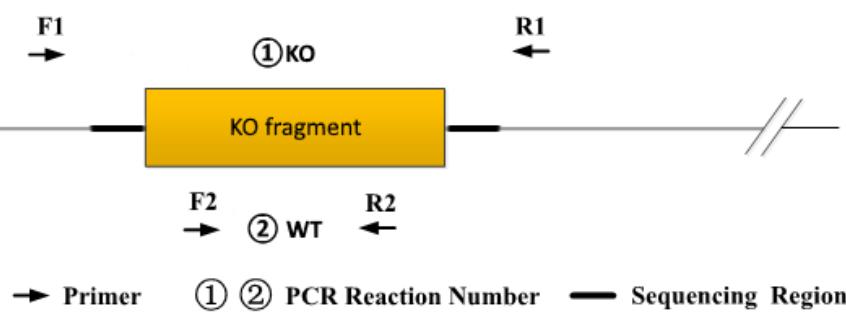




Genotyping Report

Strain ID	T013988	Strain Type	KO(Cas9)	Genetic Background	C57BL/6JGpt
Designer	Zifan Lin	Gene Name			<i>Cyp7b1</i>

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 KO band; ②PCR reaction obtains a WT band.

Homozygote: ①PCR reaction obtains a single KO band; ②PCR reaction without product.

Note: 1)The sizes of WT and Targeted band are shown below.

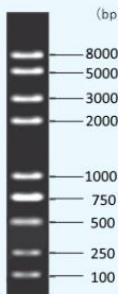
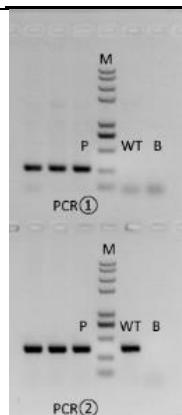
2) If the WT band is too large, it may not be possible to obtain a WT band.

2. Primer Information

PCR No.	Primer No.	Primer Name	Sequence	Band Size
PCR①	F1	JS04449-Cyp7b1-5wt-tF1	CAAACCCAACTTGATGAGAAGG AGT	WT: 1747bp KO: 254bp
	R1	JS04449-Cyp7b1-3wt-tR1	AAACTAATGCTGCCTGTTCAAGG G	
PCR②	F2	JS14449-Cyp7b1-wt-tF1	CACTTGGTACACAAACATACATG AAGGC	WT: 330bp KO:0bp
	R2	JS14449-Cyp7b1-wt-tR1	TCAACAACTGCTTGAGACCTGAG ACAC	

3. Gel Image

ttaagttagagaccatggctggccccaaccct---1493bp---cagcagcaacctctgtacattttggctaccatg



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

PCR Reaction Component

Seg.	reaction component	Volume (μl)
1	2 × Rapid Taq Master Mix (Vazyme P222)	12.5
2	ddH ₂ O	9.5
3	Primer A(10pmol/μl)	1
4	Primer B(10pmol/μl)	1
5	Template(20~80ng/μl)	1

PCR program I (priority selection)

Seg.	Temp.	Time	Cycle
1	95°C	5min	20x
2	98°C	30s	
3	65°C * (-0.5°C/cycle)	30s	
4	72°C	45s*	
5	98°C	30s	
6	55°C *	30s	
7	72°C	45s*	
8	72°C	5min	
9	10°C	hold	

PCR program II (the second choice)

Seg.	Temp.	Time	Cycle
1	95°C	5min	
2	98°C	30s	35x



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3	58°C*	30s	
4	72°C	45s*	
5	72°C	5min	
6	10°C	hold	

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