

IVD CE

bosphore

INSTRUCTIONS FOR USE

West Nile Virus Quantification Kit v2

For *In Vitro* Diagnostic Use

MB100v9f
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Anatolia
geneworks

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1. PRODUCT DESCRIPTION

Bosphore West Nile Virus Quantification Kit v2 is a Real-Time PCR kit for *in vitro* diagnostics that detects and characterizes the 3' UTR region of West Nile virus RNA from serum, plasma, urine, whole blood, and CSF samples. Fluorescence detection is performed using FAM, and HEX filters. West Nile virus RNA is amplified and fluorescence detection is performed using the FAM filter.

Component	FAM (Gene)	HEX
PCR Master Mix	West Nile virus (3' UTR)	Internal Control

Internal control has been integrated into the kit to check RNA extraction, PCR inhibition, or application problems. The amplification data of the internal control is detected with the HEX filter. The internal control can be added either during RNA extraction or the PCR step.

2. CONTENT

Bosphore West Nile Virus Quantification Kit v2 consists of the following dH₂O, Real-Time PCR Master Mix, RT mix, positive control, standards, and internal control.

Component	Reagent	100 Reactions	50 Reactions	25 Reactions
1	dH ₂ O	(1,000 µL)	(1,000 µL)	(1,000 µL)
2	PCR Master Mix	(1,640 µL)	(820 µL)	(410 µL)
3	RT Mix	(28 µL)	(14 µL)	(7 µL)
4	Internal Control	(550 µL)	(275 µL)	(275 µL)
5	Positive Control	(44 µL)	(22 µL)	(22 µL)
6	Standard 1 (1,000 copies/µL)	(88 µL)	(44 µL)	(44 µL)
7	Standard 2 (100 copies/µL)	(88 µL)	(44 µL)	(44 µL)
8	Standard 3 (10 copies/µL)	(88 µL)	(44 µL)	(44 µL)
9	Standard 4 (2 copies/µL)	(88 µL)	(44 µL)	(44 µL)

3. STORAGE

PCR reagents for Bosphore West Nile Virus Quantification Kit v2 should be stored at -20 °C. Repeated thawing and freezing (>3x) should be avoided since it may reduce sensitivity. If the components are to be used in small amounts, they should be frozen in aliquots.

While preparing the PCR, the components should not be exposed to room temperature for more than 10 minutes, and the PCR master mix components should not be exposed to light or air more than necessary. Vials must be kept closed except during pipetting. We recommend preparing the PCR on a cooling block and keeping the PCR master mix in a closed container. If the components are stored according to the recommended conditions, they will remain stable until the expiry dates on the labels.

4. REQUIRED MATERIALS AND DEVICES

- Montania 4896 Real-Time PCR Instrument – Anatolia Geneworks, CFX96 Real-Time PCR Detection System - Bio-Rad, QuantStudio 5 Real-Time PCR System – ThermoFisher, LightCycler 480 Instrument II – Roche*, Q qPCR Cycler - Quantabio, Rotor-Gene Q – QIAGEN or another Real-Time PCR system with FAM, and HEX filters**
- 0.1 mL or 0.2 mL thin-wall PCR tubes, PCR plates or strips
- Unio B24 Extraction System or Unio B48 Extraction System or Unio A24S Extraction & PCR Setup System and Unio Viral DNA/RNA Extraction Kit 600 µl, Unio 96 Extraction & PCR Setup System and Unio 96 Nucleic Acid Extraction Versatile Kit, Unio M32 Extraction System and Unio M32 Viral DNA/RNA Extraction Kit, Magrev 24 Manual Magnetic Bead Nucleic Acid Extraction Stand and Magrev Nucleic Acid Extraction Versatile Kit or Magrev Viral DNA/RNA Extraction Kit, Bosphore Viral RNA Extraction Spin Kit, Bosphore Viral DNA/RNA Extraction Spin Kit or Bosphore Nucleic Acid Extraction Versatile Spin Kit or Magnesia 16 Nucleic Acid Extraction Instrument and Magnesia Viral DNA/RNA Extraction Kit EP (Anatolia Geneworks) or other high-quality RNA extraction kits and systems
- Deep freezer (-20 °C)
- Desktop centrifuge with rotor for 2 mL or 1.5 mL microcentrifuge tubes
- DNase, RNase, pyrogen-free 1.5 mL or 2 mL microcentrifuge tubes
- Calibrated, adjustable micropipettes
- DNase, RNase, pyrogen-free micropipette tips with filters
- Disposable laboratory gloves, coats and caps

* See section 9.4 for the thermal protocol suitable for LightCycler 480 Instrument II – Roche.

