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

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530-754-MMRRC

Protocol Name: C57BL/6N-Atm1Brd Sh3bp5ltm1(KOMP)Wtsi/MbpMmucd

**MMRRC: 049640-UCD** 

Protocol:

Reagent/Constituent	Volume (µL)
Water	11.275
10x Buffer	2.5
MgCl <sub>2</sub> (stock concentration is 25mM)	1.7
Betaine (stock concentration is 5M) Optional	6.5
dNTPs (stock concentration is 10mM)	0.5
DMSO Optional	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:	25.000 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 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?	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	$\infty$	n/a

#### Primers

ners:	Electrophoresis Protocol:			
Name	Nucleotide Sequence (5' - 3')	Argarose: 1.5%	V: 90	
1. 49640-lacF	GCTACCATTACCAGTTGGTCTGGTGTC	Estimated Running:Time: 90 min.		
2. 49640-neoF	GGGATCTCATGCTGGAGTTCTTCG	<b>Primer Combination</b>	Band (bp)	Genotype
3. 49640-wtF	AGCGTGTGACGAGGCTGTGC	2 & 4	591	PreCre
4. 49640-R	CTTTGCTTCCTCATCAGTGTTCTCC	1 & 4	542	PostCre
5. 49640-F	AGTTTCCTCATCCCTGCAGTGG	3 & 4	295	Wildtype
		5 & 4	483	PostFlp

Please note, these primers are auto-designed and may not have been verified by the repository, and as such may require optimization or redesign by your facility. We recommend running primers singleplex. For screening of pups prior to any FIp or Cre recombination, the Floxed or PreCre primers may be used to identify the mutant mice. The Floxed primers test for the distal LoxP site. The PostCre primers should be used if mutant mice were crossed with a Cre recombinase line (without any FLP recombination). The PostFlp primers should be used if mutant mice were crossed with a FIp recombinase line. The PostFIp & Cre primers should be used if mutant mice were crossed with a FIp recombinase line and then a Cre recombinase line. The wildtype primers should be used for zygosity testing of all mutant mice (pre or post recombination).

# Untested Protocol: no gel image available