AmpliSens[®] EBV-screen/monitor-FRT PCR kit

use

Instruction Manual

KEY TO SYMBOLS USED

REF	Catalogue number	$\overline{\Sigma}$	Contains sufficient for <n> tests</n>
LOT	Batch code	$\mathbf{\Sigma}$	Use-by Date
IVD	<i>In vitro</i> diagnostic medical device	ī	Consult instructions for use
VER	Version	Ň	Keep away from sunlight
X	Temperature limit	NCA	Negative control of amplification
-	Manufacturer	C-	Negative control of extraction
${}^{\frown}$	Date of manufacture	C+	Positive control of amplification
EC REP	Authorized representative in the European Community	IC	Internal control
$\overline{\mathbb{V}}$	Caution	PCE	Positive Control of Extraction
		KSG1, KSG2	DNA calibrators

1. INTENDED USE

AmpliSens[®] EBV-screen/monitor-FRT PCR kit is an *in vitro* nucleic acid amplification test for qualitative and quantitative detection of the Epstein-Barr virus (EBV) DNA in the clinical material (peripheral blood plasma, amniotic fluid, cerebrospinal fluid (liquor), saliva, oropharyngeal swabs, bronchoalveolar lavage, whole human blood, white blood cells, and viscera biopsy material) by using real-time hybridization-fluorescence detection. The results of PCR analysis are taken into account in complex diagnostics of

NOTE: disease

2. PRINCIPLE OF PCR DETECTION

EBV detection by the polymerase chain reaction (PCR) is based on the amplification of pathogen genome specific region using special EBV primers. In real-time PCR the amplified product is detected using fluorescent dyes. These dyes are usually linked to oligonucleotide probes which bind specifically to the amplified product during thermocycling. The real-time PCR monitoring of the fluorescence intensities during the real-time PCR allows the detection of accumulating product without re-opening the reaction tubes after the PCR run.

AmpliSens[®] EBV-screen/monitor-FRT PCR kit is a qualitative and quantitative test which is used with two internal controls: exogenous and endogenous. The Internal Control STI-87 (IC) – the internal exogenous control – must be used in the extraction procedure of peripheral blood plasma, amniotic fluid, cerebrospinal fluid (liquor), saliva, oropharyngeal swabs, bronchoalveolar lavage in order to monitor test stages of each individual sample and to identify possible reaction inhibition.

When the clinical material containing cells (whole human blood, white blood cells, and viscera biopsy material) is extracted then the DNA fragment of β -globin gene is amplified. DNA fragment of β-globin gene is used as an internal endogenous control (a human genome fragment). Thus, the use of an endogenous internal control makes it possible not only to monitor test stages (DNA extraction and amplification) but also to assess the

only to monitor test stages (DNA extraction and amplification) but also to assess the adequacy of sampling and storage of clinical material. **AmpliSens® EBV-screen/monitor-FRT** PCR kit uses 'hot-start", which greatly reduces the frequency of nonspecifically primed reactions. "Hot-start" is guaranteed by using a chemically modified polymerase (TaqF). The chemically modified polymerase (TaqF) is activated by heating at 95 °C for 15 min. The PCR kit contains the system for prevention of contamination by amplicons using the enzyme uracil-DNA-glycosylase (UDG) and dUTP. The enzyme UDG recognizes and catalyzes the destruction of the DNA containing deoxyuridine, but has no effect on DNA postpointed polymerities. DNA writing a bacet in the authorite DNA but is a burger.

containing deoxythymidine. Deoxyuridine is absent in the authentic DNA, but is always present in amplicons, because dUTP is a part of dNTP mixture in the reagents for the amplification. Due to the deoxyuridine containing contaminating amplicons are sensitive to the destruction by UDG before the DNA-target amplification. So the amplicons cannot be amplified.

The enzyme UDG is thermolabile. It is inactivated by heating at temperature above 50 °C. Therefore, UDG does not destroy the target amplicons which are accumulated during PCR. The results of amplification are registered in the following fluorescence channels:

			I able 1
Channel for fluorophore	FAM	JOE	ROX
DNA-target	DNA (IC) Glob	Human gammaherpesvirus 4 DNA	Internal Control STI-87L (IC) DNA
Target gene	DNA fragment of the β -globin gene	DNA fragment of the LMP gene	Artificially synthesized sequence

3. CONTENT

AmpliSens[®] EBV-screen/monitor-FRT PCR kit is produced in 1 form: variant FRT-100 F REF R-V9-100-S(RG,iQ,Mx)-CE.

