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Preparation of dsRNA viruses for MinION - CLM

Forked from [Preparation of dsRNA viruses for next-generation sequencing](#)

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In Development



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ABSTRACT

Updated protocol by CLM

EXTERNAL LINK

<https://doi.org/10.1099/mgen.0.000315>

PROTOCOL CITATION

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<https://protocols.io/view/preparation-of-dsrna-viruses-for-minion-clm-bvbkn2kw>



FORK NOTE

FORK FROM

Forked from Preparation of dsRNA viruses for next-generation sequencing, Sam Diaz-Munoz

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May 26, 2021

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50252

MATERIALS TEXT

MATERIALS

[NEBNext Second Strand Synthesis \(dNTP-free\) Reaction Buffer - 0.4 ml](#) **New England**

Biolabs Catalog #B6117S

[E.coli DNA Ligase - 1,000 units](#) **New England**

Biolabs Catalog #M0205L

[DNA Polymerase I \(E.coli\) - 2,500 units](#) **New England**

Biolabs Catalog #M0209L

[DMSO](#) **Sigma**

Aldrich Catalog #D1435

[Nuclease-free water](#)

Ambion Catalog #AM9932

[Qubit dsDNA HS Assay Kit](#) **Thermo Fisher**

Scientific Catalog #Q32851

[DNase I \(e.g., NEB #M0303\) and DNase I Cleanup Reagents or Kit for Removal of DNA Prior to Depletion](#) **New England Biolabs**

[Millex-GP Syringe Filter Unit, 0.22 µm](#) **Emd**

Millipore Catalog #SLGP033RS

[Qubit RNA HS Assay Kit](#) **Thermo Fisher**

Scientific Catalog #Q32852

[RNase H](#) **New England**

Biolabs Catalog #M0297S

[RNeasy® Mini](#)

Kit Qiagen Catalog #74104

[SuperScript™ III First-Strand Synthesis System](#) **Thermo Fisher**

Scientific Catalog #18080051

[RNase AT/1 Mix](#) **Thermo**

Scientific Catalog #EN0551

[RNeasy MinElute Cleanup](#)

Kit Qiagen Catalog #74204

[Nucleospin Gel and PCR cleanup](#) **Macharey**

Nagal Catalog #740609.250

- 1 Viral lysates should be passed through a 0.22µm filter to remove host debris, then treated with nucleases to degrade extracapsular nucleic acids.

To 1ml filtrate:

25µl DNase I,

50µl RNase A/T1

107.5 DNase I Buffer

Scale down for concentrated lysates, which should be in the range of 50-500uL.

Incubate for 1 hour 30 minutes at 37°C.

Inactivate nucleases at 65°C for 10 minutes and proceed to RNA extraction.

- 2 RNA should be extracted using a commercially available kit. We used an RNeasy Mini Kit (Qiagen). Kits that require carrier RNA are not recommended because this will interfere with sequencing.

Follow manufacturer instructions with the following modifications:

1. Pass all lysate through column at the first step (i.e. add buffer + lysate mix multiple times until lysate is used up)
2. Elute with 45uL of Elution Buffer. Let column sit at 25°C for 5 minutes before spinning.
3. Split into 20uL aliquots. RNA may be frozen at -80°C for future processing or may move on to next step.

- 3 Quantify the RNA using Qubit RNA High Sensitivity Kit (HS). Note that low viral titres and/or small viral genomes may result in the amount of RNA being below the Qubit's limit of detection.
- 4 Add 50% DMSO (v/v) to RNA sample, and incubate for 1 hour 30 minutes at 65°C.
- 5 Purify RNA using a column cleanup kit (we recommend Qiagen's RNeasy MinElute Cleanup Kit) or your preferred other method (e.g. ethanol precipitation, TRIzol purification). This step removes DMSO from sample.

If using RNeasy MinElute Cleanup Kit:

1. Adjust sample volume to 100 uL using RNase free water (ex: if sample volume = 20 uL RNA + 10 uL DMSO, add 70 uL RNase free water)
2. Centrifuge 'full speed' == 21000 rcf or highest speed on centrifuge
3. Store at -80 °C or place on ice and proceed to first strand synthesis

- 6 First strand synthesis - Superscript IV (in BSC):

Calculate reagent volumes for MM:

A	B
Reagent	Vol for 1 rxn
dNTPs	1 uL
Random hexamers	1 uL
RNase free water	4 uL

Step 1

A	B
Reagent	Vol for 1 rxn
5x SSIV buffer	4 uL
100 mM DTT	1 uL
RNase out *	1 uL
SSIV RT	1 uL

Step 2: (* = RNase out instead of RNase inhibitor)

1. Mix appropriate volume (# samples+ 1) of 1st step mastermix and add 6 uL of MM to each tube
Keep MM tube on ice
Start with RNase free water
2. Add 7 uL of each RNA eluate to a tube strip and label + record scheme in notebook
keep on cooling tray, do not allow to freeze
Total vol of each tube is 13 uL
Minifuge
3. Place on thermal cycle, start incubation at 65°C for 5 minutes. At end of 5 min, place on ice for at least 1 min
4. During 5 min incubation, mix appropriate volume (# samples+ 1) of 2nd step mastermix.
Get SSIV RT on cold block from freezer
Keep MM tube on ice
Mix SSIV buffer
Add SSIV RT last (pipette slowly)

5. Add 7 uL (vol for 1 rxn) of MM to each sample on the tube strip
 - Keep tube strip on cooling tray
 - Total vol of each tube is now 20 uL
 - Minifuge
6. Place back on thermal cycler and proceed thru the following incubation steps:
 - 10 min at 23 °C
 - 10 min at 52.5 °C
 - 10 min at 80 °C
7. Store at 4 °C or place reactions on ice and proceed to second strand synthesis.

7 Second strand synthesis.

Calculate reagent volumes for MM:

A	B
Reagent	Vol for 1 rxn
dNTPs	1 uL
E. Coli DNA ligase	0.5 uL
DNA polymerase I	2 uL
RNase H	0.5 uL
5x second strand synthesis buffer (OR 10x second strand synthesis buffer)	8uL(4uL)
Nuclease free water	8 uL (use 12 uL if used 10x buffer)

1. Mix appropriate volume (# samples+ 1) of second strand mastermix.
 - Keep MM tube on ice
 - Keep enzymes on cold block
 - Start with nuclease free water
 - Add enzymes last

2. Add 20 uL (vol for 1 rxn) of MM to each sample on the tube strip
 - Keep tube strip on cooling tray
 - Total vol of each tube is now 20 uL
 - Minifuge

3. Place on thermal cycler and begin 5 hour incubation at 16 °C

4. Store at 4 °C or proceed to cleanup

- 8 Purify DNA using your preferred method (we used a Macharey-Nagal Nucleospin Gel and PCR Clean-up kit, per manufacturer instructions). Quantify DNA using Qubit.
- 9 Prepare DNA libraries for sequencing using your chosen platform and library preparation kit (we used Nextera XT, we will modify this protocol for Oxford Nanopore Native Barcode sequencing).