

GENOTYPING PROTOCOL

MUTANT MOUSE RESOURCE & RESEARCH CENTER: UC DAVIS

mmrrc@ucdavis.edu

530-754-MMRRC

Protocol Name: CR11721 Sbp 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:

Name	Nucleotide Sequence (5' - 3')	Agarose: 1.5%	V: 90
1. CR_Sbp_comF	CTGAAGTCTCAGGAGCCGATGG	Estimated Running Time: 90 min.	
2. CR_Sbp_wtR*	GGCTCCAGTGTGAGATGAGTACC	Primer Combination	Band (bp)
3. CR_Sbp_mutR	TGTAAAGGAAGAGGGTGAGCAGG	1 & 2, 1 & 3	491, 3677
		1 & 3	531
			wildtype
			mutant

Electrophoresis Protocol:

Allele Description: Exon 2-4 (ENSMUSE00001276045, ENSMUSE00001028131, ENSMUSE00000138570) and flanking splicing regions were constitutively deleted from the Sbp Gene ENSMUSG00000024128 using CRISPR Cas9 gene editing technology in mouse zygotes. The subsequent 3146bp deletions from Chr 17 (Deletion: 17:24161390 - 17:24164583 and Deletion: 17:24164273 - 17:24164844) GRM39 was sequence verified. The selected founder animal was backcrossed to C57BL6/N to produce sequence confirmed heterozygous animals for establishing and archiving the line.

*wtR primer untested (ePCR verified)

