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First-Strand cDNA Synthesis from Cultured Human Epithelial Cells (HNSCC)

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Protocol status: Working

We use this protocol and it's working well.

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Abstract

This protocol describes the synthesis of first-strand complementary DNA (cDNA) from total RNA extracted from epithelial cell cultures of HNSCC cell lines using the Bio-Rad iScript cDNA Synthesis Kit. The method is optimized for downstream quantitative real-time PCR (RT-qPCR) applications and gene expression analysis.

Guidelines

Notes and Best Practices

1. **RNA Handling:** Always work in an RNase-free environment. Use dedicated pipettes and filter tips for RNA work.
2. **Reaction Scaling:** The protocol can be scaled to 10 μ L reactions (half volumes) if RNA is limited, though 20 μ L reactions are recommended for optimal performance.
3. **Primer Design Consideration:** The iScript kit contains a blend of oligo(dT) and random hexamer primers, providing unbiased 3' and 5' coverage. Design qPCR primers accordingly.
4. **Dynamic Range:** The iScript kit supports RNA inputs from 1 pg to 1 μ g, making it suitable for both high and low-abundance transcripts.
5. **No-Template Control (NTC):** In addition to no-RT controls, include a no-template control (water only) in downstream qPCR reactions to monitor for contamination.
6. **Documentation:** Record all calculations, lot numbers of reagents, and thermal cycler used for full traceability and reproducibility.



Materials

Materials and Reagents

Equipment:

- NanoDropTM spectrophotometer (or equivalent UV-Vis spectrophotometer)
- Thermal cycler with heated lid capability
- Microcentrifuge
- Vortex mixer
- Pipettes (P2, P10, P20, P200, P1000)
- Ice bucket with ice

Consumables:

- 0.2 mL PCR tubes (thin-walled, nuclease-free)
- Nuclease-free pipette tips (filter tips recommended)
- Nuclease-free microcentrifuge tubes (1.5 mL)

Reagents:

- Bio-Rad iScript cDNA Synthesis Kit (Cat# 1708890/1708891) containing:
 1. 5× iScript Reaction Mix (4 µL per 20 µL reaction)
 2. iScript Reverse Transcriptase (1 µL per 20 µL reaction)
- Nuclease-free water (molecular biology grade)
- Total RNA (extracted from epithelial cells, stored at -80°C)



RNA Quantification and Quality Assessment

- 1 Thaw RNA samples on ice. Mix gently by pipetting or brief vortexing. 
- 2 Measure RNA concentration using NanoDrop™ spectrophotometer. 
- 3 Assess RNA purity by evaluating absorbance ratios (A260/A280 and A260/A230)
Note: Proceed if both ratios are within acceptable ranges 

RNA Normalization

- 4 **Rationale:** Since cDNA concentration is difficult to accurately quantify post-synthesis, normalize the RNA input amount to ensure consistency across samples. This standardization enables reliable comparison of gene expression levels between samples.
- 5 **Recommended:** 500 ng total RNA per reaction (adjustable based on RNA availability: 1 pg - 1 µg range) 

cDNA Synthesis Reaction Setup

- 6 Thaw all iScript kit components on ice. Mix gently by inverting tubes 5-10 times. Briefly centrifuge to collect contents. 
- 7 Calculate the number of reactions needed (include 10% overage for pipetting error):
 - 5× iScript Reaction Mix: **4 µL**
 - iScript Reverse Transcriptase: **1 µL**
 - Total per reaction: **5 µL**
 - RNA Template: **Variable**, calculated from step 5
 - Water (µL) = **20-5-RNA volume (µL)**
- 8 Mix gently by pipetting up and down 5-8 times. Avoid introducing bubbles. 
- 9 Briefly centrifuge tubes (2-3 seconds) to collect contents at the bottom. 
- 10 Program the thermal cycler with the following protocol: 



Step	Temperature	Duration	Purpose
1. Priming	25°C	5 minutes	Primer annealing
2. Reverse Transcription	46°C	20 minutes	cDNA synthesis
3. RT Inactivation	95°C	1 minute	Enzyme inactivation
4. Hold	4°C	∞	Storage

Thermocycler Program for iScript cDNA Synthesis

Post-Synthesis Processing

- 11 **Rationale for Diluting cDNA:** Diluting cDNA reduces potential PCR inhibitors from the reverse transcription reaction and extends the usable volume for multiple downstream qPCR reactions.
- 12 **Recommended Dilution:** 1:3 (v/v) or 1:4 (v/v) depending on RNA input amount and expected target gene abundance:
 - For 500 ng RNA input: 1:3 dilution is optimal
 - For lower RNA inputs (<100 ng): 1:2 dilution or use undiluted
- 13 **Dilution procedure** (1:3 dilution example):
 - Transfer the entire 20 µL cDNA reaction to a labeled 1.5 mL microcentrifuge tube
 - Add 60 µL nuclease-free water
 - Mix by pipetting or gentle vortexing
 - **Final volume: 80 µL**
- 14 Briefly centrifuge to collect contents.



Storage and Quality Control

- 15 Aliquot diluted cDNA into appropriate volumes based on anticipated use:
 - For frequent use (within 1 week): store at **4°C**
 - For long-term storage (>1 week): store at **-20°C** or **-80°C**
- 16 Minimize freeze-thaw cycles. For samples requiring multiple uses, prepare 10-20 µL aliquots to avoid repeated freezing and thawing.
- 17 Label all tubes clearly





Protocol references

1. Bio-Rad Laboratories. (2025). iScript™ cDNA Synthesis Kit Instruction Manual. Bio-Rad Laboratories, Inc.
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