

**GENOTYPING BY PCR PROTOCOL**  
**MUTANT MOUSE REGIONAL RESOURCE CENTER: UC DAVIS**  
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 530-754-MMRRC

**NAME OF PCR:** B6;129S5-Gpr56<sup>tm1Lex</sup>/Mmcd **MMRRC #** 032341-UCD

**Protocol:**

| Reagent/ Constituent                             | Volume (μL) |
|--------------------------------------------------|-------------|
| Water                                            | 10.775      |
| 10x Buffer ( 15mM MgCl <sub>2</sub> )            | 2.5         |
| MgCl <sub>2</sub> (stock concentration is 25mM)  | 1.7         |
| Betaine (stock concentration is 5M)              | 6.5         |
| dNTPs (stock concentration is 10mM)              | 0.5         |
| DMSO                                             | 0.325       |
| Primer 1 (stock concentration is 20μM) DNA085-5  | 0.5         |
| Primer 2 (stock concentration is 20μM) DNA085-14 | 0.5         |
| Primer 3 (stock concentration is 20μM) Neo3a     | 0.5         |
| Taq Polymerase 5Units/μL                         | 0.2         |
| DNA Sample                                       | 1.0         |
| <b>TOTAL VOLUME OF REACTION:</b>                 | <b>25μL</b> |

**Comments on protocol:**

**Strategy:**

| Steps                                                                | Temp (°C )            | Time (m:ss) | # of Cycles |
|----------------------------------------------------------------------|-----------------------|-------------|-------------|
| 1. Initiation/Melting HOT START? CHECK HERE <input type="checkbox"/> | 94                    | 5:00        | 1           |
| 2. Denaturation                                                      | 94                    | 0:15        |             |
| 3. Annealing } steps 2-3-4 will cycle in sequence                    | 65 to 55 (↓1°C/cycle) | 0:30        | } 40x       |
| 4. Elongation                                                        | 72                    | 0:40        |             |
| 5. Amplification                                                     | 72                    | 5:00        | 1           |
| 6. Finish                                                            | 4                     | ∞           | n/a         |

**Primers:**

| Primer Name  | Nucleotide Sequence (5' - 3') |
|--------------|-------------------------------|
| 1: DNA085-5  | CGA GAA GAC TTC CGC TTC TG    |
| 2: DNA085-14 | AAA GTA GCT AAG ATG CTC TCC   |
| 3: Neo3a     | GCA GCG CAT CGCCTT CTA TC     |

**Electrophoresis Protocol:**

**% Agarose:** 1.5 **V:** 90

**Estimated Running Time (min):** 90

| Primers | Expected Bands | Genotype  |
|---------|----------------|-----------|
| 1 and 2 | 639 bp         | Wild-type |
| 2 and 3 | 369 bp         | Mutant    |