

HBV Quantification Kit v1 USER MANUAL

For in vitro Diagnostic Use



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1. PRODUCT DESCRIPTION

Bosphore® HBV Quantification Kit v1 detects and quantitates Hepatitis B Virus DNA in human serum or plasma, encompassing all the major HBV genotypes (A-H). The linear range of quantitation is 1x10¹-1x10⁰ IU/ml and the analytic sensitivity is 10 IU/ml. A region within the S gene is amplified and fluorescence detection is accomplished using the FAM filter.

An internal control has been integrated into the kit in order to check PCR inhibition. The amplification data of the internal control is detected with the Cy5 filter. The internal control can be added either during DNA extraction or PCR step.

2. CONTENT

Bosphore® HBV Quantification Kit v1 is composed of Real-Time PCR reagents and quantitation serum standards which have been calibrated against WHO International Standard (NIBSC Code 97/750):

Component	REAGENT	100	50 Tests	25 Tests
		Tests		
1	dH₂O	(1000 µl)	(1000 µl)	(500 µl)
2	PCR Mix	(1400 µl)	(700 µl)	(350 µl)
3	Detection Mix1	(79 µl)	(39.5 µl)	(19.5 µl)
4	Detection Mix2	(95 µl)	(47.5 μl)	(23.75µl)
5	Internal Control	(560 µl)	(280 µl)	(140 µl)
6	Positive Control 1	(45 µl)	(23 µl)	(11 µl)
7	Standard 1 (1 x 10 ⁶) IU/ml	(880 µl)	(440 µl)	(440 µl)
8	Standard 2 (1 x 10⁵) IU/ml	(880 µl)	(440 µl)	(440 µl)
9	Standard 3 (1 x 10 ⁴) IU/ml	(880 µl)	(440 µl)	(440 µl)
10	Standard 4 (5 x 10 ²) IU/ml	(880 µl)	(440 µl)	(440 µl)

3. STORAGE

Bosphore® HBV Quantification Kit v1 PCR reagents 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 min. and the detection mix components should not be exposed to light more than 1-2 min. We recommend preparing the PCR on a cooling block and keeping the detection mixes within a closed container.

The components maintain their stability until the expiry dates on the labels, if they are stored at advised conditions.

4. REQUIRED MATERIALS AND DEVICES

Montania® 483 Real-Time PCR Instrument (Anatolia Geneworks), or another Real-Time PCR system with FAM and Cy5 filters (iCycler, iQ5, CFX-BioRad, LightCycler 1.5, 2.0, 480-Roche, 7500

Document Code: MB01v4f Date: April 2011 Real-Time PCR System-ABI, Stratagene Mx3005P, Mx3000P-Agilent, LineGeneK, LineGene 9600-Bioer, Rotorgene 2000, 3000, 6000, Q-Qiagen)

- 0.2 ml Thin-Wall PCR tubes or strips
- Magnesia® 16 Nucleic Acid Extraction System / Magnesia® Viral Nucleic Acid Extraction Kit (Anatolia Geneworks), or other high quality viral DNA extraction kits and systems
- Deep freezer (-20°C)
- Desktop centrifuge with rotor for 2 ml. microcentrifuge tubes
- Calibrated adjustable micropipettes
- DNAse, RNAse, pyrogen free micropipette tips with filters
- DNAse, RNAse, pyrogen free 1.5 or 2 ml. microcentrifuge tubes
- Disposable laboratory gloves

5. IMPORTANT NOTES AND SAFETY INSTRUCTIONS

Important!:

- The product should be delivered on dry ice. Check for presence of dry ice upon arrival.
- Check for the expiry dates on the box and tube labels, upon arrival. Do not use expired products or components.
- Calibrated or verified micropipettes, DNAse, RNAse, pyrogen free micropipette tips with filters, and DNAse, RNAse, 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 prior to use.
- The kit components should be kept on ice or a cooling block until the reaction is prepared, and they should be quickly returned to -20°C.
- PCR and nucleic acid isolation 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.
- Serum samples (including the standards) should be handled with extreme caution, suitable class microbiological safety cabinet should be used: Physical contact with pathogens should be avoided by; wearing lab coats and gloves, no allowance for eating or drinking within the workspace, prevention of unauthorized individuals' access to the working area.
- All the pathogenic wastes produced during the nucleic acid isolation step; including the serum samples and material contacted with them, should be discarded into medical waste and disposed safely.

