

GENOTYPING BY PCR PROTOCOL
MUTANT MOUSE REGIONAL RESOURCE CENTER: UC DAVIS
 2795 2nd Street, Suite 400, Davis, CA 95618
mmrrc@ucdavis.edu
 530-754-MMRRC

NAME OF PCR: STOCK Tg(4930572J05Rik-EGFP)MD16Gsat /Mmucd MMRRC # 031169-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. 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 (↓1°C/cycle)	0:30	} 40x
4. Elongation		72	0:40	
5. Amplification		72	5:00	1
6. Finish		4	Hold	n/a

Primers:

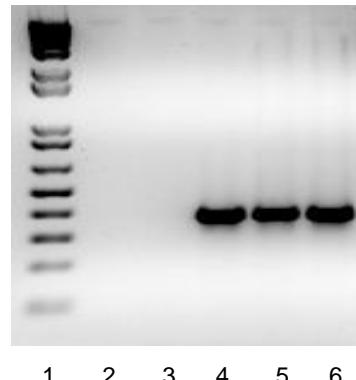
Name	Nucleotide Sequence (5' - 3')
1. 493052J05Rik (31169) F	CCCCTCGGACTGCTGGAGAG
2. GS eGFP R3	GGTCGGGGTAGCGGCTGAA

Electrophoresis Protocol:

Agarose: 1.5% V: 90

Estimated Running Time: 90 min.

Primer Combination	Band	Genotype
1 and 2	375 bp	transgenic
Tg copy # ~ 12 copies/genome		



- Lanes**
1. 1 kb+ ladder (Invitrogen, Cat. #10787-026)
 2. Non-template control
 3. Wild-type Control
 - 4, 5 & 6. 493052J05Rik tg/+