

# Clarifi COVID-19 Test Kit

For Real Time qRT-PCR Test

## Instructions for Use

Catalog #1105 and 1110

For Use under Emergency Use Authorization Only

Rx Only  
For *in vitro* diagnostic use



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## Intended Use

The Clarifi COVID-19 Test Kit (Clarifi Test) is a real-time qRT-PCR test intended for the qualitative detection of RNA from the SARS-CoV-2 in saliva swab specimens collected from individuals suspected of COVID-19 by their healthcare provider. Saliva specimens are collected by a HCP in a healthcare setting using the ORAcollect•RNA (OR-100) saliva collection device from individuals who are suspected of COVID-19 by their healthcare provider. Testing is limited to laboratories which are certified under the Clinical Laboratory Improvement Amendments of 1988 (CLIA), 42 U.S.C. §263a, and meet the requirements to perform high complexity tests.

Results are for the identification of SARS-CoV-2 RNA. The SARS-CoV-2 RNA is generally detectable in saliva specimens during the acute phase of infection. Positive results are indicative of the presence of SARS-CoV-2 RNA; clinical correlation with patient history and other diagnostic information is necessary to determine patient infection status. Positive results do not rule out bacterial infection or co-infection with other viruses. The agent detected may not be the definite cause of disease. Laboratories within the United States and its territories are required to report all results to the appropriate public health authorities.

Negative results do not preclude SARS-CoV-2 infection and should not be used as the sole basis for patient management decisions. Negative results must be combined with clinical observations, patient history, and epidemiological information. Negative results for SARS-CoV-2 RNA from saliva should be confirmed by testing of an alternative specimen type if clinically indicated.

The Clarifi COVID-19 Test Kit is intended for use by qualified clinical laboratory personnel, specifically instructed and trained in the techniques of qRT-PCR and *in vitro* diagnostic procedures. The Clarifi COVID-19 Test Kit is only for use under the Food and Drug Administration's Emergency Use Authorization.

## Summary and Explanation

An outbreak of pneumonia caused by a novel coronavirus (SAR-CoV-2) in Wuhan City, Hubei Province, China was initially reported to the World Health Organization (WHO) in early December 2019. On January 31, 2020, Health and Human Services Secretary Alex M. Azar II declared a public health emergency (PHE) for the United States to aid the nation's healthcare community in responding to SARS-CoV-2. The emergence and rapid spread of SARS-CoV-2 to numerous areas throughout the world, has necessitated preparedness and response in laboratories, as well as health care and other areas of society in general. The availability of specific and sensitive assays for the detection of the virus are essential for accurate diagnosis of cases, assessment of the extent of the outbreak, monitoring of intervention strategies and surveillance studies.

## Principles of the Procedure

The Quadrant Biosciences Clarifi COVID-19 Test Kit is a real-time reverse transcription polymerase chain reaction (qRT-PCR) test for the detection of RNA from the SARS-CoV-2. The SARS-CoV-2 primer and probe set(s) are designed to detect two specific sequences of the RdRp gene of the SARS-CoV-2 genome. RNA is extracted from saliva swabs collected from patients suspected of COVID-19 by their healthcare provider. Saliva swab collection is performed by a healthcare provider using the ORAcollect•RNA (OR-100) saliva collection device according to the Instructions for Use of the collection device.

Once received by the laboratory, the ORAcollect•RNA OR-100 saliva swab collection devices containing saliva samples are incubated at 60 °C for 2 hours. Subsequently, RNA is extracted using the Quick-RNA Viral 96 Kit R1041 for which 100 µl sample input volume is used. RNA is eluted in 30 µL of DNase/RNase-Free Water and 1.5 µL of the eluted RNA is used for the downstream qRT-PCR reaction.

The test consists of three processes in a single-well assay:

- Reverse transcription of total RNA to cDNA
- PCR amplification of target and Internal Control cDNA
- Simultaneous detection of PCR amplicons by fluorescent dye labeled probes

## Components and Storage

### Kit Components

Clarifi COVID-19 Test Kit includes the following components:

*Note: The Clarifi COVID-19 Kit is shipped at different temperatures each with a unique part number. See table below to indicate the kit part number the component is contained within. The collection devices are packaged separately from the lab components.*

Name	Part Number	Quantity	Storage Conditions
<b>Part Number 1110 (shipped at ambient temperature)</b>			
Quick-RNA Viral 96 Kit	R1041	1 unit	15°C to 30°C
ORAc collect•RNA saliva swab collection device*	OR-100	1 swab/test	15°C to 25°C
<b>Part Number 1105 (shipped on dry ice)</b>			
4x Reliance One-Step Multiplex Supermix**	12010176	1 x 1mL	-20°C
Positive Control	1111	2,000 copies/μL	-20°C
Clarifi COVID-19 Primer/Probe Set	1108	1 x 780μL	-20°C

\* The ORAc collect•RNA saliva swab collection device lot provided with the Clarifi COVID-19 Test Kit should not be exchange with ORAc collect•RNA saliva swabs from lots that were procured from other sources or with different swabs as this may impact the performance of this test.

\*\*The BioRad enzyme mix lot provided in this test kit was specifically qualified for use with this assay. Do not exchange the enzyme lot provided in this test kit with different lots of the same enzyme or with different enzymes as this may impact the performance of this test.

