

# GENOTYPING BY PCR PROTOCOL

## MUTANT MOUSE RESOURCE & RESEARCH CENTER: UC DAVIS

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530-754-MMRRC

Protocol Name: \_\_\_\_\_ FVB.Cg-Mbtm1Jsc/Mmucd \_\_\_\_\_ MMRRC: **41495-UCD**

Protocol: **GoTaq® G2 Colorless Master Mix(Promega)**

Reagent/Constituent	Volume (µL)
Water	5.6
GoTaq® G2 Colorless Master Mix,2X	7.5
Primer 1. (stock concentration is 20µM)	0.45
Primer 2. (stock concentration is 20µM)	0.45
DNA (example) extracted w/ "Qiagen DNeasy columns or other similar silica based kits"	1.0
<b>TOTAL VOLUME OF REACTION:</b>	<b>15.00 µL</b>

**Comments on protocol:**

- Protocol may work with other DNA extraction methods.

**Strategy:**

Steps	Temp (°C )	Time (m:ss)	# of Cycles
1. Initiation/Melting <span style="float: right;">HOT START? <input type="checkbox"/></span>	94	2:00	<b>1x</b>
2. Denaturation	94	0:10	
3. Annealing <span style="float: right;">steps 2-3-4 cycle in sequence</span>	65 (↓1°C/cycle)	0:30	
4. Elongation	68	2:00	
			<b>Go to 2 (9x)</b>
5. Denaturation	94	0:15	
6. Annealing <span style="float: right;">steps 5-6-7 cycle in sequence</span>	55	0:30	
7. Elongation	68	2:00 (+20s/cycle)	
			<b>Go to 5 (24x)</b>

**Primers:**

Name	Nucleotide Sequence (5' - 3')	Agarose: 1.5%	:	90
1. 41495-mutF	CTTGCAAAACCACACTGCTC	Estimated Running	90	min.
2. 41495-mutR	GAGGGAGCTGGTGTCAACAG	<b>Primers</b>	<b>Band (bp)</b>	<b>Genotype</b>
3. 41495-WtF	GAGGACCTGAAGAAGCATGG	1 & 2	~1100	mutant
4. 41495-WtR	TGAGTGGCTAAGCTGGGATT	3 & 4	198	wildtype

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## Electrophoresis Protocol:

