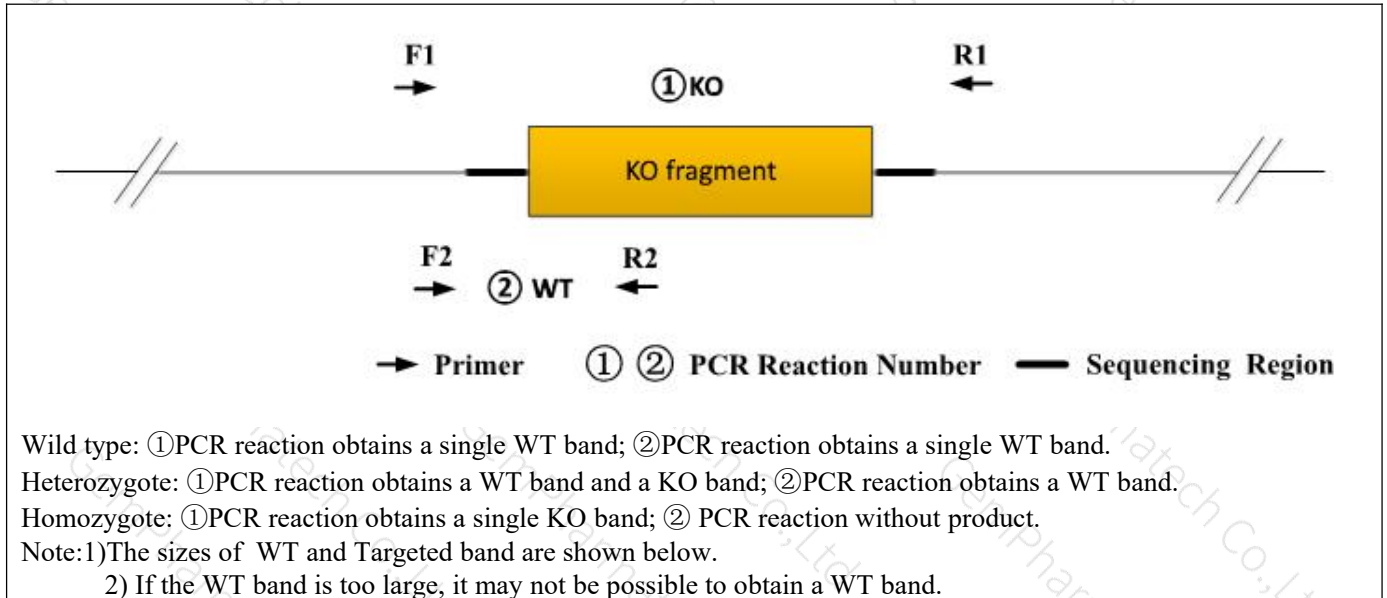


Genotyping Report

Strain ID	T045824	Strain Type	KO(Cas9)	Genetic Background	C57BL/6JGpt
Designer	Yuting Liu	Gene Name	Yod1		

1. Strategy of Genotyping



2. Primer Information

PCR No.	Primer No.	Primer Name	Sequence	Band Size
PCR①	F1	T045824(P1)-F1	GCTGTAGAATTAACCAACAGCAG	WT: 6612bp KO: 430bp
	R1	T045824(P1)-R1	TTTTCAAGATTGGCAGAGACCA	
PCR②	F1	T045824(P1)-F1A	CTCGAATCGAGCTGTCAGATGG	WT: 520bp KO: 0bp
	R2	T045824(P1)-R2A	GCACAAGCTGGATTCAAGACTCC	

3. Gel Image

catgcgaagtgcaaagaacaccccagtgac---6182bp---atctgggtgggttagatctgtttgaagat



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

(Generally recommend to use Vazyme P222; If the sequences contain special structures such as GC% ≥ 60% or GC% ≤ 40%, recommend to use Vazyme P515.)

PCR Reaction Component			
Seg.	reaction component		Volume (μl)
1	2 × Rapid Taq Master Mix(Vazyme P222) or 2 × Phanta Max Master Mix (Vazyme P515)		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	20×
2	98°C	30s	
3	65°C* (-0.5°C/cycle)	30s	
4	72°C	45s*	
5	98°C	30s	15×
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	35×
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.