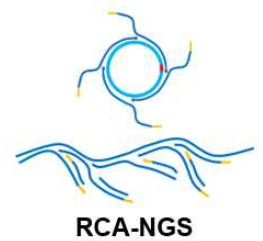


Feb 10, 2023

RCA-NGS for RNA viruses

DOI

<https://dx.doi.org/10.17504/protocols.io.rm7vzb7wrvx1/v1>



Masayasu Misu^{1,2}, Tomoki Yoshikawa¹, Satoko Sugimoto¹, Yuki Takamatsu¹, Takeshi Kurosu¹, Yukiteru Ouji², Masahide Yoshikawa², Masayuki Shimojima¹, Hideki Ebihara¹, Masayuki Saijo¹

¹Department of Virology I, National Institute of Infectious Diseases;

²Department of Pathogen, Infection and Immunity, Nara Medical University



Masayasu Misu

Department of Virology I, National Institute of Infectious D...

Create & collaborate more with a free account

Edit and publish protocols, collaborate in communities, share insights through comments, and track progress with run records.

Create free account

OPEN  ACCESS



DOI: <https://dx.doi.org/10.17504/protocols.io.rm7vzb7wrvx1/v1>

Protocol Citation: Masayasu Misu, Tomoki Yoshikawa, Satoko Sugimoto, Yuki Takamatsu, Takeshi Kurosu, Yukiteru Ouji, Masahide Yoshikawa, Masayuki Shimojima, Hideki Ebihara, Masayuki Saijo 2023. RCA-NGS for RNA viruses. **protocols.io**

<https://dx.doi.org/10.17504/protocols.io.rm7vzb7wrvx1/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: January 29, 2023

Last Modified: February 10, 2023

Protocol Integer ID: 76026

Keywords: Oxford Nanopore Technology, RNA virus, Sequence method, MinION, Nanopore sequencing, RCA-NGS, ngs for rna virus, amplification of the viral genomic cdna, viral genome sequence determination, nucleic acid amplification with virus, whole rna viral genome, terminal of viral genome sequence determination, viral genome, viral genome by nuclease treatment, viral genomic cdna, rna virus, isothermal dna amplification technique, isothermal dna amplification, nucleic acid amplification, physical viral particle enrichment, whole rna, rna, rna other than the rna, segmented genome, virus, nucleic acid, genome, genome sequence, specific pcr primer, removal of unwanted dna, amplification, rca, such as rolling circle amplification

Abstract

This RCA-NGS were optimized for an NGS machine, MinION. These methods do not require nucleic acid amplification with virus-specific PCR primers, physical viral particle enrichment, and RACE.

These methods enable whole RNA viral genome sequencing by combining the following techniques:

- 1) removal of unwanted DNA and RNA other than the RNA viral genome by nuclease treatment
- 2) the terminal of viral genome sequence determination by barcoded linkers ligation
- 3) Amplification of the viral genomic cDNA using an isothermal DNA amplification technique, such as rolling circle amplification (RCA).

This method can be exploited to determine any whole RNA viral genomes (i.e., single-stranded, double-stranded, positive-stranded, negative-stranded, non-segmented or multi-segmented genomes).



