

## Medina Lab, Penn State University

*RT-qPCR protocol (with aRNA), prepared by M. DeSalvo*

*Last updated: 29 July 2008*

### 1. cDNA synthesis

- a. Place 1 $\mu$ g aRNA into a PCR tube (volume can't exceed 11 $\mu$ L).
- b. Add 1 $\mu$ L random hexamers.
- c. Add 1 $\mu$ L 10mM dNTPs.
- d. Bring final volume to 13 $\mu$ L with dH<sub>2</sub>O.
- e. Place @ 65°C for 5min.
- f. Chill on ice for at least 1min.
- g. Prepare the RT master mix.

| Reagent         | 1x (uL) | e.g. 11x (uL) |
|-----------------|---------|---------------|
| 5x RT buffer    | 4       | 44            |
| 0.1M DTT        | 1       | 11            |
| RNaseOUT        | 1       | 11            |
| SuperScript III | 1       | 11            |

- h. Add 7 $\mu$ L RT master mix to each tube (mix by pipetting).
- i. Place @ 25°C for 5min.
- j. Place @ 50°C for 50min.
- k. Place @ 70°C for 15min.
- l. Chill on ice.
- m. Add 1 $\mu$ L RNaseH to each tube.
- n. Place @ 37°C for 20min. Test PCRs
- a. Place RT reactions into new 1.5mL tubes.
- b. Add 180 $\mu$ L dH<sub>2</sub>O.
- c. Use 2 $\mu$ L of cDNA in a PCR reaction using good qPCR primers.
- d. PCR master mix (25 $\mu$ L reaction volumes):
  - i. This will change depending on Taq being used.

| Reagent                   | 1x (uL) | e.g. 11x (uL) |
|---------------------------|---------|---------------|
| 10x ExTaq PCR buffer      | 2.5     | 27.5          |
| ExTaq dNTPs               | 2.0     | 22.0          |
| 10 $\mu$ M forward primer | 0.5     | 5.5           |
| 10 $\mu$ M reverse primer | 0.5     | 5.5           |

|                   |        |         |
|-------------------|--------|---------|
| Takara ExTaq      | 0.125  | 1.375   |
| dH <sub>2</sub> O | 17.375 | 191.125 |
| cDNA              | 2.0    |         |

**2. PCR Program**

e. PCR program as follows:

- i. 94°C for 30sec
- ii. 94°C for 15sec
- iii. 60°C for 15sec
- iv. 72°C for 15sec
- v. go to step ii. – repeat 35x
- vi. 72°C for 10min
- vii. 4°C for ever.

f. Run 5uL of each PCR reaction on a 2% agarose gel.

g. You should get a single band ~100bp.

RT-qPCR

a. Use 12.5uL reaction volumes.

b. Use 96-well Optical Reaction Plates (ABI Part #4306737).

c. Use Power SYBR Green PCR Master Mix (ABI Part #4367659).

d. Prepare a separate master mix (in a 1.5mL tube) for each gene that you will be testing.

| Reagent               | 1x (uL) |
|-----------------------|---------|
| SYBR Green            | 6.25    |
| Forward primer (10uM) | 0.25    |
| Reverse primer (10uM) | 0.25    |
| dH <sub>2</sub> O     | 3.75    |

3.

e. Calculate the # of reactions needed for each gene:

- i. (# of samples) X (3 technical rep's per sample)
- ii. e.g. (9 samples) X (3) = 27 total rxns.

3.

f. Overshoot by a few reactions to allow for slop, and to have left over master mix for your 3 non-template controls (NTCs).

- i. Thus, the total # of actual qPCR reactions cannot exceed 93.

3.

g. Bring the following to the Choi lab:

- i. Sample tubes.
- ii. Master mixes.
- iii. 96-well plate.
- iv. Plate sealers.
- v. Tube with PCR-grade water.
- vi. P200 and P2 tips.

3.

h. Start software.

i. Choose "Absolute quantification" à "Next".

j. Click "SYBR Green" à "Add" à "Next".

k. Drag the mouse over (thus highlighting) the wells you plan to fill with reactions à click "SYBR Green" à "Finish".

l. Click on the "Instrument" tab.

m. Highlight "Stage 1" (50°C for 2min) à "Delete".

n. Click "Add Dissociation Stage".

o. Enter "13" in the "Sample Volume (uL)" field.

p. Make sure "Data Collection" takes place at "Stage 2, Step 2".

q. Click on the "Setup" tab.

r. Enter names for all reactions.

s. Choose "NTC" for 3 wells.

t. Fill appropriate wells with 10.5uL of the appropriate Master Mix.

u. Add 2uL of cDNA to each well.

v. Add 2uL of dH<sub>2</sub>O to the NTC wells.

w. Seal the plate with a plate sealer and use the plastic tool to ensure that each well is sealed.

x. Spin the plate for 30sec.

y. Insert the plate into the machine.

Go to the "Instrument" tab and click "Start".