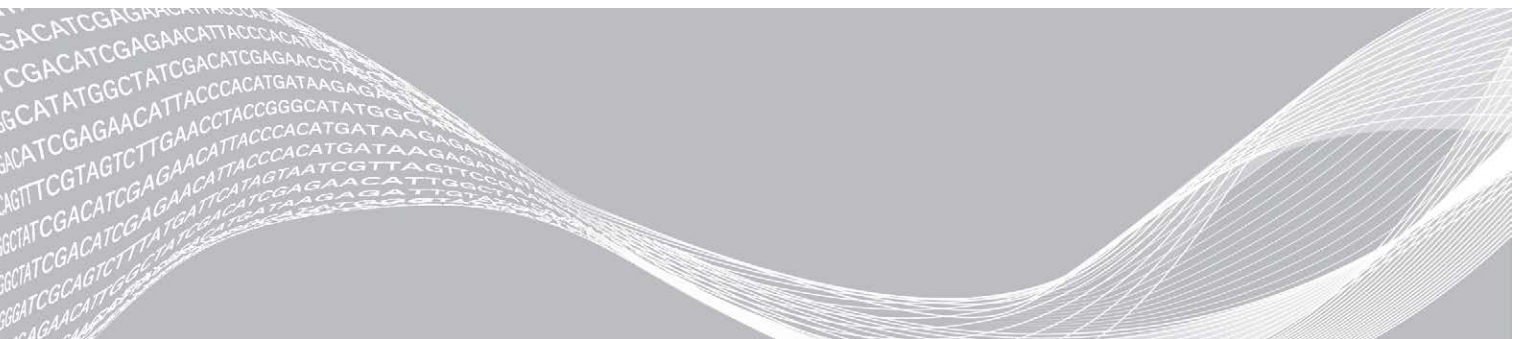


Illumina COVIDSeq RUO Kits

Reference Guide



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Revision History

| Document | Date | Description of Change |
|---------------------------------|---------------|---|
| Document # 1000000126053 v06 | July 2021 | <p>Added information for compatibility with the iSeq 100 System to appropriate topics, including the Introduction, Pool and Dilute Libraries, and the Prepare for Sequencing sections.</p> <p>Moved sample sheet information for Illumina COVIDSeq Test (RUO) to an appendix.</p> <p>Updated Illumina COVIDSeq Assay naming conventions and kit configurations to reflect current on-market use.</p> <p>Reorganized Prepare for Sequencing section for improved ease of use.</p> <p>Removed redundant sequencing consumables information from Prepare for Sequencing section. Information remains in Consumables and Equipment list in an appendix.</p> |
| Document # 1000000126053 v05 | June 2021 | <p>Renamed document Illumina COVIDSeq RUO Kits Reference Guide to account for use with the 96-sample Illumina COVIDSeq Assay.</p> <p>Added information for the 96-sample Illumina COVIDSeq Assay to the Introduction, Extract DNA, Pool and Clean Up Libraries, Pool and Dilute Libraries, and Prepare for Sequencing sections.</p> <p>Added information for compatibility with the MiSeq and MiniSeq Sequencing Systems to the Introduction, Pool and Dilute Libraries, and the Prepare for Sequencing sections.</p> <p>Added new Kit Contents section to Appendix A for the 96-sample Illumina COVIDSeq Assay.</p> <p>Moved COVIDSeq Positive Control (CPC) preparation instructions to new Appendix B and updated for Illumina COVIDSeq Assay.</p> |
| Document # 1000000126053 v04 | April 2021 | <p>Added information about variant analysis software options to the Introduction and Prepare for Sequencing section.</p> <p>Added reference to technical note for performing library prep multiple times for a low throughput batch size.</p> <p>Updated the Illumina COVIDSeq Test (3072 samples) from an eight box configuration to a four box configuration in the Kit Contents section.</p> <p>Updated the Read Length recommendations in the Prepare for Sequencing section.</p> <p>Updated Thermal Cycler recommendations for confirmation of microtiter plate compatibility.</p> |

| Document | Date | Description of Change |
|---------------------------------|------------------|---|
| Document # 1000000126053 v03 | February 2021 | <p>Added instructions to set up analysis in BaseSpace Sequence Hub for NextSeq 2000 and other NextSeq 2000-specific information throughout the protocol.</p> <p>Added Thermal Cycler recommendations.</p> <p>Updated materials, consumables, and equipment part numbers in the Prepare for Sequencing, Product Components, and Consumables and Equipment sections, including new consumables for the NextSeq 2000.</p> <p>Updated version numbers of flow cells and control software for the NovaSeq 6000 system.</p> <p>Updated the tube for Dilution 3 in Extract RNA to 5 ml LoBind tube.</p> <p>Updated the protocol options for Quick-DNA/RNA Viral MagBead Procedure for precision and clarity.</p> <p>Updated temperature in PCR program on the thermal cycler for Amplify cDNA Preparation from 65°C to 63°C.</p> |
| Document # 1000000126053 v02 | July 2020 | <p>Added instructions for extracting RNA using the Quick-DNA/RNA Viral Magbead kit.</p> <p>Added safe stopping point after pooling and cleaning up libraries.</p> <p>Updated index kit configurations to IDT for Illumina-PCR Indexes.</p> <p>Removed sequencing instructions.</p> <p>Added dilution and sequencing preparation instructions for the NovaSeq 6000 Sequencing System SP flow cell, NextSeq 500 Sequencing System, NextSeq 550 Sequencing System, and NextSeq 550Dx Instrument.</p> <p>Moved data analysis information to <i>Illumina COVIDSeq Test Pipeline Software Guide document # 1000000128122</i>.</p> |
| Document # 1000000126053 v01 | June 2020 | No content changes. |
| Document # 1000000126053 v00 | June 2020 | Initial release. |

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Chapter 1 Overview

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Introduction

This guide explains how to detect the SARS-CoV-2 virus using either of two different research use only kits: the Illumina COVIDSeq Test (RUO) or the Illumina COVIDSeq Assay (96 Samples). The Illumina COVIDSeq Test supports sample processing for high throughput (HT) sequencing, while the Illumina COVIDSeq Assay is oriented toward sample processing for low throughput (LT) sequencing.

The Illumina COVIDSeq Test offers preparation of up to 3072 samples using the NovaSeq 6000 Sequencing System or up to 384 samples using the NextSeq 500/550 Sequencing Systems, NextSeq 550Dx Instrument in RUO mode, or NextSeq 2000 Sequencing System. The Illumina COVIDSeq Assay offers preparation of up to 96 samples using the iSeq 100 Sequencing System, MiSeq Sequencing System or MiniSeq Sequencing System. [Table 1](#) displays some of the differences between these two kit options.

Table 1 Comparison of Illumina COVIDSeq (RUO) Kit Options

| Kit Name | Samples | Recommended Systems | Indexes | Positive Control |
|------------------------------|------------|--|--------------|------------------|
| Illumina COVIDSeq Test (RUO) | Up to 3072 | NovaSeq 6000, NextSeq 500/550, Next Seq 550Dx in RUO mode, or NextSeq 2000 | Not Included | Included |
| Illumina COVIDSeq Assay | Up to 96 | iSeq 100, MiSeq, or MiniSeq | Included | Not Included |

Both the Illumina COVIDSeq Test (RUO) and the Illumina COVIDSeq Assay (96 Samples) support:

- ▶ RNA extraction from decontaminated nasopharyngeal (NP), oropharyngeal (OP), and nasal swab samples, as well as mid-turbinate specimens collected from individuals who meet COVID-19 clinical or epidemiological criteria using an RNA extraction kit. Recommended kits include the QIAamp Viral RNA Mini Kit and the Quick-DNA/RNA Viral Magbead Kit.
- ▶ Qualitative detection of SARS-CoV-2 RNA using either the Illumina DRAGEN COVIDSeq Test Pipeline locally or with the Illumina DRAGEN COVIDSeq Test app on BaseSpace Sequence Hub.

With either kit, you can also perform surveillance of characteristics of the SARS-CoV-2 viral genome using either the DRAGEN COVID Pipeline with the COVID Lineage Tools locally or with the DRAGEN COVID Lineage app on BaseSpace Sequence Hub.

