

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

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530-754-MMRRC

NAME OF PCR: B6;129-ErbB4^{tm1Gr/}/Mmucd MMRRC # 010461-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 (50-200ng/ μ L) 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:

Name	Nucleotide Sequence (5' - 3')
1. 10461-ErbB4-comF	TGTGCGCAGGAACAGAGAAC
2. 10461-ErbB4-mutR	CTGCACGAGACTAGTGAGAC
3. 10461-ErbB4-wtR	CCGCAGGAAGGGAGAGGTC

Electrophoresis Protocol:

Agarose: 1.5% V: 90

Estimated Running Time: 90 min.

Primer Combination	Band	Genotype
1 and 2	320 bp	MUT
1 and 3	150 bp	WT

