Genotyping Protocol: MMRRC 16831

Assay Type: PCR (can distinguish heterozygous animals from homozygous animals)

DNA Extraction: DNA from tail snips was extracted using Sigma's Extract-N-Amp Tissue PCR Kit (Cat#XNAT2R). Kit directions for fresh or frozen tails were performed with a few minor modifications as follows: use 50 μ l of Extraction Solution and 12.5 μ l of Tissue Preparation Solution and 50 μ l of Neutralization Solution B.

Primer Information:

Name: XP26 Sequence: 5'- GTG TCA GGC ATA AGA TCT ATG ACA A -3'

Name: XP47 Sequence: 5'- AGG CAA GCA CCT GCA GCT GT -3'
Name: PGK2 Sequence: 5'- GGC CAC TTG TGT AGC GCC AA -3'

XP26 and Xp47 are located in xeroderma pigmentosum, complementation group A (*Xpa*). Xp26 is the forward primer. PGK2 is located in the PGK promoter. PGK2 is a reverse primer.

Assay Name: Xpa KO PCR

PCR Master Mix Components:

KO (Mutant) Master Mix

component	manufacturer	concentration	μl/rxn
Extract-N-Amp PCR			
Reaction Mix	Sigma (Cat#XNAT2R)	2X	10
XP26	Sigma or IDT	25 μM	0.3
PGK2	Sigma or IDT	25 μM	0.3
sterile water			5.4

Wild-type Master Mix

Tha type maeter mix				
component	manufacturer	concentration	μl/rxn	
Extract-N-Amp PCR				
Reaction Mix	Sigma (Cat#XNAT2R)	2X	10	
XP26	Sigma or IDT	25 µM	0.3	
XP47	Sigma or IDT	25 µM	0.3	
sterile water			5.4	

PCR Setup:

Final Reaction: 16µl master mix & 4µl 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) 94°C 3 minutes 2) 94°C 1 minute 3) 61°C 1 minute 4) 72°C 1 minute

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)

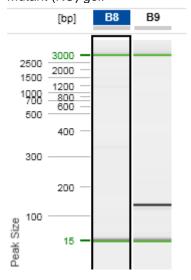
Method: AM320 Injection: 10s at 5KV Separation: 320s at 6KV

Expected products:

Wild-type PCR: 214bp Mutant (KO) PCR: 132bp

Example of Gels:

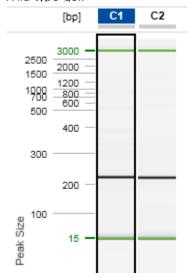
Mutant (KO) gel:



Lane B8 displays a sample negative for the KO allele (no product)
Lane B9 displays a sample positive for the KO allele (132bp product)

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

Wild-type gel:



Lanes C1 and C2 display samples positive for the WT allele (214bp product)

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

Genotype Interpretation:

Conotype interpretation.				
Genotype:	Wild-type PCR	KO (Mutant) PCR)		
Wild-type	214bp product	No product		
Heterozygous	214bp product	132bp product		
Homozygous mutant	No product	132bp product		