Variant FRT-100 F includes

Reagent		Description	Volume, ml	Quantity
PCR-mix-1-FL EB screen/monitor	/	clear liquid from colorless to light lilac colour	0.6	2 tubes
PCR-mix-2-FRT		colorless clear liquid	0.3	2 tubes
Polymerase (TaqF)	colorless clear liquid	0.03	2 tubes
RNA-buffer		colorless clear liquid	0.6	1 tube
DNA calibrators	KSG1	colorless clear liquid	0.2	1 tube
	KSG2	colorless clear liquid	0.2	1 tube
RNA-buffer		colorless clear liquid	1.2	1 tube
Negative Control (C-)*		colorless clear liquid	1.2	2 tubes
Positive Control DNA EBV and human DNA**		colorless clear liquid	0.1	2 tubes
Internal Control STI-87 (IC)***		colorless clear liquid	0.6	2 tubes

must be used in the extraction procedure as Negative Control of Extraction (C must be used in the extraction procedure as Positive Control of Extraction (PCE). add 10 μ I of Internal Control STI-87 (IC) during the DNA extraction procedure directly ***

to the sample/lysis mixture. Variant FRT-100 F is intended for 110 reactions, including controls and DNA calibrators.

- 4. ADDITIONAL REQUIREMENTS
- Hemolytic. DNA extraction kit.
- Disposable powder-free gloves and a laboratory coat
- Pipettes (adjustable)
- Sterile RNase-free pipette tips with aerosol filters (up to 200 µl).
- Tube racks. Vortex mixer
- Desktop centrifuge with rotor for 2-ml reaction tubes.
 - PCR box.
- Real-time instruments (for example, Rotor-Gene 3000/6000 (Corbett Research, Australia), Rotor-Gene Q (QIAGEN, Germany), iCycler iQ5 (Bio-Rad, USA), Mx3000P (Stratagene, USA), or equivalent).
- Disposable polypropylene PCR tubes (0.1- or 0.2-ml) when working with PCR kit variant FRT-100 F a) 0.2-ml PCR tubes with optical transparent domed or flat caps if a plate-type
 - instrument is used; b) 0.2-ml PCR tubes with flat caps or strips of four 0.1-ml Rotor-Gene PCR tubes if a
 - rotor-type instrument is used Refrigerator for 2-8 °C.
- Deep-freezer at the temperature from minus 24 to minus 16 °C. Reservoir for used tips.
- 5. GENERAL PRECAUTIONS The user should always pay attention to the following:
- - Use sterile pipette tips with aerosol barriers and use a new tip for every procedure. Store all extracted positive material (specimens, controls and amplicons) away from all other reagents and add it to the reaction mix in a distantly separated facility. Thaw all components thoroughly at room temperature before starting detection.
- When thawed, mix the components and centrifuge briefly.
- Use disposable protective gloves and laboratory cloths, and protect eyes while samples and reagents handling. Thoroughly wash hands afterward. Do not eat, drink, smoke, apply cosmetics, or handle contact lenses in laboratory work
- areas Do not use a kit after its expiration date.
- Dispose of all specimens and unused reagents in compliance with local regulations.
- Samples should be considered potentially infectious and handled in a biological cabinet
- in accordance with appropriate biosafety practices. Clean and disinfect all samples or reagent spills using a disinfectant, such as 0.5 % sodium hypochlorite, or other suitable disinfectant.
- Avoid samples and reagents contact with the skin, eyes, and mucous membranes. If these solutions come into contact, rinse the injured area immediately with water and seek medical advice immediately.
- Safety Data Sheets (SDS) are available on request.
- Use of this product should be limited to personnel trained in DNA amplification techniques. Workflow in the laboratory must be one-directional, beginning in the Extraction Area and
- moving to the Amplification and Detection Area. Do not return samples, equipment and reagents to the area in which the previous step was performed.



