

# BactoReal<sup>®</sup> Kit

## *Neisseria meningitidis*

## Manual

**CE****IVD**For *in vitro* diagnostic use**REF**

DHUB00153

**Σ**

50 reactions



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## Index

1. Intended use.....	3
2. Product description.....	3
3. Pathogen information.....	3
4. Contents of the kit, stability and storage.....	4
5. Additionally required materials and devices.....	4
6. Precautions and safety information.....	4
7. Limitations.....	5
8. Preparation of samples.....	6
9. Preparation of real-time PCR.....	6
9.1. Pipetting scheme.....	6
9.2. Programming of temperature profile.....	7
10. Interpretation of PCR-data.....	7
11. Troubleshooting.....	8
11.1. No signal in FAM channel and Cy5 channel with controls and sample:.....	8
11.2. Valid results for controls, but no signal in FAM channel and Cy5 channel with sample:.....	8
11.3. Signal in FAM channel in negative control:.....	8
11.4. Signal in FAM channel in extraction negative control:.....	8
12. Specifications and performance evaluation.....	9
12.1. Kit performance.....	9
12.2. Limit of detection and linearity.....	10
12.3. Analytical specificity.....	10
12.4. Diagnostic evaluation.....	10
13. References.....	10

## Explanation of symbols



Batch code



Catalogue number



Contains sufficient for <n> tests



This product fulfils the requirements of the European Directive 98/79 EC for *in vitro* diagnostic medical devices



Corrosion, GHS05



Use by



Manufactured by



Store at



For *in vitro* diagnostic use



Exclamation mark, GHS07

## 1. Intended use

BactoReal<sup>®</sup> Kit *Neisseria meningitidis* is an *in vitro* diagnostic test, based on real-time PCR, for the detection of *Neisseria meningitidis*.

## 2. Product description

BactoReal<sup>®</sup> Kit *Neisseria meningitidis* detects two regions of the capsular transport protein (ctrA) gene of *N. meningitidis*.

This test allows the rapid and sensitive detection of DNA of *N. meningitidis* purified from EDTA-blood, blood culture or cerebrospinal fluid (e.g. with the QIAamp DNA Mini Kit, Qiagen). This kit detects all known serotypes of *N. meningitidis*.

A probe-specific amplification-curve in the FAM channel indicates the amplification of *N. meningitidis* specific DNA. An internal DNA positive control (DNA IPC) is detected in Cy5 channel and is used as DNA extraction as well as real-time PCR inhibition control. The target for the DNA IPC is extracted with the sample.

This test has been validated with the Applied Biosystems<sup>®</sup> 7500 Fast Real-time PCR System (Thermo Fisher Scientific) and tested with a LightCycler<sup>®</sup> 480 Instrument II (Roche) and Mx3005P<sup>®</sup> QPCR System (Agilent), but is also compatible with other real-time PCR instruments which detect and differentiate fluorescence in FAM and Cy5 channel.

The test is based on real-time PCR. A specific DNA sequence of the pathogen genome is detected and amplified. The generated PCR-product is detected by an oligonucleotide-probe labelled with a fluorescent dye. This technology allows a sequence-specific detection of PCR amplicates.

Ingenetix ViroReal<sup>®</sup>, BactoReal<sup>®</sup> and ParoReal Kits are optimized to run under the same thermal cycling conditions. RNA and DNA can be analysed in one run.

## 3. Pathogen information

*Neisseria meningitidis* is a Gram-negative bacterium causing meningitis and septicemia. There are 13 clinically significant serogroups, which are classified according to the antigenic structure of their polysaccharide capsule. Five serogroups (A, B, C, Y and W135) are responsible for virtually all cases of the disease. Serogroup B is the predominant one. The capsular transport protein (ctrA) gene is unique to *N. meningitidis* and can be found in all meningococcal serogroups.

