



QUALITY CONTROL

It is necessary to confirm the correct functioning of the kit in every assay by amplifying a blank (DEPC-treated water) and the negative and positive controls included in the kit.

A blank that gives a positive result for either gene (E gene, M gene, N gene or GAPDH) in the analysis indicates a contamination problem. In this case, the process should be repeated, ensuring that the work area and equipment are adequately decontaminated and that extreme precautions are taken during the PCR.

A negative control that is negative for the GAPDH gene indicates either a negative control degradation problem or a failure to prepare the reaction. In this case, the process must be repeated.

A positive control that returns a negative result indicates a failure of the PCR reaction, either due to inadequate reagent preparation, failure of the reagents, or failure of the positive control. Before repeating the amplification process, make sure that all reagents have been stored correctly and have not expired.

TEST LIMITATIONS

As with any other diagnostic test, results of Real SARS-CoV-2/Flu/RSV kit must be interpreted by a healthcare professional, taking into account the patient's medical history, clinical symptoms and any other diagnostic tests available.

The results obtained with this product depend on the proper collection, transportation, storage and processing of samples. To avoid erroneous results, it is necessary to pay particular attention to these stages and carefully follow the instructions provided with the products for the extraction and storage of the nucleic acids.

This assay could be used with other types of samples, although it has only been validated with nasopharyngeal, oropharyngeal or nasal swabs, aspirates and saliva.

There is the possibility of false negatives or of variable results when working with samples with a low number of copies of the target template, below the detection limit of the test (see sensitivity heading).

Due to the high analytical sensitivity of the technique, there is a possibility of false positives due to cross-contamination from positive samples with high viral load, with the positive control or with the same products of the amplification reaction. These phenomena can only be avoided with good laboratory practice and carefully following product instructions.

A positive result does not necessarily indicate the presence of viable viruses and does not imply that these viruses are infectious or that they are the causative agents of clinical symptoms. However, a positive result may be indicative of the presence of the target viral sequences.

Negative results do not exclude suffering from virus infection and should not be used as the sole basis for treatment or other patient management decisions. Collecting multiple samples (sample types and at various points over time) from the same patient may be necessary to detect the virus.

FEATURES / EXTERNAL AND INTERNAL EVALUATIONS

1. Diagnostic sensitivity and specificity concordance with the CLART PneuMoVir kit (Genómica)

The presence of Flu and RSV in respiratory samples (nasopharyngeal aspirates) and nucleic acids extracted from those samples was evaluated using the Real SARS-CoV-2/Flu/RSV kits (respiratory sample and nucleic acid) and the CLART PneuMoVir kit (only nucleic acid). This evaluation was performed in Operon's facilities (Cuarte de Huerva, Spain).

Table with 2 columns: DIRECT SAMPLE (N = 62) and CLART® PneuMoVir 2. Rows include Influenza, Real SARS-CoV2/Flu/RSV, RSV, and Concordance Sensitivity/Specificity/Prevalence/PPV/NPV/Efficiency data.

Table with 2 columns: NUCLEIC ACID (N = 153) and CLART® PneuMoVir 2. Rows include Influenza, Real SARS-CoV2/Flu/RSV, RSV, and Concordance Sensitivity/Specificity/Prevalence/PPV/NPV/Efficiency data.

2. Diagnostic sensitivity and specificity concordance with the BioGX Real Flu A, B and RSV kit for the BD Max system.

The presence of Flu and RSV in respiratory samples (nasopharyngeal aspirates) and nucleic acids extracted from those samples was evaluated using the Real SARS-CoV-2/Flu/RSV kits (respiratory sample and nucleic acid) and the BioGX Real Flu A, B and RSV kit for the BD Max system (only nucleic acid). This evaluation was performed in Operon's facilities (Cuarte de Huerva, Spain).

Table with 2 columns: DIRECT SAMPLE (N = 80) and BioGX. Rows include Influenza, Real SARS-CoV2/Flu/RSV, RSV, and Concordance Sensitivity/Specificity/Prevalence/PPV/NPV/Efficiency data.

Table with 2 columns: NUCLEIC ACID (N = 90) and BioGX. Rows include Influenza, Real SARS-CoV2/Flu/RSV, RSV, and Concordance Sensitivity/Specificity/Prevalence/PPV/NPV/Efficiency data.

3. Diagnostic sensitivity and specificity concordance with the Xpert Xpress Flu/RSV kit (GeneXpert)
The presence of Flu and RSV in respiratory samples (nasopharyngeal aspirates) and nucleic acids extracted from those samples was evaluated using the Real SARS-CoV-2/Flu/RSV kits (respiratory sample and nucleic acid) and the GeneXpert Xpert Xpress Flu/RSV kit (only nucleic acid). This evaluation was performed in Operon's facilities (Cuarte de Huerva, Spain).