### Components Required but Not Provided with The Kit

Name	Manufacturer	Part Number	Storage Conditions
PCR Grade Water	Any	Any	15°C to 30°C
Deep 96 well plate (1.0 mL)	Nest	501062	N/A
Ethanol	Fisher Scientific	BP2818500	15°C to 30°C
Beta-mercaptoethanol	Fisher Scientific	03446I-100	15°C to 30°C

### Components Required but Not Provided with The Kit (Continued)

<b>Name</b>	<b>Manufacturer</b>	<b>Part Number</b>	<b>Storage Conditions</b>
MicroAmp™ Optical 96-Well Reaction Plate with Barcode, or equivalent	ThermoFisher Scientific	4306737	N/A
Hard-Shell 96-Well PCR Plates, low profile, thin wall, skirted, white/white, or equivalent	BioRad	HSP9655	N/A
Hard-Shell 384-Well PCR Plates, thin wall, skirted, clear/white, or equivalent	BioRad	HSP3805	N/A
Microseal 'B' PCR Plate Optical Sealing Film, or equivalent	BioRad	MSB1001	N/A
Single-Channel Pipette	Ranin	P-10, P-20, P-200, P-1000	N/A
Multi-Channel Pipette	Ranin	P-10, P-20, P-50, P-200	N/A
Aerosol Barrier Pipette Tips	Ranin	P-10, P-20, P-50, P-200, P-1000	N/A
MicroCentrifuge tube, 1.5 or 2mL, PCR grade	N/A	N/A	N/A

### Equipment/Instrumentation Required

<b>Name</b>	<b>Manufacturer</b>	<b>Part Number</b>
BioRad-CFX96 Touch Real Time Detection System	BioRad	1855195
BioRad-CFX384 Touch Real Time Detection System	BioRad	1855485
QuantStudio 5	Applied Biosystems	A28139
Swing Bucket Plate Centrifuge	N/A	N/A
Incubator	N/A	N/A

## Reagent Storage, Handling, and Stability

- Clarifi COVID-19 Test Kit is shipped at two temperatures (ambient temperature and on dry ice).
- All components of the kit must be stored at the appropriate storage conditions as listed in the section *Kit Components*.
- The Clarifi COVID-19 Primer/Probe Set should be stored at -20°C and protected from light.
- Do not use kit components after expiration date printed on the tube label.
- If there is damage to the packaging inside and outside or kit contents have been tempered with or storage condition failed to meet above -20°C do not use.
- Dispose of unused reagent and waster in accordance with country, federal, state, and local regulations.
- Repeated freezing and thawing may lead to inaccurate results

## Sample Collection, Handling and Storage

Proper collection of specimens is the most important step in the laboratory diagnosis of infectious disease. A specimen that is not collected correctly may lead to false negative test results. All testing for SARS-CoV-2 should be conducted in consultation with a healthcare provider. Specimens should be collected as soon as possible once a decision has been made to pursue testing, regardless of the time of symptom onset. Training in specimen collection is highly recommended due to the importance of specimen quality.

- **Collecting the Sample**
  - Refer to Interim Guidelines for Collecting, Handling, and Testing Clinical Specimens from Persons for Coronavirus Disease 2019 (COVID-19). See <https://www.cdc.gov/coronavirus/2019-ncov/lab/guidelines-clinical-specimens.html>
  - Follow the instructions of the sample collection device for proper collection methods.
- **Transporting Samples**
  - Label each sample container with patient's ID number (e.g. medical record number), date and time sample were collected.
  - Specimens must be packaged, shipped, and transported according to the current edition of the International Air Transport Association (IATA) Dangerous Goods Regulations. Follow shipping regulations for UN 3373 Biological Substance, Category B when sending potential SARS-CoV-2 specimens. Store specimens at ambient temperatures and ship overnight.
- **Storing Samples**
  - Upon receipt, samples can be stored at ambient temperatures for up to 48 hours. Within this 48 hour period the ORAcollect•RNA OR-100 collected saliva samples should be incubated at 60 °C for 2 hours.
  - Store purified nucleic acids at -80°C or below.



## Warning and Precautions

All procedures should be performed in a laboratory of adequate Bio Safety Level (BSL) as recommended by CDC and specimens handled within a Biological Safety Cabinet (BSC). Samples should always be considered potentially infectious and handled in accordance with safe laboratory procedures. Refer to Interim Laboratory Biosafety Guidelines for Handling and Processing Specimens Associated with Coronavirus disease 2019 (COVID-19).

<https://www.cdc.gov/coronavirus/2019-ncov/lab/lab-biosafety-guidelines.html>

Separate work areas should be used for:

- Reagent Preparation (e.g., preparation of RT-PCR master mix); **No** amplified reactions, target solutions, control materials or clinical specimens should be brought into this area. After working in this area, laboratory coat and gloves should be changed before moving into nucleic acid addition area).
- Nucleic acid addition
- Instrumentation (e.g. thermocyclers)

### General Handling:

Proper molecular biological, aseptic technique should always be used when working with RNA. Hands and dust particles may carry bacteria and molds and are the most common sources of RNase contamination. Always wear powder-free latex, vinyl, or nitrile gloves when handling reagents tubes and RNA samples to prevent RNase contamination from the surface of the skin or from dusty laboratory equipment. Change gloves frequently and keep tubes closed. During the procedure, work quickly and keep everything on cold blocks when possible to avoid degradation of RNA by endogenous or residual RNase.

Cleaning working surfaces, pipettes, with cleaning reagents that destroy nucleic acids and RNase. To eliminate accelerated deterioration of any plastics and metals, wipe down surfaces with 70% ethanol.

As with any testing procedure, good laboratory practices are essential to the proper performance of this assay.

- For *In Vitro* diagnostic use (IVD)
- For use under Emergency Use Authorization only
- For prescription Use only
- This test has not been FDA cleared or approved;
- This test has been authorized by FDA under an Emergency Use Authorization (EUA) for use by laboratories certified under the Clinical Laboratory Improvement Amendments (CLIA) of 1988, 42 U.S.C. §263a, that meet the requirements to perform high complexity tests.
- This test has been authorized only for the detection of nucleic acid from SARS-CoV-2, not for any other viruses or pathogens.
- This test is only authorized for the duration of the declaration that circumstances exist justifying the authorization of emergency use of in vitro diagnostics for detection and/or

diagnosis of COVID-19 under Section 564(b)(1) of the Federal Food, Drug and Cosmetic Act, 21 U.S.C. § 360bbb-3(b)(1), unless the authorization is terminated or revoked sooner.