Some components of this kit contain sodium azide as a preservative. Do not use metal tubing for reagent transfer.

For Professional Use Only

6. SAMPLING AND HANDLING

Obtaining samples of biological materials for PCR-analysis, transportation, and storage are described in the manufacturer's handbook [1]. It is recommended NOTE

that this handbook is read before starting work. AmpliSens[®] EBV-screen/monitor-FRT PCR kit is intended for the analysis of DNA extracted using DNA extraction kits from the clinical material (peripheral blood plasma, amniotic fluid, cerebrospinal fluid (liquor), saliva, oropharyngeal swabs, bronchoalveolar lavage, whole human blood, white blood cells, and viscera biopsy material). Whole peripheral and umbilical blood

Before extraction it is necessary to pretreat blood. Transfer 1.0 ml of Hemolytic (REF 137-CE, manufactured by Federal Budget Institute of Science "Central Research Institute for Epidemiology") and 0.25 ml of whole blood into 1.5-ml Eppendorf-type tube using an individual tip. Carefully vortex the content of the tube and incubate it for 10 min with periodic stirring. Centrifuge tubes at 8000 rpm for 2 min. Remove the supernatant using vacuum aspirator. Do not disturb the pellet. After washing, the pellet should be white. A small quantity of a pinkish film above the pellet (crythrocyte debris) is allowed. Washing with Hemolytic can be repeated, if necessary. The obtained pellet of leukocytes should be lysed immediately (in case of RIBO-prep extraction, add 300 µl of Solution for Lysis and then extract DNA according to RIBO-prep Instruction manual; do not add Solution for Lysis again) or it can be stored at \leq -68 °C for a long time.

again of it can be stored as 3-06 °C hor a hong time. Packed white cells of peripheral and/or umbilical blood Blood can be stored for 6 hours after sampling at room temperature. To obtain white cells, centrifuge tube with blood at 800-1600 g (3000 rpm) for 20 min. Then remove the white film formed on the surface of the blood and carry out the pretreatment as described for whole peripheral and umbilical blood. White blood cells of peripheral and umbilical blood can be stored at ≤ -68 °C for a long time.

7. WORKING CONDITIONS

AmpliSens® EBV-screen/monitor-FRT PCR kit should be used at 18-25 °C.

8. PROTOCOL

8.1. DNA extraction

- It is recommended to use the following nucleic acid extraction kits:
- RIBO-prep, REF K2-9-Et-100-CE.
- NucliSENS easyMAG automated system (for details see Guidelines [2]).
- The DNA extraction of each clinical sample is carried out in the presence of Internal Control STI-87 (add 10 µl of Internal Control STI-87 into each sample).
- Addition of Internal Control STI-87 is not necessary for the samples of whole human blood, white blood cells, and viscera biopsy material. NOTE:
- In the extraction procedure it is necessary to carry out the control reaction Add 100 µl of Negative Control (C-) to the tube labelled C- (Negative C-
- Control of Extraction). Add 90 µl of Negative Control and 10 µl of Positive Control DNA *EBV* and human DNA to the tube labeled PCE (Positive Control of Extraction). PCF
- NOTE: Extract the DNA according to the manufacturer's protocol.

8.2. Preparing PCR

The total reaction volume is 25 µl, the volume of DNA sample is 10 µl.

8.2.1 Preparing tubes for RT-PCR

- Prepare the mixture of PCR-mix-2-FRT and polymerase (TaqF). For this purpose transfer 30 μ I of polymerase (TaqF) into the tube with PCR-mix-2-FRT and vortex without foam forming. The prepared mixture is intended for analysis of 60 samples. The mixture is to
- NOTE: be stored at the temperature 2-8 °C for 3 months and used as needed. If the mixture cannot be used up for 3 months, it is necessary to prepare a
- NOTE: mixture for fewer reactions. For example, mix 150 µl of PCR-mix-2-FRT and 15 µl of polymerase (TaqF). The obtained mixture is intended for 30 reactions.
- 2. Prepare the reaction mixture.

