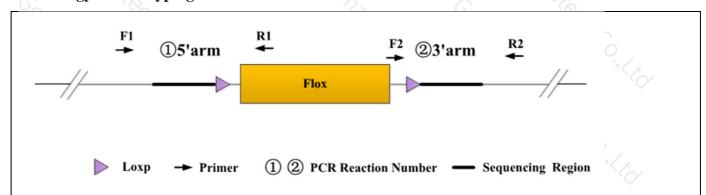


## **Genotyping Report**

Strain ID	T025050	Strain Type	CKO(Cas9)	Genetic Background	C57BL/6JGpt
Designer	Zifan Lin	Gene Name	3/2	Nsmce1	9

## 1. Strategy of Genotyping



Wild type: ①PCR reaction obtains a single WT band; ②PCR reaction obtains none band.

Heterozygote: ①PCR reaction obtains a WT band and a Targeted band; ②PCR reaction obtains a Targeted band.

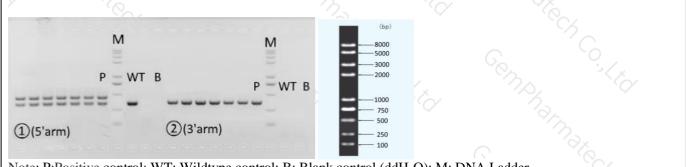
Homozygote: ①PCR reaction obtains a single Targeted band; ②PCR reaction obtains a Targeted band.

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

#### 2. Primer Information

PCR No.	Primer No.	Sequence	Band Size	
①(5'arm)	T025050-F1	GCTGTTAGTATCTGTTCACCCAAGC	WT: 340bp Targeted: 445bp	
	T025050-R1	TAGCCAGGACACATCTGAAGGC		
②(3'arm)	T025050-F2	CATCGCATTGTCTGAGTAGGTG	AGGTG WT: 0bp	
	T025050-R2	CTGTACCACTGTGACACAGGATAAGC	Targeted: 342bp	

## 3. Gel Image & Conclusion



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.
- 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 Compo	onent		3
Seg.	reaction component		Volume (μl)
10 PX	2 × Rapid Taq Master Mix (Vazyme	12.5	
2 2	ddH2O	9.5	
3	Primer A(10pmol/μl)	7	1 6
4	Primer B(10pmol/μl)	1, 34	
5	Template(≈100ng/μl)	19%	
PCR program ① prid	ority selection		3
Seg.	Temp.	Time	Cycle
1	95℃	5min	
2	98℃	30s	20×
3	65°C* (-0.5°C/cycle)	30s	9%
4	<b>72℃</b>	45s*	, %
5	98℃	30s	20×
6	55℃*	30s	3/2
7 🕝	<b>72℃</b>	45s*	(2)X
8	<b>72℃</b>	5min	^ 33
9	10°C	hold	9./
PCR program ② the	e second choice	<u></u>	m, 0
Seg.	Temp.	Time	Cycle
1	95℃	5min	3
2	98℃	30s	35×
3	58℃*	30s	79x
4	<b>72℃</b>	45s*	.62
5	72℃	5min	6
6	10℃	hold	75 34E

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