

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

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530-754-MMRRC

Protocol Name: C57BL/6N-Ergem1(IMPC)Bay/Mmucd MMRRC: 047583-UCD

**Protocol:**

Reagent/Constituent	Volume ( $\mu$ L)
Water	5.6
GoTaq® G2 Colorless Master Mix,2X	7.5
Primer 1. (stock concentration is 20 $\mu$ M)	0.45
Primer 2. (stock concentration is 20 $\mu$ M)	0.45
DNA (example) extracted w/ "Qiagen DNeasy columns or other similar silica based kits"	1.0
	<b>TOTAL VOLUME</b>
	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	<b>10x</b>
4. Elongation		68	2:00	
5. Denaturation		94	0:15	
6. Annealing	steps 5-6-7 cycle in sequence	55	0:30	<b>25x</b>
7. Elongation		68	2:00 ( $\uparrow 20\text{sec}/\text{cycle}$ )	

**Primers:**

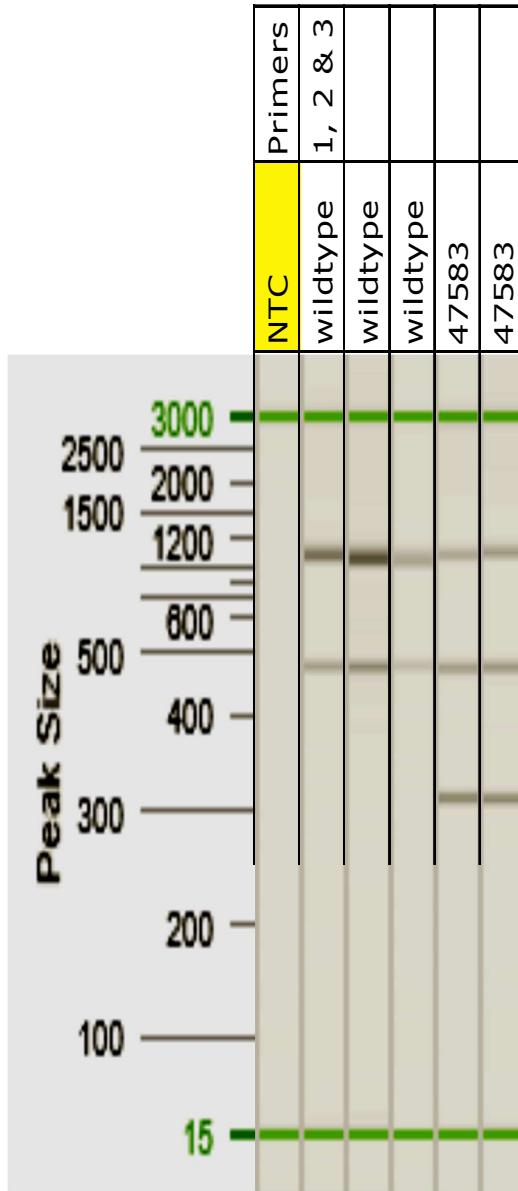
**Electrophoresis Protocol:**

Name	Nucleotide Sequence (5' - 3')	Argarose: 1.5% V: 90		
1. 47583-comF	CATGATGGTCTTCCAGGTG	Estimated Running Time: 90 min.		
Primer Combination	Band (bp)	Genotype		
2. 47583-WtR	GTAGCTGCCGTAGCTCATCC	1 & 3	300, 950	mutant
3. 47583-mutR	GGAACAGTGGAGCTTACGAA	1 & 2	480	wildtype

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