## 4. Contents of the kit, stability and storage

Labelling	Content	Amount	Storage
Neisseria meningitidis Assay Mix (green cap)	Primer and probe (FAM) for <i>N. meningitidis</i> detection	1 x 50 µl	-15 °C to -25 °C
DNA IPC-3 Assay Mix (yellow cap)	Primer and probe (Cy5) for DNA IPC detection	1 x 50 µl	-15 °C to -25 °C
DNA IPC Target (orange cap)	Target for DNA IPC (internal DNA positive control system)	1 x 100 µl	-15 °C to -25 °C
Neisseria meningitidis Positive Control (red cap)	DNA positive control (approx. 10,000 target copies/µl)	1 x 25 µl	-15 °C to -25 °C
DNA Reaction Mix (white cap)	2 x DNA reaction mix	1 x 500 µl	<b>-15 °C to -25 °C until first use, then at +4 °C</b>
Nuclease-free water (blue cap)	Nuclease-free water	1 x 1000 µl	-15 °C to -25 °C

The components of BactoReal® Kit *Neisseria meningitidis* are stable until the expiry date stated on the label.

## 5. Additionally required materials and devices

- Reagents and devices for DNA-extraction
- Nuclease-free water for dilution of DNA IPC Target
- Disposable powder-free gloves
- Pipettes (adjustable)
- Sterile pipette tips with filters
- Vortex mixer
- Desktop centrifuge with rotor for 2 ml reaction tubes
- Real-time PCR instrument which is able to detect and differentiate fluorescence in FAM and Cy5 channel
- Optical 96 well reaction plates or optical reaction tubes

## 6. Precautions and safety information

- For *in vitro* diagnostic use. The use of this kit is limited to persons instructed in the procedures of real-time PCR and *in vitro* diagnostics.
- Clean benches and devices periodically.
- Use sterile filter pipette tips.
- Specimens should be handled as if infectious in accordance with safe laboratory procedures. Wear protective disposable powder-free gloves when handling kit reagents and specimens.
- Use separated working areas for specimen preparation, reagent preparation and amplification. Supplies and equipment must be dedicated to each of these separate working areas and ensure workflow in the laboratory from pre- to post-PCR.
- Be careful when handling samples and positive control to avoid cross contamination. Change gloves after handling of samples or positive control. Store positive or potentially positive material separately from reagents
- Prevent contamination of work equipment and reagents with DNA/RNA, nuclease or amplification products by good laboratory practice.
- Quality of DNA has a profound impact on the test performance. Ensure that the used DNA extraction system is compatible with real-time PCR technology.
- Always include a negative control per PCR-run (nuclease-free water instead of sample).
- For a valid interpretation of results, a negative control should be included during DNA-extraction (e.g. extraction of water instead of sample material), in order to exclude false-positive results due to contamination with DNA of pathogen during extraction.
- Please note the expiry date of the kit.
- Do not interchange or mix reagents from kits with different lot numbers.

- Repeated thawing and freezing of kit components should be avoided. Protect kit components from light. **Caution:** DNA IPC Target is stored in RNA/DNA stabilizer which contains Guanidinium thiocyanate/Triton X-100 (see MSDS, [www.ingenetix.com](http://www.ingenetix.com)).
- Use established laboratory practices according to your local safety regulations for discarding specimens, reagents and waste.

## 7. Limitations

- Optimal performance of this test requires appropriate specimen collection, transport and storage, as well as an appropriate DNA extraction procedure.
- DNA extraction and *N. meningitidis* detection have been validated for EDTA-blood, blood culture and cerebrospinal fluid with this kit.
- A negative test result does not exclude the possibility of *N. meningitidis* infection, because test results may be affected by improper specimen collection, technical error, specimen mix-up or pathogen quantities below the assay sensitivity. The presence of PCR inhibitors may lead to invalid results.
- For this kit highly specific primers and probes have been selected. However, false-negative or less sensitive results might be obtained due to sequence heterogeneity within the target region of not yet described clinical subtypes.
- Results should be interpreted in consideration of clinical and laboratory findings.