** For other Real-Time PCR devices that can be used with Bosphore West Nile Virus Quantification Kit v2, please contact Anatolia Geneworks from the information in Section 15.

5. IMPORTANT NOTES AND SAFETY INSTRUCTIONS

- The product should be delivered on dry ice. Check for the presence of dry ice upon arrival.
- Check for the expiration dates on the box and tube labels upon arrival. Do not use expired products or components.

- Calibrated or verified micropipettes, DNase, RNase, and pyrogen-free micropipette tips with filters, and DNase, RNase, and pyrogen-free microcentrifuge tubes should be used.
- Before starting a test procedure, all components should be thoroughly thawed. After thawing, all components should be centrifuged briefly (spin-down for 3-5 seconds) and mixed well to ensure homogeneity before use.
- The kit components should be kept on ice or a cooling block until the reaction is prepared and quickly returned to -20 °C.
- PCR and nucleic acid extraction must be performed in different compartments. Samples should be stored separately to avoid contact with the kit components.
- Pathogen information should be reviewed to be aware of the health-related risks.
- Biological samples should be handled with extreme caution and in a microbiological safety cabinet of the appropriate class. Physical contact with pathogens should be avoided by wearing lab coats and gloves, making no allowance for eating or drinking within the workspace, and preventing unauthorized individuals' access to the working area.
- After working biological samples, all pathogenic wastes produced during the nucleic acid extraction step, including materials contacted with them, should be discarded into medical waste, and disposed of safely.

6. PRODUCT USE LIMITATIONS

- All the components may exclusively be used for *in vitro* diagnostics.
- This product should be used by this user manual.
- This product is to be used by personnel specially trained to perform *in vitro* diagnostic procedures.

7. INFECTION

West Nile virus (WNV) is a *Flavivirus* belonging to the *Flaviviridae* family that was first isolated in 1937 in the "West Nile" province of Uganda/Africa and was named after this region. WNV virions are approximately 50 nm in diameter and are composed of a single-stranded RNA genome of length between 11 kb and 12 kb (Brinton, 2002).

West Nile virus was first isolated in a woman in the West Nile district of Uganda in 1937. It was identified in birds (crows and Columbiformes) in the Nile Delta region in 1953. Before 1997 WNV was not considered pathogenic for birds, but at that time in Israel, a more virulent strain caused the death of different bird species presenting signs of encephalitis and paralysis. Human infections attributable to WNV have been reported in many countries in the World for over 50 years.

In 1999 a WNV circulating in Israel and Tunisia was imported in New York producing a large and dramatic outbreak that spread throughout the continental United States of America (USA) in the following years.

The WNV outbreak in the USA (1999-2010) highlighted that the importation and establishment of vector-borne pathogens outside their current habitat represent a danger to the world. The largest outbreaks occurred in Greece, Israel, Romania, Russia, and the USA. Outbreak sites are on major birds' migratory routes.

In its original range, WNV was prevalent throughout Africa, parts of Europe, the Middle East, West Asia, and Australia. Since its introduction in 1999 into the USA, the virus has spread and is now widely established from Canada to Venezuela (WHO, 2017).

Most of the WNV infections are asymptomatic without any symptoms in people. The second type of WNV infection results in mild fever (West Nile Fever) with recovery from the illness, but however, third type of WNV infection with the virus crossing the blood-brain barrier causes severe outcomes such as inflammation of the brain (encephalitis) and the tissue surrounding the brain and spinal cord (meningitis) that all resulting in long-term effects like memory loss, depression, etc. The common route of WNV transmission is shown to be the mosquitos as vectors carrying the virus across vertebrate hosts. Mosquitos receive the virus upon feeding on infected birds, and the virus is transmitted to other hosts via mosquito bites. Humans, horses, and most other mammals are widely accepted as incidental hosts as they don't transmit the virus to others. However, with increasing research after WNV outbreaks, new WNV transmission routes are transmitted via blood transfusion, organ transplantation, intrauterine exposure, and breastfeeding (Hayes et al., 2005).

