

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

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 530-754-MMRRC

NAME OF PCR: STOCK Tg(Aldh1l1-EGFP)OFC789Gsat/Mmucl MMRRC # 011015-UCD

Protocol:

Reagent/ Constituent	Volume (μ L)
Water	11.275
10x Buffer	2.5
MgCl ₂ (stock concentration is 25mM)	1.7
Betaine (stock concentration is 5M)	6.5
dNTPs (stock concentration is 10mM)	0.5
DMSO	0.325
Primer 1 (stock concentration is 20 μ M)	0.5
Primer 2 (stock concentration is 20 μ M)	0.5
Taq Polymerase (5Units/ μ L)	0.2
DNA extracted with <input type="checkbox"/> NaOH <input checked="" type="checkbox"/> Proteinase K <input type="checkbox"/> Other:	1.0
TOTAL VOLUME OF REACTION:	25μL

Comments on protocol:

- Use Touch-Down cycling protocol-first 10 cycles anneal at 65° C decreasing in temperature by 1.0 ° C; next 30 cycles anneal at 55° C.
- Betaine/DMSO is standardized due to high GC content in promoter regions and protocol may be tested without. Also, may adjust MgCl₂ to increase reaction or decrease non specific amplifications.

Strategy:

Steps	HOT START? <input type="checkbox"/>	Temp (°C)	Time (m:ss)	# of Cycles
1. Initiation/Melting		94	5:00	1
2. Denaturation		94	0:15	
3. Annealing	} steps 2-3-4 will cycle in sequence	65 to 55 ($\downarrow 1^{\circ}\text{C}/\text{cycle}$)	0:30	} 40x
4. Elongation		72	0:40	
5. Amplification		72	5:00	1
6. Finish		4	Hold	n/a

Primers:

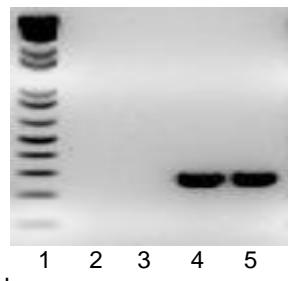
Name	Nucleotide Sequence (5' - 3')
1: Fthfd (11015) F1	CCTCTGGCTGCTCCTAACAG
2: GS eGFP R3	GGTCGGGGTAGCGGCTGAA

Electrophoresis Protocol:

% Agarose: 1.5 V: 90

Estimated Running Time (min): 90

Primer Combination	Band	Genotype
1 and 2	300 bp	transgenic



1: 1kb+ ladder (Invitrogen, Cat. #10787-026)
 2: non-template control
 3: wild-type & eGFP
 4-5: Fthfd+