## 8. Preparation of samples

Extract samples with a DNA extraction system compatible with real-time PCR technology. An extraction negative control should be included during DNA-extraction (e.g. extraction of water instead of sample material).

The **DNA IPC Target** has to be added during extraction. The DNA IPC is used as a control of DNA extraction, identifies possible PCR inhibition and confirms the integrity of kit reagents.

**Caution:** The DNA IPC Target must not be added directly to the sample material but has to be pipetted to the lysis buffer.

→ For an elution volume of 50-100 µl: Per sample, spike 1 µl DNA IPC Target into lysis buffer.

→ For an elution volume of >100 µl or when using an automated extraction system: Per sample, spike 2 µl DNA IPC Target into lysis buffer.

## 9. Preparation of real-time PCR

- Include one negative control (water), one positive control and one extraction negative control per PCR run.
- It is recommended to analyse samples in duplicates, which increases the probability of pathogen detection and facilitates interpretation of results.
- Thaw DNA samples on ice.
- Thaw kit components at room temperature. When thawed, mix components, centrifuge briefly and keep on ice.
- Mix the DNA Reaction Mix to ensure homogeneity of solution.
- **Positive Control**
  - Use 1 µl of *Neisseria meningitidis* Positive Control + 4 µl nuclease-free water. Pipette positive control at last.

### 9.1. Pipetting scheme

		<b>Per sample</b>
<b>Preparation of Master Mix</b> (mix well)	Nuclease-free Water*	3.0 µl
	DNA Reaction Mix	10.0 µl
	<i>Neisseria meningitidis</i> Assay Mix	1.0 µl
	DNA IPC-3 Assay Mix	1.0 µl
	<b>Total volume Master Mix</b>	<b>15.0 µl</b>
<b>Preparation of PCR</b>	Master Mix	15.0 µl
	DNA-Sample*	5.0 µl
	<b>Total volume</b>	<b>20.0 µl</b>

\*1-8 µl of the sample can be used. When using ≠ 5 µl sample, the volume of water has to be adjusted accordingly.

→ **If DNA IPC Target was not added during extraction:** Freshly dilute the DNA IPC Target 1:100 with nuclease-free water and add 1 µl per sample directly to the master mix.

**Caution:** The use of more than 1 µl diluted (1:100) DNA IPC Target per reaction causes inhibition of the real-time PCR reaction.

## 9.2. Programming of temperature profile

Please find further information on programming the real-time PCR instrument in the respective operator's manual. Take into consideration that some PCR-platforms have to be calibrated with the corresponding dye before performing a multiplex-PCR.

**Select detection channel:** FAM-TAMRA, 530 nm (for *N. meningitidis*)  
Cy5-NONE, 667 nm (for DNA IPC-3)

**Select reference dye (passive reference):** ROX

**Sample Volume:** 20 µl

**Temperature Profile:**

Program 1	Program 2	Program 3
Cycles: 1 Analysis: None	Cycles: 1 Analysis: None	Cycles: 45 Analysis: Quantification Acquisition at 60°
50°C *2 min	95°C 20 sec	95°C 5 sec 60°C 1 min

For ABI PRISM® 7500:  
Ramp speed: "Standard"

For LightCycler® 480 instrument:  
Detection format: 2 Color Hydrolysis Probe

**\*Note:** If viral RNA should be also detected in the same PCR run, program 1 has to be prolonged to 15 min at 50°C. This temperature profile can be used for all ingenetix ViroReal®, BactoReal®, MycoReal and ParoReal kits for the detection of DNA or RNA.

## 10. Interpretation of PCR-data

For analysis of PCR results select fluorescence display options 530 nm (FAM channel) for the *N. meningitidis* target and 667 nm (Cy5 channel) for the DNA IPC target. Samples with positive Ct or Cp-values are considered positive. Please additionally check the amplification-curves manually. Samples should be inspected both in logarithmic and linear scale view and compared with the negative control.