Even for analysis of one DNA sample in the qualitative format, it is necessary to run **two controls** of amplification: the Positive Control of Amplification (KSG2) and the Negative Control of Amplification (RNA-buffer). And even for analysis of

- one DNA sample in the quantitative format, it is necessary to run five controls of amplification: two calibrators (KSG1 and KSG2) in two replicates and the NOTE: Negative Control of Amplification (RNA-buffer). In addition, you should take reagents for one extra reaction.
- 3. Mix PCR-mix-1-FL EBV screen/monitor and the mixture of PCR-mix-2-FRT and polymerase (TaqF) prepared before in the individual tube in the following proportion: 10 µl of PCR-mix-1-FL EBV screen/monitor,

 - 5 µl of mixture of PCR-mix-2-FRT and polymerase (TaqF). Calculate the required reaction number including clinical and control samples (see Table 2)

Total reaction volume is 25 µl, including the volume of DNA sample - 10 µl					
Reagent vo	lume for 1 reaction (µl)	10.0	5.0		
Quantity of	of clinical samples				
For quantitative analysis	For qualitative analysis	PCR-mix-1-FL EBV screen/monitor ¹	Mixture of PCR-mix-2-FRT and polymerase (TaqF) ¹		
1	4	70	35		
2	5	80	40		
3	6	90	45		
4	7	100	50		
5	8	110	55		
6	9	120	60		
7	10	130	65		
8	11	140	70		
9	12	150	75		
10	13	160	80		
11	14	170	85		
12	15	180	90		
13	16	190	95		
14	17	200	100		
15	18	210	105		
16	19	220	110		
17	20	230	115		
18	21	240	120		
19	22	250	125		
20	23	260	130		
21	24	270	135		
22	25	280	140		
23	26	290	145		
24	27	300	150		
25	28	310	155		
30	33	360	180		

Scheme of reaction mixture preparation

If 60 samples are analyzed simultaneously, you can use a simplified variant of mixture preparation: transfer the content of one tube with PCR-mix-2-FRT and

NOTE: the content of one tube with polymerase (TaqF) into the tube with PCR-mix-1-FL

- EBV screen/monitor. 4. Take the required number of tubes for amplification for the clinical and control samples. Transfer 15 µl of the prepared mix into each tube
- 5. Add 10 µl of DNA obtained at the DNA extraction stage into tubes with the reaction mixture.
- 6. Carry out the control amplification reactions:
- For qualitative analysis Add **10 µI** of **RNA-buffer** to the tube labeled NCA (Negative Control of Amplification). NCA
- Add 10 µl of KSG2 to the tube labeled C+ (Positive Control of Amplification). C+ C-
- Add 10 μ I of the sample extracted from the Negative Control (C–) reagent to the tube labeled C– (Negative control of Extraction).
- PCE Add 10 ul of the sample extracted from the Positive control DNA EBV and human DNA reagent to the tube labeled PCE (Positive control of Extraction). For quantitative analysis:
- Add 10 µl of RNA-buffer to the tube labeled NCA (Negative Control of NCA Amplification)
- KSG1 - Add 10 µl of KSG1 to two tubes and 10 µl of KSG2 to other two tubes KSG2
- c– Add 10 μI of the sample extracted from the Negative Control (C–) reagent to the tube labeled C- (Negative control of Extraction).
- Add 10 µl of the sample extracted from the Positive control DNA EBV and PCE human DNA reagent to the tube labeled PCE (Positive control of Extraction).

8.2.2. Amplification

1. Create a temperature profile on your instrument as follows:

Table 3

	AmpliSens-1 amplification program						
	Rotor-type instruments ²			Plate-type instruments ³			
Step	Temperature, °C	Time	Cycles	Temperature, °C	Time	Cycles	
1	95	15 min	1	95	15 min	1	
	95	5 s		95	5 s		
2	60	20 s	5	5	60	20 s	5
	72	15 s		72	15 s		
	95	5 s		95	5 s		
3	60	20 s	40	60	30 s	40	
	72	15 s		72	15 s		

Fluorescent signal is detected in the channels for the FAM and JOE fluorophores (when another tests are performed simultaneously the detection in another channels is enabled).

Adjust the fluorescence channel sensitivity according to the Important Product Information Bulletin and Guidelines [2].

3.

Insert tubes into the reaction module of the device. Run the amplification program with fluorescence detection.