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6. PRODUCT USE LIMITATIONS

All the components may exclusively be used for in vitro diagnostics.

• This product should be used in accordance with this user manual.

This product is to be used by personnel specially trained to perform in vitro diagnostic

procedures.

7. PATHOGEN

Causative Agents

Hepatitis B virus (HBV) is one of the smallest enveloped double-stranded DNA viruses, and a member

of the Hepadnaviridae family. The replication process of HBV that takes place in liver, is unique among the

animal DNA viruses in which reverse transcription is involved. HBV may destroy the liver and cause

diseases such as cirrhosis and hepatocellular carcinoma. There are 8 distinctly classified genotypes of

hepatitis B virus and further recognized subgenotypes. [1], [2]

Epidemiology

Hepatitis B virus (HBV) infection is a worldwide health problem with the highest burden of disease in

Asia, Pacific Islands and Sub-Saharan Africa. There are 2 billion people infected worldwide (one third of

world's population), and 400 million suffering from chronic HBV infection (90% of infants and up to 50% of

young children infected with hepatitis B will develop chronic infections). HBV infections result in roughly 1

million deaths per year; including the deaths caused by HBV and its complications (HBV-related liver

diseases). [3], [4]

Modes of Transmission: Transmission of hepatitis B virus follows the same modes as HIV, but unlike

HIV, HBV is 50-100 times more infectious and survives in the open air for at least 7 days. Common modes of

transmission are; perinatal (from mothers to infants primarily at birth), early childhood infections

(inapparent infection through close contact with infected household), unsafe injection practices, blood

transfusions and sexual contact. [4]

8. METHOD

Bosphore® HBV Quantification Kit v1 is based on the Real-Time PCR method. Polymerase chain reaction

is a technique that is used for amplification of a DNA region. The reaction occurs by the repeating cycles of

heating and cooling. The main components of PCR are primers, dNTPs, Taq polymerase enzyme, buffer

solution and template. As a brief explanation, primers are small synthetic DNA those anneal to the specific

regions of the template in order to start the synthesis. dNTPs are the building blocks of the amplified

products. Tag polymerase amplifies the DNA template. Buffer solution provides the pH adjustment

required for the reaction and template, as referred, is the target region for synthesis.

In Real Time PCR technique, in contrast to conventional PCR, PCR product can be monitored during the

reaction. Therefore Real-Time PCR obviates the need for further analysis methods like gel electrophoresis,

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whereby minimizing the risk of contamination. Dual labeled probes employed in the reaction in addition to the conventional PCR reagents, enable detection of the amplified target with increased sensitivity.

The assay utilizes the 5' exonuclease activity of Tag Polymerase to cleave a dual-labeled fluorescent 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 in close proximity, and the reporter is excited by light, no reporter fluorescence can be detected. During the elongation step of PCR, Tag Polymerase encounters and cleaves the probe bound to the template. As the reporter is freed from the suppressing effect of the quencher, 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 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 starting amount of unknown templates can be determined using standard curves constructed using C_T values of the known starting amounts of target templates.

Bosphore HBV Quantification Kit v1 employs multiplex PCR, and an internal control is incorporated into the system in order to control the isolation procedure and to check for possible PCR inhibition. HBV DNA and an internal control are co-amplified in a single reaction, using sequence-specific primers. The fluorescent signal generated by the HBV amplification is detected by a probe labeled at the 3' end with FAM, through the FAM channel. The fluorescent signal generated by the internal control amplification, is detected by a second probe (labeled at the 5' end with a different reporter molecule, Cy5) through the Cy5 channel.

9. PROCEDURE

9.1. Sample Preparation, Storage and Transport

To isolate serum from the clinical specimen, the blood sample should be collected into sterile vacutainers without any anticoagulant. For venipuncture, only sterile material should be used.

The serum should be separated from blood within 6 hours after blood collection. To separate the serum, the blood container should be centrifuged at 800-1600 x g for 20 minutes. The separated serum should be transferred to polypropylene tubes and stored at -20°C or lower, until use.

The samples should be transported in containers with capacity to resist pressure. Transportation should be done according to local and national regulations for pathogen material transport.

