

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

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530-754-MMRRC

Protocol Name: C57BL/6-Clec4e^{tm1.1Cfg}/Mmucd

MMRRC: 031936-UCD

Protocol:

Reagent/Constituent	Volume (μ L)
Water	10.775
10x Buffer	2.5
MgCl ₂ (stock concentration is 25mM)	1.7
Betaine (stock concentration is 5M) <i>Optional</i>	6.5
dNTPs (stock concentration is 10mM)	0.5
DMSO <i>Optional</i>	0.325
Primer 1. (stock concentration is 20 μ M)	0.5
Primer 2. (stock concentration is 20 μ M)	0.5
Primer 3. (stock concentration is 20 μ M)	0.5
Taq Polymerase 5Units/ μ L	0.2
DNA (example) extracted w/ "Qiagen DNeasy columns or other similar silica based kits"	1.0
TOTAL VOLUME OF REACTION:	25.000 μL

Comments on protocol:

- Protocol may work with other DNA extraction methods.
- 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 and DMSO have been standardized due to high GC content. 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 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		15	∞	n/a

Primers:

Electrophoresis Protocol:

Name	Nucleotide Sequence (5' - 3')	Agarose: 1.5%	V: 90
1. 31936-P1	ATTGCCACTGACCCTCCACC	Estimated	9 min.
2. 31936-P2	CCCCTGTCACTGTTCTTGCA	Primer	Band (bp)
3. 31936-P3 (mutant)	TGCAGCCCAAGCTGATCCTC	1, 2 & 3	488/593
		1 & 2	593
			wildtype