Input Recommendations

The Illumina COVIDSeq Test (RUO) and Illumina COVIDSeq Assay (96 Samples) support patient samples derived from nasopharyngeal (NP), oropharyngeal (OP), and nasal swabs. Transport samples according to the governing regulations for the transport of etiologic agents applicable to your region.

Store samples according to the instructions from the manufacturer of tubes used for sample transport. Exceeding the storage times can negatively impact test results.

The following sample factors might affect SARS-CoV-2 detection:

- ▶ Sample collection methods, patient factors, and/or the stage of the infection.
- ▶ Viral RNA degradation during shipping and storage. RNA degradation can produce false-negative results.



CAUTION

Handle all specimens as infectious reagents.

Chapter 2 Library Prep

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Introduction

This chapter describes library preparation using either the Illumina COVIDSeq Test (RUO) or the Illumina COVIDSeq Assay (96 Samples).

- ▶ Confirm kit contents and make sure that you have the required equipment and consumables. See [Supporting Information on page 17](#).



NOTE

Reagents in the Illumina COVIDSeq Test (RUO) kit have HT on their labels to indicate use for high throughput sequencing. Reagents in the Illumina COVIDSeq Assay (96 Samples) do not.

- ▶ Follow the protocols in the order shown, using the specified volumes and incubation parameters.
- ▶ Make sure that reagents are not expired. Using expired reagents might negatively affect performance.
- ▶ When using the Illumina COVIDSeq Test (RUO) for detection, include one no template control (NTC) and one positive control per 96-well plate. The COVIDSeq Positive Control (CPC) is included in the Illumina COVIDSeq Test (RUO). See [COVIDSeq Positive Control Preparation on page 24](#) for preparation instructions.
- ▶ For the Illumina COVIDSeq Assay (96 Samples) or when using the Illumina COVIDSeq Test (RUO) for surveillance, an NTC and a positive control are recommended for quality control but are not required. The Illumina COVIDSeq Assay (96 Samples) does not include a positive control, which you can purchase separately if necessary.
- ▶ If performing library prep multiple times with the Illumina COVIDSeq Test (RUO), refer to the *Aliquot Procedure for Illumina COVIDSeq Test (RUO version) Kit Reagents* Illumina Technical Note.
- ▶ Do not allow multiple freeze-thaw cycles of the CPC. If performing library prep multiple times, aliquot CPC into low-bind tubes, and then store at -85°C to -65°C.
- ▶ For all other reagents besides the CPC, do not allow more than eight freeze-thaw cycles.
- ▶ Sequence libraries as soon as possible after pooling. Pooled libraries are stable for up to 30 days at -25°C to -15°C.

Tips and Techniques

Unless a safe stopping point is specified in the protocol, proceed immediately to the next step.

Avoiding Contamination

- ▶ Use proper laboratory practices to prevent nuclease and PCR product contamination. Nuclease and PCR product contamination can cause inaccurate and unreliable results.
- ▶ Perform library preparation in a RNase/DNase-free environment. Thoroughly decontaminate work areas with a RNase/DNase-inhibiting solution, such as RNaseZap and DNAZap.
- ▶ Use fresh tips and fresh consumable labware between samples and dispensing reagents.
- ▶ Use aerosol-resistant tips to reduce the risk of carry over and sample to sample cross contamination.
- ▶ Due to the potential for contamination, take extreme care to make sure that well contents remain fully in the well. Do not splash contents.
- ▶ Do not use aerosol bleach sprays when performing library preparation. Trace bleach contamination can lead to assay failure.
- ▶ Use a unidirectional workflow when moving from pre-amplification to post-amplification environments.
- ▶ One or more no template controls (NTCs) are recommended per plate to monitor contamination.

Sealing and Unsealing the Plate

- ▶ Always seal the 96-well plate before the following steps in the protocol:
 - ▶ Shaking steps
 - ▶ Vortexing steps
 - ▶ Centrifuge steps
 - ▶ Thermal cycling steps
- ▶ To seal the plate, apply the adhesive cover to the plate and then seal with a wedge or rubber roller.
- ▶ Make sure the edges and wells are completely sealed to reduce the risk of cross-contamination and evaporation.
- ▶ Microseal 'B' adhesive seals are effective at -40°C to 110°C, and suitable for skirted or semiskirted PCR plates. Use Microseal 'B' for shaking, centrifuging, and long-term storage.
- ▶ Before unsealing:
 - ▶ Briefly centrifuge the 96-well plate at 1000 × g for 1 minute. For bead steps, centrifuge at 500 × g for 1 minute.
 - ▶ Place the plate on a flat surface before slowly removing the seal.

Plate Transfers

- ▶ When transferring volumes between plates, transfer the specified volume from each well of a plate to the corresponding well of the other plate.
- ▶ If beads are aspirated into the pipette tips, dispense back to the plate on the magnetic stand and wait until the liquid is clear (~2 minutes).

Centrifugation

- ▶ Centrifuge as needed at any step in the procedure to consolidate liquid or beads in the bottom of the well, and to prevent sample loss.

Handling Beads

- ▶ Pipette bead suspension slowly to prevent splashing and bubbles.

- ▶ When mixing, mix thoroughly.
- ▶ To avoid sample loss, confirm that no beads remain in pipette tips after resuspension and mixing steps.
- ▶ When washing beads:
 - ▶ Use the appropriate magnet for the plate.
 - ▶ Dispense liquid so that beads on the side of the wells are wetted.
 - ▶ Keep the plate on the magnet until the instructions specify to remove it.
 - ▶ Do not agitate the plate while on the magnetic stand. Do not disturb the bead pellet.

Extract RNA

This step extracts RNA from decontaminated viral transport medium tubes. You can extract RNA using the Quick-DNA/RNA Viral MagBead, Zymo Research, part # R2141 or the QIAamp Viral RNA Mini Kit, Qiagen, part # 52906. Follow the procedure corresponding to your extraction method.

If you plan to use the COVIDSeq Positive Control, make sure to follow the appropriate preparation procedure in *COVIDSeq Positive Control Preparation* on page 24.

Consumables

- ▶ [QIAamp Viral RNA Mini Kit] 1.7 ml LoBind tubes
- ▶ [Quick-DNA/RNA Viral MagBead] 2000 µl 96 deep well plate

Quick-DNA/RNA Viral MagBead Procedure

- 1 For each sample, add 400 µl patient sample to a new deep-well plate.
If you plan to use controls, include one tube of dilution 3 CPC (positive control) and ELB (no template control) per sample batch.
- 2 To extract RNA, use the Quick-DNA/RNA Viral MagBead. For information, see *Quick-DNA/RNA Viral MagBead Instruction Manual* from Zymo Research.
Use the following protocol options:
 - ▶ Before adding MagBinding Beads, pipette up and down ten times to mix.
 - ▶ After adding 20 µl MagBinding Beads, pipette up and down ten times to mix, and then shake at 1500 rpm for 10 minutes.

QIAamp Viral RNA Mini Kit Procedure

- 1 For each sample, add 140 µl patient sample to new 1.7 ml microcentrifuge tube.
If you plan to use controls, include one tube of dilution 3 CPC (positive control) and ELB (no template control) per sample batch.
- 2 To extract RNA, use the QIAamp Viral RNA Mini Kit. For information, see *QIAamp Viral RNA Mini Handbook (document # HB-0354-006)* available on the QIAGEN website.
Use the following protocol options:
 - ▶ Purify viral RNA using the spin protocol.
 - ▶ Incubate elution for at least 1 minute.
 - ▶ Elute in 30 µl Buffer AVE instead of 60 µl.

Anneal RNA

During this process the extracted RNA is annealed using random hexamers to prepare for cDNA synthesis.

If you plan to use the COVIDSeq Positive Control and have not yet prepared the control, make sure to follow the appropriate procedure in [COVIDSeq Positive Control Preparation on page 24](#).