9.2. Interfering Substances

The following factors may have possible influences on PCR:

- Hemolytic samples,
- Samples of heparinised patients,
- Samples of patients with elevated levels of bile salts, bilirubin or lipids

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9.3. DNA Isolation

We recommend that the Magnesia® 16 Nucleic Acid Extraction System / Magnesia® Viral Nucleic Acid Extraction Kit (Anatolia Geneworks) isolation system is used with Bosphore® HBV Quantification Kit v1. The DNA isolation should be performed according to the manufacturers' instructions. The starting volume is 400 μ l, the elution volume is 60 μ l and the amount of internal control that should be used during isolation for each system is 5 μ l. The external quantitation standards are provided as serum, so that they undergo the same steps as the patient samples, starting from DNA isolation.

9.4. Kit Components

9.4.1. PCR Mix

HotStarTaq DNA Polymerase: HotStarTaq DNA Polymerase is a modified form of a recombinant 94 kDa DNA polymerase, originally isolated from *Thermus aquaticus*, cloned into E.Coli. The enzyme is provided in an inactive form. It is activated by a 15-minute 95 °C incubation step. This prevents the formation of misprimed products and primer-dimers during reaction setup and the first denaturation step, leading to high PCR specificity and accurate quantification. PCR Buffer: contains Tris-Cl, KCl, (NH₄)₂SO₄, 8 mM MgCl₂, pH 8.7 (20°C). dNTP Mix: Contains ultrapure quality dATP, dGTP, dCTP ve dTTP/dUTP.

9.4.2. Detection Mix 1

Detection Mix 1 contains HBV-specific forward and reverse primers and a dual-labeled probe.

9.4.3. Detection Mix 2

Detection Mix 2 contains internal control-specific forward and reverse primers and a dual-labeled probe.

9.4.4. Internal Control

An internal control is included in the kit to control DNA isolation and PCR inhibition. The internal control is a synthetic DNA molecule derived from human genome. It is added into the serum, proteinase K and carrier RNA mixture during DNA isolation, to control the isolation efficiency and PCR inhibition. The amount of IC that should be added during isolation is 5 µl per serum sample. Alternatively, the internal control can be added directly into the PCR master mix to control the PCR inhibition exclusively. For this purpose, 0.1 µl of internal control should be added for each reaction into the master mix. Lack of internal control amplification in the FAM negative samples, may indicate a problem in isolation or PCR inhibition. In this case, isolation and PCR should be repeated. In samples that contain a high viral load, internal control can be suppressed and no increase of the signal is detected. Please use the table below for the interpretation of internal control data:

HBV (FAM)	Internal Control (Cy5)	Interpretation
+	+	Sample positive
-	+	Sample negative
+	-	Sample positive
-	-	Repeat the test!

9.4.5. Positive Control

The positive control contains HBV DNA. It can be included in the PCR to test the efficiency of the PCR exclusively. The threshold cycle for the positive control is given in the acceptance criteria table (Section 10. Analysis). Threshold cycles higher than the acceptance criteria may indicate an efficiency loss in the reaction.

9.4.6. Quantitation Standards

The quantitation serum standards are calibrated by WHO International Standard (NIBSC Code 97/750).

9.5. Preparing the PCR

All four external quantitation standards should be added into the PCR reaction together with the samples and the negative control (PCR-grade water). Make sure that all the kit components are thawed before use. Refer to the table below for preparing the PCR. It is for only one reaction, multiply these values with the sample number to find the values required for the master mix. While preparing master mixes for more than 5 samples, an extra 10% should be added to the total sample number.

When the Internal Control is added in the extraction step:

PCR Mix	12.5 µl
Detection Mix 1	0.7 μΙ
Detection Mix 2	0.85 µl
dH2O	0.95 μl
Sample DNA	
(Standard,	10.0 μl
Negative/Positive Control)	
Total Volume	25.0 μl

When the Internal Control is added in the PCR step:

PCR Mix	12.5 µl
Detection Mix 1	0.75 µl
Detection Mix 2	0.85 µl
Internal Control	0.1 μΙ
dH2O	0.85 µl
Sample DNA	·
(Standard,	10.0 µl
Negative/Positive Control)	
Total Volume	25.0 μl

Pipette 15 μ l of the master mix into the PCR tubes or strips, and add 10 μ l of DNA (sample/standard/positive or negative control). Close the tube cap. Make sure that the solution in each tube is at the bottom of the tube. Centrifuge if necessary.

