

# GENOTYPING BY PCR PROTOCOL

## KOMP Repository: UC DAVIS

### Protocol: CR912 Prkab1

Reagent/Constituent	Volume ( $\mu$ L)
Water	10.275
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 OF REACTION:</b>	<b>25.000 <math>\mu</math>L</b>

#### Comments on protocol:

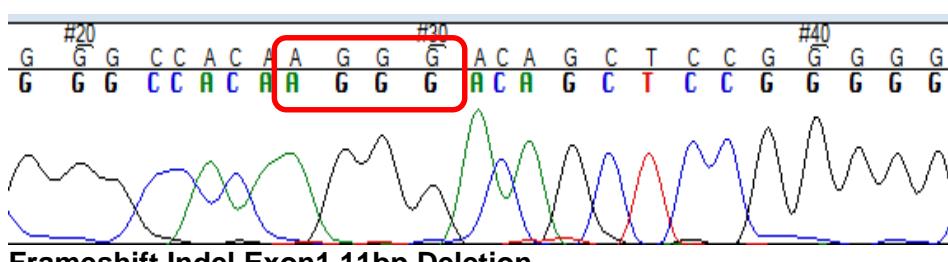
- Protocol may work with other DNA extraction methods.
- Use Touch-Down cycling protocol-first 10 cycles anneal at 65°C decreasing in temperature by 1.0°C; next 30 cycles anneal at 55°C.
- Betaine and DMSO have been standardized due to high GC content. Protocol may be tested without. Also, may adjust 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	40x
4. Elongation	72	0:40	
5. Amplification	72	5:00	1
6. Finish	15	$\infty$	n/a

#### Primers:

Name	Nucleotide Sequence (5' - 3')	Agarose: 1.5% V: 90
1. Prkab1 CR912-F	CATGGGCAACACGGAGCAGC	Estimated Running Time: 90 min
2. Prkab1 CR912-R	GGCTGTCCATGAGGATCTTGG	Primer Combination
		Band (bp)
		1 & 2
		122
		Wildtype
		1 & 2
		111
		mutant



Contig gggccacaag**acgcccgcggag**ggacagctccgggggg

**Allele Description:** Exon 1 (ENSMUSE00000375698) received a 11bp deletion (**acgcccgcggag**) from the Prkab1 gene (ENSMUST00000031486.13) using CRISPR Cas9 gene editing technology in mouse zygotes. This causes a frameshifted transcript followed by early termination signal. Subsequent founders were backcrossed to C57BL6/N to produce sequence confirmed heterozygous animals.