Consumables

- ▶ EPH3 (Elution Prime Fragment 3HC Mix)
- ▶ 96-well PCR Plate
- ▶ Microseal 'B' adhesive seals

About Reagents

- ▶ Vortex before each use

Preparation

- 1 Prepare the following consumables:

| Reagent | Storage | Instructions |
|---------|----------------|---|
| EPH3 | -25°C to -15°C | Thaw at room temperature, and then invert to mix. |

- 2 Save the following COVIDSeq Anneal program on the thermal cycler:
 - ▶ Choose the preheat lid option
 - ▶ Set the reaction volume to 17 μ l
 - ▶ 65°C for 3 minutes
 - ▶ Hold at 4°C

Procedure

- 1 Label new PCR plate CDNA1.
- 2 Add 8.5 μ l EPH3 to each well.
- 3 Add 8.5 μ l eluted sample to each well.
- 4 Seal and shake at 1600 rpm for 1 minute.
- 5 Centrifuge at 1000 \times g for 1 minute.
- 6 Place on the preprogrammed thermal cycler and run the COVIDSeq Anneal program.

Synthesize First Strand cDNA

This step reverse transcribes the RNA fragments primed with random hexamers into first strand cDNA using reverse transcriptase.

Consumables

- ▶ FSM (First Strand Mix)
- ▶ RVT (Reverse Transcriptase)
- ▶ 1.7 ml tubes (1 per 96-well sample plate)
- ▶ Microseal 'B' adhesive seal

Preparation

- 1 Prepare the following consumables:

| Reagent | Storage | Instructions |
|---------|----------------|--|
| FSM | -25°C to -15°C | Thaw and bring to room temperature. Invert to mix, and then keep on ice. |
| RVT | -25°C to -15°C | Invert to mix before use. Keep on ice. |

- 2 Save the following COVIDSeq FSS program on the thermal cycler:
 - ▶ Choose the preheat lid option
 - ▶ Set the reaction volume to 25 µl
 - ▶ 25°C for 5 minutes
 - ▶ 50°C for 10 minutes
 - ▶ 80°C for 5 minutes
 - ▶ Hold at 4°C

Procedure

- 1 In a 1.7 ml tube, combine the following volumes to prepare First Strand cDNA Master Mix. Multiply each volume by the number of samples.
 - ▶ FSM (9 µl)
 - ▶ RVT (1 µl)
 Reagent overage is included to account for small pipetting errors.
- 2 Add 8 µl master mix to each well of the CDNA1 plate.
- 3 Seal and shake at 1600 rpm for 1 minute.
- 4 Centrifuge at 1000 × g for 1 minute.
- 5 Place on the preprogrammed thermal cycler and run the COVIDSeq FSS program.

SAFE STOPPING POINT

If you are stopping, seal the plate and store at -25°C to -15°C for up to 7 days.

Amplify cDNA

This step uses two separate PCR reactions to amplify cDNA.

Consumables

- ▶ IPM (Illumina PCR Mix)
- ▶ CPP1 (COVIDSeq Primer Pool 1)
- ▶ CPP2 (COVIDSeq Primer Pool 2)
- ▶ Nuclease-free water
- ▶ 15 ml tube (2 for four 96-well sample plates)
- ▶ 96-well PCR plates (2)
- ▶ Microseal 'B' adhesive seal

Preparation

- 1 Prepare the following consumables:

| Reagent | Storage | Instructions |
|---------|----------------|--|
| CPP1 | -25°C to -15°C | Thaw at room temperature. Keep on ice until use. |
| CPP2 | -25°C to -15°C | Thaw at room temperature. Keep on ice until use. |
| IPM | -25°C to -15°C | Thaw at room temperature, and then invert to mix. Keep on ice until use. |

- 2 Save the following COVIDSeq PCR program on the thermal cycler:

- ▶ Choose the preheat lid option
- ▶ Set the reaction volume to 25 μ l
- ▶ 98°C for 3 minutes
- ▶ 35 cycles of:
 - ▶ 98°C for 15 seconds
 - ▶ 63°C for 5 minutes
- ▶ Hold at 4°C

Procedure

- 1 Label two new PCR plates COV1 and COV2.
The plates represent two separate PCR reactions on each sample and control in the CDNA1 plate.
- 2 In a 15 ml tube, combine the following volumes to prepare COVIDSeq PCR 1 Master Mix and COVIDSeq PCR 2 Master Mix. Multiply each volume by the number of samples.
Reagent overage is included to account for small pipetting errors.

| Reagent | COVIDSeq PCR 1 Master Mix (μ l) | COVIDSeq PCR 2 Master Mix (μ l) |
|---------------------|--------------------------------------|--------------------------------------|
| IPM | 15 | 15 |
| CPP1 | 4.3 | N/A |
| CPP2 | N/A | 4.3 |
| Nuclease-free water | 4.7 | 4.7 |

- 3 Add 20 μ l COVIDSeq PCR 1 Master Mix to each well of the COV1 plate corresponding to each well of the CDNA1 plate.
- 4 Add 5 μ l first strand cDNA synthesis from each well of the CDNA1 plate to the corresponding well of the COV1 plate.
- 5 Add 20 μ l COVIDSeq PCR 2 Master Mix to each well of the COV2 plate corresponding to each well of the CDNA1 plate.
- 6 Add 5 μ l first strand cDNA synthesis from each well of the CDNA1 plate to the corresponding well of the COV2 plate.
- 7 Seal and shake at 1600 rpm for 1 minute.
- 8 Centrifuge at 1000 x g for 1 minute.
- 9 Place in the preprogrammed thermal cycler and run the COVIDSeq PCR program.

SAFE STOPPING POINT

If you are stopping, seal the plate and store at -25°C to -15°C for up to 3 days.

Tagment PCR Amplicons

This step uses EBLTS to tagment PCR amplicons, which is a process that fragments and tags the PCR amplicons with adapter sequences.

Consumables

- ▶ EBLTS (Enrichment BLT)
- ▶ TB1 (Tagmentation Buffer 1)
- ▶ Nuclease-free water
- ▶ 1.7 ml tube
- ▶ 15 ml tube (1 per four 96-well sample plates)
- ▶ 96-well PCR plate
- ▶ Microseal 'B' adhesive seal

About Reagents

- ▶ Store EBLTS upright at temperatures above 2°C. Make sure beads are always submerged in the buffer.
- ▶ If beads are adhered to the side or top of the 96-well plate, centrifuge at 500 × g for 1 minute, and then pipette to resuspend.

Preparation

- 1 Prepare the following consumables:

| Reagent | Storage | Instructions |
|---------|----------------|--|
| EBLTS | 2°C to 8°C | Bring to room temperature. Vortex thoroughly before use. |
| TB1 | -25°C to -15°C | Bring to room temperature. Vortex thoroughly before use. |

- 2 If COV1 and COV2 plates were stored frozen, prepare as follows.
 - a Thaw at room temperature.
 - b Check seals, and then shake at 1600 rpm for 1 minute.
 - c Centrifuge at 1000 × g for 1 minute.
- 3 Save the following COVIDSeq TAG program on the thermal cycler:
 - ▶ Choose the preheat lid option
 - ▶ Set the reaction volume to 50 µl
 - ▶ 55°C for 5 minutes
 - ▶ Hold at 10°C

Procedure

- 1 Label a new PCR plate TAG1.
- 2 Combine COV1 and COV2 as follows.

- a Transfer 10 μ l from each well of the COV1 plate to the corresponding well of the TAG1 plate.
 - b Transfer 10 μ l from each well of the COV2 plate to each well of the TAG1 plate containing COV1.
- 3 In a 15 ml tube, combine the following volumes to prepare Tagmentation Master Mix. Multiply each volume by the number of samples.
 - ▶ TB1 (12 μ l)
 - ▶ EBLTS (4 μ l)
 - ▶ Nuclease-free water (20 μ l)
 - 4 Add 30 μ l master mix to each well in TAG1 plate.
 - 5 Seal and shake at 1600 rpm for 1 minute.
 - 6 Place on the preprogrammed thermal cycler and run the COVIDSeq TAG program.

Post Tagmentation Clean Up

This step washes the adapter-tagged amplicons before PCR amplification.

Consumables

- ▶ ST2 (Stop Tagment Buffer 2)
- ▶ TWB (Tagmentation Wash Buffer)
- ▶ Microseal 'B' adhesive seal

About Reagents

- ▶ Dispense ST2 and TWB slowly to minimize foaming.
- ▶ Dispense TWB directly onto beads.

