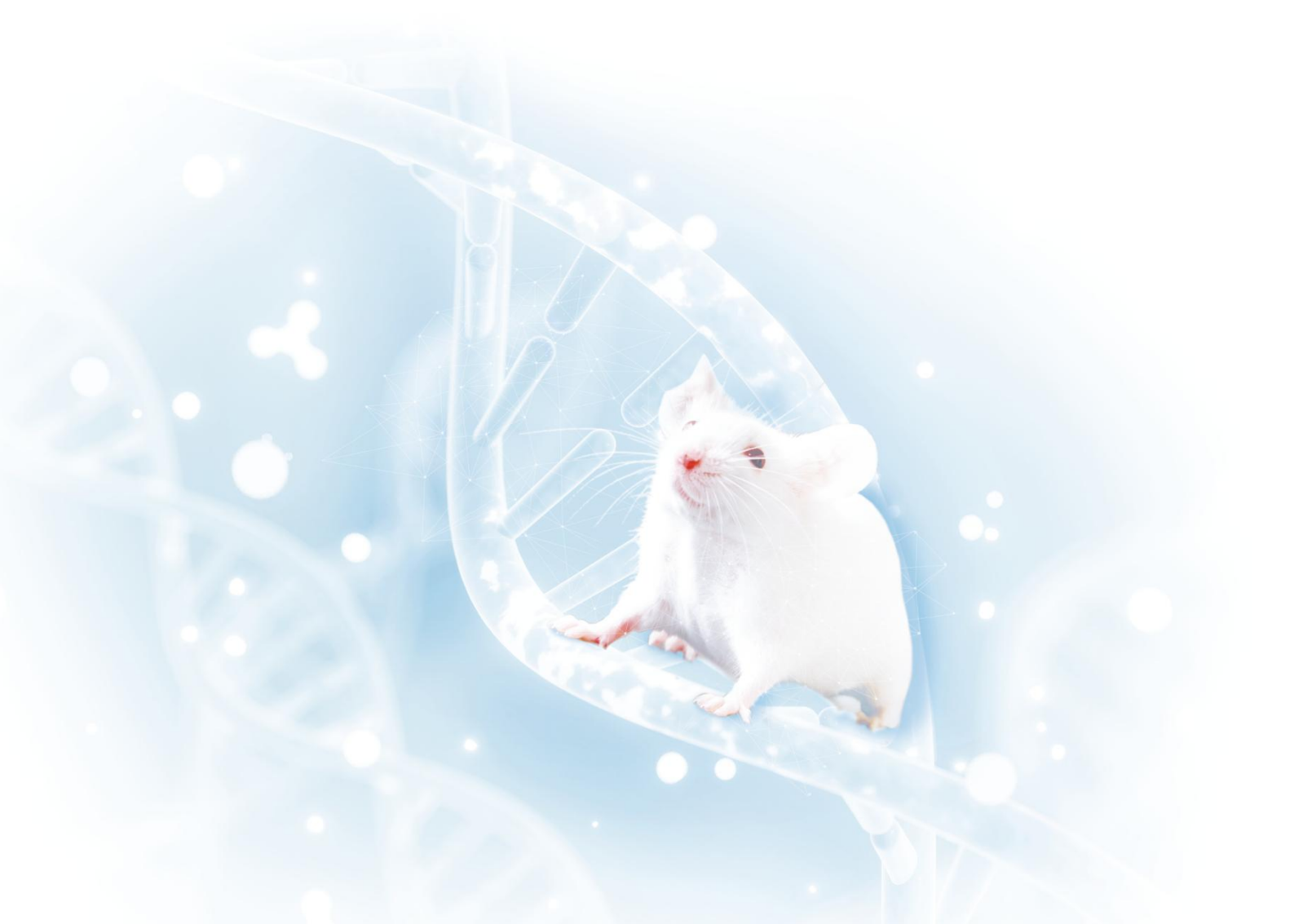


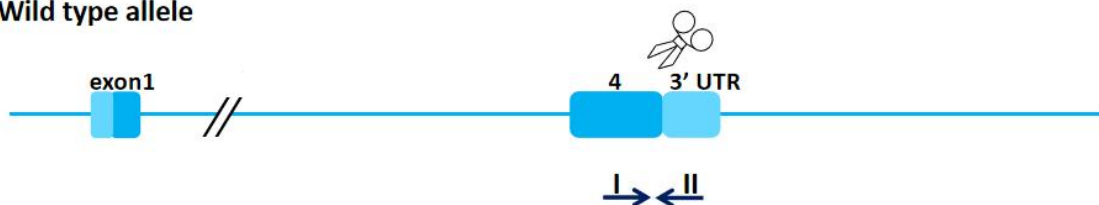
Tubb3-2A-CreERT2-2A-tdTomato Genotyping Protocol



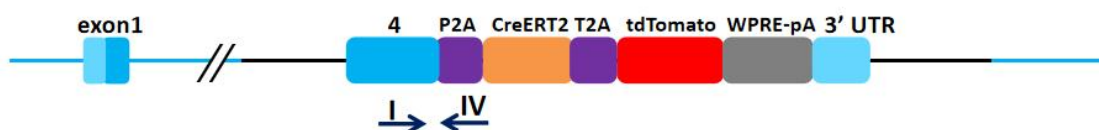
Common Name	<i>Tubb3-2A-CreERT2-2A-tdTomato</i>	Cat. NO.	NM-KI-240472
Strain of Origin	C57BL/6J	Version	V1

Genotyping strategy

Wild type allele



Targeted allele

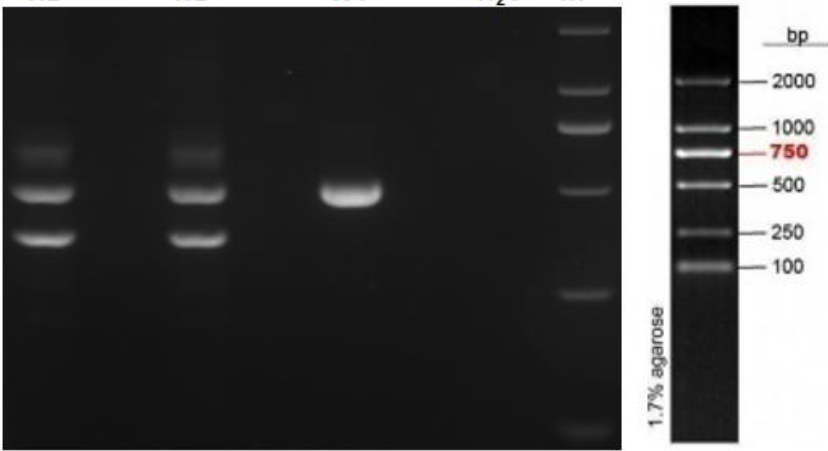


→ : primer location

Primers

Primer	Sequence (5'→3')	Primer type
P1	TCAAGGTAGCCGTGTGTGAC	Forward
P2	GTGGACTCACATGGAGTGGG	Reverse
P4	TAGGACCAGGTTCTCCTCC	Reverse

Expected results

Results	P1+P2+P4  490bp 372bp DL2000 bp 2000 1000 750 500 250 100 1.7% agarose
Genotype	Wild type: P1P2 =490 bp Heterozygote: P1P2 =490 bp; P1P4 =372 bp Homozygote: P1P4 =372 bp

Reaction &Cycling

PCR Reaction System	Reaction Component	Volume (μl)
	ddH ₂ O	7.5
	2×Rapid Taq Master Mix	10.0
	P1(10 pmol/μl)	0.5
	P2(10 pmol/μl)	0.5
	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	