Materials

-  Micrococcal Nuclease - 320,000 gel units **New England Biolabs Catalog #M0247S**
-  High Pure Viral RNA Kit **Roche Catalog #11858882001**
-  Turbo DNA-free Kit **Invitrogen - Thermo Fisher Catalog #AM1907**
- **NucleoSpin RNA Clean-up XS - Takara, Catalog #740903.10**
-  T4 RNA Ligase 2, truncated KQ - 2,000 units **New England Biolabs Catalog #M0373S**
- **The barcode-polyA linker DNA (e.g., The cSP6-polyA linker DNA)**
-  Superscript IV **Thermo Fisher Scientific Catalog #18090050**
- **SP6 primer (e.g., 5' phosphorylated SP6 primer)**
-  Deoxynucleotide (dNTP) Solution Mix **New England Biolabs Catalog #N0447S**
-  Superscript-In RNase Inhibitor **ThermoFisher Catalog #AM2694**
- **Dr.GentLE Precipitation Carrier - Takara Catalog #9094**
-  RNase H - 250 units **New England Biolabs Catalog #M0297S**
-  Agencourt AMPure XP **Beckman Coulter Catalog #A63880**
- **CircLigase II ssDNA Ligase - Biosearch Technologies Catalog #CL9021K**
- **GenomiPhi V3 Ready-To-Go DNA Amplification Kit - Cytiva Catalog #25-6601-24**
-  T7 Endonuclease I - 250 units **New England Biolabs Catalog #M0302S**
-  NEBNext FFPE DNA Repair Mix - 24 rxns **New England Biolabs Catalog #M6630S**
-  NEBNext Ultra II End Repair/dA-Tailing Module - 24 rxns **New England Biolabs Catalog #E7546S**
-  Blunt/TA Ligase Master Mix - 50 rxns **New England Biolabs Catalog #M0367S**
-  NEBNext Quick Ligation Module - 20 rxns **New England Biolabs Catalog #E6056S**
- **Ligation Sequencing Kit - Oxford Nanopore Technologies Catalog #SQK-LSK109**
- **Native Barcoding Expansion - Oxford Nanopore Technologies Catalog #EXP-NBD104, #EXP-NBD114**
-  Qubit 4 Fluorometer **Thermo Fisher Scientific Catalog #Q33238**
-  Qubit 1X dsDNA HS Assay Kit **Thermo Fisher Scientific Catalog #Q33230**
-  DNA LoBind Tube 1.5ml **Eppendorf Catalog #022431021**
-  0.2 ml PCR Tube strips **Eppendorf Catalog #0030124359**
- **100 % ethanol**
- **70 % ethanol**
- **TE(pH8.0)**
- **nuclease-free H₂O**



Protocol materials

- ⊗ Micrococcal Nuclease - 320,000 gel units **New England Biolabs Catalog #M0247S**
- ⊗ Qubit 1X dsDNA HS Assay Kit **Thermo Fisher Scientific Catalog #Q33230**
- ⊗ T7 Endonuclease I - 250 units **New England Biolabs Catalog #M0302S**
- ⊗ RNase H - 250 units **New England Biolabs Catalog #M0297S**
- ⊗ Blunt/TA Ligase Master Mix - 50 rxns **New England Biolabs Catalog #M0367S**
- ⊗ NEBNext Quick Ligation Module - 20 rxns **New England Biolabs Catalog #E6056S**
- ⊗ Qubit 4 Fluorometer **Thermo Fisher Scientific Catalog #Q33238**
- ⊗ Deoxynucleotide (dNTP) Solution Mix **New England Biolabs Catalog #N0447S**
- ⊗ NEBNext Ultra II End Repair/dA-Tailing Module - 24 rxns **New England Biolabs Catalog #E7546S**
- ⊗ Agencourt AMPure XP **Beckman Coulter Catalog #A63880**
- ⊗ Turbo DNA-free Kit **Invitrogen - Thermo Fisher Catalog #AM1907**
- ⊗ T4 RNA Ligase 2, truncated KQ - 2,000 units **New England Biolabs Catalog #M0373S**
- ⊗ Superscript IV **Thermo Fisher Scientific Catalog #18090050**
- ⊗ Superase-In RNase Inhibitor **ThermoFisher Catalog #AM2694**
- ⊗ High Pure Viral RNA Kit **Roche Catalog #11858882001**
- ⊗ NEBNext FFPE DNA Repair Mix - 24 rxns **New England Biolabs Catalog #M6630S**
- ⊗ DNA LoBind Tube 1.5ml **Eppendorf Catalog #022431021**
- ⊗ 0.2 ml PCR Tube strips **Eppendorf Catalog #0030124359**

Safety warnings

- ⚠ Follow your facility's regulations and biosafety practices.

Before start

This method was only confirmed to work with the working stocks that contain isolated RNA viruses at least 3.0×10^5 TCID₅₀ per ml.