Preparation

- 1 Prepare the following consumables:

| Reagent | Storage | Instructions |
|---------|------------------|--------------------|
| ST2 | Room temperature | Vortex before use. |
| TWB | 2°C to 8°C | Vortex before use. |

Procedure

- 1 Centrifuge the TAG1 plate at 500 x g for 1 minute.
- 2 Add 10 μ l ST2 to each well of the TAG1 plate.
- 3 Seal and shake at 1600 rpm for 1 minute.
- 4 Incubate at room temperature for 5 minutes.
- 5 Centrifuge at 500 x g for 1 minute.
- 6 Place on the magnetic stand and wait until the liquid is clear (~3 minutes).
- 7 Inspect for bubbles on the seal. If present, centrifuge at 500 x g for 1 minute, and then place on the magnetic stand (~3 minutes).
- 8 Remove and discard all supernatant.

- 9 Wash beads as follows.
 - a Remove from the magnetic stand.
 - b Add 100 µl TWB to each well.
 - c Seal and shake at 1600 rpm for 1 minute.
 - d Centrifuge 500 × g for 1 minute.
 - e Place on the magnetic stand and wait until the liquid is clear (~3 minutes).
 - f For first wash only, remove and discard all supernatant from each well.
- 10 Wash beads a **second** time.
Leave supernatant in plate for second wash to prevent beads from overdrying.

Amplify Tagmented Amplicons

This step amplifies the tagmented amplicons using a PCR program. The PCR step adds prepared 10 base pair Index 1 (i7) adapters, Index 2 (i5) adapters, and sequences required for sequencing cluster generation.

Consumables

- ▶ EPM (Enhanced PCR Mix)
- ▶ Index adapters (IDT for Illumina-PCR Indexes Set 1, 2, 3, 4)
- ▶ Nuclease-free water
- ▶ 15 ml tubes (1 per two 96-well sample plates)
- ▶ 96-well PCR plate

About Reagents

- ▶ Index adapter plates
 - ▶ Do not add samples to the index plate wells.
 - ▶ Index plate wells are considered single use and should not be reused.

Preparation

- 1 Prepare the following consumables:

| Reagent | Storage | Instructions |
|----------------|----------------|--|
| EPM | -25°C to -15°C | Invert to mix. Keep on ice until use. |
| Index adapters | -25°C to -15°C | Thaw at room temperature. Vortex to mix, and then centrifuge at 1000 × g for 1 minute. |

- 2 Open each prepared index adapter plate seal as follows. Use a new PCR plate for each different index set.
 - a Align a new 96-well PCR plate above the index adapter plate, and then press down to puncture the foil seal.
 - b Discard the PCR plate.

- 3 Save the following COVIDSeq TAG PCR program on the thermal cycler:
 - ▶ Choose the preheat lid option and set to 100°C
 - ▶ Set the reaction volume to 50 µl
 - ▶ 72°C for 3 minutes
 - ▶ 98°C for 3 minutes
 - ▶ 7 cycles of:
 - ▶ 98°C for 20 seconds
 - ▶ 60°C for 30 seconds
 - ▶ 72°C for 1 minute
 - ▶ 72°C for 3 minutes
 - ▶ Hold at 10°C

Procedure

- 1 In a 15 ml tube, combine the following volumes to prepare PCR Master Mix. Multiply each volume by the number of samples.
 - ▶ EPM (24 µl)
 - ▶ Nuclease-free water (24 µl)
- 2 Vortex PCR Master Mix to mix.
- 3 Keep the TAG1 plate on magnetic stand and remove TWB.
- 4 Use a 20 µl pipette to remove any remaining TWB.
- 5 Remove the TAG1 plate from the magnetic stand.
- 6 Add 40 µl PCR Master Mix to each well.
- 7 Add 10 µl index adapters to each well of the PCR plate.
- 8 Seal and shake at 1600 rpm for 1 minute.
- 9 If liquid is visible on the seal, centrifuge at 500 x g for 1 minute.
- 10 Inspect to make sure beads are resuspended. To resuspend, set your pipette to 35 µl with the plunger down, and then slowly pipette to mix.
- 11 Place on the preprogrammed thermal cycler and run the COVIDSeq TAG PCR program.

Pool and Clean Up Libraries

This step combines libraries from each 96-well sample plate into one 1.7 ml tube. Libraries of optimal size are then bound to magnetic beads, and fragments that are too small or large are washed away.

Consumables

- ▶ ITB (Illumina Tune Beads)
- ▶ RSB (Resuspension Buffer)
- ▶ Freshly prepared 80% ethanol (EtOH)
- ▶ 1.7 ml tube (2 per 96-well sample plate)
- ▶ [Illumina COVIDSeq Test (RUO)] PCR 8-tube strip

About Reagents

- ▶ ITB
 - ▶ Vortex before each use.
 - ▶ Vortex frequently to make sure that beads are evenly distributed.
 - ▶ Aspirate and dispense slowly due to the viscosity of the solution.

Preparation

- 1 Prepare the following consumables:

| Reagent | Storage | Instructions |
|---------|------------------|--|
| ITB | Room temperature | Vortex thoroughly to mix. |
| RSB | 2°C to 8°C | Let stand for 30 minutes to bring to room temperature. Vortex and invert to mix. |

- 2 Prepare 2.5 ml 80% EtOH from absolute EtOH for each tube of pooled libraries.

Procedure

- 1 Centrifuge the TAG1 plate at 500 × g for 1 minute.
- 2 Place on the magnetic stand and wait until the liquid is clear (~3 minutes).
- 3 To pool libraries, complete the following steps appropriate for your kit. Repeat the steps for each additional sample plate.
 - a Label a new 1.7 ml tube Pooled ITB.
 - b **[Illumina COVIDSeq Assay (96 Samples)]** Transfer 5 µl library from each well of the TAG1 plate into the Pooled ITB tube.
 - c **[Illumina COVIDSeq Test (RUO)]** Use a 20 µl eight-channel pipette to transfer 5 µl library from each well of the TAG1 plate to a PCR 8-tube strip, resulting in 60 µl pooled library per row. Change tips after each column.
 - d **[Illumina COVIDSeq Test (RUO)]** Transfer 55 µl pooled library from each well of the PCR 8-tube strip into the Pooled ITB tube. For each sample plate, these volumes results in 440 µl pools of pooled libraries.
- 4 Vortex the Pooled ITB tubes to mix, and then centrifuge briefly.
- 5 Vortex ITB to resuspend.
- 6 Add ITB using the resulting volume of Pooled ITB tube volume multiplied by 0.9. For example, for 96 samples, add 396 µl ITB to each tube.
- 7 Vortex to mix.
- 8 Incubate at room temperature for 5 minutes.
- 9 Centrifuge briefly.
- 10 Place on the magnetic stand and wait until the liquid is clear (~5 minutes).
- 11 Remove and discard all supernatant.
- 12 Wash beads as follows.
 - a Keep on the magnetic stand and add 1000 µl fresh 80% EtOH to each tube.
 - b Wait 30 seconds.

- c Remove and discard all supernatant.
- 13 Wash beads a **second** time.
- 14 Use a 20 µl pipette to remove all residual EtOH.
- 15 Add 55 µl RSB.

**NOTE**

Due to library yield excess, the RSB volume does not impact batches with a small number of samples.

- 16 Vortex to mix, and then centrifuge briefly.
- 17 Incubate at room temperature for 2 minutes.
- 18 Place on the magnetic stand and wait until the liquid is clear (~2 minutes).
- 19 Transfer 50 µl supernatant from each Pooled ITB tube to a new microcentrifuge tube.

SAFE STOPPING POINT

If you are stopping, cap the tube and store at -25°C to -15°C for up to 30 days.

Quantify and Normalize Libraries

- 1 Analyze 2 µl library pool using a Qubit dsDNA HS Assay kit.
If libraries are outside the standard range, dilute to 1:10 concentration, and analyze again.
- 2 Calculate the molarity value using the following formula.
 - ▶ Use 400 bp as the average library size.

$$\frac{\text{Library concentration ng/}\mu\text{l}}{660 \frac{\text{g}}{\text{mol}} \times \text{average library size (bp)}} \times 10^6 = \text{Molarity (nM)}$$

- 3 Dilute each library pool to a minimum of 30 µl at a normalized concentration 4 nM using RSB.

Pool and Dilute Libraries

After diluting to the starting concentration of 4 nM, libraries are ready to be denatured and diluted to the final loading concentration.