- Positive results are indicative of the presence of SARS-CoV-2 RNA
- Laboratories within the United States and its territories are required to report all results to the appropriate public health authorities.
- All human-sourced materials should be considered potentially infectious and should be handled with universal precautions. If spillage occurs, immediately disinfect with freshly prepared solution of 0.5% sodium hypochlorite (20% v/v bleach). Dispose of cleaning materials in a biohazard waste container.
- Do not use bleach in areas where DNA RNA Shield is used.
- Proper personal protective equipment including lab coats, gowns, gloves, eye protection, and biological safety cabinet are recommended for manipulation of clinical specimens. Refer to Microbiologic and Biomedical Laboratories (BMBL), 5<sup>th</sup> Edition- CDC.
- Do not eat, drink, smoke, apply cosmetics or handle contact lenses in areas where reagents and human specimens are handled.
- Handle all specimens as if infectious using safe laboratory procedures. Refer to Interim Laboratory Biosafety Guidelines for Handling and Processing Specimens Associated with Coronavirus Disease 2019 (COVID-19). See <https://www.cdc.gov/coronavirus/2019-ncov/lab/lab-biosafety-guidelines.html>
- Specimen processing should be performed in accordance with national biological safety recommendations. See <https://www.cdc.gov/labs/BMBL.html>
- If infection with SARS-CoV-2 is suspected based on current clinical and epidemiological screening criteria recommended by public health authorities, specimens should be collected with appropriate infection control precautions.
- Process human clinical specimens within a Class II (or higher) biological safety cabinet (BSC).
- Closely follow procedures and guidelines provided to ensure that the test is performed correctly. Any deviation from the procedures and guidelines may affect the test performance.
- Avoid over exposure of the primer-probe mixes to light for optimal fluorescent signal.
- If exposure of biological materials to skin or mucous membranes occurs, immediately wash the area with large amounts of water. Seek medical advice immediately.
- Do not use components beyond the expiration the date printed on the kit boxes.
- Reagents supplied are formulated specifically for use with this kit. Make no substitutions in order to ensure optimal performance of the kit. Do not mix reagents from different lots.
- Return all components to the appropriate storage conditions after preparing working reagents.
- Keep all the materials on ice when in use.

## Quality Control

Patient samples must be collected according to CDC guidelines.

The following controls are required to be included in every run to accurately interpret patient test results.

- a) A negative control (processing blank) containing PCR grade RNase-Free water is needed to monitor contamination of extraction and PCR reagent reagents with amplifiable material and 1.5  $\mu\text{L}$  is used in each extraction run. This control is then included as template at least once per RT-PCR plate (i.e. for each qRT-PCR run). This control should be included in the sample RNA extraction step.
- b) A positive control is needed to monitor integrity of reagents, screen for improper assay set up and qRT-PCR reagent failure. Recommended concentration for the control included in the Kit is 30 copies per reaction (1.5 $\mu\text{L}$  per reaction). The control needs to be pre-diluted by the user with PCR-grade RNase-Free water to a concentration of 20 copies/ $\mu\text{L}$ . At least one well should be used per qRT-PCR run.
- c) A no template (negative) control containing PCR grade RNase-Free water is needed to determine if amplicon contamination occurred during the qRT-PCR step. The control consists of 1.5 $\mu\text{L}$  of water (in place of sample) least once per plate (i.e. for each qRT-PCR run).
- d) An internal (sample) control is used to monitor poor specimen quality, extraction, reverse transcription and PCR processes and reagent failures. The internal control is the amplification of the human Peptidylprolyl Isomerase A (PPIA) gene with its own primer and probe set included in the Clarifi COVID-19 Test Kit. Failure to amplify the human gene indicates a failed test.

### Clarifi COVID-19 Positive Control Preparation:

**Caution:** The positive control should be handled in a separate area from samples and from PCR set up and amplification with caution to prevent possible contamination. Freeze-thaw cycles should be avoided. Maintain on cold block when thawed.

The positive control is needed to monitor integrity of reagents, screen for improper assay set up and PCR reagent failure. Recommended concentrations for the control included in the Clarifi COVID-19 Test Kit is 30 copies per reaction. The control is provided as 100  $\mu\text{L}$  at a concentration of 200 copies/ $\mu\text{L}$  and needs to be diluted by adding 900  $\mu\text{L}$  nuclease free water to the positive control vial for a concentration of 20 copies/ $\mu\text{L}$  (10:1 dilution). At least one well should be used per batch of specimens.

## Procedure

### Sample Collection:

The workflow begins with saliva collection using the ORAcollect•RNA OR-100 saliva swab collection device provided with the Clarifi COVID-19 Test Kit. Saliva swab collection is performed by a healthcare provider according to the Instructions for Use of the collection device.

These collection devices contain a stabilizing solution that stabilizes collected RNA at room temperature when used according to the provided instructions.

### Assay:

After samples are received in the laboratory for processing, the following steps are to be followed:

**Note: Each Extraction Run requires one negative control to be included.**

### Reagent Preparation

1. Clarifi COVID-19 Test Kit saliva swab collection devices containing saliva samples should be incubated at 60 °C for 2 hours. After incubation, vials can be stored at room temperature prior downstream processing.

**NOTE: Prepare the following reagents fresh on the day of use. Calculate the appropriate volume of reagents needed based on the number of samples to be processed.**

2. Prepare a Beta-mercaptoethanol/Viral RNA Buffer solution by adding beta-mercaptoethanol to the Viral RNA Buffer (Cat. No: 1119) to a final dilution of 0.5 % (v/v) (i.e. 500  $\mu$ L per 100 mL). The final volume should be enough to ensure 400  $\mu$ L of the solution can be added to each sample.
3. Prepare an Ethanol/Viral Wash Buffer Solution by adding 4 mL of 100% ethanol to every 1 mL Viral Wash Buffer (Cat. No: 1117) for a final volume of 5 mL. Volumes may be adjusted to ensure there is enough volume to add 1mL to every sample.

