

GENOTYPING BY PCR PROTOCOL

MUTANT MOUSE RESOURCE & RESEARCH CENTER: UC DAVIS

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530-754-MMRRC

Protocol Name: STOCK Inpp5btm2Nbm Ocr1tm1Nbm Tg(INPP5B)CNbm/Mmucd **MMRRC:** 050927-UCD

Protocol:

Reagent/Constituent	Volume (μL)
Water	5.15
GoTaq® G2 Colorless Master Mix, 2X	7.5
Primer 1. (stock concentration is 20μM)	0.45
Primer 2. (stock concentration is 20μM)	0.45
Primer 3. (stock concentration is 20μM)	0.45
DNA (example) extracted w/ "Qiagen DNeasy columns or other similar silica based kits"	1.0
TOTAL VOLUME	
15	

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.

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	30X
7. Elongation	68 (+20s/cycle)	2:00	

Primers:

Electrophoresis Protocol:

Name	Nucleotide Sequence (5' - 3')	Argarose: 1.5% V: 90		
1. 50927_Ocr1_wtF	CCCTTTTCATCTGTTAGGAGAAATC	Estimated Running Time: 90 min.		
2. 50927_Ocr1_wtR	GCATGGTTAAACGCACTATGTGG	Primer Combination	Band (bp)	Genotype
3. 50927_Ocr1_koF	GCCCTTTGATTCTAATCCCTTTTCATC	1 & 2	227	wildtype
4. 50927_Ocr1_koR	TCTGAGCCCAGAAAGCGAA	3 & 4	425	mutant
5. 50927_BAC-INPP5B_tgF	CCACCCACGATTGACTC	5 & 6	225	mutant
6. 50927_BAC-INPP5B_tgR	GGTGTCCCAGCCCTCAG	7 & 8	146/200	WT/floxed
7. 50927_Inpp5b_LoxF	CTCATTTCTCCTTGATTCCAAT			
8. 50927_Inpp5b_LoxR	TAAAGTCTGAAAATCCAAGGC			

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