- 1 Transfer the designated volume of normalized libraries containing the appropriate index adapter sets to a new microcentrifuge tube for each number of samples specified in [Table 2](#).
If you have multiple normalized pools, combine the designated volume of each normalized pool in the tube. Doing so produces a final pool of samples diluted to a starting concentration of 4 nM. Do not combine pools with the same index adapter set.

Table 2 Normalized Pool Volumes and Sample Numbers for Denature and Dilution by Instrument

| Sequencing System | Volume of Normalized Libraries Used per Run (µl) | Samples per Final Pool of Normalized Libraries | Samples per Flow Cell |
|-----------------------------|--|--|-----------------------|
| iSeq 100 v1 or v2 Flow Cell | 2 | 4 | 4 |
| MiSeq v2 Flow Cell | 5 | 15 | 15 |
| MiSeq v3 Flow Cell | 5 | 24 | 24 |

| Sequencing System | Volume of Normalized Libraries Used per Run (µl) | Samples per Final Pool of Normalized Libraries | Samples per Flow Cell |
|---------------------------------------|--|--|----------------------------------|
| MiniSeq HO Flow Cell | 25 | 24 | 24 |
| NextSeq 500/550 or 550Dx HO Flow Cell | 25 | 384 | 384 |
| NovaSeq 6000 SP Flow Cell | 25 | 384 | 384 per lane, 768 per flow cell |
| NovaSeq 6000 S4 Flow Cell | 25 | 384 | 384 per lane, 1536 per flow cell |
| NextSeq 1000/2000 P2 Flow Cell | 25 | 384 | 384 |

- Follow the denature and dilute instructions for your system to dilute to the final loading concentration.
 - ▶ For the iSeq Sequencing System, see the *iSeq 100 Sequencing System Guide (document # 1000000036024)*.
 - ▶ For the MiSeq Sequencing System, see the *MiSeq System Denature and Dilute Libraries Guide (document # 15039740)*.
 - ▶ For the MiniSeq Sequencing System, see the *MiniSeq System Denature and Dilute Libraries Guide (document # 1000000002697)*.
 - ▶ For the NextSeq 500/550 Sequencing System and NextSeq 550Dx Sequencing System, see the *NextSeq System Denature and Dilute Libraries Guide (document # 15048776)*.
 - ▶ For the NovaSeq 6000 Sequencing System, see the *NovaSeq 6000 Denature and Dilute Libraries Guide (document # 1000000106351)*.
 - ▶ For the NextSeq 2000 Sequencing System, see the *NextSeq 1000/2000 Sequencing System Guide (document # 1000000109376)*.
- Use the following loading concentrations for your system.

Table 3 Loading Concentrations by Instrument

| Sequencing System | Starting Concentration (nM) | Final Loading Concentration (pM) |
|---------------------------------------|-----------------------------|----------------------------------|
| iSeq 100 v1 or v2 Flow Cell | 4 | 75 |
| MiSeq v2 Flow Cell | 4 | 10 |
| MiSeq v3 Flow Cell | 4 | 12 |
| MiniSeq HO flow cell | 4 | 1.2 |
| NextSeq 500/550 or 550Dx HO flow cell | 4 | 1.4 |
| NovaSeq 6000 SP Flow Cell | 4 | 100 |
| NovaSeq 6000 S4 Flow Cell | 4 | 100 |
| NextSeq 1000/2000 P2 Flow Cell | 4 | 1000 |

Adjustments to final loading concentration should follow the denature and dilute instructions for your sequencing system.

Prepare for Sequencing

The Illumina COVIDSeq Assay (96 Samples) is compatible with the iSeq 100 i1 reagents v2, the MiSeq reagent kits v2 and v3, and the MiniSeq High Output (HO) reagent kit.

The Illumina COVIDSeq Test (RUO) is compatible with the NovaSeq 6000 Sequencing System SP and S4 flow cells, the NextSeq 2000 Sequencing System, the NextSeq 500/550 Sequencing Systems, and the NextSeq 550DX instrument.

Use the `samplesheet.csv` file for your sequencing system available on the Illumina COVIDSeq Research Use Only (RUO) Kits support site as a template to create the sample sheet. For specific sample sheet requirements for the Illumina DRAGEN COVIDSeq Test Pipeline, refer to *Illumina DRAGEN COVIDSeq Test Pipeline Sample Sheet on page 26*.

Set Up Sequencing Run for Illumina COVIDSeq Assay (96 Samples)

Refer to the documentation for your sequencing system and the following information to set up your run.

- ▶ For read length recommendations, refer to the Illumina Technical Note *Sequencing Guidelines for COVID-19 Surveillance Using the Illumina COVIDSeq Test (RUO Version)* for guidance.
- ▶ If using the iSeq System, refer to the *iSeq System Guide (document # 1000000036024)*.
- ▶ If using the MiSeq, refer to the *MiSeq System Guide (document # 15046563)*.
- ▶ If using the MiniSeq System, refer to the *MiniSeq System Guide (document # 1000000002695)*.
- ▶ Set up your sequencing run in manual mode unless you are performing surveillance and use Local Run Manager.
 - ▶ If using Local Run Manager, refer to the *Local Run Manager Software Guide (document # 1000000111492)*.
- ▶ If using a BaseSpace Sequence Hub app, make sure to enable monitoring and storage as needed for your instrument.
 - ▶ For the iSeq System, select or enable **Run Analysis, Collaboration, and Storage** in the system settings.
 - ▶ For the MiniSeq System, select **Run Monitoring and Storage** as the Configuration option.
- ▶ Enter **Paired End** as the Read Type.
- ▶ Enter **10** as the value for Index 1 and Index 2.

Set Up Sequencing Run for Illumina COVIDSeq Test (RUO)

Refer to the documentation for your sequencing system and the following information to set up your run.

For read length recommendations, refer to the Illumina Technical Note *Sequencing Guidelines for COVID-19 Surveillance Using the Illumina COVIDSeq Test (RUO Version)* for guidance.

- 1 If using the NextSeq 500/550 or NexSeq 550Dx, refer to the *NextSeq 500 System Guide (document # 15046563)*, *NextSeq 550 System Guide (document # 15069765)*, or *NextSeq 550Dx Instrument Reference Guide (document # 1000000009513)*.
 - ▶ Use v4.0 of the NextSeq Control Software (NCS).
 - ▶ If using the NextSeq 550Dx, use RUO mode.
 - ▶ Set up your sequencing run in manual mode unless you are performing surveillance and use Local Run Manager.
 - ▶ If using Local Run Manager, refer to the *Local Run Manager Software Guide (document # 1000000111492)*.
 - ▶ If using a BaseSpace Sequence Hub app, select **Run Monitoring and Storage** as the Configuration option.
 - ▶ Enter **Single-Read** as the Read Type. For surveillance, enter **Paired End** as the Read Type.
 - ▶ Enter **10** as the value for Index 1 and Index 2.
- 2 If using the NovaSeq 6000 system, refer to the *NovaSeq 6000 Sequencing System Guide (document # 1000000019358)* for sequencing instructions.
 - ▶ Use v1.7 of the NovaSeq Control Software (NVCS).
 - ▶ If using a BaseSpace Sequence Hub app, select **Run Monitoring and Storage** as the Configuration option.

- ▶ Enter **10** as the value for Index 1 and Index 2.
- 3 If using the NextSeq 2000, refer to the *NextSeq 1000/2000 Sequencing System Guide (document # 1000000109376)*.
 - ▶ When creating a run in BaseSpace Sequence Hub, make sure to do the following:
 - ▶ Select **BaseSpace** for analysis location.
 - ▶ Select **Illumina DRAGEN COVIDSeq Test** for analysis type.
 - ▶ If Illumina DRAGEN COVIDSeq Test does not appear as an analysis type, contact Illumina Technical Support.
 - ▶ Set up the analysis as described in the following *Set Up Analysis in BaseSpace Sequence Hub for NextSeq 2000* section.
 - ▶ Use v1.2 of the NextSeq 1000/2000 Control Software.
 - ▶ Make sure **Online Run Setup** and **Proactive, Run Monitoring, and Storage** are selected in the Settings screen to enable Cloud mode.

Set Up Analysis in BaseSpace Sequence Hub for NextSeq 2000

Use the following steps to configure the Illumina DRAGEN COVIDSeq Test analysis in BaseSpace Sequence Hub when using a NextSeq 2000 instrument.

