

GENOTYPING BY PCR PROTOCOL
MUTANT MOUSE RESOURCE & RESEARCH CENTER: UC DAVIS

mmrrc@ucdavis.edu

530-754-MMRRC

NAME OF PCR: STOCK Tg(Syt6-cre)KI109Gsat/Mmucl MMRRC # 036088-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	∞	n/a

Primers:

Name	Nucleotide Sequence (5' - 3')
1: Syt6 (36088) F	GAGGGTCTTCTGGGGCCTT
2: CreGS R2	CCCCAGAAATGCCAGATTACGTAT

Electrophoresis Protocol:

Agarose: 1.5% V: 90

Estimated Running Time: 90 min.

Primer Combination	Band	Genotype
1 and 2	300 bp	transgenic



Lanes

1. 1 kb+ ladder (Invitrogen, Cat. #10787-026)
2. Non-template control
3. Wild-type & Cre
- 4 & 5. Syt6