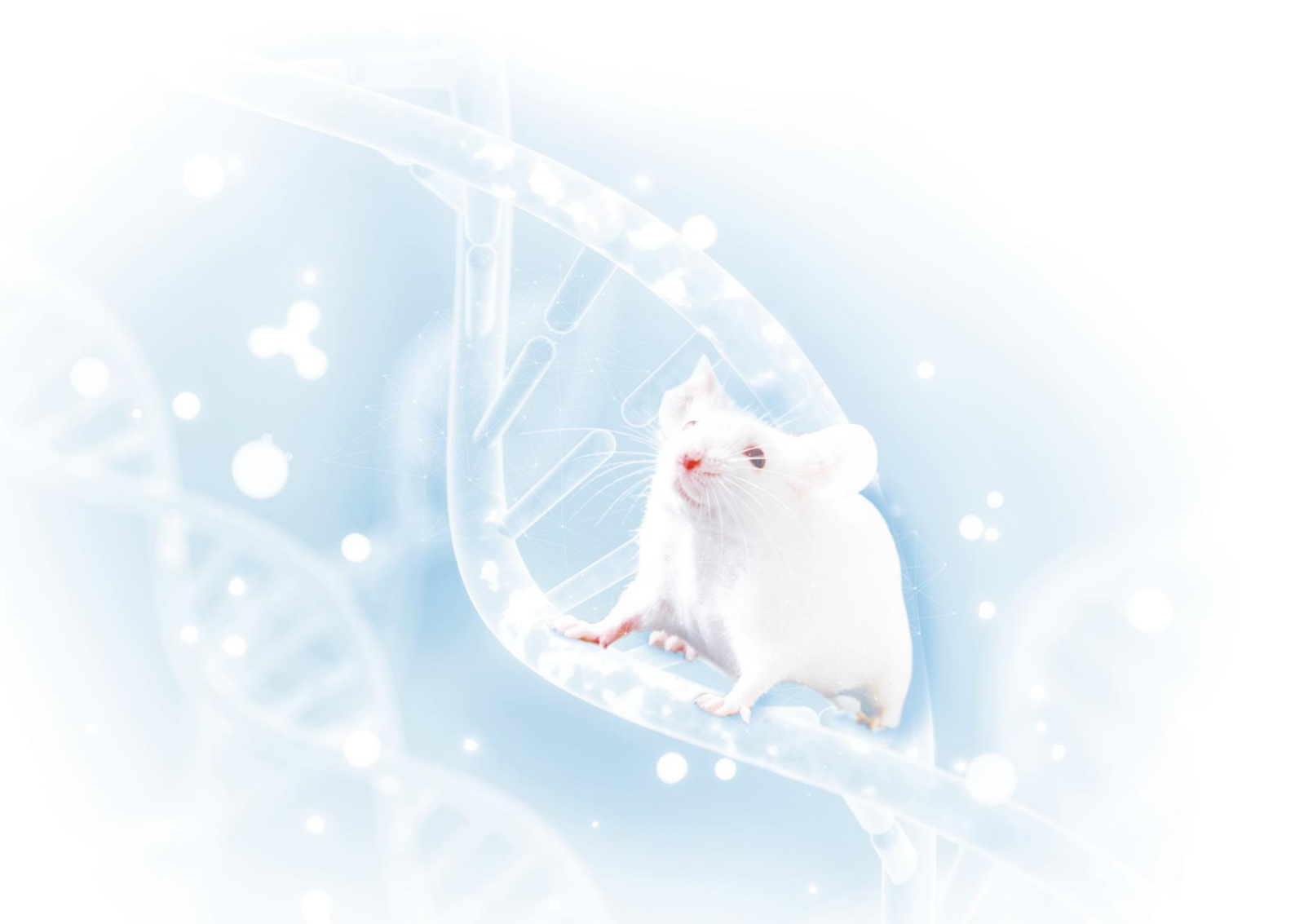


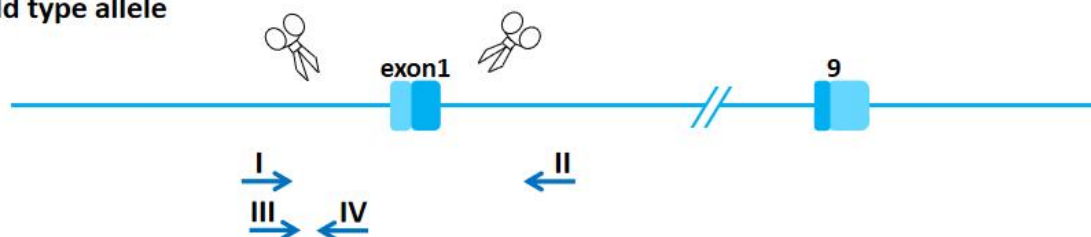
Trim29-KO Genotyping Protocol



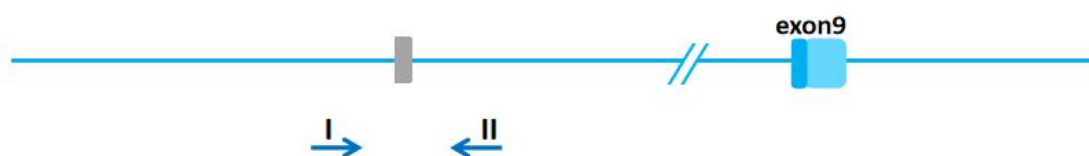
Common Name	Trim29-KO	Cat. NO.	NM-KO-241983
Strain of Origin	C57BL/6J	Version	V1

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	CCCAAAGGCTTTCCTCCCAT	Forward
P2	GCTCAGGAAGGGACAAGACC	Reverse
P3	CTTTCCTCCCTGCTTCTCC	Forward
P4	ATGAGGCTTCTGGGTGTTGGG	Reverse

Expected results

Results	
Genotype	Knockout type: -3909 bp Wild type: P1P2 =4341 bp; P3P4 =392 bp Heterozygote: P1P2 =4341 bp and 432 bp; P3P4 =392 bp Homozygote: P1P2 =432 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

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	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	