It is recommended to check no bacterial contamination(e.g., *Mycoplasma* spp.).



Preparation for virus supernatant

- 1 Centrifuge the working stock virus to remove debris.

6000 x g, Room temperature, 00:10:00

10m

- 2 Transfer 180 μ L virus supernatant to a 1.5ml screw cap tube.

- 3 Unwanted DNA and RNA mainly originating from the virus-infected cells are digested using



Micrococcal Nuclease - 320,000 gel units **New England Biolabs** Catalog #M0247S

.

- 3.1 Total 201 μ l reaction

1h

- 180 μ L virus supernatant
- 20 μ L 10X Micrococcal Nuclease Reaction Buffer
- 1 μ L Micrococcal nuclease

Mix by pipetting and spin down.

37 °C water bath 01:00:00

The viral genomic RNA extraction

- 4 The viral genomic RNA extraction is performed using



High Pure Viral RNA Kit **Roche** Catalog #11858882001 .

- 4.1 Add 400 μ L of binding buffer (with 4 μ L PolyA carrier RNA).

10m

Mix gently by ~5 times pipetting and flicking thoroughly the tube, and spin down.

Room temperature 00:10:00

- 4.2 Transfer the sample to a High Pure Filter Tube.

1m

8000 x g, Room temperature, 00:01:00



Discard the flow-through liquid and Collection Tube, and insert the Filter Tube into a new Collection Tube.

- 4.3 Add  500 μL of inhibitor removal bo transfer the sample to a High Pure Filter Tube.

1m

 8000 x g, Room temperature, 00:01:00

Discard the flow-through liquid and Collection Tube, and insert the Filter Tube into a new Collection Tube.

- 4.4 Add  450 μL of wash buffer.

1m

 8000 x g, Room temperature, 00:01:00

Discard the flow-through liquid and Collection Tube, and insert the Filter Tube into a new Collection Tube.

- 4.5 Add  450 μL of wash buffer.

1m

 13000 x g, Room temperature, 00:01:00 and discard the flow-through liquid.

Discard the Collection Tube and insert the Filter Tube into a 1.5 ml tube -

 DNA LoBind Tube 1.5ml **Eppendorf Catalog #022431021** .

- 4.6 Add  50 μL Elution Buffer.

1m

 13000 x g, Room temperature, 00:01:00

Note

The eluted RNA can be stored at -80°C .

Remove unwanted DNA

- 5 Unwanted DNA mainly from the virus-infected cells in the RNA sample is digested using a  Turbo DNA-free Kit **Invitrogen - Thermo Fisher Catalog #AM1907** .

- 5.1 Total 56 μL reaction

30m

-  50 μL the eluted RNA
-  5 μL 10X reaction buffer



-  1 μL DNase I

Mix gently by pipetting and spin down.

 37 °C  00:30:00

- 6 The viral RNA is purified using **NucleoSpin RNA Clean-up XS - Takara, Catalog #740903.10**.

- 6.1 Add an equal volume  56 μL of Buffer RCU and mix gently.

- 6.2 Transfer the sample to a NucleoSpin RNA XS Column.

 11000 x g, Room temperature, 00:01:00

1m

- 6.3 Wash the column by  400 μL Buffer RA3.

 11000 x g, Room temperature, 00:01:00

1m

Discard the flow-through liquid and Collection Tube, and insert the NucleoSpin RNA XS Column into a new Collection Tube.

- 6.4 Wash the column by  200 μL Buffer RA3.

 11000 x g, Room temperature, 00:02:00

2m

Discard the flow-through liquid and Collection Tube, and insert the NucleoSpin RNA XS Column into a Nuclease-free Collection Tube(1.5 ml).

- 6.5 Add  10 μL RNase-free H_2O .

 11000 x g, Room temperature, 00:01:00

1m

Transfer the sample to a 0.2 ml PCR tube -

 0.2 ml PCR Tube strips **Eppendorf Catalog #0030124359** .

cSP6-polyA Linker DNA ligation

7

The viral RNA is ligated with cSP6-polyA Linker DNA using

 T4 RNA Ligase 2, truncated KQ - 2,000 units **New England Biolabs Catalog #M0373S**

.