5. Analyze results after the amplification program is completed.

¹ Values are given with account of one extra reaction and five controls (2 DNA-calibrators KSG1 and KSG2 (in two replicates), negative control (RNA-buffer) for quantitative analysis of EBV DNA, and two repiredies), regardle control (RNA-bullet) for qualitative analysis of EBV DNA, and two controls (positive and negative) for qualitative analysis of EBV DNA. ² For example, Rotor-Gene 3000, Rotor-Gene 6000 (Corbett Research, Australia), Rotor-Gene Q (QIAGEN, Germany).

9. DATA ANALYSIS

Analysis of results is performed by the software of the real-time PCR instrument used by measuring fluorescence signal accumulation in three channels

- The signal of β -globin gene DNA (IC Glob) amplification product is detected in the channel for the FAM fluorophore,
- The signal of EBV DNA (Positive Control DNA EBV and human DNA) amplification product is detected in the channel for the JOE fluorophore
- The signal of Internal Control STI-87 (IC) DNA amplification product is detected in the channel for the ROX fluorophore.

The results are interpreted by the software of the used instrument by the crossing (or not crossing) of the fluorescence curve with the threshold line.

9.1. Interpretation of results for DNA extracted from cell suspension (whole human blood, white blood cells, viscera biopsy material)

- The results are analysed in two channels:
 the signal of *B*-globin gene DNA (IC Glob) amplification product is detected in in the channel for the FAM fluorophore,
 the signal of *EBV* DNA (Positive Control DNA *EBV* and human DNA) amplification
- product is detected in the in the channel for the JOE fluorophore. If the total DNA is extracted from cell suspension (whole human blood, white blood
- <u>cells, viscera biopsy material)</u> the principle of interpretation is the following: 1. *EBV* DNA is **detected** if the *Ct* value determined in the results grid in the channel for the JOE fluorophore does not exceed the threshold value of the positive result (for details see Guidelines [2]). Moreover, the fluorescence curve of the sample should cross the
- threshold line in the area of typical exponential growth of fluorescence. *EBV* DNA is **not detected** if the *Ct* value is not determined (absent) in the results grid in the channel for the JOE fluorophore (the fluorescence curve does not cross the threshold line), whereas the *Ct* value in the channel for the FAM fluorophore does not exceed the boundary *Ct* value specified in the *Important Product Information Bulletin* (for qualitative analysis) or the quantity of IC Glob DNA is more than 2000 copies/reaction (for quantitative analysis).
- The result is **invalid** if the *Ct* value is not determined (absent) in the channel for JOE fluorophore, whereas the *Ct* value in the channel for the FAM fluorophore is greater than 3 the boundary Ct value specified in the *Important Product Information Bulletin* (for qualitative analysis) or the quantity of IC Glob DNA is less than 2000 copies/reaction (for quantitative analysis). In such case the PCR analysis should be repeated for required sample.
- The result is **equivocal** if the Ct value in the channel for JOE fluorophore exceeds the boundary Ct value specified in the Important Product Information Bulletin. In that case, it is necessary to conduct additional analysis for that DNA sample with two repeats. If the repeated positive *Ct* value is obtained, the result is considered positive. If the positive *Ct* value can't be reproduced in two repeats, the result is considered **equivocal**.
- The negative result is considered unreliable if the *Ct* value in the channel for **FAM** fluorophores is greater than the boundary value specified in the *Important Product Information Bulletin* (for qualitative analysis). The quantitative positive or negative results are considered **unreliable** if the quantity of IC Glob DNA is less than 2000 copies/reaction.

The result of the analysis is considered reliable only if the results obtained for Positive and Negative Controls of amplification as well as for the Negative Control of extraction are correct (see Table 4). For quantitative analysis the results for Positive Control should fall in the concentration range specified in the *Important Product* Information Bulletin

Table 4 Results for controls for DNA extracted from cell suspension (whole human blood, white blood cells, and viscera biopsy material)

