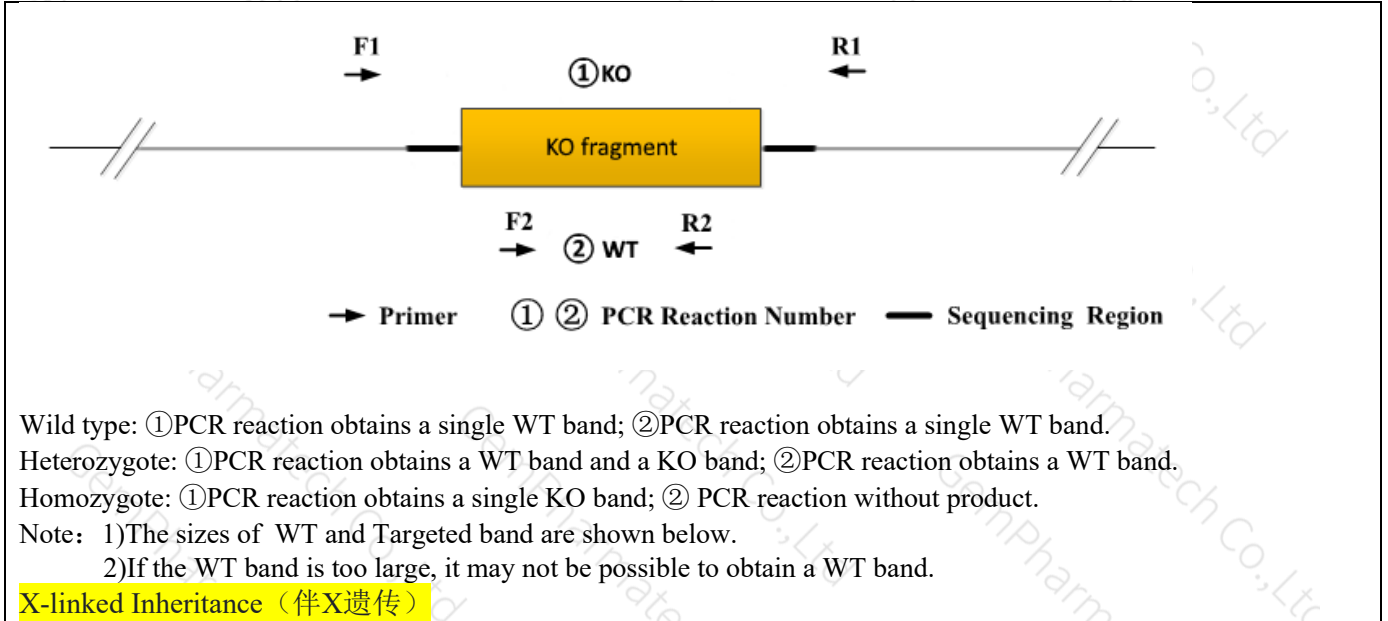


Genotyping Report

Strain ID	T006559	Strain Type	KO(Cas9)	Genetic Background	C57BL/6JGpt
Designer	Zifan Lin	Gene Name	Cxcr3		

1. Strategy of Genotyping

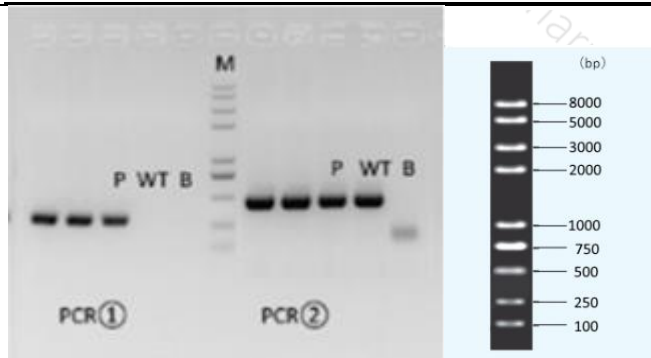


2. Primer Information

PCR No.	Primer No.	Primer Name	Sequence	Band Size
PCR①	F1	JS10636-Cxcr3-KO-tF1	AGGTGTAGAGCTGAAGTGGAAAGTG	WT:2728bp KO: 270bp
	R1	JS10636-Cxcr3-KO-tR1	CCAAAGCATTACTGTCTTCAGCC	
PCR②	F2	JS10636-Cxcr3-wt-tF1A	CCAAGTCCTAACACACTCCAAGTG	WT:426bp KO:0bp
	R2	JS10636-Cxcr3-wt-tR1A	CCAGACCTGGTTCAAAGAATGAC	

3. Gel Image

CCCAAGAGTTTCAAATCAGCCTGGCCATCTTGGGAAGA---2458bp--- TCAGCAAGATGGCAAGATCTATCTGAAATGGCTTTCTGG



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℃	5min	
2	98℃	30s	20×
3	65℃*(-0.5℃/cycle)	30s	
4	72℃	45s*	
5	98℃	30s	15×
6	55℃*	30s	
7	72℃	45s*	
8	72℃	5min	
9	10℃	hold	

PCR program II the second choice

Seg.	Temp.	Time	Cycle
1	95℃	5min	
2	98℃	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.

Reviewer:Ting Sun

Date:2020-7-1