8. METHOD

Bosphore West Nile Virus Quantification Kit v2 is based on the Real-Time PCR method. The polymerase chain reaction is a technique that is used for the amplification of an DNA region. The reaction occurs through repeated cycles of heating and cooling. The main components of PCR are primers, dNTPs, *Taq* DNA Polymerase (with hot-start property), buffer solutions, and templates. As a brief explanation, primers are small synthetic DNA that anneals to the specific regions of the template to start the synthesis, dNTPs are the building blocks of the amplified products, and *Taq* DNA Polymerase amplifies the RNA template. Finally, buffer solutions provide the pH adjustment required for the reaction, and the template, as referred to, is the target region for synthesis. In addition to these components, in RT-PCR reverse transcriptase is added to the reaction, and cDNA synthesis from the RNA template is acquired.

In the Real-Time PCR technique, in contrast to conventional PCR, PCR products can be monitored during the reaction. Therefore, Real-Time PCR obviates the need for further analysis methods like gel electrophoresis, thereby minimizing the risk of contamination.

Dual-labeled probes employed in the reaction, in addition to the conventional PCR reagents, enable the detection of the amplified target with increased sensitivity. The assay utilizes the 5' exonuclease activity of *Taq* DNA Polymerase to cleave a dual-labeled fluorescent hydrolysis probe during the extension phase of PCR.

The probe is labeled at the 5' end with a fluorescent 'reporter' molecule, and at the 3' end with another fluorescent molecule that acts as a 'quencher' for the 'reporter'. When the two fluorophores are nearby, and the reporter is excited by light, no reporter fluorescence can be detected. During the elongation step of PCR, *Taq* DNA Polymerase encounters and cleaves the probe bound to the template.

As the reporter is freed from the suppressing effect of the quencher, a fluorescence signal can be detected. The fluorescence generated by the reporter increases as the PCR product is accumulated; the point at which the signal rises above the background level and becomes distinguishable is called the threshold cycle (C_T). There is a linear relationship between the log of the starting amount of a template and its threshold cycle. Thus, the starting number of unknown templates can be determined using standard curves constructed using C_T values of the known starting amounts of target templates.

Bosphore West Nile Virus Quantification Kit v2 uses multiplex qPCR and internal control is included in the system to control the extraction procedure, PCR inhibition, and application issues.

The reaction is performed in one PCR tube containing PCR Master Mix. In PCR Master Mix, West Nile virus genome amplification is screened using FAM filter, and the fluorescent signal produced by the internal control amplification is detected in PCR Master Mix via the HEX filter.

9. PROCEDURE

9.1. RNA Extraction

We recommend that the Unio B24 Extraction System or Unio B48 Extraction System or Unio A24S Extraction & PCR Setup System and Unio Viral DNA/RNA Extraction Kit 600 µl, Unio 96 Extraction & PCR Setup System and Unio 96 Nucleic Acid Extraction Versatile Kit, Unio M32 Extraction System and Unio M32 Viral DNA/RNA Extraction Kit, Magrev 24 Manual Magnetic Bead Nucleic Acid Extraction Stand and Magrev Nucleic Acid Extraction Versatile Kit or Magrev Viral DNA/RNA Extraction Kit, Bosphore Viral RNA Extraction Spin Kit, Bosphore Viral DNA/RNA Extraction Spin Kit or Bosphore Nucleic Acid Extraction Versatile Spin Kit or Magnesia 16 Nucleic Acid Extraction Instrument and Magnesia Viral DNA/RNA Extraction Kit EP (Anatolia Genetworks) or other high-quality RNA extraction kits and systems or other high-quality RNA extraction kits and systems are used with Bosphore West Nile Virus Quantification Kit v2. The RNA extraction should be performed according to the manufacturer's instructions.

9.2. Kit Components

9.2.1. Negative Control

The negative control is an essential component of Bosphore West Nile Virus Quantification Kit v2 for Real-Time PCR. It consists of distilled water (dH₂O) and serves as a reference sample to ensure accurate results by detecting any contamination or errors that could lead to false-positive outcomes. Including the negative control and carefully monitoring it will enable you to confidently interpret your data.

9.2.2. PCR Master Mix

PCR Master Mix contains a highly specific and accurate *Taq* DNA Polymerase (with hot-start property), PCR buffers, and dNTPs mix. PCR Master Mix also contains forward and reverse primers and dual-labeled probes specific for West Nile virus and internal control.

