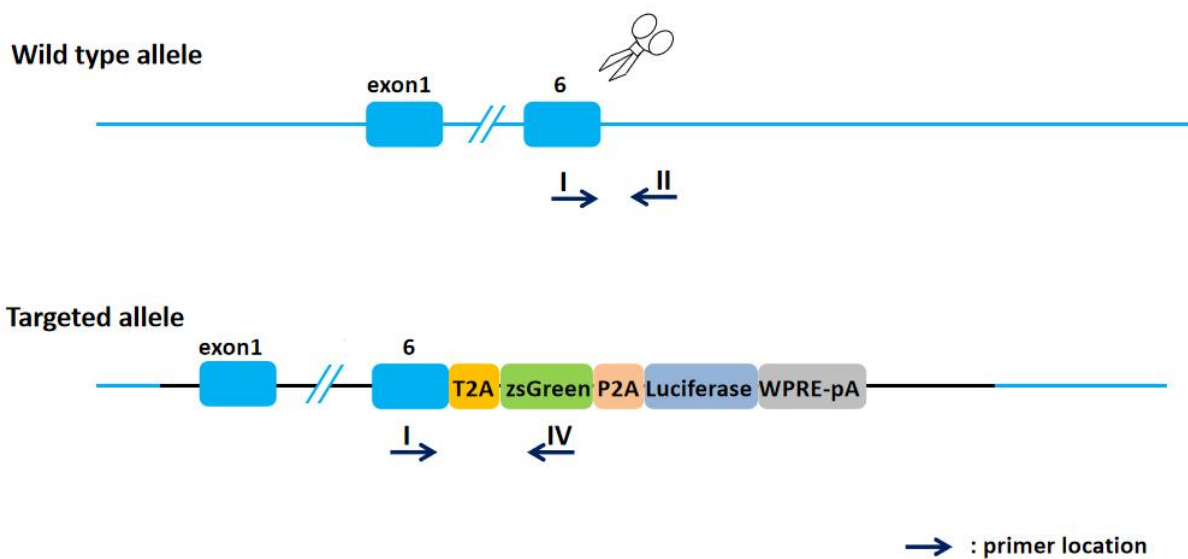


Cd68-2A-zsGreen-2A-Luc Genotyping Protocol



Common Name	<i>Cd68-2A-zsGreen-2 A-Luc</i>	Cat. NO.	<i>NM-KI-240468</i>
Strain of Origin	<i>C57BL/6J</i>	Version	<i>V1</i>

Genotyping strategy



Primers

Primer	Sequence (5'→3')	Primer type
P1	TGCCTCTCATCATTGGCCTG	Forward
P2	TCTTGCCAGCTGTCTCTTG	Reverse
P4	GACAAGATGTCCTCGGCGAA	Reverse

Expected results

Results	<p style="text-align: center;">P1+P2+P4</p> <p style="text-align: center;">HE HE WT H₂O M DL2000</p> <p style="text-align: center;">410bp 346bp</p> <p style="text-align: right;">bp — 2000 — 1000 — 750 — 500 — 250 — 100</p> <p style="text-align: right;">1.7% agarose</p>
Genotype	<p>Wild type: P1P2 =410 bp</p> <p>Heterozygote: P1P2 =410 bp; P1P4=346 bp</p> <p>Homozygote: P1P4 =346 bp</p>

Reaction & Cycling

	Reaction Component	Volume (μ l)
PCR Reaction System	ddH ₂ O	7.5
	2xRapid Taq 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

<i>2xRapid Taq Master Mix from Vazyme(Code Number: P222-01)</i>				
	<i>Step</i>	<i>Temp</i>	<i>Time</i>	<i>Note</i>
<i>Cycling Reaction</i>	<i>1</i>	<i>95°C</i>	<i>5 min</i>	
	<i>2</i>	<i>95°C</i>	<i>20 sec</i>	
	<i>3</i>	<i>60°C</i>	<i>20 sec</i>	
	<i>4</i>	<i>72°C</i>	<i>20 sec</i>	<i>35 repeats to 2</i>
	<i>5</i>	<i>72°C</i>	<i>5 min</i>	
	<i>6</i>	<i>12°C</i>	<i>Hold</i>	