

Genotyping Protocol: **MMRRC 36698**

Assay Type: Precision Melt SNP analysis – can distinguish between heterozygous and homozygous samples.

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.

Strain Description: This strain carries a G to A single nucleotide change in Exon 39 of the mouse *Myh9* gene. This results in a glutamic acid to lysine amino acid change at codon 1841 (E1841K). Details can be found in Zhang et al (2012) Blood 119(1): 238–250.

Primer Information:

- 1) Name: M36698 melt F Sequence: 5'- CCC CTT TCA GTC AGC ATT TC-3'
 2) Name: M36698 melt R Sequence: 5'- GCA CGT CCT TCA GCT TCT TC-3'

Primer Location: M36698 melt F and melt R are located on either side of the SNP: M36698 melt F is in Intron 38 and M36698 melt R is in Exon 39 of mouse *Myh9*.

WT Mouse *Myh9* sequence (red letters with tan highlights are Exon 39):

CAGCAGCTACTGAGTGTCCAGGAAGACTAGCACCTTAGCTTAGCTCCAGGGGCCCCAGGAAGCGGAAGCTCCGGTTGCCATCTTGCTGGCTGAGCTA
 TCTGGTGGAGGGTGC **CCCCTTTTCAGTCAGCATTTC** TCTCCTACCCAG **GGAGCGCCAGGCAGCCTCCAAGCAGGTGCGCCGGACG** **AGAAGAAGCTGAAG**
GACGTGCTGCTGCAGGTGGAGGACGAGCGGAGGAACGCGGAACAGTTCAAGGACCAG GTCTGTGCTCTGCAGGGCCTGGGTTCTGTCTGTGCTCAGTAG
 AGCCTTGGGCTGCACTCAGTGTCTCACCAGCGTCCCTGGCCACCTGTTTCTGCAG

M36698 melt F

M36698 melt R

Nucleotide which changes from G to A in mutant allele

Codon 1841

Assay Name: M36698 SNP PCR**Master Mix Components:**

component	manufacturer	concentration	µl/rxn
Precision Melt Supermix	BioRad (Cat# 172-5110)	2X	10
M36698 melt F	Sigma or IDT	2µM	1
M36698 melt R	Sigma or IDT	2µM	1
sterile water			6

PCR Setup:

Final Reaction: 18µl master mix & 2µl DNA template (DNA is diluted to 5ng/µl, for 10ng total for each reaction)

All reactions were performed in either BioRad Hard-Shell PCR Plates (Cat#HSP9601) with Optical Tape (Cat#2239444) covers or Life Technologies' Optical 8-tube strips (Cat#4316567) with Optical Caps (Cat#4323032). All were run in a BioRad CFX96 RealTime System utilizing BioRad CFX Manager software.

