

# GENOTYPING BY PCR PROTOCOL

## MUTANT MOUSE RESOURCE & RESEARCH CENTER: UC DAVIS

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530-754-MMRRC

**Protocol Name:** B6J.Cg-Ptprctm1Holm Tg(ITGAL-Ptprca)BRasch/Mmucd **MMRRC: 044051-UCD**

**Protocol:**

Reagent/Constituent	Volume (µL)
Water	5.6
GoTaq® G2 Colorless Master Mix, 2X	7.5
Primer 1. (stock concentration is 20µM)	0.45
Primer 2. (stock concentration is 20µM)	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.
- Strains 44051, 46054, 46256, 46257, 46258, 46266, 46268, 46269, 46270 share similarities. PstI digest distinguishes between Ptprca and Ptprca\*. Primers 3 and 4 are to detect ΔC strain which is present in 26258.

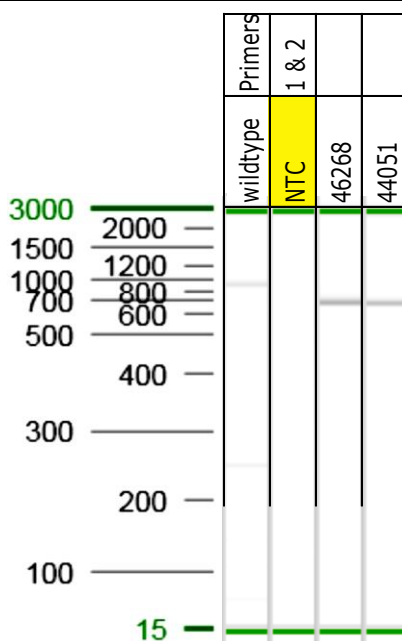
**Strategy:**

Steps	Temp (°C)	Time (m:ss)	# of Cycles
1. Initiation/Melting <span style="float: right;">HOT START? <input type="checkbox"/></span>	94	2:00	<b>1x</b>
2. Denaturation	94	0:10	
3. Annealing <span style="float: right;">steps 2-3-4 cycle in sequence</span>	65 (↓1°C/cycle)	0:30	<b>10x</b>
4. Elongation	68	2:00	
5. Denaturation	94	0:15	
6. Annealing <span style="float: right;">steps 5-6-7 cycle in sequence</span>	55	0:30	<b>30X</b>
7. Elongation	68 (+20s/cycle)	2:00	

**Primers:**

**Electrophoresis Protocol:**

Name	Nucleotide Sequence (5' - 3')	Argarose: 1.5%	V: 90
1. 44051-CD363-F	GCCAGCTACATTGATGGCTTC	Estimated Running Time: 90 min.	
2. 44051-CD365-R	CCTGTATGAAGGAAGTCTCTGG	<b>Primer Combination</b>	<b>Band (bp)</b>
3. 44051-321-F	GGTCACTGGAATGAAAACCTCCCG	1 & 2	734
4. 44051-332-R	GCATAGGAAATGGCCATAGTC	3 & 4	0
			<b>Genotype</b>
			mutant



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PstI digest of primers 1 & 2 PCR product

