

GENOTYPING PROTOCOL

MUTANT MOUSE RESOURCE & RESEARCH CENTER: UC DAVIS

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530-754-MMRRC

Protocol Name: CR11694 1110032F04Rik EXDEL

Protocol: GoTaq® G2 Colorless Master Mix(Promega)

Reagent/Constituent	Volume (µL)
Water	4.5
GoTaq® G2 Colorless Master Mix,2X	7.5
Primer 1. (stock concentration is 20µM) comF	0.5
Primer 2. (stock concentration is 20µM) wtR	0.5
Primer 3. (stock concentration is 20µM) mutR	0.5
DNA (example) extracted w/ "Qiagen DNeasy columns or other similar silica based kits"	1.5
TOTAL VOLUME OF REACTION:	15.00 µL

Comments on protocol:

- Protocol may work with other DNA extraction methods.

Strategy:

Steps	Temp (°C)	Time (m:ss)	# of Cycles
1. Initiation/Melting HOT START? <input type="checkbox"/>	94	2:00	1x
2. Denaturation	94	0:10	
3. Annealing steps 2-3-4 cycle in sequence	65 (↓1°C/cycle)	0:30	10x
4. Elongation	68	2:00	
5. Denaturation	94	0:15	
6. Annealing steps 5-6-7 cycle in sequence	55	0:30	25x
7. Elongation	68	2:00 (↑20sec/cycle)	
8. Finish	4	∞	n/a

Primers:

Electrophoresis Protocol:

Name	Nucleotide Sequence (5' - 3')	Agarose: 1.5%	V: 90
1. CR_1110032F04Rik_comF	ACACCTACGCTAGTGAGAGAGC	Estimated Running Time: 90 min.	
2. CR_1110032F04Rik_wtR*	AGGTACTIONGACCTCCTCGTAGC	Primer Combination	Band (bp)
3. CR_1110032F04Rik_mutR	CAGGCACTTACAGTGAACACAAGG	1 & 2, 1 & 3	1092, 1551
		1 & 3	738
			wildtype
			mutant

Allele Description: Exon 1 ENSMUSE00000396332 and flanking splicing regions were constitutively deleted from the 1110032F04Rik Gene ENSMUSG00000046999 using CRISPR Cas9 gene editing technology in mouse zygotes. The subsequent 813bp deletion from Chr 3: 68777009 - 68777821 GRCh39 was sequence verified. The selected founder animal was backcrossed to C57BL6/N to produce sequence confirmed heterozygous animals for establishing and archiving the line.

Note: Non-coding exon 1 ENSMUSE00000898859 of Gm17641 gene is also deleted.

*wtR primer untested (ePCR verified)