9.6. Programming the Montania® 483 Real-Time PCR Instrument

The thermal protocol for Bosphore® HBV Quantification Kit v1 is composed of an initial denaturation for activation the HotStarTaq DNA Polymerase, a two-step amplification cycle and a terminal hold. The real-time data is collected at the second step of the amplification cycle.

Denaturation	97°C	00:30 min.	٦
Annealing and Synthesis	54°C	01:30 min.	} 50 cycles
(Data Collection)			
Hold	22°C	05:00 min.	

Montania® 483 Real-Time PCR Instrument is installed and calibrated as it is delivered to the end user.

In order to establish an appropriate link between the system components, first the thermal cycler and the optical module, and then the PC and the software should be started.

Before starting a Real-Time PCR reaction using the Bosphore® Kits, the following steps should be completed:

- Choose the filter pairs to be used (FAM and Cy5),
- Identify unknown samples, standards, positive and negative controls, assign quantitative values to the standards,
- Select the correct thermal protocol.

These steps are described below:

From the main menu of the Montania® 483 Real-Time PCR Instrument, "File" and then "New" is selected. "Create a new Experiment" is selected. In the "Select Channel" window channels 1 (FAM) and 3 (Cy5) are selected (Fig. 1). Standards, samples and negative controls are identified in the "Module Edit" menu (Fig. 2). Standards should only be defined for the FAM channel and their concentration/viral load should be entered. To select the thermal protocol "Gene Amplification" menu is used. The "Open" button in the "Experiment Program" is clicked and the appropriate thermal protocol is selected. (Fig. 3a). The thermal cycles of the selected protocol is displayed. The experiment starts by clicking the "Start" button (Fig. 3b).

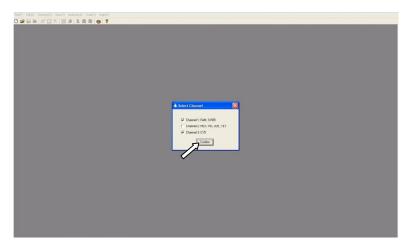


Fig. 1: Filter Selection in Montania® 483

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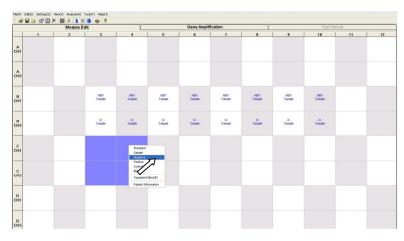


Fig. 2: Sample Location and Identification

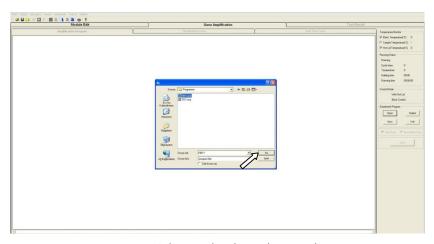


Fig. 3a: Selecting the Thermal Protocol

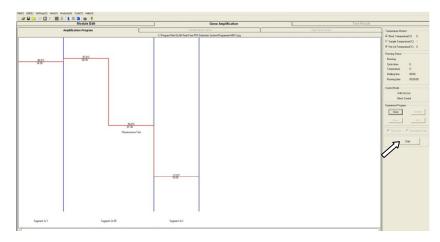


Fig. 3b: Starting the Experiment

10. ANALYSIS

By the end of the thermal protocol, the Montania® 483 Real-Time PCR Instrument software automatically calculates the baseline cycles and the threshold.

Example of an amplification curve is given in Fig. 4.

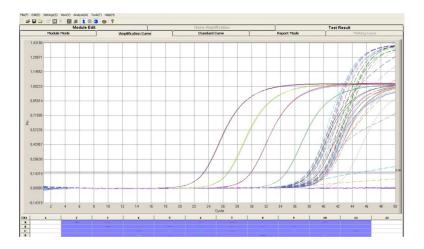


Fig. 4: Amplification Curve of a Bosphore® HBV v1 test

The standard curve is plotted using the data obtained from the defined standards, with the axes Ct-Threshold Cycle and Log Starting Quantity. Example of a standard curve is given in Fig. 5

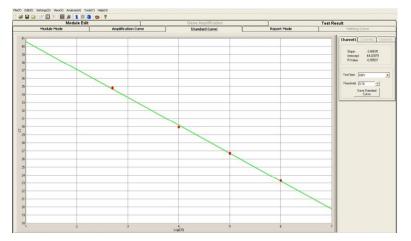


Fig.5: Standard Curve of a Bosphore® HBV v1 test

Analysis of the results should be performed by trained personnel who have received the required training for analysing Real-Time PCR data. We recommend that the test results must 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 manufacturer, consider it as necessary, the system allows pulling down the threshold as much as possible in order to detect low positive samples. In this case, attention should be paid to keep the threshold line above the background and to keep the correlation coefficient at the maximum possible value (and within its acceptance criteria.