### RNA Extraction (manual)

**NOTE: Perform all steps at room temperature. All centrifugation steps are performed on swinging bucket plate centrifuge at 3000-5000g for 5 minutes. Use 96- deep well plates or PCR tubes (1.0-2.0mL per well) for steps 1 and 2 since the total volume of each well can be up to 600  $\mu$ L.**

**NOTE: Each Extraction Run requires one negative control to be included.**

1. Add 100 $\mu$ L of each saliva sample to each well.

**NOTE: Add 100 $\mu$ L of nuclease free water to one well to be used as a negative control and process it in parallel with each batch of samples.**

2. Add 100 $\mu$ L of DNA/RNA Shield™ (Cat. No: 1118) to each 100 $\mu$ L of saliva sample. Mix well by pipetting.

3. Add 400µL of Beta-mercaptoethanol /Viral RNA Buffer solution prepared above to each 200µL sample solution. Mix well by pipetting.
4. Transfer the sample mixture into each well of the I-96 Plate (Cat. No: 1124) mounted on a Collection Plate (Cat. No: 1123) and centrifuge. Discard the flow-through from the collection plate.
5. Add 500µL of Ethanol/Viral Wash Buffer Solution prepared above in Reagent Preparation to each well and centrifuge. Discard the flow-through from the collection plate. Repeat this step on each well.
6. Add 500µL ethanol (95-100%) to each well and centrifuge. Discard the flow-through from the collection plate.
7. Mount the plate onto an Elution Plate (Cat. No: 1122).
8. To elute RNA, add 30 µL of DNase/RNase-Free Water (Cat. No: 1120) directly to the column matrix of each well and centrifuge. Once RNA is eluted it should be kept on ice during processing.
9. Cover the elution plate with cover foil (Cat. No: 1121) to prevent evaporation.  
*NOTE: The eluted RNA can be used immediately or stored frozen.*

### **PCR Reagent Preparation**

1. Place BioRad Reliance (4x Reliance One-Step Multiplex Supermix) and Clarifi Probe/Primer Mix (both provided in the kit) on a lab bench on ice and protected from light and equilibrate until they reach ambient temperature, vortex for 2-3 seconds, and then spin briefly to collect the reagent
2. Prepare Clarifi qRT-PCR Mix in bulk according to the following table. This should be done on ice.
  - a. To determine the total volume to prepare, use the volume per test below multiplied by the number of tests plus 2 (positive and negative control)
  - b. It is recommended that 10-15% extra qRT-PCR Mix is prepared to account for loss that may occur during pipetting of mix to individual tubes/plate wells, numbers in the table are actual volumes required for each sample
  - c. Add appropriate reagent amounts in the order of: PCR Grade Water, BioRad Reliance and Clarifi Probe/Primer Mix.
  - d. Mix reagents by pipetting up and down 10 times.
  - e. Centrifuge at 3000g for 10 seconds.
  - f. Keep reagents on ice and protected from light until ready to use

### **Clarifi qRT-PCR Mix**

Reagent	Volume/test
BioRad Reliance (4x Reliance One-Step Multiplex Supermix)	3.75 µL
Clarifi Probe/Primer Mix	7.0 µL
PCR Grade Water	2.75 µL
Total volume	13.5 µL

## **qRT-PCR Run**

Plate setup:

1. The assay has been optimized for 15µL reactions in either 96-well or 384-well plates. For the total number of specimens and controls to be run, add 13.5µL Clarifi qRT-PCR Mix to a separate PCR reaction well on the PCR plate.

***Note: Each PCR plate requires 1 no template control (water), 1 negative control (from extraction run) and 1 positive control well.***

2. Add 1.5µL of each sample (and each control) from the sample RNA preparation to each well on the PCR plate. ***Note: The final volume in each well is 15 µL/test.***
3. Seal the plate with optical film.
4. Using a plate spinning rotor, Centrifuge the plate for approximately 1 minute at 2000 - 3000 rpm.
5. Load the plate onto Bio-Rad CFX96 or CFX384 Touch Real-Time PCR Detection System, or Applied Biosystems™ QuantStudio 5 Real-Time PCR System, as desired

## **qRT-PCR Instrument Setup and Operation**

Initial setup:

1. Follow instructions on qRT-PCR equipment to create the following new qRT-PCR protocol for the Clarifi Test:

<b>Step</b>	<b>Cycles</b>	<b>Temp.</b>	<b>Time</b>
Reverse transcription (RT)	1	50°C	20 minutes
Enzyme Inactivation, Polymerase activation	1	95°C	10 minutes
Amplification	45	95°C	10 seconds
		59°C	20 seconds

1. Select 15µL reaction volume
2. Select 99°C lid temperature
3. Save the protocol for future runs
4. Load the plate on the instrument
5. Start run. Note: when using BioRad CFX96 or CFX384, you will receive a prompt, select “all channel”

Repeat setup:

1. Open the previously saved setup file
2. Load the plate on the instrument
3. Start run. Note: when using BioRad CFX96 or CFX384, you will receive a prompt, select “all channel”

### **qRT-PCR Results Export - BioRad CFX96 or CFX384**

1. Export the run file from the instrument. Files are stored in a folder named “Real-Time Data”
2. Using BioRad CFX Maestro Software, open the BioRad compressed file (.zpcr) for the desired run
3. Go to “Settings” > “Plate Setup” > “View/Edit Plate”. Label positive control, negative control and test samples according to your plate layout. Choose HEX and Cy5 channels and give them “Target Name” as HEX and Cy5 respectively. Hit “OK>Apply” to finish your plate setup
4. Threshold setup: under the “Quantification” tab, drag and set both threshold bars to 500 RFU

### **qRT-PCR Results Export - Applied Biosystems QuantStudio 5**

1. Export the run file from the instrument
2. Using QuantStudio software, open the .eds file for the desired run
3. Under the “Plate” tab > “Advanced Setup”, add HEX and Cy5 channels to the “Targets” section and give them names as HEX and Cy5 respectively. Label positive control, negative control and test samples according to your plate layout in the “Samples” section. Hit “Next” > “Next” > “Analyze” to view your results
4. Export your results to excel files under the “Export” tab.

