

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

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530-754-MMRRC

Protocol Name: B6N.129S5-Layn<sup>tm1Lex</sup>/Mmucd

MMRRC: 032426-UCD

**Protocol:**

Reagent/Constituent	Volume ( $\mu$ L)
Water	9.775
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
Primer 3. (stock concentration is 20 $\mu$ M)	0.5
Primer 4. (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</b>	<b>24.5</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. Also, may adjust MgCl<sub>2</sub> 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	
3. Annealing steps 2-3-4 cycle in sequence	65 to 55 ( $\downarrow 1^{\circ}\text{C}/\text{cycle}$ )	0:30	<b>40x</b>
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. DNA026-15	TAGGCAAACCTCAGAACATCTCCC	Estimated	9 min.
2. GT IRES	CCCTAGGAATGCTCGTCAAGA	Primer	Band (bp)
3. DNA026-5	TTGGCTAACCGAGAGGAGC	1 & 2	572
4. DNA026-6	CAGACTGCCAACGAGAAAGC	3 & 4	271

