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Ncaph (MCEE; EPD0156_5_B09)

Allele: Ncaphtm1a(EUCOMM)Wtsi

Genotyping Information

These mice may be genotyped through a combination of separate PCR reactions that detect *LacZ*, the gene-specific wild type allele, and a mutant allele-specific short range PCR. Interpretation of the consolidated results produces the genotype of the mice.

For example: LacZ positive, mutant positive, wild type positive = heterozygous.

PCRs primer pairs and expected size bands

PCR type	Forward primer	Reverse primer	Expected size band (bp)
Mutant PCR	Ncaph_48020_F	CAS_R1_Term x	216
Wild type PCR	Ncaph_48020_F	Ncaph_48020_R	400
LacZ PCR	LacZ_2_small_F	LacZ_2_small_R	108

> Primer sequences

Primer name	Primer sequence (5' > 3')		
CAS_R1_Term	TCGTGGTATCGTTATGCGCC		
Ncaph_48020_F	TGTACGACACCAGGTCCAGC		
Ncaph_48020_R	CACTCCCCTTTACAGCAGGG		
LacZ_2_small_F	ATCACGACGCGCTGTATC		
LacZ_2_small_R	ACATCGGGCAAATAATATCG		

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Reaction

Reagent	μΙ
DNA (~50-100 ng)	1.0
10x Buffer	2.0
MgCl ₂ (50 mM)	0.6
PtTaq (Platinum Taq (Invitrogen))	0.2
dNTPs (100 mM)	0.2
Primer 1 (10 μM)	0.4
Primer 2 (10 μM)	0.4
H_20	<u>15.2</u>
Total	20.0

Cycling conditions

Wild type and mutant PCRs

Go to '2' + 34 cycles

5 min

forever

72 °C

12 °C

Cycle			Cycle		
1	94 °C	5 min	1	94 °C	5 min
2	94 °C	30 sec	2	94 °C	30 sec
3	58 °C	30 sec	3	60 °C	30 sec
4	72 °C	45 sec	4	72 °C	30 sec

LacZ PCR

Go to '2' + 34 cycles

5 min

forever

72 °C

12 °C

5

6

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