

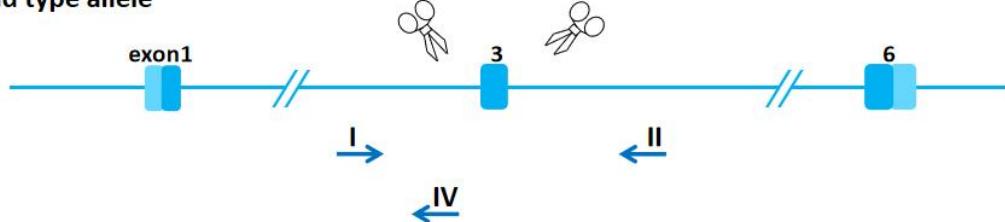
# C1qbp-KO Genotyping Protocol



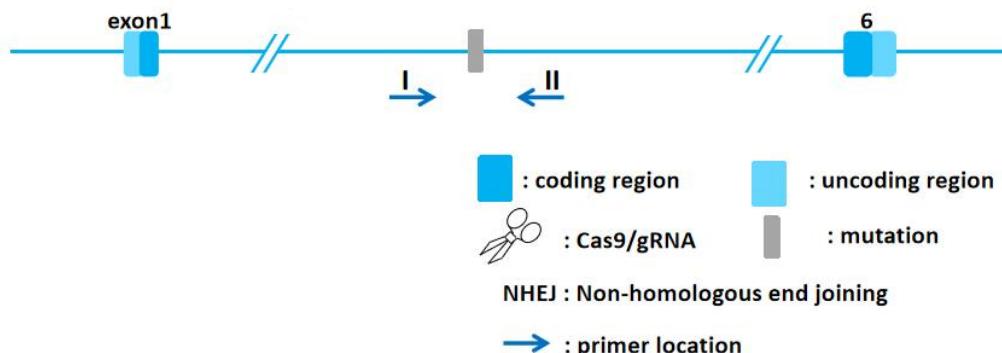
Common Name	C1qbp-KO	Cat. NO.	NM-KO-235080
Strain of Origin	C57BL/6J	Version	V1

### Genotyping strategy

#### Wild type allele



#### knockout allele



■ : coding region      ■ : uncoding region  
 : Cas9/gRNA      ■ : mutation

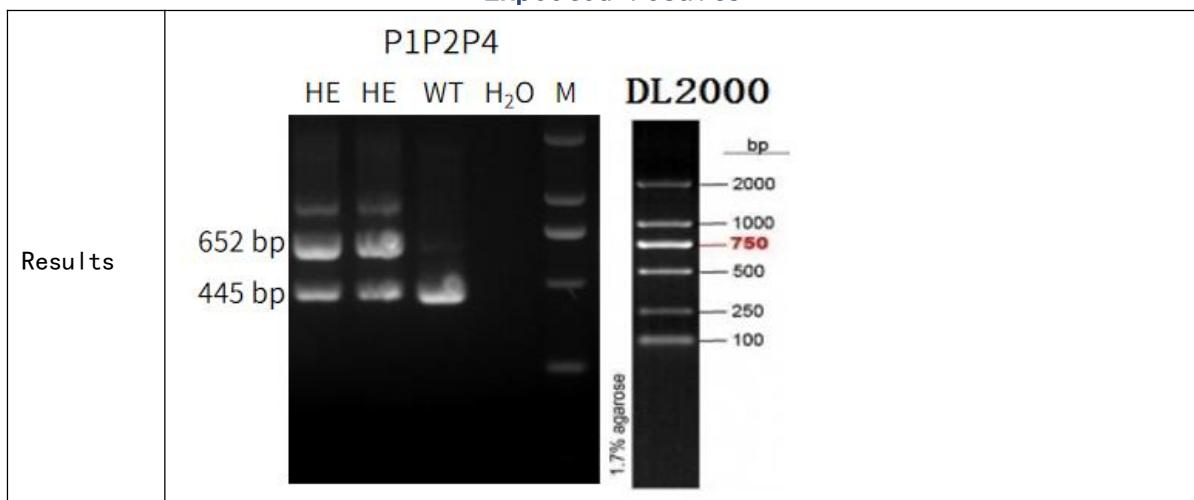
NHEJ : Non-homologous end joining

→ : primer location

### Primers

Primer	Sequence (5' → 3' )	Primer type
P1	GCCATCAGACAAAGGTAGAAGGTT	Forward
P2	GGCTGTGGATAAGGCCTCAA	Reverse
P4	CCTCTGCCTGGCTGGGATT	Reverse

### Expected results



	Knockout type: -1976bp
Genotype	Wild type: P1P4=445 bp Heterozygote: P1P2 =652 bp; P1P4=445 bp Homozygote: P1P2 =652 bp

Note: In both wild-type and heterozygous mice, whether the P1 and P2 primers can amplify larger bands does not affect the interpretation of the results, because the purpose of designing this pair of primers is to amplify K0 band

### Reaction &Cycling

PCR Reaction System	Reaction Component		Volume ( $\mu$ l)
	ddH2O		7.5
	2 $\times$ Taq Plus Master Mix		10.0
	P1(10 pmol/ $\mu$ l)		0.5
	P2(10 pmol/ $\mu$ l)		0.5
	P4(10 pmol/ $\mu$ l)		0.5
	Genomic DNA		1.0
	Total		20
2 $\times$ Taq Plus Master Mix from Vazyme (Code Number: P222-1)			
Cycling Reaction	Step	Temp	Time
	1	95° C	5 min
	2	95° C	20 sec
	3	60° C	20 sec
	4	72° C	20 sec
	5	72° C	5 min
	6	12° C	Hold
			Note
			35 repeats to 2