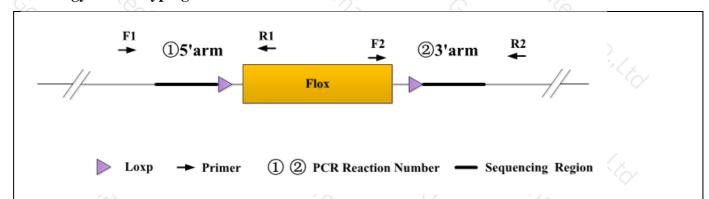


# **Genotyping Report**

Strain ID	T027127	Strain Type	CKO(Cas9)	Genetic Background	C57BL/6JGpt
Designer	Ya'nan Xu	Gene Name	3/2	Coro2b	6

# 1. Strategy of Genotyping



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

1 1/2			7/1
PCR No.	Primer No.	Sequence	Band Size
①(5'arm)	T027127-F1	TTGGCCCATCTTTCTCTCCACAG	WT: 265bp
	T027127-R1 AAATCTGCTTCTCCTCCTTCTGCC		Targeted:370bp
②(3'arm)	T027127-F2	AAGAGCAGCTTCCATCCCAGTTG	WT: 292bp
	T027127-R2	TGAGTAGGCAGAGCTTGCACAGTG	Targeted:398bp

# 3. Gel Image & Conclusion



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 C	77.			
Seg.	r	reaction component		
1 70,	2 × Rapid Taq Master M	2 × Rapid Taq Master Mix (Vazyme P222)		
2	ddH2O	70, 70	9.5	
3	Primer A(10pmol/μl)	79/2 3/x	12	
4	Primer B(10pmol/μl)	Primer B(10pmol/μl)		
5	Template(≈100ng/μl)	Template(≈100ng/μl)		
PCR program	1) priority selection	%, (C)	G., 3/x	
Seg.	Temp.	Time	Cycle	
1	95℃	5min	1977	
2	98℃	30s	20×	
3	65°C* (-0.5°C/cycle)	30s	30 S	
4	72℃	45s*	9/2	
5	98℃	30s	20×	
6	55℃*	30s	C YOU	
7	<b>72</b> ℃	45s*	7/20 · // C	
8	72℃	5min	3/x 3/x	
9	<b>10</b> ℃	hold	- 73×	
PCR program (	2) the second choice			
Seg.	Temp.	Time	Cycle	
1 3/7	95℃	5min		
2	98℃	30s	35×	
3	58℃*	30s	3	
4	<b>72</b> ℃	45s*	G	
5	<b>72</b> ℃	5min	70/	
6	10℃	hold	70A.	

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