## Interpretation of Results

All test controls must be examined prior to interpretation of patient results. If the controls are not valid, the patient results are invalid and cannot be interpreted. If the external controls are invalid a root-cause investigation should be performed and the entire run has to be repeated after the root cause is eliminated. Results are interpreted based on a cutoff of  $Ct \leq 40.00$ .

### **Examination and Interpretation of Control Results:**

- 1. Positive control review:** To be valid, the positive control should demonstrate amplification below the designated Ct cutoff ( $\leq 40$ ) in the HEX channel. If the positive control is not valid, stop here. The entire plate is invalid and the assay will need to be repeated.
- 2. No template Control review:** To be valid, the new template control should demonstrate no amplification (exceeding background levels) below the designated Ct cutoff ( $\leq 40$ ) in the HEX channel and no amplification (exceeding background levels) below the designated Ct cutoff ( $\leq 35$ ) in the Cy5 channel.
- 3. Negative control review:** To be valid, the negative control should demonstrate no amplification (exceeding background levels) below the designated Ct cutoff ( $< 40$ ) in the HEX channel. If the negative control is not valid, stop here. The entire plate is invalid and the assay will need to be repeated.
- 4. Internal control review:** once both the positive and negative control pass, examine each sample for the internal control. Each sample should demonstrate amplification below the designated Ct cutoff ( $\leq 35$ ) in the Cy5 channel. If the internal control is not valid, the sample should be excluded from further interpretation. The sample is invalid and will need to be repeated. If the internal control fails a second time, the sample will need to be recollected.

### **Expected performance of the controls in the Clarifi COVID-19 Test Kit**

Control	Expected Ct Value (SARS-CoV-2; HEX)	Expected Ct Value (Internal Control; Cy5)
Positive Control	$\leq 40.00$	N/A
Negative Control (processing blank)	$> 40.00$	N/A
No Template Control	$> 40.00$	$> 35.00$
Internal Control	N/A	$\leq 35.00$

### **Examination and Interpretation of Patient Specimen Results:**

Assessment of clinical specimen test results should be performed after the positive, negative and internal controls have been examined and determined to be valid and acceptable. If the controls are not valid, the patient results cannot be interpreted.



1. **Internal Control:** The results of the sample control (internal control) should be examined prior to the release of clinical specimen results. The internal control should produce a positive within each sample analysis. If the internal control is not valid, the clinical specimen results cannot be released, and the sample will need to be rerun. If the internal control fails a second time, the sample needs to be recollected.
  
2. **SARS-CoV2 targets:** To interpret clinical specimen results, the Ct cutoff value for SARS-CoV-2 is 40.00 or less, and the amplification curve shape must be typical in appearance.
  - i. **Positive:** A Ct value  $\leq 40.00$  indicates that the specimen contains the target nucleic acid sequence and is positive (+) for SARS-CoV-2.
  
  - ii. **Negative:** A Ct value  $> 40.00$  indicates that the specimen does not contain the target nucleic acid sequence and is negative (-) for SARS-CoV-2.

HEX channel (SARS-CoV-2)	Cy5 channel (Internal Control)	Result
Early amplification below Ct cutoff (Ct $\leq$ 35)	Amplification below Ct cutoff (Ct $\leq$ 35)	Positive for SARS-CoV-2
	No amplification below Ct cutoff (Ct $>$ 35)	Positive for SARS-CoV-2
Amplification below Ct cutoff (35 $\leq$ Ct $\leq$ 40)	Amplification below Ct cutoff (Ct $\leq$ 35)	Positive for SARS-CoV-2
	No amplification below Ct cutoff (Ct $>$ 35)	SARS-CoV-2 presumptive positive; retest specimen*
No amplification below Ct cutoff (Ct $>$ 40)	Amplification below Ct cutoff (Ct $\leq$ 35)	Negative for SARS-CoV-2
	No amplification below Ct cutoff (Ct $>$ 35)	Invalid - failed internal control, retest specimen**

\*Results for SARS-CoV-2 RNA is Presumptive Positive. Samples with lower target signal (Ct  $> 35.00$ ,  $\leq 40.00$ ) and no Cy5 signal, may be retested by re-extracting RNA from the same specimen. For samples with a repeated Presumptive Positive result, additional confirmatory testing may be conducted.

\*\*First conduct the repeat test by re-isolating RNA phase from the same specimen. If the test fails again, collect a new specimen from the patient and repeat the test.

## Limitations

- All users, analysts, and any person reporting diagnostic results should be trained to perform this procedure by a person with documented competency. Ability to competently perform the test and interpret the results should be documented prior to performing the assay independently. Quadrant Biosciences will limit the distribution of this device to only those users who have documented successful completion of appropriate training.
- Saliva testing is limited to individuals with signs and symptoms of COVID-19.
- The test was validated for use only with oral saliva swabs. Performance with other specimen types is unknown.
- Performance with other sample matrices is unknown.
- Negative results do not preclude SARS-Cov-2 infection and should not be used as the sole basis for treatment or other patient management decisions. Optimum specimen types and timing for peak viral levels during infections caused by SARS-CoV-2 have not been determined. Additional testing of other sample types from the same patient may be necessary to detect the virus.
- A false negative result may occur if a specimen is improperly collected, transported or handled. False negative results may also occur if amplification inhibitors are present in the specimen or if inadequate numbers of organisms are present in the specimen.
- Positive and negative predictive values are highly dependent on prevalence. False negative test results are more likely when prevalence of disease is high. False positive test results are more likely when prevalence is moderate to low.
- If the virus mutates in the RT-PCR target region, SARS-CoV-2 may not be detected or may be detected less predictably. Inhibitors or other types of interference may produce a false negative result. An interference study evaluating the effect of common cold medications was not performed.
- Detection of viral RNA may not indicate the presence of infectious virus or that SARS-CoV-2 is the causative agent for clinical symptoms.
- The performance of this test has not been established for monitoring treatment of SARS-CoV-2 infection.
- The performance of this test has not been established for screening blood or blood product for the presence of SARS-CoV-2.
- This test cannot rule out diseases caused by other bacterial or viral pathogens.