The table below displays the acceptance criteria for Bosphore® HBV v.1.

Component/Parameter	Cycle Threshold (C _T)
Standard 1	23±2.5
Standard 2	27±2
Standard 3	30±2
Standard 4	35±2
Positive Control	33±4
Correlation Coefficient	>0.950
PCR Efficiency*	>%60

^{*} PCR efficiency is calculated by the following formula: 10^(-1/slope)-1 x100

Test results should not be reported unless the assay results meet the criteria stated above. Please contact the manufacturer if an impairment in the product's performance is observed (See the last page for contact information).

The quantitative results of the test are displayed on the "Report Mode" screen. A spread sheet containing the calculated starting quantities of the unknown samples in each tube is shown. The samples that cross the threshold in Channel 1 (FAM) are displayed with a calculated starting quantity, samples that do not cut the threshold are displayed as "No Ct". These samples are regarded as negative or having a viral load below the detection limit of the assay. For these undetectable samples, the Cy5 data of the internal control should also be checked to avoid false negative results (Fig. 6).

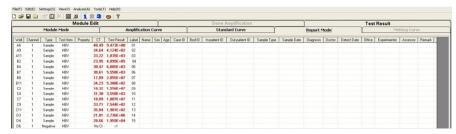


Fig. 6: A Report Mode Screen Showing the Results

The following table shows the possible results and their interpretation:

Signal detected in FAM	The sample contains	No need to check the internal control since the sample
filter pair	HBV DNA, the result	is positive (high positive samples may suppress the
	is positive	signal from the internal control)
No signal in FAM, signal	The HBV DNA in the	Signal from Cy5 filter pair rules out the possibility of
in Cy5	sample is not	PCR inhibition
	detectable	
No signal in FAM and	The diagnosis is	No signal in Cy5 points out to PCR inhibition or to a
Cy5	inconclusive	problem in DNA isolation (See 11.Troubleshooting)

11. TROUBLESHOOTING

Please contact the manufacturer in case of a problem during a run.

Late or no signal from the FAM filter

Wrong thermal protocol is chosen	Make sure that the correct thermal protocol is chosen.
----------------------------------	--

Late or weak signal from the standards

Deterioration of the standards or	Don't use expired standards or kit components. Follow the
the core kit components	instructions for the storage of kit components (Section 3. Storage).

No signal from the internal control

Deterioration of the internal	Follow the instructions for the storage of kit components (See
control or detection mix 2	Section 3. Storage).
PCR inhibition	Make sure that you use the recommended DNA isolation method
	(See 9.3 DNA isolation).
DNA lost during isolation	Make sure that you use the recommended DNA isolation method
_	(See 9.3 DNA isolation).

Signal from FAM Filter in the Negative Control

_	-g				
	Contamination	Use filter-tips. Repeat PCR with new kit components.			

The Threshold is Above Low Signals

The threshold should be manually	Using the mouse pull the threshold down until it cuts the low	
adjusted	signals. Avoid the background and the signal from negative	
	control.	

12. SPECIFICATIONS

12.1. 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 cutoff value.

The analytical detection limit for Bosphore® HBV v1 was found to be 1x10¹ IU/ml (p=0.05). The sensitivity was determined using serial dilutions of DNA calibrated with the WHO International Standard for HBV DNA NAT assays, (NIBSC Code 97/750). The dilutions were tested in different runs in replicates. The results were analyzed by probit method.

12.1.1. Genotype Detection and Quantitation Efficiency

Efficiency of detecting and quantitating different genotypes were ensured both by sequence comparison analysis and a Real-Time PCR assay using HBV DNA Genotype Performance PHD 350 (Seracare). The following genotypes were tested and found positive:

Panel Member	Genotype	HBV (FAM)
1	D	+
2	D	+
3	F	+
4	C	+
5	Α	+
6	E	+
7	E	+
8	E	+
9	С	+
10	F	+
11	G	+
12	Н	+
13	Α	+
14	В	+
15	В	+

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12.2. Linear Range

The linear range of Bosphore® HBV Quantification Kit v1 was determined to be 1x10¹-1x109 IU/ml.