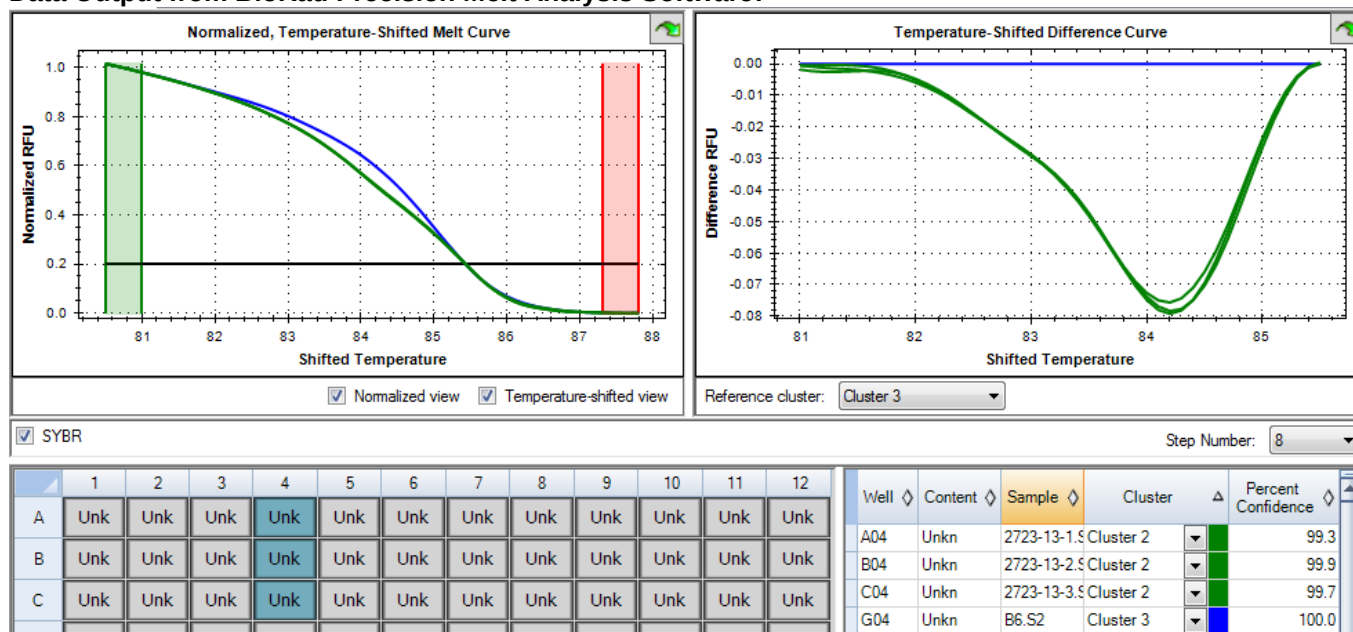
Cycle Parameters:

- 1) 95°C 2 minutes
- 2) 95°C 10 seconds
- 3) 60°C 30 seconds (+plate read)
- 4) 72°C 30 seconds
- 5) Repeat steps 2-4 39 times for a total of 40 cycles
- 6) 95°C 30 seconds
- 7) 60°C 1 minute
- 8) 65-95°C (in 0.2°C increments) 10 seconds/step (+plate read)

Product Analysis:

Results were analyzed with BioRad Precision Melt Analysis Software.

If products are run on a gel, a 91bp product can be expected.

Expected Results:**Data Output from BioRad Precision Melt Analysis Software:**

In this example, Cluster 2 (green) are heterozygous samples and Cluster 3 (blue) is a WT control.

For Genotype Confirmation via Sequencing (Optional):**Primer Information:**

1) Name: M36698 551 F

Sequence: 5'-TTG CCA TCT TGC TGC TGA GCT ATC TG-3'

2) Name: M36698 552 R

Sequence: 5'-ACC TGC AGC AGC ACT TCA GCT TCT TC-3'

Primer location: Primers are located on either side of the G to A single nucleotide change in Exon 39 of *Myh9*.

PCR Master Mix Components:

component	manufacturer	concentration	µl/rxn
Buffer with MgCl ₂ (green cap)	Roche	10X	2
dNTPs	Promega (Cat# U1515)	1.25mM	3.2
M36698 551 F	Sigma or IDT	25µM	0.3
M36698 552 R	Sigma or IDT	25µM	0.3
FastStart Tag	Roche (Cat# 12032953001)	5 U/µl	0.2
sterile water			13

PCR Setup:

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

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) | 95°C | 3 minutes |
| 2) | 94°C | 20 seconds |
| 3) | 66°C | 25 seconds |
| 4) | 72°C | 30 seconds |
| 5) | Repeat steps 2-4 34 times for a total of 35 cycles | |
| 6) | 72°C | 10 minutes |
| 7) | 4°C | hold until refrigerate product |

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)

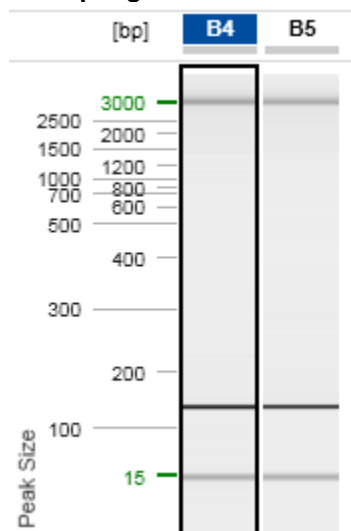
Size Marker: QX DNA Size Marker 100-3Kb (Cat# 929553)

Method: AM320 Injection: 10s at 5KV

Separation: 320s at 6KV

Expected product: 138bp

Example gel:



Lanes B4 and B5 display the 138bp product from the sequencing PCR (M36698 551 F and M36698 552 R). All samples will produce this 138bp product.

