

# Genotyping protocol

## General information:

Strain name	B6J.B6N(Cg)-Htr3a<tm1c(EUCOMM)Wtsi>/Cnbc
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## Primers:

Name	Sequence	Primer type
Htr3a-5'_ALT_Fw	GGAGCAAAGCAAACGTCAGA	
Htr3a-3'_ALT_Rv	CACTGTTATCGAGGCTGTGC	

In case more than two primers are introduced, please indicate how they should be combined:

	Forward primer	Reverse primer
Wt / Mutant Floxed	Htr3a-5'_ALT_Fw	Htr3a-3'_ALT_Rv

## Reaction mix:

ddH <sub>2</sub> O	18,5	μl
PCR Buffer 15mM MgCl <sub>2</sub>	2,5	μl
dNTPs 2,5mM	2,5	μl
Primer 100μM	0,125	μl
Primer 100μM	0,125	μl
Taq Polymerase 5U/μl	0,250	μl
DNA	1	μl
Final volume	25	μl

## PCR program:

95	°C	5	min	X35
95	°C	30	sec	
63	°C	30	sec	
72	°C	60	sec	
72	°C	10	min	

## Expected fragment size:

Wt	248	bp
Mutant Floxed	453	bp

## Comments/Additonal information:

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