

Genotyping Protocol: MMRRC 11588

Strain Characteristics: Acadm (acetyl-Coenzyme A dehydrogenase, medium chain) knockout

Assay Type: PCR to confirm presence of Neomycin gene. Can only distinguish positive from negative animals. Cannot distinguish heterozygous from homozygous mutant animals.

Mice have a knockin of redundant exons 8-10 inverted between exon 9 and 10 of the wild type gene with a neomycin gene in between redundant exons 10 and 8 (see figure below). Details can be found in Tolwani et al (2005) PLoS Genet. 1(2): e23.



DNA Extraction: DNA from tail snips was extracted using Qiagen's DNeasy Blood and Tissue kit (Cat# 69506). Kit directions for animal tissues were performed with a few minor modifications as follows: repeat AW1 and AW2 wash steps one time, elute in 200µl of AE buffer once.

Primer Information:

- | | |
|----------------|---|
| 1) Name: Neo F | Sequence: 5'-CATTTCGACCACCAAGCGAAACATC-3' |
| 2) Name: Neo R | Sequence: 5'-ATATCACGGGTAGCCAACGCTATG-3' |

Assay Name: Neomycin PCR

PCR Master Mix Components:

| Component | Manufacturer | Concentration | µl/rxn |
|---|--------------------------|---------------|--------|
| Buffer with MgCl ₂ (green cap) | Roche | 10x | 2 |
| dNTPs | Promega (Cat# U1515) | 1.25 mM | 3.2 |
| Neo F | Sigma | 25 µM | 0.3 |
| Neo R | Sigma | 25 µM | 0.3 |
| FastStart Taq | Roche (Cat# 12032953001) | 5 U/µl | 0.2 |
| Sterile Water | | | 13 |

PCR Setup:

Final Reaction: 19 µl master mix & 1µl DNA template (10-20ng/µl of genomic DNA)

All reactions were performed in 200µl thin walled PCR tubes and were run in an Applied Biosystems 2700 thermocycler.

Cycle Parameters:

- | | | |
|----|--|------------|
| 1) | 95°C | 5 minutes |
| 2) | 94°C | 30 seconds |
| 3) | 68°C | 30 seconds |
| 4) | 72°C | 1 minute |
| 5) | Repeat steps 2-3 34 times for a total of 35 cycles | |
| 6) | 72°C | 10 minutes |
| 7) | 4°C | hold |

Product Analysis:

All products were analyzed on the Qiaxcel (instrument and all supplies from Qiagen) with the Qiaxcel DNA Screening Kit (Cat# 929004).

Alignment Marker: QX Alignment Marker 15bp/3Kb (Cat# 929522)

Size Marker: QX DNA Size Marker 100-3Kb (Cat# 929553)

Method: AM320 Injection: 10s at 5KV
Separation: 320s at 6KV

09.10.10 MS
08.07.13 MLS

PCR Setup:

Final Reaction: 48µl master mix & 2µl RNA template

All reactions were performed in 200µl thin walled PCR tubes and were run in Perkin Elmer 2400 thermocycler or Applied Biosystems 2700 thermocycler.

Cycle Parameters:

- | | | |
|----|--|--------------------------------|
| 1) | 50°C | 30 minutes |
| 2) | 95°C | 15 minutes |
| 3) | 94°C | 30 seconds |
| 4) | 64°C | 1 minute |
| 5) | 72°C | 2 minute |
| 6) | Repeat steps 3-5 39 times for a total of 40 cycles | |
| 7) | 72°C | 10 minutes |
| 8) | 4°C | hold until refrigerate product |

Product Analysis:

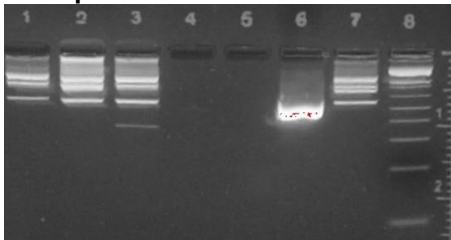
All products were analyzed on a 3% agarose gel with ethidium bromide staining.

Wild type mRNA product: 550 bp

Knockout gene product: ladder effect: weak or absent band at 550 bp, strong bands at 700, 800, 1000-bp and larger, easily detected in homozygous animals.

A mixture of 50% wildtype RNA with 50% homozygous RNA results in only amplification of the wild type transcript (550bp product) and this has been confirmed in known heterozygous animals (see Case #13316). Because of this, the assay cannot be used to distinguish between wild type and heterozygous animals.

Example of Gel:



Lanes 1-3 display homozygous samples (ladder of products: 700bp, 800bp, 1000bp bands).
Lanes 4 and 5 are extraction and PCR blanks, respectively.
Lane 6 displays a WT control (550bp band).
Lane 7 displays a homozygous control (ladder of products).
Lane 8 displays 1Kb+ Ladder size marker (Invitrogen Cat# 10787-018).

Interpretation if both assays are used:

Wild type: negative for neo and 550 bp RT-PCR product

Heterozygous: positive for neo and 550 bp RT-PCR product

Homozygous mutant: positive for neo and ladder effect in RT-PCR assay