- 1 To enable fast mode, set the Fast Mode option to **True**.
Fast mode turns off alignment, variant calling, and consensus sequence FASTA generation to analyze results.
- 2 To exclude run logs, QC metric files, and other file types, set the Metrics and Logs Datasets option to **False**.
Setting this option to false improves analysis speed, but the Logs_Intermediates_Lane_* folder is not generated.
- 3 Identify the location for your positive and no template controls using either the sample ID or well position.
- 4 Enter the positive control and no template control for each index set.
 - ▶ If you used the index set during library preparation, enter the sample ID or well position for the positive and no template controls.
 - ▶ If you did not use the index set, enter NA.
- 5 Select **Submit Run**.

Analysis Software

After sequencing completes, analysis either takes place locally using installed pipeline software or in BaseSpace Sequence Hub.

- ▶ Local analysis for qualitative detection of SARS-CoV-2 RNA uses the Illumina DRAGEN COVIDSeq Test Pipeline.
- ▶ Local analysis for surveillance uses the Illumina DRAGEN COVID Pipeline with COVID Lineage Tools.
- ▶ Cloud-based analysis in BaseSpace Sequence Hub can use the Illumina DRAGEN COVIDSeq Test for qualitative detection of SARS-CoV-2 or the DRAGEN COVID Lineage app for surveillance.

Refer to one of the following resources for additional information.

- ▶ *Illumina DRAGEN COVIDSeq Test Pipeline Software Guide (document # 1000000128122)*
- ▶ *Illumina DRAGEN COVIDSeq Test App Guide (document # 1000000129548)*
- ▶ *Illumina DRAGEN COVID Pipeline Software Guide (document # 1000000158680)*

Appendix A Supporting Information

| | |
|--|----|
| Illumina COVIDSeq Assay Kit Contents (96 Samples) | 17 |
| Illumina COVIDSeq Test (RUO) Kit Contents (3072 Samples) | 18 |
| Consumables and Equipment | 20 |

Illumina COVIDSeq Assay Kit Contents (96 Samples)

The Illumina COVIDSeq Assay (96 Samples) for low throughput sequencing has four different kit options. Each kit option contains a different set of IDT for Illumina-PCR Indexes.

The Illumina COVIDSeq Assay (96 Samples) does not contain the COVIDSeq Positive Control, but you can purchase it separately. See [COVIDSeq Positive Control for 96 Samples \(Optional\)](#) on page 18

| Kit | Catalog # |
|--|-----------|
| Illumina COVIDSeq Assay (96 Samples) including Index Set 1 | 20049393 |
| Illumina COVIDSeq Assay (96 Samples) including Index Set 2 | 20051772 |

Illumina COVIDSeq Assay (96 Samples)

Promptly store reagents at the indicated temperature to ensure proper performance.

Table 4 Illumina COVIDSeq Assay Box 1 – 96 Samples, Part # 20051272

| Quantity | Label Volume (ml) | Reagent | Description | Storage |
|----------|-------------------|---------|-----------------------|--|
| 1 | 15 | ITB | Illumina Tune Beads | Room temperature, post-amp environment |
| 1 | 2 | ST2 | Stop Tagment Buffer 2 | Room temperature, post-amp environment |

Table 5 Illumina COVIDSeq Assay Box 2 – 96 Samples, Part # 20051273

| Quantity | Label Volume (ml) | Reagent | Description | Storage |
|----------|-------------------|---------|--------------------------|----------------------------------|
| 2 | 2 | EBLTS | Enrichment BLT | 2°C to 8°C, post-amp environment |
| 3 | 2 | ELB | Elution Buffer | 2°C to 8°C, pre-amp environment |
| 2 | 2 | RSB | Resuspension Buffer | 2°C to 8°C, post-amp environment |
| 1 | 50 | TWB | Tagmentation Wash Buffer | 2°C to 8°C, post-amp environment |

Table 6 Illumina COVIDSeq Assay Box 3 – 96 Samples, Part # 20051274

| Quantity | Label Volume (ml) | Reagent | Description | Storage |
|----------|-------------------|---------|--------------------------------|--------------------------------------|
| 1 | 2 | CPP1 | COVIDSeq Primer Pool 1 | -25°C to -15°C, pre-amp environment |
| 1 | 2 | CPP2 | COVIDSeq Primer Pool 2 | -25°C to -15°C, pre-amp environment |
| 4 | 0.5 | EPH3 | Elution Prime Fragment 3HC Mix | -25°C to -15°C pre-amp environment |
| 3 | 2 | EPM | Enhanced PCR Mix | -25°C to -15°C, pre-amp environment |
| 3 | 0.5 | FSM | First Strand Mix | -25°C to -15°C, pre-amp environment |
| 4 | 2 | IPM | Illumina PCR Mix | -25°C to -15°C, pre-amp environment |
| 2 | 0.5 | RVT | Reverse Transcriptase | -25°C to -15°C, pre-amp environment |
| 6 | 0.5 | TB1 | Tagmentation Buffer 1 | -25°C to -15°C, post-amp environment |

Table 7 Illumina COVIDSeq Assay Box 4 – 96 Samples, Indexes

| Quantity | Description | Storage |
|----------|--|----------------|
| 8 | One of the following sets: <ul style="list-style-type: none"> • IDT for Illumina- PCR Indexes Set 1 (96 Indexes) • IDT for Illumina- PCR Indexes Set 2 (96 Indexes) • IDT for Illumina- PCR Indexes Set 3 (96 Indexes) • IDT for Illumina- PCR Indexes Set 4 (96 Indexes) | -25°C to -15°C |

COVIDSeq Positive Control for 96 Samples (Optional)

The COVIDSeq Positive Control (CPC) is optional for the Illumina COVIDSeq Assay (96 Samples). It is sold separately from the Illumina COVIDSeq Assay (96 Samples). Store the CPC at -85°C to -65°C in the pre-amp environment.

| Quantity | Label Volume | Reagent | Part Number |
|----------|--------------|---------------------------|-------------|
| 1 | 100 µl | COVIDSeq Positive Control | 20051775 |

Illumina COVIDSeq Test (RUO) Kit Contents (3072 Samples)

The Illumina COVIDSeq Test (RUO) for high throughput sequencing requires the Illumina COVIDSeq Test (3072 Samples) and 8 IDT for Illumina-PCR Indexes.

| Component | Kit | Catalog # |
|----------------------------|---|-----------|
| Library Preparation | Illumina COVIDSeq Test (3072 Samples) | 20043675 |
| Indexes | IDT for Illumina-PCR Indexes Sets 1–4 (384 Indexes) | 20043137 |

Illumina COVIDSeq Test (RUO)

Promptly store reagents at the indicated temperature to ensure proper performance.

Table 8 Illumina COVIDSeq Test Box 1 – 3072 Samples, Part # 20044405

| Quantity | Label Volume (ml) | Reagent | Description | Storage |
|----------|-------------------|---------|--------------------------|--|
| 1 | 233 | ITB | Illumina Tune Beads | Room temperature |
| 1 | 56 | ST2 HT | Stop Tagment Buffer 2 HT | Room temperature, post-amp environment |

Table 9 Illumina COVIDSeqTest Box 2 – 3072 Samples, Part # 20044406

| Quantity | Label Volume (ml) | Reagent | Description | Storage |
|----------|-------------------|----------|-----------------------------|----------------------------------|
| 2 | 6.1 | EBLTS HT | Enrichment BLT HT | 2°C to 8°C, post-amp environment |
| 1 | 114 | ELB HT | Elution Buffer HT | 2°C to 8°C, pre-amp environment |
| 1 | 10 | RSB HT | Resuspension Buffer HT | 2°C to 8°C, post-amp environment |
| 1 | 845 | TWB HT | Tagmentation Wash Buffer HT | 2°C to 8°C, post-amp environment |

