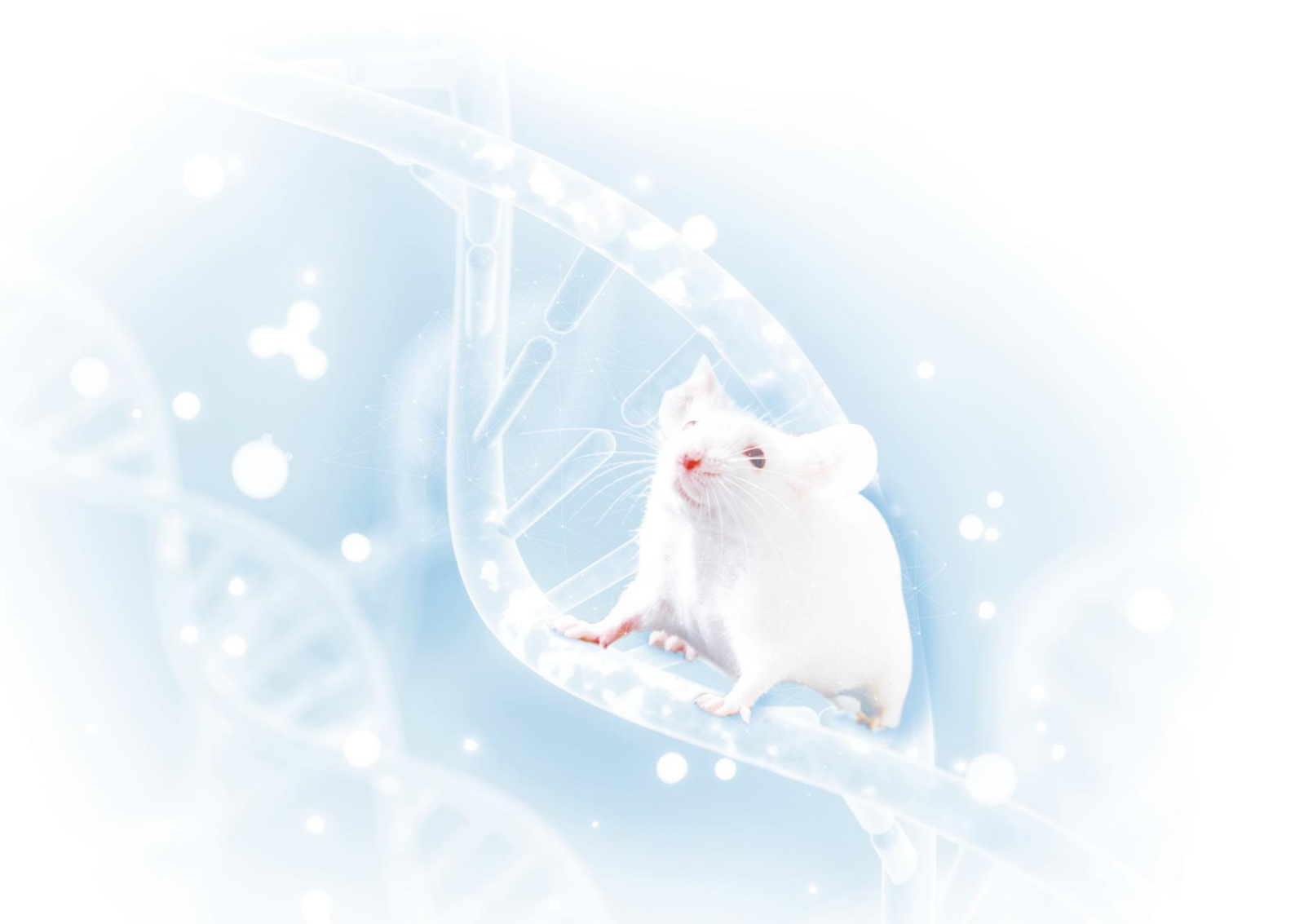
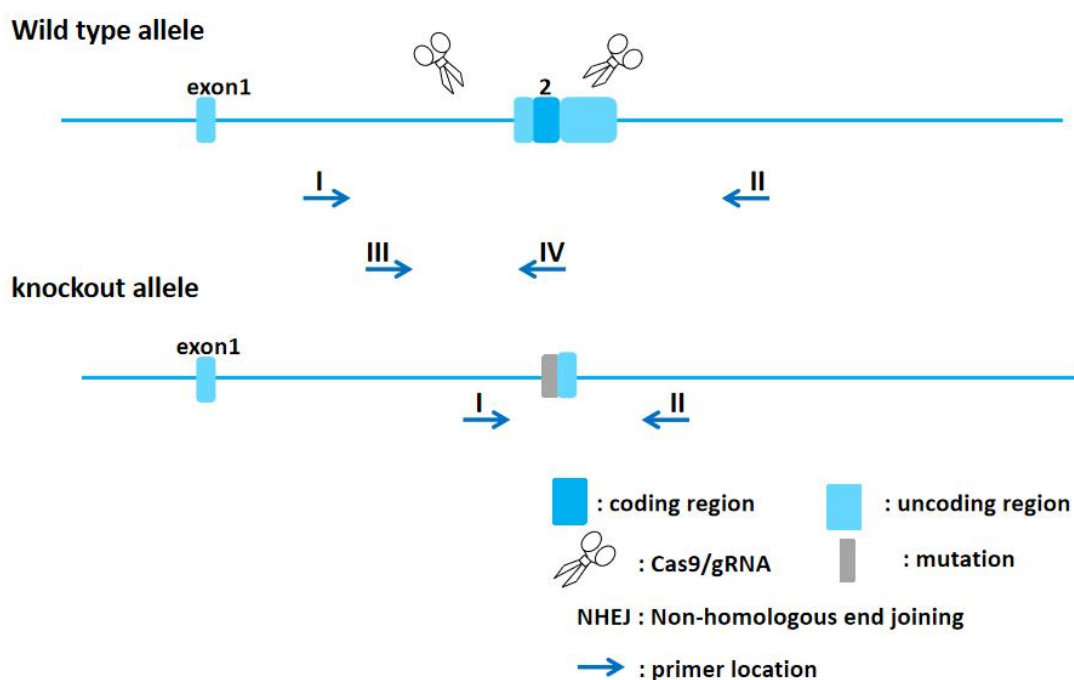