The RNA is ligated to the 3' end with the barcoded (complementary sequence of SP6 (cSP6)) polyA linker DNA. It is able to identify the 3' terminal viral genome sequence. The PolyA sequence is required for reverse transcription for ONT kit (SQK-PBK004/ PCS109).

Note

The cSP6-polyA linker DNA (5'-5rApp-CTATAGTGTACCTAAATCAAAAAAAAAAAAAAAAAAAAA-3ddC-3'), which is pre-adenylated at the 5' terminal (5rApp), and consists of the complementary sequence of SP6 (CTATAGTGTACCTAAATC), oligo (dA) 20, and dideoxycytidine (3ddC) at the 3' terminal, was synthesised for 3' linker ligation by Integrated DNA Technologies (Coralville, IA).

7.1 Total 20 µl reaction

15m

-  10 µL Purified RNA
-  1 µL 10 µM the cSP6-polyA linker DNA
-  2 µL 10X T4 RNA Ligase Reaction Buffer
-  6 µL 50% PEG8000 solution
-  1 µL T4 RNA Ligase 2, truncated KQ

Mix gently by pipetting and spin down.

Incubate  25 °C  00:15:00

8 The linker-ligated viral RNA is purified using **NucleoSpin RNA Clean-up XS - Takara, Catalog #740903.10**

 [go to step #6](#)

Fill the sample to 100 µl with 80 µl TE (pH 8.0) and add 100 µl (equal volume) of Buffer RCU.

Eluted the RNA in  10 µL RNase-free H₂O and transfer the sample to a 0.2 ml PCR tube.

Reverse transcription



9 The viral RNA is reverse transcribed using

Superscript IV Thermo Fisher Scientific Catalog #18090050 .

5' phosphorylated SP6 primer is used for reverse transcription.

Note

SP6 primer (5' phosphorylated SP6 primer); 5' [Phos]GATTTAGGTGACACTATAG 3'
5' phosphorylation is due to circularization.

9.1 Set up pre-mixture

6m

- 10 μ L RNA (~ 50ng)
- 1 μ L 50 μ M SP6 primer
- 1 μ L nuclease-free H₂O
- 1 μ L 10mM dNTP -

Deoxynucleotide (dNTP) Solution Mix New England Biolabs Catalog #N0447S

Mix gently by flicking the tube, and spin down.

65 °C 00:05:00 and 4 °C 00:01:00

9.2 Total 20 μ l reaction

20m

- 13 μ L pre-mixture sample
- 4 μ L 5X SSIV Buffer
- 1 μ L 100mM DTT
- 1 μ L RNase OUT -
- Superscript-III RNase Inhibitor Thermofisher Catalog #AM2694
- 1 μ L SuperScript IV Reverse Transcriptase

Mix gently by flicking the tube, and spin down.

55 °C 00:10:00



80 °C

00:10:00

RNase H treatment and ethanol precipitation

20m

10 Add 1 µL RNase H - 250 units **New England Biolabs Catalog #M0297S** .

20m

37 °C

00:20:00

11 Ethanol precipitation using **Dr.GenTLE Precipitation Carrier - Takara Bio Catalog #9094**.

11.1 Add 20 µL TE(pH8.0) to fill up the sample to 40 µL .

11.2 Add 4 µL 3M CH₃COONa (pH5.2) .

11.3 Add 1 µL Dr.GenTLE Precipitation Carrier.

11.4 Add 100 µL 100% ethanol.

11.5 13000 x g, Room temperature, 00:15:00

15m

11.6 Discard the supernatant.

11.7 Wash the pellet with 500 µL 70% ethanol.

5m

13000 x g, Room temperature, 00:05:00

11.8 Discard the supernatant and dry.

11.9 Dissolve the pellet in 12 µL nuclease-free H₂O.



(Optional step) Short cDNA fragment removal instead of Ethanol precipitation (#11)

12 Short cDNA fragment is removed from the viral RNA sample using

 Agencourt AMPure XP **Beckman Coulter Catalog #A63880**

Prepare AMPure XP beads for use; resuspend by vortexing.

