Primer Normalization

1. Make cDNA standards A→ E

2. Select the cycle and fix the annealing temperature.

3. Run Q-PCR with standards and Blank

4. Check the melting curve, whether it has single product or not.

5. Check 3.3 CT value difference between 1:10 dilutions

6. Check whether blanks amplified or not.

7. Check whether it has non-specific amplification.

8. Confirm the single product by running gel.