

GENOTYPING BY PCR PROTOCOL

KOMP Repository: UC DAVIS

Protocol: CR956 Gja10

Reagent/Constituent	Volume (μL)
Water	10.275
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
Primer 3. (stock concentration is 20μM)	0.5
Primer 4. (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 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:

Name	Nucleotide Sequence (5' - 3')	Electrophoresis Protocol:		
1. SEQ-Gja10-F	GCTTCTGTATGTTTCAGATGTGACTTCC	Agarose: 1.5%	V: 90	
2. SEQ-Gja10-R	GGACTGCTCATCATCCAGACG	Estimated Running Time: 90 min		
		Primer Combination	Band (bp)	Genotype
		1 & 2	183	wildtype
		1 & 2	160	mutant

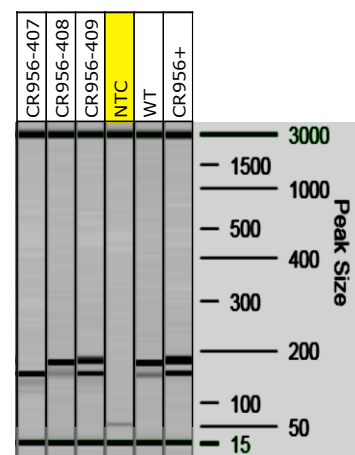
Frameshift Indel

Gja10 Exon 1

23bp Deletion



Contig gaagtccactccactccactatagtggggaagatctggctgaccatcctctcatct



Allele Description: Exon 1 ENSMUSE00001407006 had 23 bp (ccactatagtggggaagatctgg) deleted from the Gja10 gene ENSMUST00000219644.1 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.