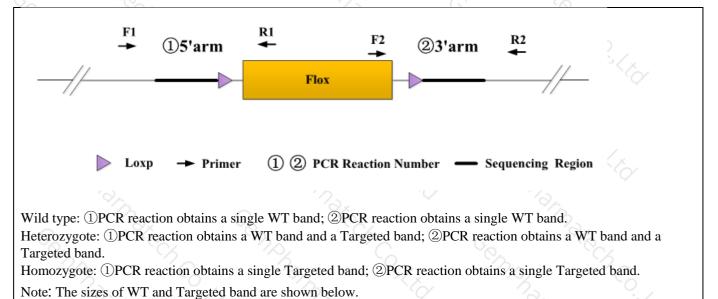


		Genotyp	ing Report		~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~
Strain ID	T008226	Strain Type	CKO(Cas9)	Genetic Background	C57BL/6JGpt
Designer	Ya'nan Xu	Gene Name	~<~~	Traf6	C
9/2		~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~	Ý	Ph	34

## 1. Strategy of Genotyping



## 2. Primer Information

PCR No.	Primer No.	Sequence	Band Size	
(1)(5'arm)	T008226-F1	CGTTCATGGAGATTGGTTGCA	WT: 272bp Targeted: 377bp	
	T008226-R1	T008226-R1 GCCATGTGGCAGAACTTAGATGA		
2)(3'arm)	T008226-F2	TTTGAGTTCAAGGCCAGCCTT	WT: 298bp	
	m) T008226-R2 AGTCTCTGACAAGGAGGCTCTCATT		- Targeted: 404bp	

## 3. Gel Image & Conclusion





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Note: P: Positive control; 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.

<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

PCR Reaction Com	ponent	AL COL	20°	
Seg.	reaction co	reaction component		
1 20	2 × Rapid Taq Master Mix (Vazyme	2 × Rapid Taq Master Mix (Vazyme P222)		
2 2	ddH2O	· · · · · · · · · · · · · · · · · · ·	9.5	
3	Primer A(10pmol/µl)	Primer A(10pmol/µl)		
4	Primer B(10pmol/µl)	1 6		
5	Template(≈100ng/µl)	Template(≈100ng/µl)		
PCR program $\textcircled{1}$ ,	priority selection			
Seg.	Temp.	Time	Cycle	
1 6	95°C	5min	× 500	
2	98°C	30s	20×	
3	65°C* (-0.5°C/cycle)	30s		
4	<b>72℃</b>	45s*	A A A	
5	98°C	30s	20×	
6	55℃*	30s	$\overline{\mathcal{A}}$	
7 75	72°C	45s*	on str	
8	72℃	5min	192	
9	10°C	hold		
PCR program ②	the second choice		×. 6.	
Seg.	Temp.	Time	Cycle	
1	95℃	5min	197 <sub>8</sub> 04	
2	98°C	30s	35×	
3	58°C*	30s 🕤		
4 7	72°C	45s*		
5	72°C	5min	and a	
6	10℃	hold	AX.	

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



