

GENOTYPING BY PCR PROTOCOL

MUTANT MOUSE RESOURCE & RESEARCH CENTER: UC DAVIS

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530-754-MMRRC

Protocol Name: C57BL/6-Pol^{tm1Pig}/Mmucd

MMRRC: 042060-UCD

Protocol:

Reagent/Constituent	Volume (µL)
Water	11.275
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 20µM)	0.5
Primer 2. (stock concentration is 20µM)	0.5
Taq Polymerase 5Units/µL	0.2
DNA (example) extracted w/ "Qiagen DNeasy columns or other similar silica based kits"	1.0
TOTAL VOLUME OF REACTION:	25 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	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 (↓1°C/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: 2%	V: 90
1.42060-F	ACACACACAATACTACACA	Estimated Running Time: 90 min.	
2.42060-R	AAGCTGCTGGAGTCTTTT	Primer Combination	Band (bp)
		1 & 2	320
		1 & 2	320+270+50
		1 & 2	270+50
			Genotype
			Wt
			Wt/KI
			KI/KI

Protocol / Gel Comments: After the PCR, set up the digestion using NEB.com: buffer3 3 µl, ApeK1 1 µl, PCR product 16µl, dH₂O 10 µl, for a total of 30 µl. Put the tube on the 75C heating block for 1 to 2 hrs. Spin down and add loading dye. Run 2% agarose gel for 60 min or longer. ApeK1 will only digest KI DNA, yielding 270 and 50 bp bands. Wild type DNA will not be digested. **Note:** ApeK1 has the same cutting pattern as Tse1 (65C).