Table 10 Illumina COVIDSeq Test Box 3 – 3072 Samples, Part # 20044407

| Quantity | Label Volume (ml) | Reagent | Description | Storage |
|----------|-------------------|---------|-----------------------------------|--------------------------------------|
| 1 | 14.4 | CPP1 HT | COVIDSeq Primer Pool 1 HT | -25°C to -15°C, pre-amp environment |
| 1 | 14.4 | CPP2 HT | COVIDSeq Primer Pool 2 HT | -25°C to -15°C, pre-amp environment |
| 1 | 45 | EPH3 HT | Elution Prime Fragment 3HC Mix HT | -25°C to -15°C pre-amp environment |
| 1 | 79 | EPM HT | Enhanced PCR Mix HT | -25°C to -15°C, pre-amp environment |
| 1 | 41 | FSM HT | First Strand Mix HT | -25°C to -15°C, pre-amp environment |
| 1 | 100 | IPM HT | Illumina PCR Mix HT | -25°C to -15°C, pre-amp environment |
| 1 | 4.6 | RVT HT | Reverse Transcriptase HT | -25°C to -15°C, pre-amp environment |
| 1 | 38 | TB1 HT | Tagmentation Buffer 1 HT | -25°C to -15°C, post-amp environment |

Table 11 Illumina COVIDSeq Positive Control HT, Part # 20043401

| Quantity | Label Volume | Reagent | Description | Storage |
|----------|--------------|------------------------------|------------------------------|-------------------------------------|
| 1 | 100 µl | COVIDSeq Positive Control HT | COVIDSeq Positive Control HT | -85°C to -65°C, pre-amp environment |

IDT for Illumina- PCR Indexes , Store at -25°C to -15°C

The Illumina COVIDSeq Test (RUO) requires 8 IDT for Illumina PCR Indexes Sets 1–4 (384 Indexes).

| Quantity | Description | Part Number |
|----------|--|-------------|
| 8 | IDT for Illumina- PCR Indexes Set 1 (96 Indexes) | 20043132 |
| 8 | IDT for Illumina- PCR Indexes Set 2 (96 Indexes) | 20043133 |
| 8 | IDT for Illumina- PCR Indexes Set 3 (96 Indexes) | 20043134 |
| 8 | IDT for Illumina- PCR Indexes Set 4 (96 Indexes) | 20043135 |

Consumables and Equipment

In addition to your kit (Illumina COVIDSeq Test (RUO) or Illumina COVIDSeq Assay (96 Samples)) and IDT for Illumina-PCR Indexes, make sure that you have the required consumables and equipment before starting the protocol.

Consumables

| Consumable | Supplier |
|--|---|
| 10 µl pipette tips | General lab supplier |
| 20 µl pipette tips | General lab supplier |
| 200 µl pipette tips | General lab supplier |
| 200 µl pipette tips | General lab supplier |
| 1000 µl pipette tips | General lab supplier |
| Hard-Shell 96-Well PCR Plates | Bio-Rad, catalog # HSP-9601 or equivalent |
| 96 deep-well plate, 2000 µl | Eppendorf, catalog # 951033707 |
| 1.7 ml LoBind microcentrifuge tubes | Eppendorf, catalog # 022431021 |
| 5 ml LoBind microcentrifuge tube | Eppendorf, catalog # 0030122348 |
| 15 ml tubes | General lab supplier |
| Lab tissue, low-lint | VWR, catalog # 21905-026, or equivalent |
| Lint-free alcohol wipe | General lab supplier |
| Microseal 'B' adhesive seals | Bio-Rad, part # MSB-1001 |
| RNase/DNase-free Disposable Pipetting Reservoirs | VWR, part # 89094-658 |
| One of the following, depending on the extraction method used: | <ul style="list-style-type: none"> • Qiagen, catalog # 52906 • Zymo Research, catalog # R2141 |
| Qubit dsDNA HS Assay Kit | One of the following, depending on kit size: <ul style="list-style-type: none"> • ThermoFisher Scientific, part # Q32851 • ThermoFisher Scientific, part # Q32854 |
| Qubit Assay Tubes | ThermoFisher Scientific, catalog # Q32856 |

| Consumable | Supplier |
|--|--|
| If using the iSeq 100 System: • iSeq 100 i1 Reagent v2 (300 Cycles) | • Illumina, catalog # 20031371 |
| If using the MiSeq System v2 reagent kit: • MiSeq Reagent Kit v2 (300 Cycles) | • Illumina, catalog # MS-102-2002 |
| If using the MiSeq System v3 reagent kit: • MiSeq Reagent Kit v3 (600 Cycles) | • Illumina, catalog # MS-102-3003 |
| If using the MiniSeq System: • MiniSeq High Output Reagent Kit (300 Cycles) | • Illumina, catalog # FC-420-1003 |
| If using the NovaSeq 6000 Sequencing System S4 flow cell: • 2 NovaSeq 6000 Sequencing System S4 Reagent Kit v1.5 (35 cycles) • 2 NovaSeq Xp 4-Lane Kit v1.5 | • Illumina, catalog # 20044417 • Illumina, catalog # 20043131 |
| If using the NovaSeq 6000 Sequencing System SP flow cell: • 4 NovaSeq 6000 Sequencing System SP Reagent Kit v1.5 (100 cycles) • 4 NovaSeq Xp 2-Lane Kit v1.5 | • Illumina, catalog # 20028401 • Illumina, catalog # 20043130 |
| If using the NextSeq 500/550 System or the NextSeq 550Dx instrument: • 8 NextSeq 500/550 High Output Kit v2.5 (75 Cycles) | • Illumina, catalog # 20024906 |
| If using the NextSeq 2000 System • 8 NextSeq 1000/2000 P2 Reagents (100 cycles) | • Illumina, catalog # 20046811 |

Equipment Required, Not Provided

| Equipment | Supplier |
|--|--------------------------------|
| 10 µl single-channel pipettes | General lab supplier |
| 20 µl single-channel pipettes | General lab supplier |
| 200 µl single-channel pipettes | General lab supplier |
| 1000 µl single-channel pipettes | General lab supplier |
| 10 µl 8-channel pipettes | General lab supplier |
| 20 µl 8-channel pipettes | General lab supplier |
| 200 µl 8-channel pipettes | General lab supplier |
| 1000 µl 8-channel pipettes | General lab supplier |
| 20 µl 12-channel pipettes | General lab supplier |
| 200 µl 12-channel pipettes | General lab supplier |
| 10 ml serological pipettes | General lab supplier |
| 25 ml serological pipettes | General lab supplier |
| 50 ml serological pipettes | General lab supplier |
| BioShake iQ | QInstruments, part # 1808-0506 |
| DRAGEN Bio-IT Platform or BaseSpace Sequence Hub | Illumina |

| Equipment | Supplier |
|---|--|
| Equipment for one the following extraction methods (as needed): <ul style="list-style-type: none"> • Quick-DNA/RNA Viral MagBead equipment • QIAamp Viral RNA Mini Kit equipment | <ul style="list-style-type: none"> • See <i>Quick-DNA/RNA Viral MagBead Instruction Manual</i>, Zymo Research • See <i>QIAamp Viral RNA Mini Handbook (document # HB-0354-006)</i>, Qiagen |
| Freezer, -25°C to -15°C | General lab supplier |
| Freezer, -85°C to -65°C | General lab supplier |
| Magnetic Stand-96 | Thermo Fisher Scientific, catalog # AM10027 |
| One of the following magnetic stands: <ul style="list-style-type: none"> • Dynabeads MPC-S (Magnetic Particle Concentrator) • MagnaRack Magnetic Separation Rack | <ul style="list-style-type: none"> • Thermo Fisher Scientific, catalog #A13346 • Thermo Fisher Scientific, catalog # CS15000 |
| Microcentrifuge | General lab supplier |
| Microplate Centrifuge | General lab supplier |
| One of the following sequencing systems: <ul style="list-style-type: none"> • iSeq 100 • MiSeq • MiniSeq • NextSeq 500 • NextSeq 550 • NextSeq 550Dx • NextSeq 2000 • NovaSeq 6000 | Illumina |
| NovaSeq Xp Flow Cell Dock | Illumina, # 20021663 |
| Pipette Aid | General lab supplier |
| Qubit Fluorometer 3.0 | Thermo Fisher, catalog # Q33216, Q33217, or Q33218 |
| Refrigerator, 2°C to 8°C | General lab supplier |
| One of the following thermal cyclers: <ul style="list-style-type: none"> • C1000 Touch™ Thermal Cyclers with 96-Well Fast Reaction Module • C1000 Touch™ Thermal Cyclers with 96-Deep Well Reaction Module • Veriti 96-well Thermal Cycler • GeneAmp PCR System 9700 Fast Thermal Cycler • Thermal cycler that meets the minimum specification requirements. See <i>Recommended Thermal Cycler Specifications</i> on page 22 | <ul style="list-style-type: none"> • Bio-Rad, part # 1851196 • Bio-Rad, part # 1851197 • Thermo Fisher, catalog # 4375786 • Thermo Fisher, catalog # 4339386 |
| Sealing wedge or roller | General lab supplier |
| Vortexer | General lab supplier |