	FAM channel <i>N. meningitidis</i> target	Cy5 channel DNA IPC target	Interpretation
Negative control	Negative	Negative / Positive <sup>1</sup>	Valid
Positive control	Positive	Negative / Positive <sup>1</sup>	Valid
Extraction negative control	Negative	Positive	Valid
Sample	Positive	Positive / Negative <sup>2</sup>	Positive
Sample	Negative	Positive	Negative <sup>3</sup>
Sample	Negative	Negative	Invalid

<sup>1</sup>Only positive if the DNA IPC Target was added at a 1:100 dilution directly to the master mix

<sup>2</sup>High pathogen load in the sample can lead to reduced or absent signal of DNA IPC

<sup>3</sup>The positive signal of the DNA IPC excludes a possible PCR inhibition. However, IPC Ct-values should show comparable results. A shift of Ct-values can indicate a partial inhibition of PCR.

In case of invalid data, analysis has to be repeated with the remaining or newly extracted DNA sample (see 11. Troubleshooting).

## 11. Troubleshooting

### 11.1. No signal in FAM channel and Cy5 channel with controls and sample:

- Incorrect programming of the temperature profile or detection channels of the real-time PCR instrument.  
→ Compare temperature profile and programming of detection channels with the protocol.
- Incorrect configuration of PCR reaction.  
→ Check your work steps (see pipetting scheme) and repeat PCR, if necessary.

### 11.2. Valid results for controls, but no signal in FAM channel and Cy5 channel with sample:

- Incorrect programming of detection channels with the sample.  
→ Compare programming of detection channels with protocol.
- If the DNA IPC Target was added during extraction:
  - PCR reaction has been inhibited.
  - DNA extraction has failed.
  - The undiluted DNA IPC Target has not been added to lysis buffer of sample.
  - The extracted sample has not been added to PCR reaction.  
→ No interpretation can be made. Make sure you use a recommended method for DNA isolation and stick closely to the manufacturer's instructions. Check your work steps.

### 11.3. Signal in FAM channel in negative control:

- A contamination occurred during preparation of PCR.  
→ Repeat PCR with new reagents in replicates.  
→ Strictly pipette positive control at last.  
→ Make sure that work space and instruments are cleaned at regular intervals.