## Conditions of Authorization for the Laboratory

The Clarifi COVID-19 test Letter of Authorization, along with the authorized Fact Sheet for Healthcare Providers, the authorized Fact Sheet for Patients and authorized labeling are available on the FDA website: <https://www.fda.gov/medical-devices/coronavirus-disease-2019-covid-19-emergency-use-authorizations-medical-devices/vitro-diagnostics-euas>.

However, to assist clinical laboratories running the Clarifi COVID-19 test, the relevant Conditions of Authorization are listed below.

- a) Authorized laboratories<sup>1</sup> using the Clarifi COVID-19 test will include with test result reports, all authorized Fact Sheets. Under exigent circumstances, other appropriate methods for disseminating the Fact Sheets may be used, which may include mass media.
- b) Authorized laboratories using the Clarifi COVID-19 test will use the Clarifi COVID-19 test as outlined in the authorized labeling. Deviations from the authorized procedures, including the authorized instruments, authorized extraction methods, authorized clinical specimen types, authorized control materials, authorized other ancillary reagents and authorized materials required to perform the Clarifi COVID-19 are not permitted.
- c) Authorized laboratories that receive the Clarifi COVID-19 test will notify the relevant public health authorities of their intent to run the test prior to initiating testing.
- d) Authorized laboratories using the Clarifi COVID-19 test will have a process in place for reporting test results to healthcare providers and relevant public health authorities, as appropriate.
- e) Authorized laboratories will collect information on the performance of the Clarifi COVID-19 test and report the DMD/OHT7-OIR/OPEQ/CDRH (via email:CDRH-EUA-Reporting@fda.hhs.gov) and Quadrant Biosciences Technical Support (via email: [operations@quadrantbiosciences.com](mailto:operations@quadrantbiosciences.com) or by phone: 1-866-205-7336) any suspected occurrence of false positive or false negative results and significant deviations from the established performance characteristics of the test of which they become aware.
- f) All laboratory personnel using the Clarifi COVID-19 test must be appropriately trained in RT-PCR techniques, the specific processes and instruments used in the Clarifi COVID-19 Test Kit, and use appropriate laboratory and personal protective equipment when handling this kit and use the test in accordance with the authorized labeling.
- g) Quadrant Biosciences, its authorized distributor(s) and authorized laboratories using the Clarifi COVID-19 test will ensure that any records associated with this EUA are maintained until otherwise notified by FDA. Such records will be made available to FDA for inspection upon request.

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<sup>1</sup> For ease of reference, this letter will refer to, “laboratories certified under the Clinical Laboratory Improvement Amendments of 1988 (CLIA), 42 U.S.C. § 263a, that meet requirements to perform high complexity tests” as “authorized laboratories.”

## Performance Characteristics

### Limit of Detection:

Limit of detection (LoD) studies determine the lowest detectable concentration of the SARS-CoV-2 virus at which approximately 95% of all (true positive) replicates test positive. LoD estimates were first determined using a dilution series with 5 different concentrations of heat-inactivated virus (BEI Resources, NR-52286). Saliva specimens were collected from confirmed negative subjects using the ORAcollect•RNA saliva swab collection device (approximately 250µl are absorbed by this swab). The swabs are then inserted into the transport media (1mL) included in the ORAcollect•RNA Kit which stabilizes the sample. For testing, 100 µL aliquots of this sample matrix were prepared and individually spiked with a range of 0, 20, 50, 100, 150, and 200 copies of heat-inactivated virus (BEI Resources, NR-52286) for final virus concentrations of 0, 0.2, 0.5, 1, 1.5 and 2 copies per µL. Each concentration was tested in 5 individually processed replicates following the Clarifi COVID19 Test Kit procedure (RNA eluted in 30 µL RNase-Free Water, qPCR on the BioRad CFX96 Touch Real Time Detection System) using 1.5 µL of extracted RNA per qPCR reaction.

Target Level [cp/µL*]	Valid tested replicates	SARS-CoV-2 Positive (Ct ≤ 40.0 cycles)			Internal Control Positive (Ct ≤ 35.0 cycles)		
		n	Mean Ct	Detection Rate	n	Mean Ct	Detection Rate
2.0	10	10	33.23	100%	10	28.22	100%
1.5	10	10	33.75	100%	10	27.61	100%
<b>1.0</b>	<b>10</b>	<b>10</b>	<b>34.42</b>	<b>100%</b>	<b>10</b>	<b>27.95</b>	<b>100%</b>
0.5	10	9	36.04	90%	10	28.00	100%
0.2	10	6	36.63	60%	10	27.31	100%
0	10	0	NA	0%	10	17.49	100%

\* copies/µL original saliva specimen

The tentative LoD was identified as 1 copy/µL original patient saliva. To confirm that the LoD of the Clarifi COVID-19 Test Kit is 1 copy/µL, saliva specimens were collected from 20 clinical negative subjects with ORAcollect•RNA (OR-100) saliva collection swabs. 100 µL of each sample was spiked with 100 copies of heat-inactivated SARS-CoV-2 virus. Samples were processed according to the Clarifi COVID-19 Test Kit procedure using 1.5 µL of eluted RNA. This was examined in all 20 samples reactions using the Clarifi COVID-19 Test Kit on three instrument: the BioRad-CFX96 Touch Real Time Detection System, the BioRad CFX384 Touch Real Time Detection System, and the Applied Biosystems QuantStudio 5 instruments.

This study confirmed the LoD of the Clarifi Test to be 1 copy/µL with saliva swab specimen for all tested instruments.

