

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

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NAME OF PCR: B6.129S4-Dnmt1<sup>tm2Jae</sup>/Mmucd MMRRC: 014114-UCD

**Protocol:**

Reagent/Constituent	Volume ( $\mu$ L)
Water	11.275
10x Buffer	2.5
MgCl <sub>2</sub> (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 $\mu$ M)	0.5
Primer 2. (stock concentration is 20 $\mu$ M)	0.5
Taq Polymerase 5Units/ $\mu$ L	0.2
DNA (example) extracted w/ "Qiagen DNeasy columns or other similar silica based kits"	1.0
<b>TOTAL VOLUME OF REACTION:</b>	<b>25.000 <math>\mu</math>L</b>

**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.
  - Note: non specific band overserved ~430bp on Het samples.

**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	$\infty$	n/a

**Primers:**

**Electrophoresis Protocol:**

Name	Nucleotide Sequence (5' - 3')	Agarose: 1.5% V: 90
1. Dnmt1-1	GGGCCAGTTGTGACTTGG	Estimated Running Time: 90 min.
2. Dnmt1-2	CTGGGCCTGGATCTGGGGA	
	Primer Combination	Band (bp)
	1 and 2	334 bp
	1 and 2	368 bp