Table with 2 columns: DIRECT SAMPLE (N = 93) and GeneXpert. Rows include Influenza, Real SARS-CoV2/Flu/RSV, RSV, and Concordance Sensitivity/Specificity/Prevalence/PPV/NPV/Efficiency data.

Table with 2 columns: NUCLEIC ACID (N = 90) and GeneXpert. Rows include Influenza, Real SARS-CoV2/Flu/RSV, RSV, and Concordance Sensitivity/Specificity/Prevalence/PPV/NPV/Efficiency data.

4. Diagnostic sensitivity and specificity concordance with the Multiple Real Time PCR kit for detection of 2019-CoV (XABT).
The presence of SARS-CoV-2 in respiratory samples (oropharyngeal swabs, nasopharyngeal swabs, nasal swabs, and saliva) and nucleic acids extracted from those samples was evaluated with the Real SARS-CoV-2/Flu/RSV kits (respiratory sample and nucleic acid) and the Multiple Real Time PCR Kit for detection of 2019-CoV (XABT, Beijing Applied Biological Technologies Co, Ltd; only nucleic acid). This evaluation was performed in Operon's facilities (Cuarte de Huerva, Spain).

Table with 2 columns: DIRECT SAMPLE (N = 446) and XABT. Rows include SARS-CoV2, Real SARS-CoV2/Flu/RSV, and Concordance Sensitivity/Specificity/Prevalence/PPV/NPV/Efficiency data.

5. Diagnostic sensitivity and specificity concordance with the Allplex 2019-nCoV assay (Seegene) kit
The presence of SARS-CoV-2 in respiratory samples (nasopharyngeal and oropharyngeal swabs) and nucleic acids extracted from those samples was evaluated with the kits 'Allplex 2019-nCoV assay' (Seegene, only nucleic acid) and 'Real SARS-CoV-2 Flu RSV' (Operon, respiratory sample and nucleic acid). This evaluation was performed in Operon's facilities (Cuarte de Huerva, Spain).

Table with 2 columns: DIRECT SAMPLE (N = 71) and Allplex 2019-nCoV assay. Rows include SARS-CoV2, Real SARS-CoV2/Flu/RSV, and Concordance Sensitivity/Specificity/Prevalence/PPV/NPV/Efficiency data.

6. Diagnostic sensitivity and specificity concordance with the Allplex Respiratory Panel (Seegene) kit
The presence of Flu and RSV in respiratory samples (nasopharyngeal swabs) and nucleic acids extracted from those samples was evaluated with the kits 'Allplex Respiratory Panel'

(Seegene, only nucleic acid) and 'Real SARS-CoV-2 Flu RSV' (Operon, respiratory sample and nucleic acid). This evaluation was performed by the Microbiology Service of the Donostia University Hospital.

Table with 2 columns: DIRECT SAMPLE (N = 112) and Allplex Respiratory. Rows include Influenza, Real SARS-CoV2/Flu/RSV, RSV, and Concordance Sensitivity/Specificity/Prevalence/PPV/NPV/Efficiency data.

Table with 2 columns: NUCLEIC ACID (N = 116) and Allplex Respiratory. Rows include Influenza, Real SARS-CoV2/Flu/RSV, RSV, and Concordance Sensitivity/Specificity/Prevalence/PPV/NPV/Efficiency data.

7. Diagnostic sensitivity and specificity concordance with the LightMix® SarbecoV-E gene plus EAV control (Roche) and the VIASURE SARS-CoV-2 Real Time PCR Detection Kit (Ceresist).
The presence of SARS-CoV-2 in respiratory samples (nasopharyngeal swabs) and nucleic acids extracted from those samples was evaluated with the kits LightMix® SarbecoV-E gene plus EAV control (Roche, only nucleic acid) and VIASURE SARS-CoV-2 Real Time PCR Detection Kit (Ceresist, only nucleic acid) and 'Real SARS-CoV-2 Flu RSV' (Operon, respiratory sample and nucleic acid). This evaluation was performed by the Microbiology Service of the Donostia University Hospital.

Table with 2 columns: SARS-CoV2 and SarbecoV-E gene plus. Rows include NUCLEIC ACID (N = 118), Real SARS-CoV2/Flu/RSV, and Concordance Sensitivity/Specificity/Prevalence/PPV/NPV/Efficiency data.

Table with 2 columns: SARS-CoV2 and VIASURE SARS-CoV-2. Rows include NUCLEIC ACID (N = 118), Real SARS-CoV2/Flu/RSV, and Concordance Sensitivity/Specificity/Prevalence/PPV/NPV/Efficiency data.