Transfer amplified DNA sample to 1.5ml low binding tube.

Note

If a significant proportion of the reads obtained from an NGS run fail to match with the NCBI-nr database (i.e., no hits), it could indicate a large number of short cDNA fragments in the sample. In such instances, re-performing the optional step instead of step 11, such as ethanol precipitation could significantly enhance the outcomes.

12.1 Add  36 μL (X1.8 volume) AMPure XP reagent and mix by pipetting.

Incubate on rotor mixer.

 00:05:00

 Room temperature

12.2 Spin down and pellet on a magnet.

Wait for  00:01:00 and pipette off the supernatant.

12.3 Wash twice by  200 μL 70 % ethanol and remove the ethanol using a pipette and discard.

12.4 Spin down and pipette off any residual ethanol.

12.5 Resuspend pellet in  20 μL TE(pH 8.0).

 37 °C

 00:03:00

and tapping occasionally.

Incubate on a rotor mixer.

 00:07:00

12.6 Spin down and pellet the beads on the magnet until the elute is clear and colourless.

12.7 Remove retain  20 μL elute into a new tube.



- 13 Size selection of the cDNA sample is performed using  Agencourt AMPure XP **Beckman Coulter Catalog #A63880** .
X0.8 volume of AMPure beads recovers more than 200 bp of nucleic acids.
- 13.1 Add  16 μL (X0.8 volume) AMPure beads and mix by pipetting.
Incubate on rotor mixer.
 00:05:00  Room temperature
- 13.2 Spin down and pellet on a magnet.
Wait for  00:01:00 and pipette off the supernatant.
- 13.3 Wash twice by  200 μL 70 % ethanol and remove the ethanol using a pipette and discard.
- 13.4 Spin down and pipette off any residual ethanol.
- 13.5 Resuspend pellet in  12 μL nuclease-free water.
 37 °C  00:03:00 and tapping occasionally.
Incubate on a rotor mixer.
 00:07:00
- 13.6 Spin down and pellet the beads on the magnet until the elute is clear and colourless.
- 13.7 Remove retain  12 μL elute into a new tube.

Circularization of cDNA

1h 10m

- 14 The cDNA is circularized using **CircLigase II ssDNA Ligase - Biosearch Technologies Catalog #CL9021K**.
- 14.1 Total 20 μL reaction 1h 10m
-  12 μL cDNA
 -  2 μL 10X reaction buffer
 -  1 μL 50 mM MnCl_2
 -  4 μL 5M Betaine



- 1 μL CircLigase II

Mix by pipetting and spin down.

60 °C 01:00:00

80 °C 00:10:00

15 Ethanol precipitation using **Dr.GenTLE Precipitation Carrier - Takara Catalog #9094.**

go to step #11

Dissolve the pellet in 10 μL nuclease-free H_2O .

Amplification of cDNA by rolling circle amplification (RCA)

16 cDNA is amplified by Rolling circle amplification (RCA) using **GenomiPhi V3 Ready-To-Go DNA Amplification Kit - Cytiva Catalog #25-6601-24.**

16.1 Total 20 μl reaction

3m

- 10 μL cDNA
- 10 μL 2X denaturation buffer

Mix by pipetting and spin down.

95 °C 00:03:00

4 °C on ice

16.2 Add 20 μl denatured sample to Ready to go GenomiPhi cake.

4h 10m

30 °C 04:00:00

65 °C 00:10:00

17 The cDNA is purified by

Agencourt AMPure XP **Beckman Coulter Catalog #A63880**

Prepare AMPure XP beads for use; resuspend by vortexing.

Transfer amplified DNA sample to 1.5ml low binding tube.

17.1 Add 36 μL (X1.8 volume) AMPure beads and mix by pipetting.

Incubate on rotor mixer.



