

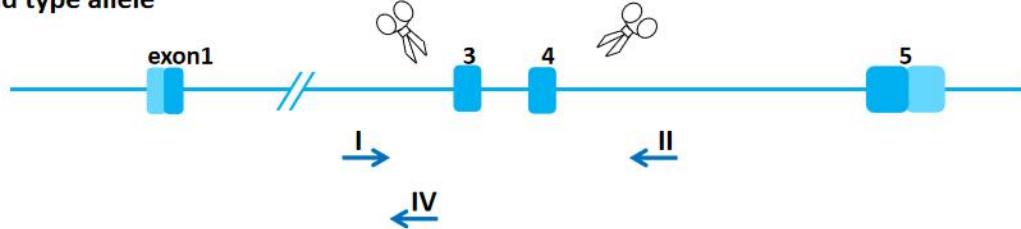
Srd5a2-KO Genotyping Protocol



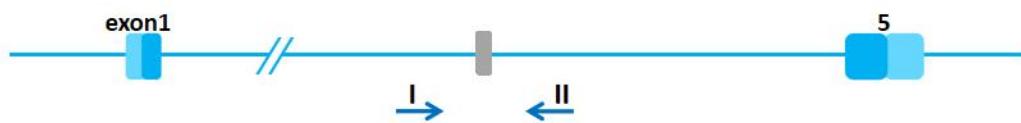
Common Name	Srd5a2-KO	Cat. NO.	NM-KO-233734
Strain of Origin	C57BL/6J	Version	V1

Genotyping strategy

Wild type allele



knockout allele



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

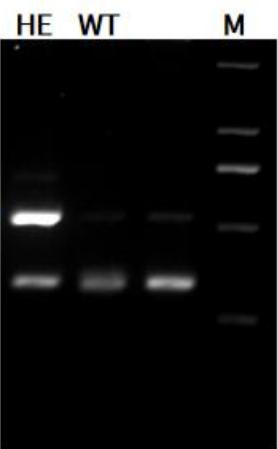
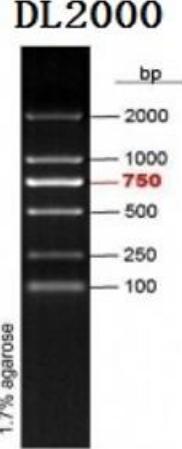
NHEJ : Non-homologous end joining

→ : primer location

Primers

Primer	Sequence (5' → 3')	Primer type
P1	TGCCTATTGGCTCACTACTCCTG	Forward
P2	CTCATCTGCTTCCTTGCTTGGT	Reverse
P4	GGGGTTATTTGTTGCTTGTTCGTC	Reverse

Expected results

	P1P2P4  DL2000 
Results	<p>Knockout type: -3676 bp</p> <p>Wild type: P1P4=327 bp</p> <p>Heterozygote: P1P2 =522 bp; P1P4=327 bp</p> <p>Homozygote: P1P2 =522 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 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