	With	Ct in the channel for fluorophore					
Control	Stage for control	F	AM		JOE		
		Qualitativ e format	Quantitativ e format	Qualitative format	Quantitative format		
C-	DNA extraction, PCR	Absent	Absent	Absent	Absent		
PCE	DNA extraction, PCR	< boundary value	< boundary value	< boundary value	concentration value falls in the range specified in the Important Product Information Bulletin		
NCA	PCR	Absent	Absent	Absent	Absent		
C+	PCR	< boundary value	_	< boundary value	_		
KSG1, KSG2	PCR	_	Ct value and calculated concentration are defined	_	Ct value and calculated concentration are defined		

For quantitative analysis the concentration in logarithm of EBV DNA copies per standard cell quantity (10⁵) in control and clinical samples (whole human blood, white blood cells, and viscera biopsy material) is calculated according to the following formula:

number of EBV DNA copies in PCR

sample number of Glob DNA copies in — x 2·10⁵}= Ig {EBV DNA copies/10⁵ cells} lg { · PCR sample

To express relative concentration of EBV DNA in copies per standard cells quantity (for example, 10^5), use the scaling ratio: 10^5 of cells = 2.10⁵ human genomes

9.2. Interpretation of results for DNA extracted from peripheral blood plasma, amniotic fluid, cerebrospinal fluid (liquor), saliva, oropharyngeal swabs, bronchoalveolar lavage with internal control sample

The results are analysed in two channels:

- the signal of the *EBV* DNA (Positive Control DNA *EBV* and human DNA) amplification product is detected in the channel for the JOE fluorophore,
- the signal of the Internal Control STI-87 (IC) DNA amplification product is detected in the channel for the ROX fluorophore.
- If the total DNA is extracted from peripheral blood plasma, amniotic fluid, cerebrospinal fluid (liquor), saliva, oropharyngeal swabs, bronchoalveolar lavage with internal control sample, principle of interpretation is the following: 1. EBV DNA is detected if the Ct value determined in the results grid in the channel for the
- JOE fluorophore does not exceed the threshold value of the positive result (for details

see Guidelines [2]). Moreover, the fluorescence curve of the sample should cross the

- Construction of the series of typical exponential growth of fluorescence.
 EBV DNA is not detected if the Ct value is not determined (absent) in the results grid in the channel for the JOE fluorophore (the fluorescence curve does not cross the threshold line), whereas the Ct value in the channel for the ROX fluorophore does not cross the threshold line), whereas the curve does not cross the threshold line by detection of the location exceed the boundary Ct value specified in the Important Product Information Bulletin.
- The result is invalid if the Ct value is not determined (absent) in the channel for the JOE fluorophore, whereas the Ct value in the channel for the ROX fluorophore is not determined (absent) or greater than the boundary *Ct* value specified in the *Important Product Information Bulletin*. In such cases, the PCR analysis should be repeated for required sample.
- 4. The result is considered to be equivocal if the Ct value in the channel for the JOE fluorophore exceeds the boundary Ct value specified in the Important Product Information Bulletin. In that case, it is necessary to conduct additional analysis for that DNA sample with two repeats. If the repeated positive Ct value is obtained, the result is considered positive. If the positive Ct value can't be reproduced in two repeats, the result is considered equivocal.

The result is considered equivocal. The result of the analysis is considered reliable only if the results obtained for Positive and Negative Controls of amplification as well as for the Negative Control of extraction are correct (see Table 5). For quantitative analysis the results for Positive Control should fall in the concentration range specified in the *Important Product* Information Bulletin

Table 5 Results for controls for DNA extracted from peripheral blood, amniotic fluid, cerebrospinal fluid (liquor), saliva, oropharyngeal swabs, bronchoalveolar lavage with internal control

		Ct in the channel for fluorophore					
Control	Stage for control		JOE	ROX			
		Qualitative format	Qualitative format	Qualitative format	Qualitative format		
C-	DNA extraction, PCR	Absent	Absent	< boundary value	< boundary value		
PCE	DNA extraction, PCR	< boundary value	Ct value falls in the range specified in the Important Product Information Bulletin	< boundary value	< boundary value		
NCA	PCR	Absent	Absent	Absent	Absent		
C+	PCR	< boundary value	-	< boundary value	-		
KSG1, KSG2	PCR	_	Ct value and calculated concentration are defined		Ct value and calculated concentration are defined		