9.2.3. RT Mix

RT Mix contains a high-quality and performance reverse transcriptase and RNase inhibitor mix.

9.2.4. Internal Control

The internal control included in the kit to check for RNA extraction, PCR inhibition, and application errors, the internal control is a synthetic DNA molecule. The internal control is added to the mixture of sample, proteinase K, and Carrier RNA at the beginning of RNA extraction to check extraction efficiency and application errors. If the internal control is to be added during nucleic acid extraction, 5 µL is added per sample; and when added directly to the PCR Master Mix to control PCR inhibition, 0.2 µL is added. We recommend adding an internal control to the negative control to evaluate the efficiency of the extraction system.

Caution! It is not necessary to include the internal control in the PCR Master Mix if it has already been added during the extraction step. The absence of internal control amplification in the HEX channel in negative samples may indicate a problem in extraction or application, or that the PCR reaction is inhibited. In this case, extraction and PCR should be repeated. In samples with high viral load, including the positive control, the internal control may be suppressed and an increase in fluorescent signal may not be detected. Therefore, internal control amplification should be evaluated according to the table in Section 10.

9.2.5. Positive Control

The kit contains 1 positive control containing synthetic West Nile virus DNA. It must be included in the PCR reaction to test the reaction efficiency.

Positive Control: Synthetic West Nile virus DNA

The threshold C_T value of the positive control is given in the acceptance criteria table (Section 10). A threshold C_T value of the positive control greater than the upper limit of the range in the table may indicate a loss of yield in the reaction.

9.2.6. Quantitation Standards

The quantitation standards are calibrated plasmid standards of 1×10^3 , 1×10^2 , 1×10^1 , and 2 copies/ μL .

9.3. Preparing the PCR

The standards, positive and negative controls should be added into the PCR reaction together with the samples. Ensure that all kit components are dissolved before use and refer to the table below to prepare the PCR reaction mix. The volumes given are for one reaction only, multiply these volumes by the number of samples to find the volumes required for the master mix. When preparing the mix for more than 5 samples, 10% should be added to the total number of samples.

Components	Volume
PCR Master Mix	14.75 μL
RT Mix	0.25 μL
Internal Control*	0.2 μL^*
Sample RNA (Standards/Positive Control/ Negative Control)	10 μL
Total Volume	25 μL

*Internal control should not be added to the reaction if it has already been added during the extraction step.

Pipette 15 μL of the PCR Master Mix into the PCR tubes or strips and add 10 μL of template (sample/standards/positive or negative control). Close the tube cap. Make sure that the mix in each tube is at the bottom of the tube. Centrifuge if necessary.

9.4. Programming the Real-Time PCR Instrument

The thermal protocol for Bosphore West Nile Virus Quantification Kit v2 consists of reverse transcription, initial denaturation for activation of the *Taq* DNA Polymerase (with hot-start property), a two-step amplification cycle, and a terminal hold. The Real-Time data is collected at the second step of the amplification cycle. The thermal protocol to be applied for the reaction is indicated on the next page.

Steps	Temperature	Time	
Reverse Transcription	50 °C	30:00 min	
Initial Denaturation	95 °C	14:30 min	
Denaturation	97 °C	00:30 min	} 50 Cycles
Annealing (Data Collection)	55.5 °C	01:10 min	
Hold	32 °C	02:00 min	

The thermal protocol suitable for LightCycler 480 Instrument II – Roche is given below:

Steps	Temperature	Time	
Reverse Transcription	50 °C	30:00 min	
Initial Denaturation	95 °C	14:30 min	
Denaturation	97 °C	00:30 min	} 50 Cycles
Annealing (Data Collection)	55 °C	01:10 min	
Hold	32 °C	02:00 min	

Before starting to work with Bosphore West Nile Virus Quantification Kit v2, the following steps must be completed and checked:

- Choose all the filters to be used (FAM, and HEX),
- Identify unknown samples, positive and negative controls, standards,
- Select the correct thermal protocol,
- Start the experiment.

10. ANALYSIS

By the end of the thermal protocol, the Real-Time PCR Instrument software automatically calculates the baseline cycles and the threshold. Analysis of the results should be performed by trained personnel who have received the required training for analyzing Real-Time PCR data. The standard curve is plotted using the data obtained from the defined standards, with the axes C_T -Threshold Cycle and Log Starting Quantity.