DETECTION CAPABILITY
The analytical sensitivity, or the test limit, is defined as the minimum amount of RNA that can always be detected with the test. The limit of the test was defined by analysing a series of dilutions in triplicate of an RNA preparation of the different viruses for detection, using three product batches.

Table with 2 columns: Real SARS-CoV-2/Flu/RSV product is: and a table with columns: RN Control, Flu A, Flu B, RSV A, RSV B, CoV 2. Rows show Copies per PCR for each component.

On the other hand, the detection capability of the test was evaluated by spiking of negative samples of the different specimens compatible with the test with the WHO international standard for SARS-CoV-2 (ref. NIBSC 20/146), for Flu from ATCC: The A strain Wisconsin/67/2005 (ref. VR-1881) and B strain Florida/07/04 (ref. VR-1804) and for RSV: RSV A 2000/3-4 (ref. NR-28530) and RSV B1 (ref. NR 4052) from Bei Resources.

Table with 2 columns: RNA specimen and a table with columns: Flu A, Flu B, RSV A, RSV B, CoV 2. Rows show detection results for various specimens.

SPECIFICITY

The presence of cross reaction with the following microorganisms was ruled out by spiking mixes of negative respiratory samples with quantified cultures of the different microorganisms:

- Bacteria/yeasts: Bordetella holmesii, Bordetella pertussis, Bordetella pertussis, Candida albicans, Haemophilus influenzae, Haemophilus parahaemolyticus, Haemophilus parainfluenzae, Klebsiella pneumoniae, Legionella pneumophila, Moraxella catarrhalis, Moraxella lacunata, Mycoplasma tuberculosis, Neisseria gonorrhoeae, Neisseria meningitidis, Neisseria subflava, Proteus vulgaris, Serratia marcescens, Staphylococcus haemolyticus, Staphylococcus saprophyticus, Staphylococcus aureus, Streptococcus A, B, C, D, F and G, Streptococcus mitis, Streptococcus mutans, Streptococcus oralis, Streptococcus pneumoniae, Streptococcus salivarius, Streptococcus sanguinis, Pseudomonas aeruginosa, Chlamydia pneumoniae and Mycoplasma pneumoniae.

Virus: coronavirus OC43, NL63, 229E y MERS, the parainfluenza viruses (HPV) 1, 2, 3, 4A and 4B, Adenovirus, Enterovirus, Metapneumovirus, Rhinovirus. The lack of cross reaction was confirmed for all cases by "in silico" analysis of every available sequence in GenBank.

HOOK EFFECT

The presence of a Hook effect was ruled out by adding increasing amounts of a positive RNA control for the different viruses to the RT-PCR reaction.

For Flu A, Flu B, and SARS-CoV-2, a control RNA from Twist Bioscience was used, up to 5 x 10^6 copies/PCR. For RSV A and B, a control RNA from ATCC was used, up to 3.25 and 1.65 x 10^5 copies/PCR.

INTRA-ASSAY PRECISION

Analysing 5 replicates of 7 series of dilutions of a positive RNA control, 4 respiratory samples and their corresponding nucleic acids, a positive control, a negative control, and a blank with three different product batches, evidence was provided for a high intra-assay precision for the Real SARS-CoV-2/Flu/RSV test, obtaining RSDs under 3% between the Cq values obtained in the different replicates.

INTER-DAY PRECISION

Analysing 2 replicates of 7 series of dilutions of a positive RNA control, 4 respiratory samples and their corresponding nucleic acids, a positive control, a negative control, and a blank over 5 consecutive days, evidence was provided for a high inter-day precision for the Real SARS-CoV-2/Flu/RSV test, obtaining RSDs under 4% between the Cq values obtained in the different replicates.

INTER-LABORATORY PRECISION

Analysing 2 replicates of 7 series of dilutions of a positive RNA control, 4 respiratory samples and their corresponding nucleic acids, a positive control, a negative control, and a blank by 3 different operators and different equipment (micropipettes, cabinet, Real Time PCR thermal cycler), evidence was provided for a high inter-laboratory precision for the Real SARS-CoV-2/Flu/RSV test, obtaining RSDs under 6% between the Cq values obtained in the different replicates.

INTER-LOT PRECISION

Analysing 2 replicates of 7 series of dilutions of a positive RNA control, 15 respiratory samples and their corresponding nucleic acids, and a blank with three different product batches, evidence was provided for a high inter-batch precision for the Real SARS-CoV-2/Flu/RSV test, obtaining RSDs under 4% between the Cq values obtained in the different replicates.

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Product information section including icons for expiration date, lot number, catalog number, and manufacturer, along with CE and ISO 9001 certifications.

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