00:05:00

Room temperature

- 17.2 Spin down and pellet on a magnet and wait for 00:01:00 and pipette off the supernatant.
- 17.3 Wash twice by 200 μL 70 % ethanol and remove the ethanol using a pipette and discard.
- 17.4 Spin down and pipette off any residual ethanol.
- 17.5 Resuspend pellet in 40 μL nuclease-free H_2O .
 37 $^{\circ}\text{C}$ 00:03:00 and tapping occasionally.
Incubate on a rotor mixer.
 00:07:00
- 17.6 Spin down and pellet the beads on the magnet until the elute is clear and colourless.
- 17.7 Remove retain 40 μL elute into a new tube.
- 18 DNA concentration is measured using a Qubit 4 Fluorometer with
 Qubit 1X dsDNA HS Assay Kit **Thermo Fisher Scientific Catalog #Q33230** .
- 199 μL 1X working solution
 - 1 μL DNA
- Mix by vortexing.
- Incubate 00:02:00 Room temperature and measure.

Note

Confirm the total amplified cDNA to be over 1500 ng, as confirmed using, for instance, a Qubit 4 Fluorometer and Qubit 1X dsDNA HS Assay Kit.



T7 endonuclease treatment

19 The amplified cDNA by RCA is digested using

 T7 Endonuclease I - 250 units **New England Biolabs Catalog #M0302S** to remove branching.

The following protocol is modified based on the Native barcoding amplicons (with EXP-NBD104, EXPNBD114, and SQK-LSK109) protocol (NBA_9093_v109_revA_12Nov2019) provided by Oxford Nanopore Technologies website.

19.1 Total 30 µl reaction

30m

-  x µL (1.0 µg) DNA
-  3 µL NEBuffer 2
-  1.5 µL T7 endonuclease I
-  25-x µL nuclease-free H₂O

Mix by pipetting and spin down.

 37 °C  00:30:00

20 The cDNA is purified using

 Agencourt AMPure XP **Beckman Coulter Catalog #A63880** .

 (Add  54 µL (X1.8 volume) AMPure beads)

Resuspend pellet in  24 µL nuclease-free H₂O.

DNA repair and end-prep

21 The purified cDNA is end-prepped using

 NEBNext FFPE DNA Repair Mix - 24 rxns **New England Biolabs Catalog #M6630S**

and

 NEBNext Ultra II End Repair/dA-Tailing Module - 24 rxns **New England Biolabs Catalog #E7546S**

21.1 Total 30 µl reaction

35m

-  24 µL DNA
-  1.75 µL NEB Next FFPE DNA repair buffer
-  1 µL NEB Next FFPE DNA repair Mix



- 1.75 µL Ultra II end-prep reaction buffer
- 1.5 µL Ultra II end-prep reaction Mix

Mix by pipetting and spin down.

20 °C 00:30:00

65 °C 00:05:00

22 The cDNA is purified using

Agencourt AMPure XP **Beckman Coulter Catalog #A63880** .

(Add 54 µL (X1.8 volume) AMPure beads)

Resuspend pellet in 30 µL nuclease-free H₂O.

23 DNA concentration is measured using a Qubit 4 Fluorometer with

Qubit 1X dsDNA HS Assay Kit **Thermo Fisher Scientific Catalog #Q33230** .

2m

- 199 µL 1X working solution
- 1 µL DNA

Mix by vortexing.

Incubate 00:02:00 Room temperature and measure.

Note

Confirm the purified cDNA to be approximately 700 ng or more using, for instance, Qubit 4 Fluorometer with a Qubit 1X dsDNA HS Assay Kit.

Native barcode ligation

24 The end-prepped cDNA is ligated with native barcode using **Native Barcoding Expansion - Oxford Nanopore Technologies Catalog #EXP-NBD104, #EXP-NBD114 and**

Blunt/TA Ligase Master Mix - 50 rxns **New England Biolabs Catalog #M0367S** .

