	
8	72℃	5min	134 24	
9 7	10°C	hold		
PCR program ②	the second choice			
Seg.	Temp.	Time	Cycle	
1	95°C	5min	197 ₀ 06	
2	98°C	30s	35×	
3	58°C*	30s 🔾		
4	72°C	45s*		
5	72°C	5min	19/2	
6 6	10°C	hold	97	

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



