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WORKS FOR ME

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Synthesis of double-strand cDNA (ds-cDNA) from viral dsRNA by using Random primers

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dx.doi.org/10.17504/protocols.io.81wgbyrxnvpk/v1[Vahid Jalali Javaran](#)¹¹Département de Biologie, Centre SÈVE, Université de Sherbrooke, Sherbrooke, QC J1K 2R1, Canada

Nanovirseq



Vahid Jalali Javaran

COMMENTS 0

ABSTRACT

Double-stranded cDNA synthesis from viral dsRNAs:

For dsRNA sequencing by nanopore sequencing, this protocol was used. Before treating samples with RNase T1, you should measure the total concentration of RNAs in the samples by using a nanodrop or Qubit device, as RNase T1 has the ability to partially digest double-stranded RNAs in the absence of single-stranded RNA.

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RNase T1 and DNase I digestion

- DNase I (RNase-free)
- DNase I Reaction Buffer with MgCl₂ (10X)
- RNase T1

Synthesize the first strand of cDNA (Reverse Transcription)

- Random primers (60 µM)
- dNTP (10 mM)
- H₂O
- First Strand cDNA synthesis Buffer
- RNase out or RNasin® Ribonuclease Inhibitor (40 u/µl)
- Superscript III or Maxima H minus

Removal of the residual RNA by RNase H

- RNase H

Synthesize the second strand of cDNA

- Klenow 10X buffer+NAD,
- dNTP (10 mM),
- Klenow DNA Polymerase I,
- E. coli DNA Ligase I

cDNA purification by AMPure XP beads

- Agencourt AMPure XP
- Magnet plate
- Ethanol 80%
- Nuclease free water
- 1.5 ml eppendorf tube

1h 20m

RNase T1 and DNase I digestion

1

- Add 10X DNase Buffer with MgCl₂ (final concentration should be 1X).
- Add 50 units RNase T1 per 1 µg of total RNA and 1 unit DNase I per 2 µg of total RNA.
- Incubate at 37 degrees C for 20 min.

20m

1.1

Stop reaction

Do a phenol-chloroform extraction and Ethanol precipitation of dsRNA.

1h

Synthesis of the first strand of cDNA

- 2 Mix well below components by pipetting and centrifuge or spin briefly.

A	B
Treated dsRNA	5 µl
Random primers (60 µM)	2 µl
dNTP (10 mM)	1 µl
H ₂ O	6 µl
Total	14 µl

- Incubate at 99C for 5 mins and put tubes **immediately on water ice** (it is better to do this step with a thermocycler).

- 2.1 Add below components and mix well by pipetting and centrifuge or spin briefly.

A	B
First Strand Buffer	4 µl
RNase out or RNasin® Ribonuclease Inhibitor (40 u/µl)	1 µl
Maxima H minus (Point: just use 200 unit)	1 µl
Total	20 µl

- 2.2 Incubation step.

A	B
25 C	20 mins
55 C	90 mins
85 C (Inactivation step)	5 mins

Removal of residual RNA by RNase H

- 3 Mix well below components by pipetting and centrifuge or spin briefly.

A	B
cDNA	20 µl
RNase H	1 µl

A	B
Total	21 μ l

Incubate at 37 C for 20 mins

2h 40m

Synthesis of the second strand of cDNA

- 4 Mix well below components by pipetting and centrifuge or spin briefly.

A	B
cDNA	21 μ l
Klenow 10X buffer	2.7 μ l
dNTP (10 mM)	1 μ l
Klenow DNA Polymerase I	0.7 μ l
E. coli DNA Ligase I	1 μ l
H ₂ O	0.6 μ l
Total	27 μ l

2h 40m

- 4.1 Incubation step.

A	B
16 C	150 mi
75 C; (Inactivation step) Point: check this part based on enzyme brand.	10 min

17m

cDNA purification by AMPure XP

- 5 Mix well below components by pipetting and incubate at room temperature for 5 mins.
Point: Put bead stock in room temperature for 20 min (before using). Vortex the bead stock very well (before using)

A	B
double stranded-cDNA	27 μ L
AMPure XP	49 μ L

Place the reaction tube onto a magnet plate for 2-5 mins

IMPORTANT: Wait for the solution to clear before proceeding to the next step.

- 5.1 This step must be performed while the reaction tube is situated on the magnet plate:

- Aspirate the cleared solution from the reaction tube and discard. Leave 5 μ L of supernatant

behind, otherwise, beads are drawn out with the supernatant.

IMPORTANT: Do not disturb the ring of separated magnetic beads.

- 5.2 This step must be performed while the reaction tube is situated on the magnet plate:
Dispense 200 μ L of 80% ethanol to the reaction tube and incubate for 30 seconds at room temperature.
Aspirate out the ethanol and discard.
Repeat previous step. 1m
- 5.3 After removing and discarding all supernatant, **A dry time is optional (2 to 5 mins)** to ensure all traces of Ethanol are removed.
NOTE: For fragments 10 kb and larger, do not over-dry the bead ring (bead ring appears cracked if over dried) as this will significantly decrease elution efficiency. 2m
- 5.4 Remove the reaction tube from the magnet plate, and then add 40 μ L of nuclease-free water to the reaction tube and pipette mix 10 times. Incubate for 2 minutes.
NOTE The liquid level will be high enough to contact the magnetic beads at a 40 μ L nuclease-free water. A greater volume of nuclease-free water can be used, but using less than 40 μ L will require extra mixing (to ensure the liquid comes into contact with the beads), and may not be sufficient to elute all cDNAs. 2m
- 5.5 Place the reaction tube onto the magnet plate for 5 minute to separate beads from the solution.
IMPORTANT Wait for the solution to clear before proceeding to the next step.
Transfer the eluate to a new plate. 5m
- Use 1 μ L of eluted ds-cDNA for concentration measurement by Qubit.