Ang2-KO Genotyping Protocol



Common Name	Ang2-KO	Cat. NO.	NM-KO-240420
Strain of Origin	C57BL/6J	Version	V1

Genotyping strategy



Primers

Primer	Sequence (5' → 3')	Primer type
P1	CATCTTGCCCTCAGGAACCC	Forward
P2	CCCTGACTAGCAGGTAACGA	Reverse
P3	ACACATACTGCACCTGTCCTT	Forward
P4	TCACAACCAGACCCAACAGG	Reverse

Expected results

Results	GeneRuler™ 1 kb DNA Ladder 1299bp 582bp P1/P2 HE HE WT M 423bp P3/P4 HE HE WT H M DL2000 bp 2000 1000 750 500 250 100 1.7% agarose 0.5 µg/lane, 8 cm length gel, 1XTAE, 7 V/cm, 45 min
Genotype	Knockout type: -717 bp Wild type: P1P2 =1299 bp; P3P4 =423 bp Heterozygote: P1P2 =1299 bp and 582 bp; P3P4 =423 bp Homozygote: P1P2 =582 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 KO band.

Reaction &Cycling

	Reaction Component	Volume (µl)
PCR	ddH2O	8.0
Reaction	2 × Rapid Taq Master Mix	10.0
System	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	Note
	1	95° C	5 min	
	2	95° C	20 sec	
	3	60° C	20 sec	
	4	72° C	20 sec	35 repeats to 2
	5	72° C	5 min	
	6	12° C	Hold	