In order to assess the linear range, a dilution series of a member of HBV DNA Genotype Performance PHD 350 (Seracare) which has been calibrated against the WHO International Standard for HBV DNA NAT assays, (NIBSC Code 97/750) was analyzed by testing each dilution in 4 replicates (Fig. 7a and 7b). The standard curve correlation coefficient was found to be 0.99968.

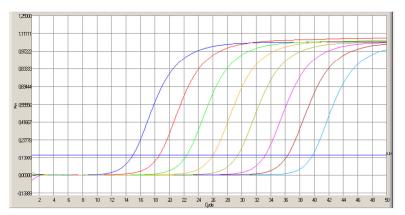


Fig. 7a: Linear Range Amplification Curve

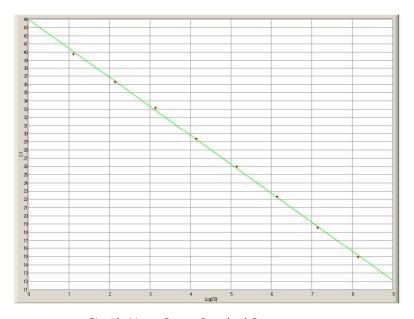


Fig. 7b: Linear Range Standard Curve

12.3. 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. Samples of CMV, EBV, MTBC, Parvovirus B19, BKV with known high positivity were tested, and found negative.

12.4. Reproducibility

Reproducibility data (on C_T value basis) were obtained by the analysis of one of the quantitation standards of the Bosphore® HBV Quantification Kit v1. Test was performed in at least 4 replicates by 3 different operators, on multiple days, using 3 different lots. The resulting data is given in Table 1:

Table 1: Reproducibility Data.

HBV (10 ⁴ IU/ml)	Standard deviation	Variance	Coefficient of variation [%]
Intra-assay Variability N=4	0.07	0.005	0.23
Inter-lot Variability N=3	0.10	0.01	0.34
Inter-operator Variability N=3	0.28	0.08	0.91
Total Inter-assay Variability N=5	0.29	0.08	0.93

HBV Quantitation Standards were calibrated against the WHO International Standard for HBV DNA NAT assays, (NIBSC Code 97/750) by performing two assays using various dilutions and testing them in triplicates. 1 IU was found to be equal to 4.5 ± 0.2 copies/ml.

12.5. Diagnostic Evaluation

The diagnostic evaluation was performed by testing 100 HBV negative and 7 HBV positive serum samples which have been previously analyzed using Roche Diagnostics Elecsys 2010. All of the negative samples were found negative, and all of the positive samples were found positive with Bosphore® HBV Quantification Kit v1. 15 HBV positive serum samples which have been previously analyzed using HBV DNA Amplicor Monitor Test v2.0 Quantitative, COBAS HBV DNA Amplicor Monitor Quantitative and Diasorin ETI-MAK2 HBsAg Plus Assay were tested with Bosphore® HBV Quantification Kit v1. All the samples were found to be positive, including the sample that could only be detected as positive by Diasorin ETI-MAK2 HBsAg Plus Assay, but was noted as below detection limit using HBV DNA Amplicor Monitor Test v2.0 Quantitative, and COBAS HBV DNA Amplicor Monitor Quantitative.

12.6. Calibration Against WHO Standard

Quantitation Standards were calibrated against the WHO International Standard for HBV DNA NAT assays (NIBSC Code 97/750). 1 IU was found to be equal to 4.5 ± 0.2 copies/ml.

13. REFERENCES

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- 3. Hepatitis B Fact Sheet No. 204. 2008, World Health Organization.
- 4. Jinlin Hou, Zhihua Liu, and Fan Gu, Epidemiology and Prevention of Hepatitis B Virus Infection, Int. J. Med. Sci. 2005 2(1),p: 50-57

14. SYMBOLS



Use by



Lot/Batch



Catalog number



Temperature limitation

Caution, consult accompanying documents



Manufacturer



In Vitro Diagnostic Medical Device

15. CONTACT INFORMATION



Anatolia Tanı ve Biyoteknoloji A.Ş.

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