

**GENOTYPING BY PCR PROTOCOL
MUTANT MOUSE REGIONAL RESOURCE CENTER: UC DAVIS**

mmrrc@ucdavis.edu
530-754-MMRRC

Please provide the following information required for genetic analysis of your mutant mice.

Note to MAC users: to ensure your graphic can be viewed on a PC please follow the steps below when inserting the graphic into this document. **DO NOT** drag and drop or copy/paste the graphic into this document.

- Open the original graphic in the program that created it
- Choose File, Save As
- Select No Compression in the save options.
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- Switch to Word, choose Insert, Picture, From File and choose the newly saved picture.

These instructions are very generic. The menu options for your graphics program may be different.

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Strain Name		MMRRC Stock Number
B6.Cg-Aldh1a2tm1Jln/Mmucd		041423

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NAME OF PCR: Raldh2 Floxed Genotyping MMRRC: 041423

Protocol: *(PCR protocol provided by Donating Investigator)*

Reagent/Constituent	Volume (µL)
Water	14.80
10x Buffer	2.00
MgCl ₂ (stock concentration is 50 mM)	0.60
Betaine (stock concentration is 5M) <i>Optional</i>	0
dNTPs (stock concentration is 10mM)	0.40
DMSO <i>Optional</i>	0
Primer 1. (stock concentration is 20µM)	0.40
Primer 2. (stock concentration is 20µM)	0.40
Primer 3. (stock concentration is 20µM)	
Primer 4. (stock concentration is 20µM)	
Taq Polymerase 5Units/µL	0.20
DNA (50-200ng/ µL) extracted w/ "Qiagen DNeasy columns or other similar silica based kits"	1.2
<i>The total volume is auto-calculated based on volumes entered, right click the total and update field to show/recalculate the total volume.</i>	TOTAL VOLUME OF REACTION: 20.0 µL

Comments on protocol:

- Cut off ~2mm of tail specimen into 1.5mL capped tube; add 500µL of 0.05M NaOH solution into tube.
- Place on heat block at 95°C for 20min. Add 50µL of 1M TRIS/EDTA solution. Vortex till disintegration.

Strategy:

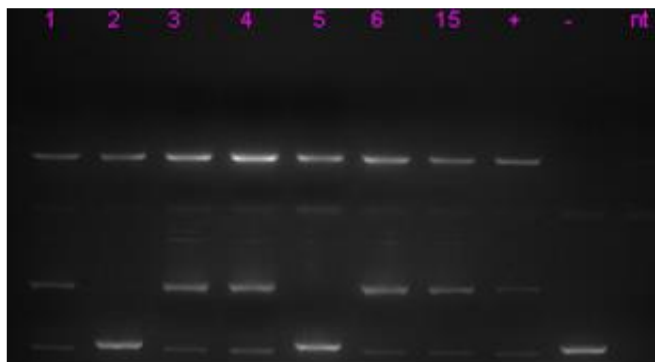
Steps	Temp (°C)	Time (m:ss)	# of Cycles
1. Initiation/Melting HOT START? <input type="checkbox"/>	94	5:00	1
2. Denaturation	94	0:15	
3. Annealing steps 2-3-4 cycle in sequence	65	0:30	36x
4. Elongation	72	0:40	
5. Amplification	72	5:00	1
6. Finish	4.0	∞	n/a

Primers:

Name	Nucleotide Sequence (5' - 3')	Argarose: 2.5% V: 125 volts		
1. Common Rev	TGAGCTGGCTAAAGGCATTTGTAGTC	Estimated Running:Time: 90 min.		
2. WT Forward	GACGGAGGACAGAGCCAACTTACTC	Primer Combination	Band (bp)	Genotype
3. Flox Forward	GGAATGTGGGACTCTGCCAGAAG	1&2	294	WT
4. Del Forward	CGACTCACTATAGGGCGAATTGGG	1&3	419	Floxed
5.		1&4	239	Deletion
6.				
7.				

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1 Protocol / Gel Comments:

Mouse 1 is +/del; Mouse 2 and 5 are +/fl; Mouse 3, 4, 6 and 15 are +/+. + and – are WT and fl/fl controls.

Note: Flox primers don't identify deletion and vice versa.

WT mice using primers 1 and 3 yield a non-specific band.