24.1 Total 25 µl reaction

20m

- x µL DNA(about 400ng)



- 1.5 µL native barcode
- 12.5 µL Blunt/TA ligase master mix
- 11-x µL nuclease-free H₂O

Mix by pipetting and spin down.

25 °C 00:20:00

25 Add 25 µL TE(pH8.0).

26 The cDNA is purified using

Agencourt AMPure XP **Beckman Coulter Catalog #A63880** .

26.1 Add 40 µL (X0.8 volume) AMPure XP reagent and mix by pipetting.

Incubate on a rotor mixer.

00:05:00 Room temperature

26.2 Spin down and pellet on a magnet. wait for 00:01:00 .

Pipette off the supernatant.

26.3 Wash twice by 200 µL 70 % ethanol and remove the ethanol using a pipette and discard.

26.4 Spin down and pipette off any residual ethanol.

26.5 Resuspend pellet in 20 µL nuclease-free water.

37 °C 00:03:00 and tapping occasionally.

Incubate on a rotor mixer.

00:07:00

26.6 Spin down and pellet the beads on the magnet until the elute is clear and colourless.

26.7 Remove retain 20 µL elute into a new tube.



27 DNA concentration is measured using a Qubit 4 Fluorometer with

Qubit 1X dsDNA HS Assay Kit **Thermo Fisher Scientific Catalog #Q33230** .

- 199 μL 1X working solution
- 1 μL DNA

Mix by vortexing.

Incubate 00:02:00 Room temperature and measure.

Convert nanogram(ng) into femtomole(fmol) by a calculator.

Note

The molar concentration of the cDNA sample can be converted based on the length of the major band confirmed by electrophoresis after T7 endonuclease treatment. Typically, the fragment lengths are around 2000 bases pairs.

Adaptor ligation

20m

28 Pool each barcoded sample into a 0.2ml PCR tube (Total 100–200 fmol).

29 Adaptor Ligation with pooled samples is performed using
Ligation Sequencing Kit - Oxford Nanopore Technologies Catalog #SQK-LSK109,
Native Barcoding Expansion - Oxford Nanopore Technologies Catalog #EXP-NBD104,
#EXP-NBD114 and

NEBNext Quick Ligation Module - 20 rxns **New England Biolabs Catalog #E6056S**

.

29.1 Total 50 μL reaction

20m

- x μL DNA (100–200 fmol)
- 2.5 μL Adaptor Mix II
- 10 μL NEB Next Quick Ligation Reaction Buffer(5X)
- 5 μL Quick T4 DNA ligase
- 32.5-x μL nuclease-free H_2O



mix gently and incubate.



25 °C



00:20:00

30 The adaptor-ligated cDNA is purified using

Agencourt AMPure XP **Beckman Coulter Catalog #A63880** .

Prepare AMPure XP beads for use; resuspend by vortexing.

Transfer amplified DNA sample to 1.5ml low binding tube.

30.1 Add  80 µL AMPure XP reagent and mix by pipetting.

Incubate on a rotor mixer.



00:05:00



Room temperature

30.2 Spin down and pellet on a magnet. Wait for  00:01:00 and pipette off the supernatant.

30.3

- Wash twice by  200 µL . Short Fragment Buffer(SFB) and remove the ethanol using a pipette and discard.

30.4 Spin down and pipette off any residual SFB.

30.5

- Resuspend pellet in  12 µL Elution Buffer (EB)



37 °C



00:03:00

and tapping occasionally.

Incubate on a rotor mixer.



00:07:00

30.6 Spin down and pellet the beads on the magnet until the elute is clear and colourless.

30.7 Remove retain  12 µL elute into a new tube.

31 DNA concentration is measured using a Qubit 4 Fluorometer with

Qubit 1X dsDNA HS Assay Kit **Thermo Fisher Scientific Catalog #Q33230** .

-  199 µL 1X working solution

-  1 µL DNA



Mix by vortexing.

Incubate  00:02:00  Room temperature and measure.

Sequencing by MinION

32 Sequencing according to the manufacturer's instructions.