### 11.4. Signal in FAM channel in extraction negative control:

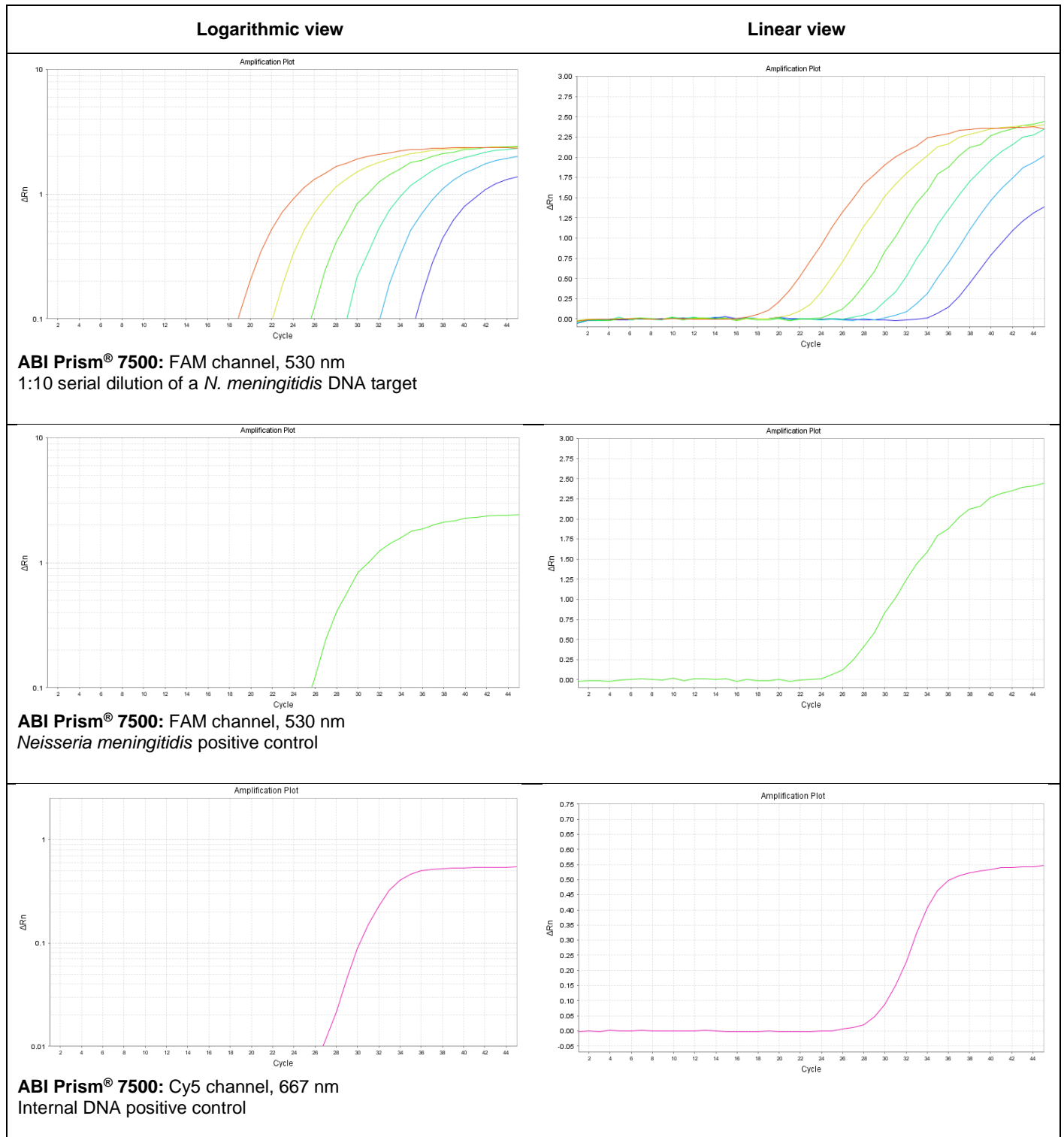
- A contamination occurred during extraction.  
→ Repeat extraction and PCR using new reagents.  
→ Make sure that work space and instruments are cleaned at regular intervals.



## 12. Specifications and performance evaluation

### 12.1. Kit performance

Performance of BactoReal® Kit *Neisseria meningitidis* with an Applied Biosystems® 7500 Fast Real-time PCR System (Thermo Fisher Scientific) is shown in Figure 1.



**Figure 1** Performance of BactoReal® Kit *Neisseria meningitidis*

## 12.2. Limit of detection and linearity

BactoReal® Kit *Neisseria meningitidis* was tested with a 10-fold dilution series of plasmids containing two fragments (region 1 and region 2) of *Neisseria meningitidis* DNA.

The **limit of detection** (LoD95 = smallest number of copies of target DNA which can be detected in 95% of cases) is 1.7 target copies/reaction.

The assay shows **linearity** for region 1 over the range of 10 to 1,000,000 target copies/reaction with a slope of  $-3.35 \pm 0.05$  and a  $R_2$  of  $> 0.99$ . For region 2 it shows linearity over the range of 100 to 1,000,000 target copies/reaction with a slope of  $-3.25 \pm 0.04$  and a  $R_2$  of  $> 0.99$  as shown in Figure 2.