Caution! Since the quantitation standards are provided as plasmid samples and as copies/ μ L, the following formula should be applied to the resulting copies/ μ L values obtained for the samples, to assess the quantitation values of the samples in copies/mL.

This mathematical factor takes the starting volume of RNA extraction and the elution volume into consideration, to ensure the correct quantitation of the samples:

$$\frac{(\text{Result in copies}/\mu\text{L}) \times (\text{Elution Volume in } \mu\text{L})}{(\text{Starting Extraction Volume in mL})} = \text{Result in copies/mL}$$

For example, if a sample's result from the automated Standard Curve was calculated as 1,000 copies/ μL , considering that the starting extraction volume is 400 μL and the elution volume is 60 μL , applying the formula; $1,000 \times 60 \div 0.4 = 150,000$ copies/mL is calculated as the West Nile virus RNA that the sample material contains.

We recommend that the test results be evaluated by an expert clinician, taking the patient's clinical findings and the results of other tests into consideration. All analysis is done automatically in routine use. However, when the trained personnel who have received the required training from the manufacturer, consider it necessary if the system allows pulling down the threshold as much as possible to detect low amplifications, attention should be paid to keep the threshold line above the background.

The negative control is essential for accurate result analysis. Please check the negative control and ensure it shows no amplification outside the associated internal control filter. If the negative control has a signal outside the filter associated with the internal control, please do not report the results. Repeat the experiment after taking the necessary precautions against contamination. If the same result is encountered again, please contact the manufacturer.

Internal control, standards, and positive control of Bosphore West Nile Virus Quantification Kit v2 are essential for accurate result analysis. The cycle threshold acceptance criteria for the internal control, standards, and positive control are listed below:

Component / Parameter	Threshold Value (C_T)
Positive Control	30 \pm 4
Internal Control	\leq 32
Standard 1	23.5 \pm 2
Standard 2	26.5 \pm 2
Standard 3	30 \pm 2
Standard 4	32.5 \pm 2
Correlation Coefficient	>0.970
PCR Efficiency	>80%

In quantitative test results, examples that cross the threshold in the FAM channel; it is evaluated as "Positive", samples that do not cross the threshold are shown as "No C_T" or "Negative". These samples are considered to have a negative or viral load below the detection limit of the assay. The internal control data in the HEX channel of these "undetected" samples should also be checked to avoid false negative results.

The delayed amplification of the internal control may indicate a problem in nucleic acid extraction / PCR inhibition or application failure. In this case, extraction and PCR should be repeated. Please consider that in the samples that contain high viral concentration, internal control can be suppressed therefore delayed or no increase in internal control signal may be detected.

Please note that this product only provides testing pathogen shown below. Experimental results from the tube must be considered when providing a result for the patients, also in consideration of the patient's clinical findings, and the guidelines of the relevant health authorities. The table below shows the possible results and their interpretation:

PCR Master Mix	West Nile virus (FAM)	Internal Control (HEX)	Result
	+	+/-	The sample is West Nile virus positive
	-	+	Sample is negative
	-	-	The test should be repeated!

LightCycler 480 Instrument II – Roche: Please use a **white plate** for Roche LightCycler 480. "**Abs Quant/Fit Points**" should be chosen as the analysis type for the appropriate threshold level selection. "**Abs Quant/2nd Derivative Max**" should be chosen as the analysis type for the quantitation results. Analysis should be performed with the '**Cycle Range**' option. **For LC480, color compensation protocol must be performed Bosphore Color Compensation Plate Set F could be used for PCR Master Mix.**

CFX96 Real-Time PCR Detection System - Bio-Rad: Use of a **white plate** and turning "**Apply Fluorescence Drift Correction**" on in "**Baseline Setting**" for the analysis is recommended.

Rotor-Gene Q – QIAGEN: Please use "**Outlier removal**" up to **15%** if necessary.

In rare cases of PCR inhibition due to medication or other PCR inhibitors in the sample, we recommend repeating the test of inhibited samples, by freezing and thawing the RNA samples and using them in the PCR after diluting them 1:2 with dH₂O.

Caution! The dilution factor must be considered while reporting the Real-Time PCR quantitative results.

