

GENOTYPING PROTOCOL

MUTANT MOUSE RESOURCE & RESEARCH CENTER: UC DAVIS

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530-754-MMRRC

Protocol Name: CR11275 Pgap6 (Tmem8) 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? ☐	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:

Name	Nucleotide Sequence (5' - 3')	Agarose: 1.5%	V: 90
1. CR_Tmem8_comF	CTACATCTTCCAGCCTGATATGCTG	Estimated Running Time: 90 min.	
2. CR_Tmem8_wtR*	GCTCTCAAATGTCACTTGTAAACCACC	Primer Combination	Band (bp)
3. CR_Tmem8_mutR	GTCAGTCTCTCCACCAAACTGCC	1 & 2, 1 & 3	525, 1558
4. attP_comKI-R	TGTACCGTACACCACTGAGACC	1 & 3, 1 & 4	466, 237
			Genotype
			wildtype
			mutant

Electrophoresis Protocol:

Allele Description: Exon 5 to 6 (ENSMUSE00000139196, ENSMUSE00000139184) and flanking splicing regions were constitutively deleted from the Pgap6 (Tmem8) gene ENSMUSG00000024180 and replaced with Bxb1 attP KI (58bp sequence) using CRISPR Cas9 gene editing technology in mouse zygotes. Subsequent founders were backcrossed to C57BL6/N to produce sequence confirmed heterozygous animals.

*wtR primer untested (ePCR verified)