For quantitative analysis the concentration of EBV DNA (CS EBV DNA) per ml of sample (peripheral blood plasma, amniotic fluid, cerebrospinal fluid (liquor), saliva, oropharyngeal swabs, urine samples, and bronchoalveolar lavage) is calculated according to the following formula:

CS EBV DNA = [KEBV DNA/ KSTI-87] x IC coefficient (copies/ml)

KEBV DNA - quantity of EBV DNA copies in DNA-sample;

K_{STI-87} – quantity of STI-87 DNA copies in DNA-sample;

IC coefficient – quantity of Internal Control STI-87 DNA copies in DNA-sample. IC coefficient, Positive Control DNA *EBV* and human DNA, Internal Control STI-87 and DNA calibrators' concentrations as well as boundary *Ct* values are specified in the Important Product Information Bulletin.

10. TROUBLESHOOTING

- Results of analysis are not taken into account in the following cases:
- If any Ct value appears in the channel for ROX fluorophore for the Negative Control of Amplification (NCA), in the channels for FAM and JOE fluorophores for Negative Control of Amplification (NCA) and Negative Control of Extraction (C-). The results testify the presence of contamination of reagents or samples. In that case the PCR analysis (beginning with the extraction stage) should be repeated for all samples, in which DNA vas found,
- 2. If Ct value is absent or greater than the threshold in the results grid for the Positive Control of Amplification (C+) KSG2 in the channel for JOE (EBV), FAM or ROX fluorophores, the amplification should be repeated for all samples where EBV DNA was not detected.
- If the Ct value is absent or greater than threshold for the Positive Control of Extraction (PCE) Positive Control DNA EBV and human DNA in the channel for JOE (EBV), FAM or ROX fluorophores, the results of analysis must be considered as invalid for all 3. samples. PCR should be repeated for all samples. 4. If the Ct value is absent or is greater than the specified boundary value in channel for
- JOE and the Ct value in the channels for FAM or ROX fluorophore is greater than the maximal value for ${\rm IC},$ the experiment should be repeated starting from DNA extraction stage
- If the Ct value is greater than the specified boundary value in channel for JOE and the Ct value in the channels for FAM or ROX is less than the specified boundary value, the results of analysis must be considered as equivocal. In that case, it is necessary to conduct additional analysis for that DNA sample with two repeats. If the repeated positive *Ct* value is obtained, the result is considered positive. If the positive *Ct* value can't be reproduced in two repeats, the result is considered **equivocal**.
- If in quantitation analysis the copies/reaction values in DNA calibrators differ by more than for 30 % from the set values, it is necessary to check the tube order in the rotor (calibrators should be placed in the wells indicated as **Standard** in sample table, concentration specified in the *Important Product*
- Information Bulletin, well no.1 must be filled with some test tube (not empty)). If the correlation coefficient R in **Standard Curve** window is less than 0.9 (in case of quantitative analysis), it means that calibration failed. Check the settings of calibrators and correct inaccuracies, if no effect, repeat PCR for all samples and calibrators

If you have any further questions or if you encounter problems, please contact our Authorized representative in the European Community.

11. TRANSPORTATION

AmpliSens® EBV-screen/monitor-FRT PCR kit should be transported at 2-8 °C for no longer than 5 days

12. STABILITY AND STORAGE

All components of the AmpliSens[®] EBV-screen/monitor-FRT PCR kit are to be stored at

AmpliSens® EBV-screen/monitor-FRT PCR kit REF R-V9-100-S(RG,iQ,Mx)-CE / VER 02.04.19-11.03.21 / Page 3 of 4

2-8 °C when not in use (except for PCR-mix-1-FL EBV screen/monitor, PCR-mix-2-FRT, and polymerase (TaqF)). All components of the AmpliSens® EBV-screen/monitor-FRT PCR kit are stable until the expiration date on the label. The shelf life of opened reagents is the same as that of unopened reagents, unless otherwise stated. NOTE- PCR-mix-1-FL *EBV* screen/monitor, PCR-mix-2-FRT and polymerase (TaqF) are

NOTE: to be stored at temperature from minus 24 to minus 16 °C

NOTE: PCR-mix-1-FL EBV screen/monitor is to be kept away from light.

