# 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.



# Neomvcin PCR

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'-CATTCGACCACCAAGCGAAACATC-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 <sub>2</sub> (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:**

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

### Expected products:

Positive: 289 bp Negative: no product

#### Example gel:



Lane A1 (highlighted in blue) displays a 15bp-3kb size marker. Lane A1 (unhighlighted) displays a sample negative for the neomycin gene (no product). Lane A2 displays a sample positive for the neomycin gene (289bp product).

\*Please note: the 15bp and 3kb bands are reference markers specific to the Qiaxcel method and do not represent expected products.\*

# MMRRC 11588 rtPCR: (alternative assay)

**Assay Type**: RT-PCR to detect wild type transcript. Wild type and heterozygous animals produce a single 550 bp product and therefore this assay cannot distinguish between wild type and heterozygous animals.

Because of the redundant exons, numerous RT-PCR products appear to be formed in homozygous mutant animals giving a specific pattern of products resulting in a ladder effect with product sizes of ~700 bp, 800bp, 1000 bp, and larger. This array of products is not seen in wild type or heterozygous animals.

**RNA Extraction**: RNA from ear punches or liver was extracted using Qiagen's RNeasy Mini kit (Cat# 74104). Kit directions for animal tissues were performed following kit directions with a single elution of 40µl.

### Primer Information:

1) Name: M11588 Forward	Sequence: 5'-ATG TGG CGG CCA TTA AGA CCA AAG-3'
2) Name: M11588 Reverse	Sequence: 5'-GCT GAT TGG CAA TGT CTC CAG CAA-3'

Primer location: Forward primer binds to exon 7 and reverse primer binds to exon 11 to produce 550 bp product

### Assay Name: MMRRC 11588 rtPCR

#### RT-PCR Master Mix Components: Master Mix for RT-PCR uses Qiagen OneStep RT-PCR Kit (Cat# 210212):

		1	,
component	manufacturer	concentration	µl/rxn
Rnase free water	Qiagen		21
5X RT-PCR Buffer	Qiagen	5X	10
dNTP mix	Qiagen		2
5X Q-Solution	Qiagen	5X	10
M11588 Forward	IDT	20µM	1.5
M11588 Reverse	IDT	20µM	1.5
One Step Enzyme Mix	Qiagen		2

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)  $50^{\circ}$ C 30 minutes
- 2)  $95^{\circ}$ C 15 minutes
- 3)  $94^{\circ}$ C 30 seconds
- 4)  $64^{\circ}$ 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