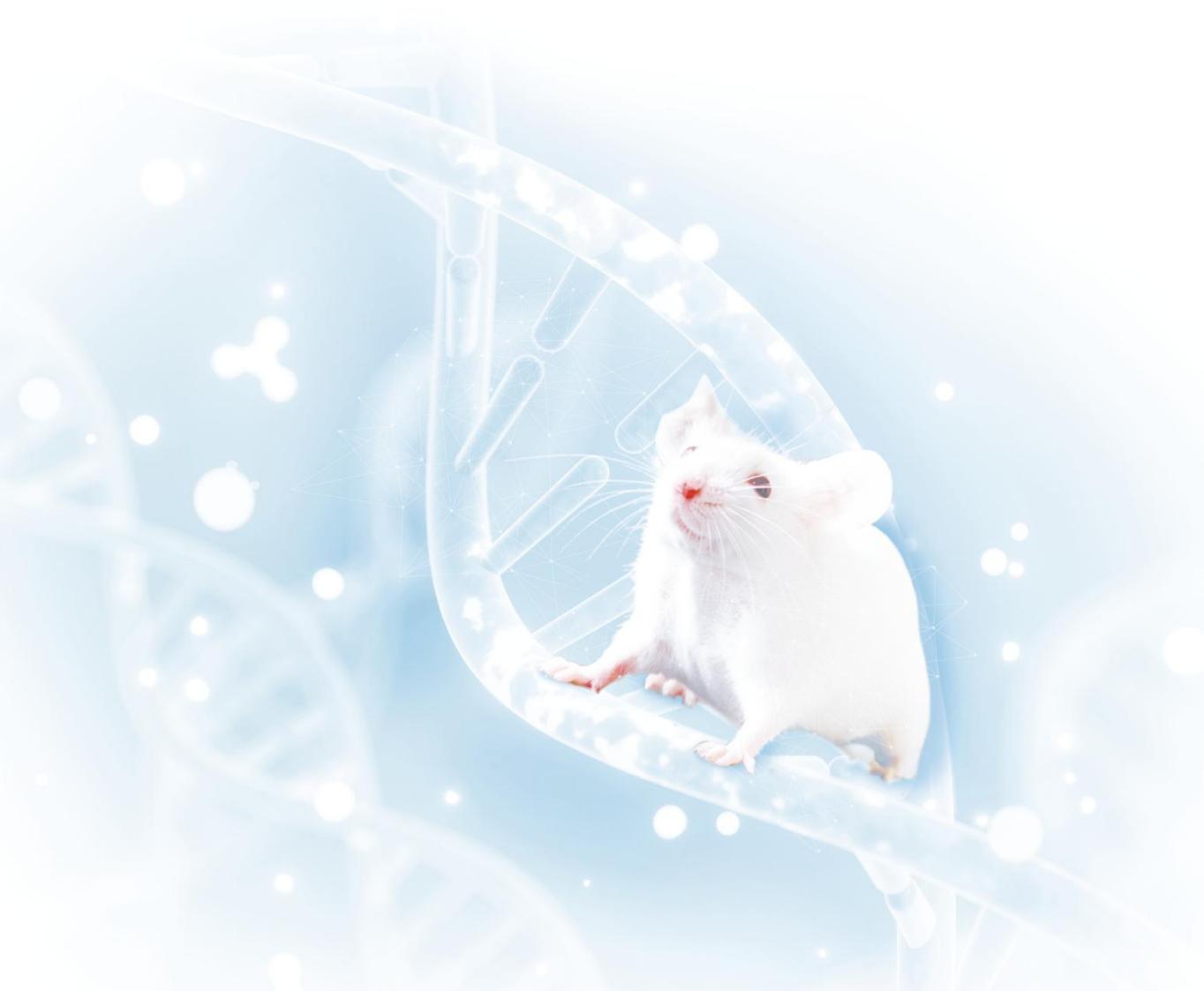


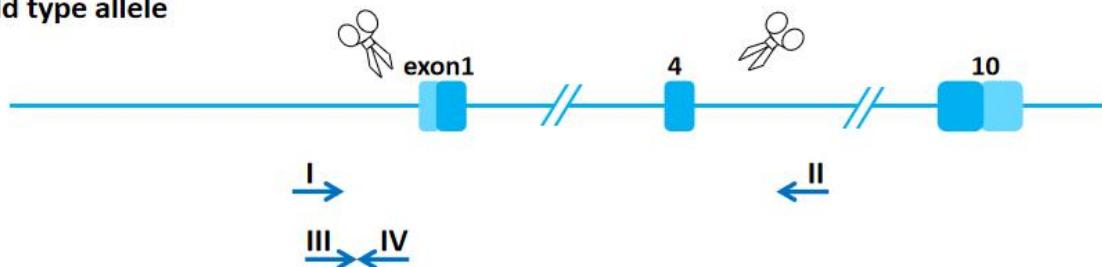
Akr1b7-KO Genotyping Protocol



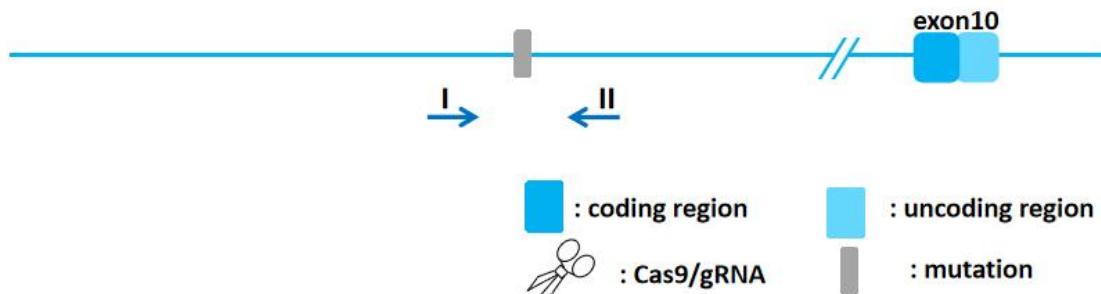
Common Name	Akr1b7-KO	Cat. NO.	NM-KO-234845
Strain of Origin	C57BL/6J	Version	V1

Genotyping strategy

Wild type allele



knockout allele



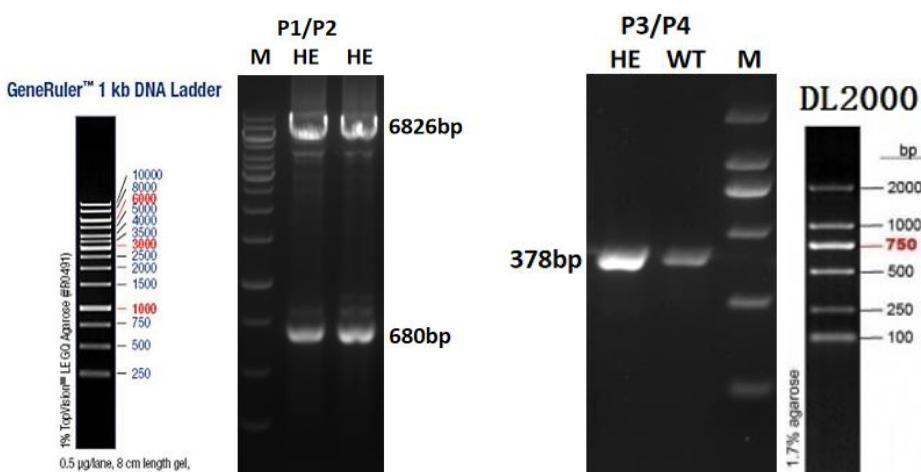
NHEJ : Non-homologous end joining

→ : primer location

Primers

Primer	Sequence (5' → 3')	Primer type
P1	AACCCTCTGCACATACTGGG	Forward
P2	AGAGCCTCTGACTTGCAGTA	Reverse
P3	CTGCTCTAACCGAGGGACCCA	Forward
P4	CTGTGAAACGCAAGGATGGT	Reverse

Expected results

Results	 <p>GeneRuler™ 1 kb DNA Ladder</p> <p>P1/P2</p> <p>M HE HE</p> <p>6826bp</p> <p>680bp</p> <p>P3/P4</p> <p>HE WT M</p> <p>378bp</p> <p>DL2000</p> <p>1.7% agarose</p> <p>bp</p> <p>10000 8000 6000 4000 3500 3000 2500 2000 1500 1000 750 500 250</p> <p>0.5 μg/lane, 8 cm length gel, 1xTAE, 7 V/cm, 45 min</p>
Genotype	<p>Knockout type: -6146bp</p> <p>Wild type: P1P2 =6826 bp; P3P4=378 bp</p> <p>Heterozygote: P1P2 =6826 bp and 680 bp; P3P4=378 bp</p> <p>Homozygote: P1P2 =680 bp</p>

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 Component	Volume (μl)
Reaction	ddH ₂ O	8. 0
System	2×Taq Plus Master Mix	10. 0

	P1(10 pmol/ μ l) or P3(10 pmol/ μ l)		0.5
	P2(10 pmol/ μ l) or 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