

AUG 09, 2023

qRT-PCR sample preparation

Louise Uoselis¹

¹Lazarou Lab, WEHI



Louise Uoselis WEHI

ABSTRACT

Protocol for qRT-PCR sample preparation for analysis using a RotorGeneQ machine (Qiagen).

OPEN ACCESS



DOI:

dx.doi.org/10.17504/protocol s.io.4r3l22713l1y/v1

Protocol Citation: Louise Uoselis 2023. qRT-PCR sample preparation. protocols.io https://dx.doi.org/10.17504/p rotocols.io.4r3l22713l1y/v1

License: This is an open access protocol distributed under the terms of the Creative Commons Attribution License. which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited

Protocol status: Working We use this protocol and it's working

Created: Aug 09, 2023

Last Modified: Aug 09,

2023

PROTOCOL integer ID:

86219

Keywords: ASAPCRN

- 1 Synthesise cDNA libraries from total RNA from each sample using a High-Capacity cDNA Reverse Transcription Kit (Applied Biosystems), using Oligo(dT)₂₀ primers.
- Place the synthesised libraries on ice, and dilute the libraries 1:3 by adding \square 60 μ L of DEPC-treated H2O to each sample, pipetting up and down gently to mix \sim 8 times.
- Dilute primer stocks by adding \square 2 μ L of the forward primer, and \square 2 μ L of the reverse primer, to \square 46 μ L of DEPC-treated H2O on ice to make a final combined forward and reverse primer stock of \square 4 micromolar (μ M) forward primer and \square 4 micromolar (μ M) reverse primer for each gene target to be analysed.
- 4 Thaw the 2x QuantiNova SYBR green master mix (Qiagen) on ice.
- Depending on the number of samples being analysed, assemble a sample master mix (not containing the cDNA library) containing the following:

Reagent	Volume (1x)
2x SYBR Green Master Mix	5 uL
4 uM forward and 4 uM reverse primer master mix	1 uL
cDNA library (to be added individually to each tube)	4 uL
Total volume	10 uL

To qRT-PCR tubes sitting on ice (Qiagen – RotorGeneQ compatible), add \bot 6 μ L of the master mix for each target gene to the desired tubes, and add \bot 4 μ L of the appropriate cDNA library to each sample.

7	Tap the rack holder to ensure all solution is at the bottom of the tube before capping each sample.(NOTE: samples do not need to be mixed prior to analysis)
8	Load and run the samples on a RotorGeneQ (Qiagen) machine using the desired parameters for the gene targets you are analysing.