

# Mutant Mouse Regional Resource Center: UC Davis

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530-754-MMRRC

NAME OF PCR: STOCK Tg(Pde1c-cre)106Gsat/Mmcd MMRRC # 030728-UCD

## Protocol:

| Reagent/ Constituent                                                                      | Volume (μL)      |
|-------------------------------------------------------------------------------------------|------------------|
| Water                                                                                     | 11.275           |
| 10x Buffer                                                                                | 2.5              |
| MgCl <sub>2</sub> (stock concentration is 25mM)                                           | 1.7              |
| Betaine (stock concentration is 5M) Optional                                              | 6.5              |
| dNTPs (stock concentration is 10mM)                                                       | 0.5              |
| DMSO <i>Optional</i>                                                                      | 0.325            |
| Primer 1 (stock concentration is 20μM)                                                    | 0.5              |
| Primer 2 (stock concentration is 20μM)                                                    | 0.5              |
| Taq Polymerase 5Units/μL                                                                  | 0.2              |
| DNA (50-200ng/μL) extracted w/ "Qiagen DNeasy columns or other similar silica based kits" | 1.0              |
| <b>TOTAL VOLUME OF REACTION:</b>                                                          | <b>25.000 μL</b> |

## 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.
- Betaine and DMSO have been standardized due to high GC content. Protocol may be tested without. Also, may adjust MgCl<sub>2</sub> to increase reaction or decrease non specific amplifications.

## Strategy:

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

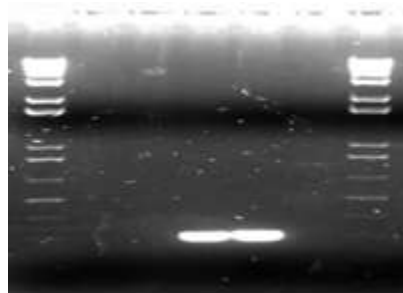
## Primers:

| Name        | Nucleotide Sequence (5' - 3') |
|-------------|-------------------------------|
| 1. Pde1c F1 | AGGGGAAGTGACAAAGTGACTGAAA     |
| 2. CreGS R1 | CGGCAAACGGACAGAAGCATT         |

## Electrophoresis Protocol:

Agarose: 1.5% V: 90  
 Estimated Running Time: 90 min.

| Primer Combination | Band   | Genotype   |
|--------------------|--------|------------|
| 1 and 2            | 300 bp | transgenic |
|                    |        |            |



1 2 3 4 5 6 7

### Lanes

- 1 & 7: 1 kb+ ladder (Invitrogen, Cat. #10787-026)
- 2: H<sub>2</sub>O
- 3: Wild-type Control
- 4 & 5: Pde1c tg/+
- 6: Other GENSAT line