

**GENOTYPING BY PCR PROTOCOL**  
**MUTANT MOUSE REGIONAL RESOURCE CENTER: UC DAVIS**  
[mmrrc@ucdavis.edu](mailto:mmrrc@ucdavis.edu)  
 530-754-MMRRC

NAME OF PCR: STOCK *Ntf3<sup>tm1Lfr</sup>*/Mmucd

MMRRC # 000191-UCD

**Protocol:**

| Reagent/ Constituent  | Volume ( $\mu$ L)          |
|---|----------------------------|
| Water   | 11.275                     |
| 10x Buffer (contains / without 15mM MgCl <sub>2</sub> )   | 2.5                        |
| MgCl <sub>2</sub> (stock concentration is 25mM)   | 1.7                        |
| Betaine (stock concentration is 5M) <i>Optional</i>   | 6.5                        |
| dNTPs (stock concentration is 10mM)   | 0.5                        |
| DMSO <i>Optional</i>  | 0.325                      |
| Primer 1 (stock concentration is 20 $\mu$ M) MNT3-IF  | 0.5                        |
| Primer 2 (stock concentration is 20 $\mu$ M) MNT3-IR  | 0.5                        |
| Primer 3 (stock concentration is 20 $\mu$ M) PGK2-mut   | 0.5                        |
| Primer 4 (stock concentration is 20 $\mu$ M) SV40F-mut  | 0.5                        |
| Taq Polymerase 5Units/ $\mu$ L  | 0.2                        |
| DNA extracted with <input type="checkbox"/> NaOH <input checked="" type="checkbox"/> Proteinase K <input type="checkbox"/> Other: | 1.0                        |
| <b>TOTAL VOLUME OF REACTION:</b>  | <b>25<math>\mu</math>L</b> |

**Comments on protocol:**

- 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.
- Betaine/DMSO is standardized due to high GC content in promoter regions and protocol may be tested without. Also, may adjust MgCl<sub>2</sub> to increase reaction or decrease non specific amplifications.

**Strategy:**

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

**Primers:**

| Name         | Nucleotide Sequence (5' - 3')               |
|--------------|---|
| 1: MNT3-IF   | ACT ACG GCA ACA GAG ACG CTA C               |
| 2: MNT3-IR   | ACA GGC TCT CAC TGT CAC ACA C               |
| 3: PGK2-mut  | GTG CCA GCG GGG CTG CTA AAG CGC             |
| 4: SV40F-mut | CTG CAT TCT AGT TGT GGT TTG TCC AAA CTC ATC |

**Electrophoresis Protocol:**

Agarose: 2% V: 80 Estimated Running Time: 90 min

| Primer Combination | Band   | Genotype |
|--------------------|--------|----------|
| 1 and 2            | 200 bp | WT +/+   |
| 3 and 4            | 150 bp | KO -/-   |