Recommended Thermal Cycler Specifications

The following are the recommended minimum requirements for a thermal cycler used in the Illumina COVIDSeq Test (RUO) or the Illumina COVIDSeq Assay (96 Samples). Make sure to also confirm compatibility of your PCR plate with the specific thermal cycler you use.

| Specification | Minimum Requirement |
|------------------------|--|
| Lid type | Heated |
| Temperature range | 4°C to 99°C |
| Format | 0.2 mL tubes, 96-well plate |
| Temperature accuracy | ±0.25°C (35°C to 99.9°C) |
| Temperature uniformity | ±0.5°C well-to-well within 30 seconds of arrival at target temperature |
| Peak ramp rate | At least 1.5°C |
| Sample ramp rate | ± 1.25°C |

Appendix B COVIDSeq Positive Control Preparation

COVIDSeq Positive Control Preparation24

COVIDSeq Positive Control Preparation

This procedure dilutes the COVIDSeq Positive Control (CPC) and prepares it for use with the Illumina COVIDSeq Assay (96 Samples) and Illumina COVIDSeq Test (RUO) kits.

Consumables

- ▶ 1.7 ml LoBind tubes
- ▶ 5 ml LoBind tubes
- ▶ COVIDSeq Positive Control

About Reagents

- ▶ Aliquot CPC into low-bind tubes. Store at -85°C to -65°C.
- ▶ Vortex before each use.

Preparation for Illumina COVIDSeq Assay (96 Samples)

The following steps describe the procedure for the Illumina COVIDSeq Assay (96 Samples) kit. For the Illumina COVIDSeq Test (RUO) kit, refer to *Preparation for Illumina COVIDSeq Test (RUO)* on page 25.

Use of the COVIDSeq Positive Control (CPC) with the Illumina COVIDSeq Assay (96 Samples) is recommended but not required.

- 1 Prepare the following consumables:

| Reagent | Storage | Instructions |
|---------|----------------|--|
| ELB | 2°C to 8°C | Thaw at room temperature, and then invert to mix. Keep on ice until use. |
| CPC | -85°C to -65°C | Dilute to 5 copies per μl using the following instructions. Keep diluted positive control on ice. |

- 2 Dilute CPC as follows.
 - a Label a 1.7 ml tube Dilution 1.
 - b Add the following volumes to the tube *in the order listed*.
 - ▶ CPC (1 μl)
 - ▶ ELB (99 μl)These volumes produce 10000 copies per μl .
 - c Pulse vortex to mix.

- 3 Dilute CPC a second time as follows.
 - a Label a 1.7 ml tube Dilution 2.
 - b Add the following volumes to the tube *in the order listed*.
 - ▶ Dilution 1 (1 µl)
 - ▶ ELB (99 µl)

These volumes produce 100 copies per µl.
 - c Pulse vortex to mix.

Preparation for Illumina COVIDSeq Test (RUO)

The following steps describe the preparation procedure for the Illumina COVIDSeq Test (RUO) kit. For the Illumina COVIDSeq Assay (96 Samples) kit, refer to [Preparation for Illumina COVIDSeq Assay \(96 Samples\) on page 24](#).

Use of the COVIDSeq Positive Control (CPC) HT with the Illumina COVIDSeq Test (RUO) is required for detection and recommended for surveillance.

- 1 Prepare the following consumables:

| Reagent | Storage | Instructions |
|---------|----------------|---|
| ELB HT | 2°C to 8°C | Thaw at room temperature, and then invert to mix. Keep on ice until use. |
| CPC HT | -85°C to -65°C | Dilute to 5 copies per µl using the following instructions. Keep diluted positive control on ice. |

- 2 Dilute CPC HT as follows.
 - a Label a 1.7 ml tube Dilution 1.
 - b Add the following volumes to the tube *in the order listed*.
 - ▶ CPC HT (5 µl)
 - ▶ ELB HT (495 µl)

These volumes produce 10000 copies per µl.
 - c Pulse vortex to mix.
- 3 Dilute CPC HT a second time as follows.
 - a Label a 1.7 ml tube Dilution 2.
 - b Add the following volumes to the tube *in the order listed*.
 - ▶ Dilution 1 (5 µl)
 - ▶ ELB HT (495 µl)

These volumes produce 100 copies per µl.
 - c Pulse vortex to mix.
- 4 Dilute CPC HT a third time as follows.
 - a Label a 5 ml tube Dilution 3.
 - b Add the following volumes to the tube *in the order listed*.
 - ▶ Dilution 2 (200 µl)
 - ▶ ELB HT (3.8 ml)

These volumes produce 5 copies per µl.
 - c Pulse vortex to mix.

Appendix C Illumina DRAGEN COVIDSeq Test Pipeline Sample Sheet

Sample Sheet Requirements 26

Sample Sheet Requirements

The Illumina DRAGEN COVIDSeq Test Pipeline requires a sample sheet for each run analysis. This requirement does not apply to the NextSeq 2000, which uses the Illumina DRAGEN COVIDSeq Test in BaseSpace Sequence Hub.

Use the `samplesheet.csv` file for your sequencing system included in the installer packager or available on the Illumina COVIDSeq Research Use Only (RUO) Kits support site as a template to create the sample sheet.

Make sure your sample sheet meets the following requirements.

- 1 Save the sample sheet with the name `SampleSheet.csv` in the sequencing run folder.
- 2 In Settings, enter the following value for the `AdapterRead1` parameter.

```
CTGTCTCTTATACACATCT
```

- 3 In the Data section, enter the following required parameters.

Make sure that there no empty rows between samples.

| Field | Description | Requirements |
|-----------|--|--|
| Sample_ID | The ID used to identify the samples in the test reports and included in the output file names. | Sample IDs are not case-sensitive. Make sure Sample IDs contain the following: <ul style="list-style-type: none">• Unique for the run.• ≤ 100 characters with no spaces.• Alphanumeric characters, underscores, and dashes only. An alphanumeric character must be added before and after an underscore or dash. |
| Index_ID | The IDT for Illumina-PCR Indexes index name associated with the sample. | See <i>Illumina Adapter Sequences (document # 100000002694)</i> for index names and additional information. The name must be unique for each flow cell lane. If the Index_ID is not specified, the Index Set field is derived from Index and Index2. If specifying all three, the index names and associated sequences must match. |
| Index | IDT for Illumina-PCR Indexes i7 index sample sheet bases | See <i>Illumina Adapter Sequences (document # 100000002694)</i> for sample sheet bases for your sequencing system and additional information. If Index_ID is specified, Index is not required. |
| Index2 | IDT for Illumina-PCR Indexes i5 index sample sheet bases. | See <i>Illumina Adapter Sequences (document # 100000002694)</i> for sample sheet bases for your sequencing system and additional information. If Index_ID is specified, Index2 is not required. |
| Lane | The flow cell lane for the sample. | If using the NovaSeq 6000 System, enter one of the following values: 1, 2, 3, or 4. If using the NextSeq 500/550, NextSeq 500Dx, or NextSeq 2000, this field is not included. |

| Field | Description | Requirements |
|-------------|----------------------------------|--|
| Sample_Type | The sample type for each sample. | Enter one of the following case-sensitive values: PatientSample, NTC, PositiveControl. If using the NovaSeq 6000 System, there must be one NTC sample and one PostiveControl sample for each Index Set/Lane combination in the sample sheet. If using the NextSeq 500/550, NextSeq 500Dx, or NextSeq 2000 there must be one NTC sample and one PostiveControl sample for each Index Set combination in the sample sheet |

- 4 **[Optional]** Enter any additional data parameters, such as `Sample_Name`.
- 5 Save your sample sheet.

Technical Assistance

For technical assistance, contact Illumina Technical Support.

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Safety data sheets (SDSs)—Available on the Illumina website at support.illumina.com/sds.html.

Product documentation—Available for download from support.illumina.com.



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