

GENOTYPING BY PCR PROTOCOL
MUTANT MOUSE RESOURCE & RESEARCH CENTER: UC DAVIS

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530-754-MMRRC

Protocol Name: C57BL/6NCrl-Rab39bem1(IMPC)Tcp/TcpMmucd MMRRC: 066531-UCD

Protocol:

Reagent/Constituent	Volume (μ L)
Water	5.6
GoTaq® G2 Colorless Master Mix,2X	7.5
Primer 1. (stock concentration is 20 μ M)	0.45
Primer 2. (stock concentration is 20 μ M)	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.
- The mutant PCR is a general LacZ PCR. The wild type is specific for this strain.

Strategy:

Steps	HOT START? <input type="checkbox"/>	Temp (°C)	Time (m:ss)	# of Cycles
1. Initiation/Melting		94	2:00	1x
2. Denaturation		94	0:10	
3. Annealing	steps 2-3-4 cycle in sequence	65 ($\downarrow 1^{\circ}\text{C}/\text{cycle}$)	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 ($\uparrow 20\text{sec}/\text{cycle}$)	

Primers:

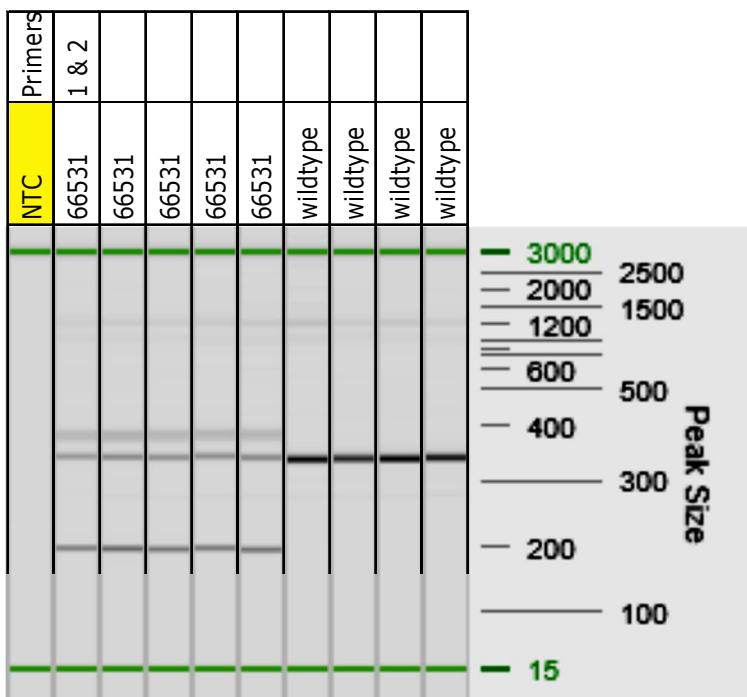
Electrophoresis Protocol:

Name	Nucleotide Sequence (5' - 3')	Argarose: 1.5% V: 90		
1. 66531-F	GAAACGGGTTGATCTGTGCTAAGG	Estimated Running Time: 90 min.		
2. 66531-R	CCCAAAGACCCTCCCACTGC	Primer Combination	Band (bp)	Genotype
		1 & 2	218	mutant
		1 & 2	351	wildtype

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Sample ID	Assay Well	Sample ID	Assay Well
ntc	A1	10466	B1
wt	B1		A2
	B2	2	
	A3	3	
	B3	4	
	A4	5	
	B4	6	
	A5	7	
	B5	8	
ntc	A6	32358	B6
wt	A7		B7
	B8	3	
	A8	4	
	B8	5	
	A9	6	
	B9	7	
	A10	8	
	B10	9	
	A11	10	
	B11	11	
	A12	12	
	B12	13	
	C1	14	
	D1	15	
ntc	C2	D2	
wt	C3		
	D3	31665	
	C4	3	
	D4	4	
	C5	5	
	D5	6	
	C6	7	
ntc	D6		
wt	C7		
	D7	13807-REG	
	C8	15	
	D8	16	
	C9	17	
	D9	18	
	C10	19	
	D10	20	
	C11	21	
	D11	22	
	C12	23	
ntc	D12		