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1 Step RT- QPCR

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Protocol status: Working

We use this protocol and it's working

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Abstract

description of how to do a 1 step rt qpcr



Organoid RNA extraction

- 1 All RNA was extracted using the following protocol as described in our paper [dx.doi.org/10.17504/protocols.io.81wgbr95qlpk/v1](https://doi.org/10.17504/protocols.io.81wgbr95qlpk/v1)

1 Step RT-QPCR

- 2
 - Use RNase-free technique throughout.
 - Decide your multiplex groups (up to 4 targets per well, including GAPDH), making sure fluorophores do not overlap.
 - Thaw master mix, probes, and RNA on ice. Mix gently and quick-spin.
- 3 Measure RNA concentration for each sample.
- 4 Dilute each RNA sample with nuclease-free water so all samples have the same concentration (so that 2 μ L gives the same RNA input for every sample)
- 5 Keep normalized RNA on ice
- 6 Create a plate map:
List each sample.
For each probe set/condition, assign 2 technical replicates per sample.
Total wells = samples $\times$ conditions $\times$ replicate
- 7 Label your 384-well plate orientation and corresponding wells
- 8 For each multiplex condition (each “probe set”), make a probe mix so pipetting is consistent. In a labeled tube for Probe Set 1, combine the probes for that multiplex (up to 4 probes). Repeat for Probe Set 2, 3, and 4
- 9 Make one “reaction mix” per probe set that contains everything except RNA. Per well (10 μ L total):
 - 5.0 μ L IDT PrimeTime One-Step RT-qPCR Master Mix
 - 0.5 μ L of each probe (up to 4 probes total; 2.0 μ L max probe volume)
 - Nuclease-free water to bring the non-RNA volume to 8.0 μ L
 - RNA will be 2.0 μ L added later
- 10 For each probe set, calculate number of reactions needed:
(number of wells for that probe set) $\times$ (1.1 or 1.2) to account for pipetting loss



- 11 In a chilled tube, combine for that probe set:
PrimeTime One-Step RT-qPCR Master Mix (5 μ L $\times$ number of reactions)
Probe mix components (0.5 μ L each probe $\times$ number of reactions)
Nuclease-free water to reach 8 μ L per reaction
- 12 Mix gently by pipetting, then quick-spin.
Keep mixes on ice and protected from light.
- 13 Add 8.0 μ L of the appropriate probe-set reaction mix into each assigned well of the 384-well plate.
Add 2.0 μ L of the corresponding normalized RNA sample into each well.
- 14 For no-template controls (NTCs), add 2.0 μ L nuclease-free water instead of RNA.
Seal the plate with an optical seal. Quick-spin the plate (brief spin) to remove bubbles and collect liquid at the bottom.
- 15 Load the plate into the Bio-Rad CFX Opus 384.
Set up the run as a one-step RT-qPCR multiplex assay:
 - Assign fluorophores/channels matching your probes.
 - Confirm plate type (384-well) and reaction volume (10 μ L).Use the cycling program recommended by IDT for PrimeTime One-Step RT-qPCR Master Mix
Start the run.

Analysis

- 16 After the run, verify amplification curves and baseline/threshold settings
- 17 Confirm NTCs show no amplification.
- 18 Export Cq values and analyze using your preferred method