Instrument	Target Level* [copies/μL SALIVA]	Valid tested replicates	SARS-CoV-2 Positive			Internal Control Positive		
			n	Mean Ct	Detection Rate	n	Mean Ct	Detection Rate
CFX96	1 cp/μL	20	20	33.85	100%	20	27.49	100%
QS5	1 cp/μL	20	19	34.31	95%	20	25.11	100%
CFX384	1 cp/μL	20	20	35.92	100%	20	29.26	100%

\* copies/μL original saliva specimen

#### Inclusivity (Analytical Sensitivity):

Analytical inclusivity of the primer probe sets was assessed through *in silico* analysis of the Clarifi COVID-19 Test primers and probes. The complete set of available SARS-CoV-2 sequences as of June 30, 2020 was downloaded from the NCBI sequence repository. All sequences longer than 500 bases (n=8,282, average length=29,052 bases) were subsequently aligned with the BWA aligner (version 0.7.15) to the SARS-CoV-2 reference sequence (NC\_045512) using the BWA-MEM algorithm. This resulted in 100% alignment of all sequences, with an average coverage depth of 8,040 bases across 100% of the genome and an average alignment length of 27,666 bases.

Sequence mismatches between the sequence set of the BWA aligner and the assays primer and probe sequences were then identified using the LoFreq program (version 2.1.3a) This resulted in the identification of 531 total sequence variants with quality scores ranging from 68 - 49,314. Within the RdRp Target 1 primer and probe regions, we identified a total of 2 sequences which had single base pair mismatched located in the RdRp Target 1 reverse primer region as shown in the table below, and none of them occurred in the same location.

SARS-CoV-2 Region	Mismatch	Position	Sequences	Frequency	Effect on T <sub>m</sub>
Forward Primer 1	None	-	-	-	-
Probe 1	None	-	-	-	-
Reverse Primer 1	C/T G/A	12,781 12,795	11/8045 9/8044	0.001367 0.001118	- 5.5 °C -5.0 °C
Forward Primer 2	None	-	-	-	-
Probe 2	None	-	-	-	-

Reverse Primer 2	None	-	-	-	-
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Notably, none of the mutations that were found in the RdRp Target 1 reverse primer are located in the 3' terminal base. Thus, the PCR amplification component of the assay is tolerant to all identifiable variants in publicly available sequence data. Even if these rare variants occurred, the RdRp Target 1 reverse primer, it will still be possible for PCR to continue for that amplicon, albeit with less efficiency.

Given that no sequence changes were found for the primers and probes covering the RdRp Target 2 amplicon, and the rare frequency of variants seen in the Clarifi RdRp Target 1 primer and probe binding sites in database, it is highly unlikely that subject samples would incur a signal loss for the SARS-CoV-2 should they contain these variants. Moreover, the dual target design of the Clarifi Test completely mitigates the likelihood of assay failure due to random mutations in the target sequences by having both fluorescent probes in the same channel.

#### Cross-Reactivity (Analytical Specificity):

Wet testing of cross-reactivity of the SARS-CoV-2 primers and probes of the Clarifi COVID-19 Test Kit was documented by laboratory testing with specimens known to be positive for various other respiratory viruses (human coronaviruses 229E, OC43, HKU1, and NL63, human influenza A and B viruses, rhinovirus, respiratory syncytial virus, parainfluenza virus, adenovirus, metapneumovirus, and picornavirus). Testing for some of these organisms was performed by Quadrant Biosciences in the context of the Microbial Interference study per the tables in this section.

For the *in silico* analysis, all of the Clarifi COVID-19 Test Kit primer and probe sequences were queried individually against organisms and pathogens potentially found in upper respiratory specimens, including other coronaviridae, shown below. This was performed using NCBI BLASTN with default alignment parameters.

The *in silico* results indicate that, across all organisms analyzed, there were 11 hits obtained that had >80% homology in one of the primers or probes in the Clarifi COVID-19 test. These 11 hits were obtained from 6 full length sequences that belong to 3 distinct organisms, namely human coronavirus OC43 strain, *Candida albicans*, and *Streptococcus salivarius* ASM78551v1. Of the 11 total hits, six sequences had 3' terminal mismatches with the primers with no risk of amplification since DNA polymerase cannot extend on a mismatch. The remaining 5 sequence hits all belong to the organism *Candida albicans*. However, the primers would align poorly with these sequences and binding would be between 6.5 kilobases to 1.7 mega basepairs away from any upstream or downstream match; based on the 20 second extension time per cycle and 59°C annealing temperature the primers of the Clarifi COVID-19 Test Kit would be unable to exponentially amplify these sequences of the non-targeted organism. Alignments data is shown in the table below.

High priority organisms	%Homology Forward and Reverse Primers and Probes*
Severe acute respiratory syndrome coronavirus 2 isolate Wuhan-Hu-1, complete genome	100%
Human coronavirus 229E, complete genome	None
Human coronavirus OC43 strain ATCC VR-759, complete genome	<b>(84%)</b>
Human coronavirus HKU1, complete genome	None
Human coronavirus NL63, complete genome	None
SARS coronavirus Tor2, complete genome	None
Middle East respiratory syndrome-related coronavirus isolate HCoV-EMC/2012, complete genome	None
Human adenovirus type 1, complete genome	None
Human metapneumovirus isolate 00-1, complete genome (hMPV)	None
Human parainfluenza virus 4a viral cRNA, complete genome, strain: M-25	None
Influenza A virus (A/California/07/2009(H1N1))	None
Influenza B virus	None
Human enterovirus, complete	None
Respiratory syncytial virus, complete genome	None
Human rhinovirus A, strain ATCC VR-1559, complete genome	None
Human rhinovirus B, complete genome	None
Human rhinovirus C, complete genome	None
<i>Chlamydia pneumoniae</i> TW-183	None
<i>Haemophilus influenzae</i> ASM76707v1	None
<i>Legionella pneumophila</i> ASM194158v1	None
<i>Mycobacterium tuberculosis</i> H37Rv	None
<i>Streptococcus pneumoniae</i> R6	None
<i>Streptococcus pyogenes</i> M1 GAS	None
<i>Bordetella pertussis</i> 18323	None
<i>Mycoplasma pneumoniae</i> FH	None
<i>Pneumocystis jirovecii</i> (PJP) RU7	None
<i>Candida albicans</i> SC5314, complete genome	<b>(89%)</b>
<i>Staphylococcus epidermidis</i> , complete genome	None
<i>Pseudomonas aeruginosa</i> strain PA0750 chromosome, complete genome	<b>(89%)</b>

