

GENOTYPING BY PCR PROTOCOL
MUTANT MOUSE REGIONAL RESOURCE CENTER: UC DAVIS
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 530-754-MMRRC

NAME OF PCR: STOCK Tg(Crym-EGFP)GF82Gsat/Mmcd **MMRRC #** 012003-UCD

Protocol:

Reagent/ Constituent	Volume (µL)
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 20µM)	0.5
Primer 2 (stock concentration is 20µM)	0.5
Taq Polymerase (5Units/µL)	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. Also, may adjust MgCl₂ to increase reaction or decrease non specific amplifications.

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	} 40x
3. Annealing } steps 2-3-4 will cycle in sequence	65 to 55 (↓1°C/cycle)	0:30	
4. Elongation	72	0:40	
5. Amplification	72	5:00	1
6. Finish	4	Hold	n/a

Primers:

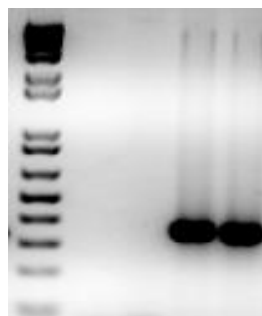
Name	Nucleotide Sequence (5' - 3')
1. Crym (012003)	GCTACTCAGGCAGTCCGCTCAT
2. GS eGFP R3	GGTCGGGGTAGCGGCTGAA

Electrophoresis Protocol:

Agarose: 1.5% **V:** 90

Estimated Running Time: 90 min.

Primer Combination	Band	Genotype
1 and 2	350 bp	transgenic



1 2 3 4 5

- Lanes
- 1 kb+ ladder (Invitrogen, Cat. #10787-026)
 - Non-template control
 - Wild-type & eGFP
 - 4-5. Crym tg/+