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

Sequencing: To confirm that the correct point mutation is present and zygosity.

PCR product is purified with the QIAquick PCR Purification Kit (Qiagen Cat# 28104). Kit directions are performed, and DNA is eluted with 50µl nuclease-free water. 15µl of the DNA is then added to 1µl M36698 551 F primer (25µM) in a 1.5ml tube. The tube is submitted to the sequencing core.

Analysis: In mutant allele, G changes to A

Example Chromatograms:

Green: A

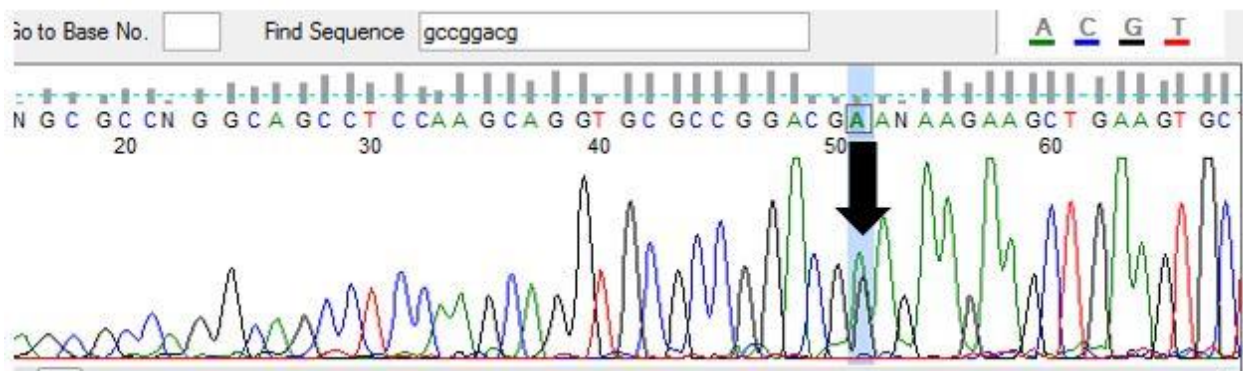
Red: T

Blue: C

Black: G

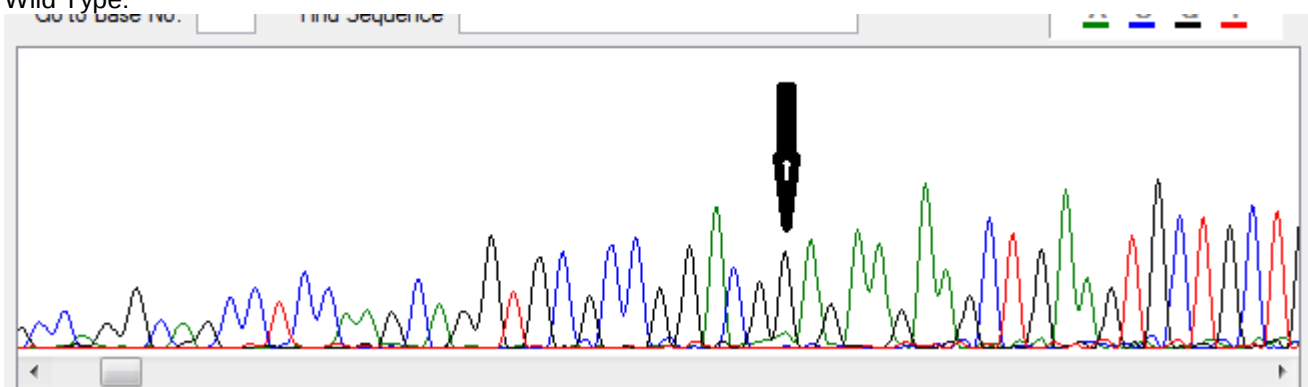
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Heterozygous Mutant:



GCCGGACG A/G

Wild Type:



GCCGGACG G