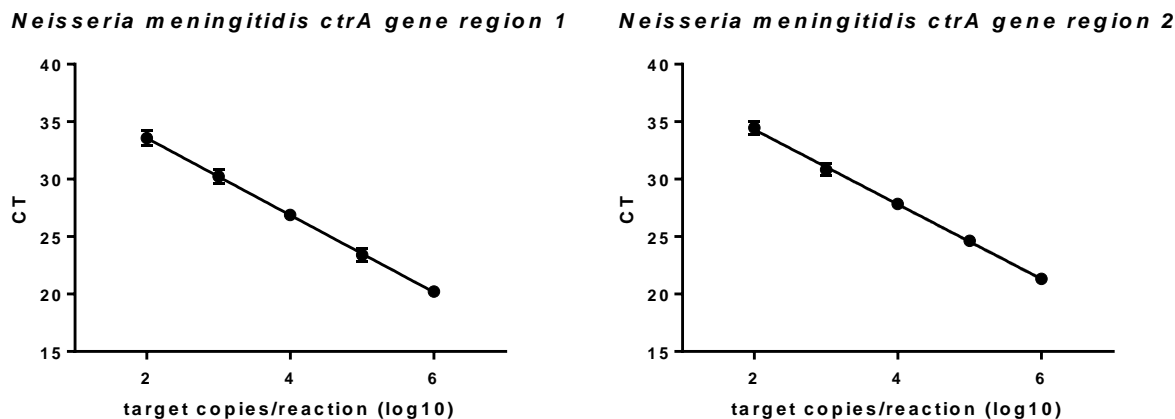


Figure 2 Ten-fold dilution series of a *N. meningitidis* DNA region 1 and region 2 standard plotted against CT

## 12.3. Analytical specificity

The selection of highly specific primers and probes ensures analytical specificity. Primers and probes have been checked for possible homologies to currently published sequences by sequence comparison analysis. This validates the detection of so far known *Neisseria meningitidis* strains.

Analytical specificity has been further evaluated by testing genomic DNA of virus (adenovirus) and bacteria (*Neisseria gonorrhoeae*, *Neisseria lactamica*, *Neisseria mucosa*, *Neisseria perflora*, *Neisseria polysaccharea*, *Neisseria subflora*, *Bordetella pertussis*, *Bordetella parapertussis*, *Haemophilus influenzae*, *Mycoplasma pneumoniae*, *Streptococcus pneumoniae*). No cross-reactions have been observed.

Furthermore, 19 *Neisseria meningitidis* isolates (seven isolates of serogroup B, five isolates of serogroup C, one isolate of serogroup W135 and six undefined isolates) were tested and correctly analysed.

With this kit two different regions of the *ctrA* gene are amplified in order to minimize false-negative results due to sequence heterogeneity within the target region. It includes also the detection of the three *N. meningitidis* strains described as *ctrA* mutants by Cavrini et al. (2010) and Jatón et al. (2010) (accession numbers GU391296, GU391295 and HQ156899).

## 12.4. Diagnostic evaluation

Diagnostic evaluation was performed with 30 DNA isolates of clinical specimens (cerebrospinal fluid n=27, secrete n=1, EDTA-blood n=1 and blood-culture n=1). Results of BactoReal® Kit *Neisseria meningitidis* were compared with an accredited reference method detecting the insertion sequence IS1106. With the reference method 15 samples were positive for *N. meningitidis*. With BactoReal® Kit *Neisseria meningitidis* 14 out of the 15 positive samples were also positive, while one sample was negative, most likely due to differences in the limit of detection between both methods. The remaining 15 samples were negative with both methods.

## 13. References

- Corless CE, Guiver M, Borrow R, Edwards-Jones V, Fox AJ, Kaczmarski EB. 2001. Simultaneous detection of *Neisseria meningitidis*, *Haemophilus influenzae*, and *Streptococcus pneumoniae* in suspected cases of meningitis and septicemia using real-time PCR. J. Clin. Microbiol. 39:1553-1558.
- Jatón K, Ninet B, Bille J, Greub G. 2010. False-negative PCR result due to gene polymorphism: the example of *Neisseria meningitidis*. J. Clin. Microbiol. 48:4590-4591.
- Cavrini F, Liguori G, Andreoli A, Sambri V. 2010. Multiple nucleotide substitutions in the *Neisseria meningitidis* serogroup C *ctrA* gene cause false-negative detection by real-time PCR. J. Clin. Microbiol. 48:3016-3018.