11. SPECIFICATIONS

11.1. Sensitivity

Analytical sensitivity may be expressed as the limit of detection: i.e., the smallest amount of the target marker that can be precisely detected. The detection limit of an individual analytical procedure is the lowest amount of nucleic acid in a sample which can be detected but not necessarily quantitated as an exact value. The analytical sensitivity or detection limit for NAT assays is expressed by the 95% positive cut-off value. The analytical detection limit for Bosphore West Nile virus Quantification v2 was found to be 50 copies/mL.

The sensitivity was determined using serial dilutions of RNA calibrated with the previously quantitated West Nile virus RNA Control. The dilutions were tested in different runs in replicates. The results were analyzed by the probit method.

11.2. Linear Range

The linear range of Bosphore West Nile Virus Quantification Kit v2 was determined to be 1.2×10^2 - 4×10^9 copies/mL. The standard curve correlation coefficient was found to be 0.999 for the linearity run with a dilution series of the West Nile virus Positive Control.

11.3. Cross-Reactivity

To eliminate potential cross-reactivity, both assay design evidence and experimental studies were employed. Primer and probe sequences were checked for possible homology to other known pathogen sequences by sequence comparison analysis using database alignment. To eliminate the risk of cross-reactivity; Zika virus, Dengue virus, Chikungunya virus, HSV1 (Herpes Simplex Virus 1), HSV2 (Herpes Simplex Virus 2), Hepatitis B Virus (HBV), Hepatitis C Virus (HCV), Human Immunodeficiency Virus (HIV), *Neisseria gonorrhoeae*, *Trichomonas vaginalis*, *Gardnerella vaginalis*, *Mycoplasma genitalium*, *Chlamydia trachomatis*, *Plasmodium malariae*, *Salmonella enterica*, *Plasmodium ovale*, *Listeria monocytogenes*, *Streptococcus pneumoniae*, *Streptococcus agalactiae*, and *Haemophilus influenzae* samples with known high positivity were tested and found negative. The experimental results indicated that the kit detects specifically and only Bosphore West Nile Virus Quantification Kit v2 pathogen that it intends to detect, but not the others.

11.4. Reproducibility

Reproducibility data (on a C_T value basis) were obtained by the analysis of the previously quantitated West Nile virus positive samples. The test was performed in at least 4 replicates by 3 different operators, on multiple days, using 3 different lots. The resulting data is given on the next page for 37,500 copies/mL.

Variability (West Nile virus)	Standard Deviation	Variance	Coefficient of Variation [%]
Intra-assay (n=4)	0.01708	0.00029	0.18624
Inter-lot (n=3)	0.02622	0.00069	0.10320
Inter-operator (n=3)	0.03126	0.00098	0.12308
Total Inter-assay (n=5)	0.03008	0.00090	0.11841

12. REFERENCES

1. Brinton M. A. (2002). The molecular biology of West Nile virus: a new invader of the western hemisphere. Annual review of microbiology, 56, 371–402. <https://doi.org/10.1146/annurev.micro.56.012302.160654>
2. World Health Organization. (3 October 2017) West Nile virus. <https://www.who.int/news-room/fact-sheets/detail/west-nile-virus>
3. Hayes EB, Komar N, Nasci RS, Montgomery SP, O'Leary DR, Campbell GL (2005). "Epidemiology and transmission dynamics of West Nile virus disease". Emerging Infect. Dis. 11 (8): 1167–73.

13. SYMBOLS



Use-by Date



Caution, consult accompanying documents.



Batch Code



Manufacturer



Catalog Number



In vitro Diagnostic Device

14. ORDERING INFORMATION

ABWNV6 (100 rxn/box)
Catalog Number: **ABWNV5 (50 rxn/box)**
ABWNV4 (25 rxn/box)

15. CONTACT INFORMATION



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Document Revision History

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V1	01	04 th December 2018	First Publishing
V2	02	12 th January 2020	Partial content correction
V3	03	06 th April 2021	The general content and type check
V4	04	15 th February 2022	The content has been updated and checked
V5	05	06 th September 2022	The general content and type check
V6	06	17 th November 2023	The content has been updated and checked
V7	07	30 th November 2023	The general content and type check
V8	08	17 th April 2024	Positive control acceptance criteria changed
V9	09	14 th November 2024	PCR Master Mix volume changed