13. SPECIFICATIONS

13.1. Analytical sensitivity The linear range of **AmpliSens®** *EBV-screen/monitor-FRT* PCR kit is **500** – **10.000.000 copies/ml**. If the result is more than 10.000.000 copies/ml, it is indicated as *the* result is more than 10.000.000 *EBV* DNA copies/ml. If the result is less than 500 copies/ml, it is indicated as the result is less than 500 EBV DNA copies/ml.

Clinical material	DNA extraction kit	Analytical sensitivity
Peripheral blood plasma, amniotic fluid, cerebrospinal fluid (liquor), saliva, oropharyngeal swabs, bronchoalveolar lavage	RIBO-prep	400 copies/ml
Whole human blood, white blood cells, viscera biopsy material	RIBO-prep	5 <i>EBV</i> DNA copies per 10 ⁵ cells

13.2. Analytical specificity

The analytical specificity of **AmpliSens®** *EBV-screen/monitor-FRT* PCR kit is ensured by the selection of specific primers and probes as well as stringent reaction conditions. The primers and probes have been checked for possible homologies to all sequences published in gene banks by sequence comparison analysis. AmpliSens[®] *EBV*-screen/monitor-FRT PCR kit is intended for *Epstein-Barr virus* DNA

fragment detection. The specific activity of AmpliSens® *EBV-screen/monitor-FRT* PCR kit is proved by analyzing QCMD panels as well as by analyzing clinical material with subsequent confirmation of results by sequencing the amplification fragments. The activity of PCR kit components with respect to DNA of other viruses (*human cytomegalovirus*, herpes simplex virus types 1 and 2, human herpes virus types 6 and 8, Varicella Zoster Virus, Parvovirus B19 and others), bacterial pathogens (Staphylococcus aureus, Streptococcus pyogenes, Streptococcus agalactiae and others), and human DNA is absent. The clinical specificity of AmpliSens[®] EBV-screen/monitor-FRT PCR kit was confirmed in laboratory clinical trials

14. REFERENCES

- Handbook "Sampling, Transportation, and Storage of Clinical Material for PCR diagnostics", developed by Federal Budget Institute of Science "Central Research Institute for Epidemiology" of Federal Service for Surveillance on Consumers' Rights Protection and Human Well-Being.
 Guidelines to the AmpliSens[®] EBV-screen/monitor-FRT PCR kit for qualitative detection and quantitation of Epstein-Bar virus (EBV) DNA in the clinical materials by using real-time hybridization-fluorescence detection developed by Federal Budget Institute of Science "Central Research Institute for Epidemiology".

15. QUALITY CONTROL

In compliance with Federal Budget Institute of Science "Central Research Institute for Epidemiology" ISO 13485-Certified Quality Management System, each lot of the AmpliSens® *EBV*-screen/monitor-FRT PCR kit has been tested against predetermined specifications to ensure consistent product quality.

List of Changes Made in the Instruction Manual

VER	Location of changes	Essence of changes	
17.06.11 RT	Cover page, text	The name of Institute was changed to Federal Budget Institute of Science "Central Research Institute for Epidemiology"	
	Through the text	Corrections according to the template. Grammar corrections	
	2. Principle of PCR detection	The additions about using endogenous and exogenous internal controls	
25.05.15 ME	8.1. DNA extraction	The phrase: "Addition of Internal Control STI-87 is not necessary for the samples of whole human blood, white blood cells, and viscera biopsy material" was added	
	8.2.1 Preparing tubes for PCR	Appendix 1 was integrated into the text of the instruction manual as Table 1	
	9. Data analysis	The sections were rewritten	
	10. Troubleshooting		
28.01.19 DV	2. Principle of PCR detection	The information about the enzyme UDG was added. The information about "hot-start" was corrected	
13.02.19 DV	3. Content	The color of the reagent was specified	
	Through the text	The text formatting was changed	
13.07.20 MM	2. Principle of PCR detection	The table with targets was added	
	Footer	The phrase "Not for use in the Russian Federation" was added	
19.01.21 EM	10. Troubleshooting	The information for Negative Control of Amplification (NCA) and Negative Control of Extraction (C–) was corrected	
11.03.21 EM	_	The name, address and contact information for Authorized representative in the European Community was changed	

AmpliSens[®]



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