\*For organism with multiple homologies, only the worst homology is presented in this table



### Microbial Interference Studies:

Because this test is a dual target assay the risk of false negative results due to competitive interference of the identified organisms with the primers of this assay is mitigated. In addition, because of the lack of substantial homology (>80%) of organisms noted above, a comprehensive microbial interference study was not performed. However, the ability to detect SARS-CoV-2 in a low titer sample (Ct > 36.0 cycles) was examined using the Clarifi COVID-19 Test Kit for some high priority organisms. This was determined by adding 1.5 µL of purified RNA from saliva samples containing SARS-CoV-2 alone or in combination with hCoV-NL63, HRV/enterovirus, *Mycoplasma pneumoniae*, hMPV/metapneumovirus, or *Chlamydia pneumoniae*. Results indicate no difference in the cycles to threshold for detection of SARS-CoV-2, and thus no interference in the assay performance.

Sample ID	SARS-CoV-2 patient sample	Known Pathogen Present	COVID (HEX)	PPIA (Cy5)
379-0907	-	Human coronavirus NL63 (hCoV-63)	NA	27.48
470-326	-	Rhinovirus	NA	29.08
132-8996	-	<i>Mycoplasma pneumoniae</i> /Human metapneumovirus	NA	28.67
106-2898	-	<i>Chlamydia pneumoniae</i>	NA	29.56
379-0907	+	Human coronavirus NL63 (hCoV-63)	36.10	26.99
470-326	+	Rhinovirus	36.34	27.48
132-8996	+	<i>Mycoplasma pneumoniae</i> /Human metapneumovirus	36.00	27.54
106-2898	+	<i>Chlamydia pneumoniae</i>	37.13	27.74

### Interfering Substances Study:

The potential interfering effects of 7 substances on the RT-PCR reactions were tested in accordance with FDA guidance. These substances included: Mucin (bovine submaxillary gland, type I-S, 2.5 mg/ml), human blood (2.5% v/v), Afrin original nasal spray (15% v/v), Basic Care allergy relief nasal spray (Glucocorticoid, 5% v/v), Cepacol Sore Throat (benzocaine/menthol lozenges, 5mg/mL), toothpaste (5% v/v), and mouthwash (5% v/v). Saliva specimens were collected from three confirmed negative subjects with the ORAcollect•RNA (OR-100) saliva collection swabs. Each subject was tested with the substances indicated in Table 17 below. Eight aliquots (100 µl) were made from each subject's sample and spiked with one of the 7 potential interfering substances or PCR grade water, as well as 200 copies of heat-inactivated SARS-CoV-2 (2x LOD). The samples were processed according to the Clarifi COVID-19 Test Kit procedure. Results indicated no differences in the cycles to threshold for detection of SARS-CoV-2.

Potential Interfering Substance	Valid Tested replicates	Valid Positive replicates	Mean Ct SARS-Cov-2	Mean Ct Internal Control
Mucin: bovine submaxillary gland, type I-S	3	3	36.62	28.55
Blood (human)	3	3	37.36	29.38



Afrin Original nasal spray	3	3	37.63	29.61
Basic Care allergy relief nasal spray (Gluococorticoid)	3	3	36.74	26.94
Cepacol Sore Throat (benzocaine/menthol lozenges)	3	3	36.53	27.07
Toothpaste	3	3	35.53	32.02
Mouthwash	3	3	35.67	26.91
PCR grade water	3	3	36.16	27.22

**Clinical Evaluations:**

A total of 63 Clarifi COVID-19 saliva swab specimens were collected within 0-5 days following collection of the comparative nasopharyngeal swabs (32 negative, 31 positive). Clinical nasopharyngeal specimen results were obtained from an EUA authorized SARS-CoV-2 RT-PCR assay. Saliva samples were processed using the Clarifi COVID-19 Test Kit as per the Instructions for Use. The Clarifi COVID-19 Test Kit resulted in a positive agreement of 100% (31/31) and a negative agreement of 100% (32/32).









		EUA authorized Comparator (NP Swab)		Total
		POSITIVE	NEGATIVE	
Clarifi COVID-19 Test Kit (Saliva)	POSITIVE	31	0	<b>31</b>
	NEGATIVE	0	32	<b>32</b>
	Total	<b>31</b>	<b>32</b>	<b>63</b>
<b>Positive Percent Agreement (PPA): 31/31 = 100% (95% CI)</b>				
<b>Negative Percent Agreement (NPA): 32/32 = 100% (95% CI)</b>				

## Additional Information

### Symbols

The following symbols are used in labeling for Quadrant Biosciences PCR diagnostic products.

Symbols used in labeling for Quadrant Biosciences PCR diagnostic products.

	In Vitro Diagnostics
	Manufactured By
	Batch Code
	Consult Instruction for use
	Catalog Number
	Temperature Limitation
	Single Use
	Date of Manufacturer
DD-MM- YYYY	Date Format (Date-Month-Year)

### Manufacturer and distributors

Quadrant Biosciences Inc.

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Syracuse, NY 13210

[www.quadrantbiosciences.com](http://www.quadrantbiosciences.com)

Technical Support: [operations@quadrantbiosciences.com](mailto:operations@quadrantbiosciences.com)

To Report an Adverse Event: 1-866-205-7336

### References

1. Center for Disease Control and Prevention. Biosafety in Microbiological and Biomedical Laboratories, 5<sup>th</sup> ed. U.S. Department of Health and Human Services, Public Health Services, Centers for Disease Control and Prevention, National Institutes of Health HHS Publications No. (CDC) 21-112, revised December 2009.
2. Clinical and Laboratory Standards Institute (CLSI). Protection of laboratory workers from occupationally acquired infections. Approved Guidelines – Fourth Edition. CLSI Document M29-A4: Wayne, PA; CLSI, 2014

## Document History

<b>Version No.</b>	<b>Version Date</b>	<b>Description of Change</b>
A	18Sep2020	First Issuance