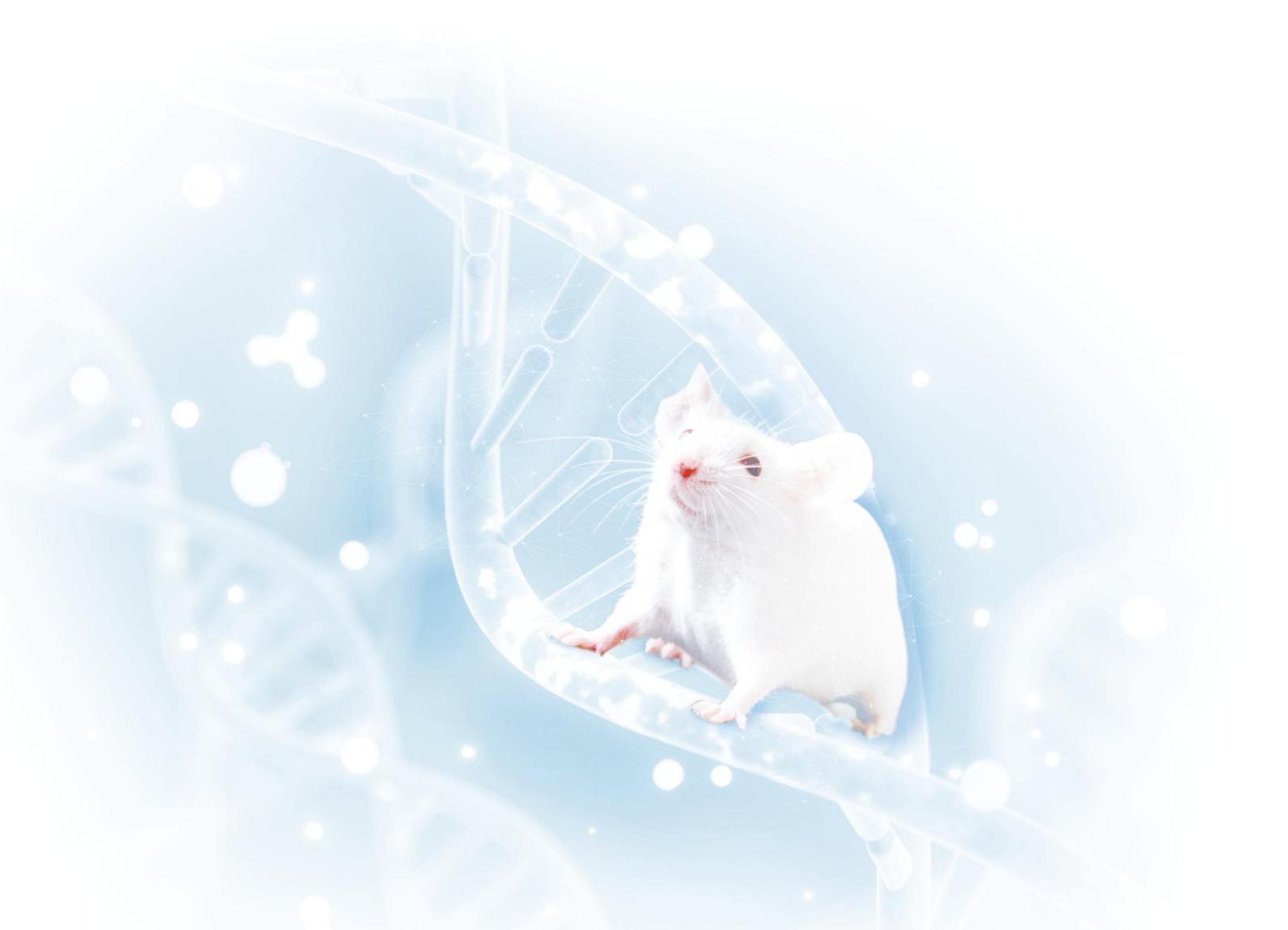


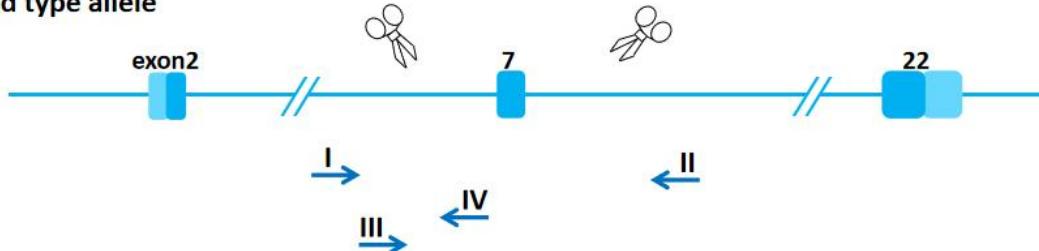
Thbs1-KO Genotyping Protocol



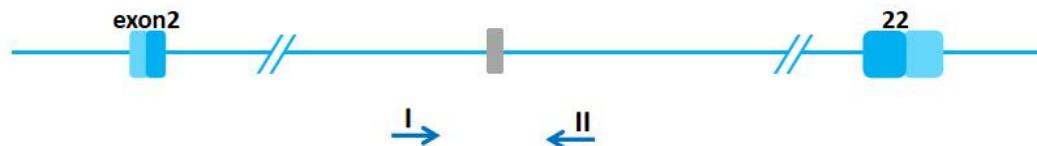
Common Name	Thbs1-KO	Cat. NO.	NM-KO-226309
Strain of Origin	C57BL/6J	Version	V1

Genotyping strategy

Wild type allele



knockout allele



█ : coding region █ : uncoding region
scissors : Cas9/gRNA █ : mutation

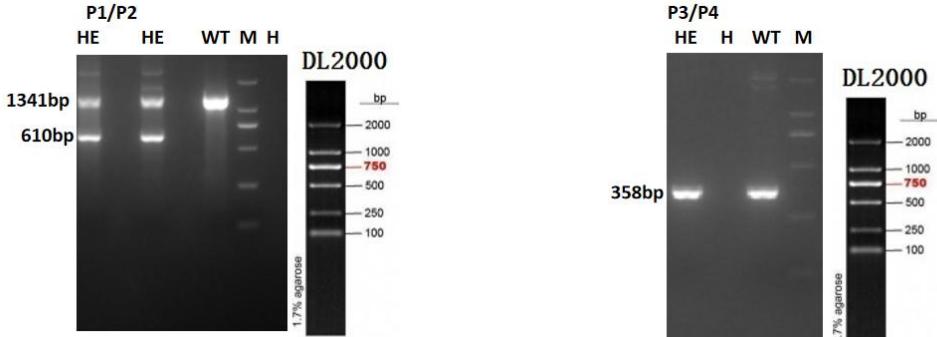
NHEJ : Non-homologous end joining

→ : primer location

Primers

Primer	Sequence (5' → 3')	Primer type
P1	TCTGGCCTGCTACCAAAGTG	Forward
P2	CTGAGTCGTGGCTCATGTGT	Reverse
P3	CGGTTTGGCCAGCCCTAAT	Forward
P4	AGGAGGGATGTCTCTGCTGG	Reverse

Expected results

Results	 <p>P1/P2</p> <p>HE HE WT M H</p> <p>1341bp 610bp</p> <p>P3/P4</p> <p>HE H WT M</p> <p>358bp</p> <p>DL2000</p> <p>1.7% agarose</p>
Genotype	<p>Knockout type: -732+1bp</p> <p>Wild type: P1P2 =1341 bp; P3P4 =358 bp</p> <p>Heterozygote: P1P2 =1341 bp and 610 bp; P3P4 =358 bp</p> <p>Homozygote: P1P2 =610 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 KO band.

Reaction &Cycling

PCR Reaction System	Reaction Component		Volume (μl)
	ddH ₂ O		8.0
	2 × Rapid Taq Master Mix		10.0
	P1(10 pmol/μl) or P3(10 pmol/μl)		0.5
	P2(10 pmol/μl) or P4(10 pmol/μl)		0.5
	Genomic DNA		1.0
	Total		20
	2 × Rapid Taq Master Mix from Vazyme(Code Number: P222-01)		
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