

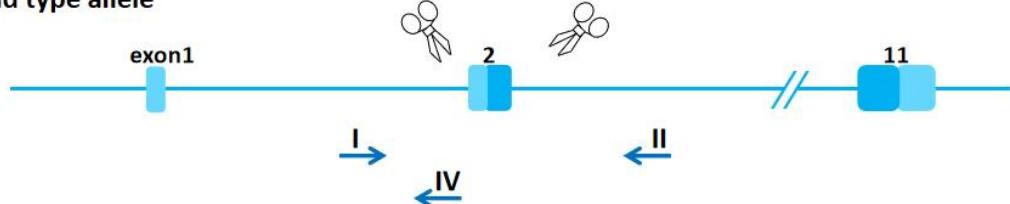
Akap1-KO Genotyping Protocol



Common Name	Akap1-KO	Cat. NO.	NM-KO-2103716
Strain of Origin	C57BL/6J	Version	V1

Genotyping strategy

Wild type allele



knockout allele



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

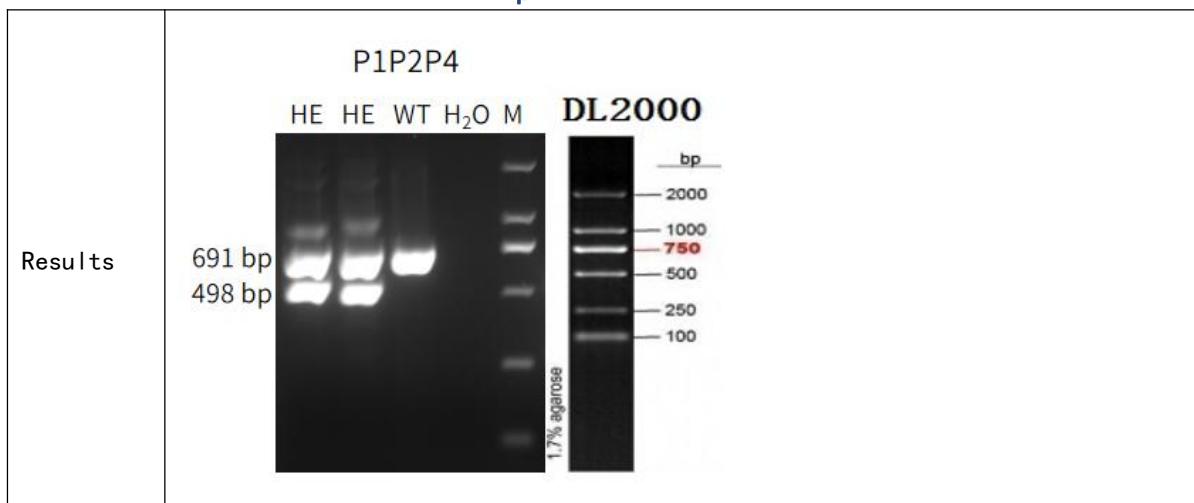
NHEJ : Non-homologous end joining

→ : primer location

Primers

Primer	Sequence (5' → 3')	Primer type
P1	AGACGCTGGAGATCCTGCTTGAAT	Forward
P2	CTGCGGCTGTCCCTGGGTGAA	Reverse
P4	AGTACGCTGGGCATGGTGAT	Reverse

Expected results



	Knockout type: -2324bp
Genotype	Wild type: P1P4=691 bp Heterozygote: P1P2 =498 bp; P1P4=691 bp Homozygote: P1P2 =498 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 (μ l)
	ddH2O		7.5
	2×Taq Plus Master Mix		10.0
	P1(10 pmol/ μ l))		0.5
	P2(10 pmol/ μ l)		0.5
	P4(10 pmol/ μ l)		0.5
	Genomic DNA		1.0
	Total		20
2×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