# virotype® BTV pan/8 2.0 RT-PCR Kit

Validation Report

For the detection of RNA from the *Bluetongue Virus* (BTV) *and the European serotype BTV-8* 





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## 1 Introduction

#### 1.1 Intended use

The virotype BTV pan/8 2.0 RT-PCR Kit is intended for the detection of *Bluetongue Virus* (BTV) RNA and differentiation of European serotype BTV-8 specifically, in ruminant whole blood and blood pools (preferred with anticoagulants, for example EDTA-blood) and tissue samples (spleen, lymph nodes) from cattle, sheep and goats.

The kit is approved by the Friedrich-Loeffler-Institut and licensed in accordance with § 11 (2) of the German Animal Health Act (FLI-C 114) for use in Germany for veterinary diagnostic procedures.

For veterinary use only.

### 1.2 General information

Bluetongue is an infectious, non-contagious disease of ruminants. The agent is the *Bluetongue Virus* (BTV), a double-stranded RNA virus of the genus *Orbivirus* of the family *Reoviridae* which includes 36 known serotypes including atypical BTV. BTV is widely distributed around the world. Sheep, cattle and goats are mainly affected by the disease. Clear clinical signs are usually seen only in sheep. In severe cases the tongue may show intense hyperemia and becomes cyanotic (Bluetongue).

BTV serotype 8 (BTV-8) is of epidemiological importance in Central Europe and cause of recent major Bluetongue Disease outbreaks. The virus is transmitted by certain midges of the genus *Culicoides*. Furthermore, the virus can be spread by contaminated needles and surgery equipment.

# 1.3 Description of the test principle

Polymerase chain reaction (PCR) is based on the amplification of specific regions of the pathogen genome. In real-time PCR, the amplified product is identified using fluorescent dyes. These are usually linked to oligonucleotide probes that bind specifically to the amplified product. Monitoring the fluorescence intensities during the PCR run (i.e., in real time) allows detection of the accumulating product without the need to re-open the reaction tubes afterward.

The virotype BTV pan/8 2.0 RT-PCR Kit contains all of the necessary reagents for the detection of BTV RNA, including a Positive and Negative Control. With this kit, both, reverse transcription and PCR are performed in one reaction tube, reducing the risk of contamination.

The virotype BTV pan/8 2.0 RT-PCR Kit uses three specific primer/probe combinations:

- FAM™ fluorescence for RNA of all known BTV serotypes (pan BTV)
- Cy®5 fluorescence for RNA of European serotype 8 (BTV-8)
- JOE™ fluorescence for the endogenous Internal Control (EC; β-actin present within the sample)

A Positive Control contains BTV-8 RNA and allows the control of the denaturation step since the successful denaturation of the viral double-stranded RNA is a prerequisite for amplification.

#### 1.4 Kit contents

virotype BTV pan/8 2.0 RT-PCR Kit	(100)		
Cat. no.	VT280465		
Number of reactions	100		
Master Mix (tube with orange cap), includes enzymes, primers, and probes	1 x 800 μl		
Positive Control (tube with red cap)	1 x 150 μl		
Negative Control (tube with blue cap)	1 x 150 μl		
Handbook	1		

# 1.5 Storage

The components of the virotype BTV pan/8 2.0 RT-PCR Kit should be stored at -30°C to -15°C and are stable until the expiration date stated on the label. Avoid repeated thawing and freezing (> 3x), as this may reduce assay sensitivity. Freeze the components in aliquots if they will only be used intermittently.

# 1.6 Equipment and reagents to be supplied by user

When working with chemicals, always wear a suitable lab coat, disposable gloves and protective goggles. For more information, consult the appropriate safety data sheets (SDSs), available from the product supplier.

- Pipets
- Nuclease-free, aerosol-resistant pipet tips with filters
- Sterile 1.5 ml Eppendorf® tubes
- Nuclease-free (RNase/DNase-free) consumables. Special care should be taken to avoid nuclease contamination of all reagents and consumables used to set up PCR for sensitive identification of viral nucleic acids
- Cooling device or ice
- Benchtop centrifuge with rotor for 1.5 ml tubes
- Real-time cycler with appropriate fluorescent channels
- Appropriate software for chosen real-time cycler
- Appropriate strip tubes and caps or 96-well optical microplate with optical sealing film or cover for chosen real-time cycler

## 1.7 RNA extraction

The virotype BTV pan/8 2.0 RT-PCR Kit can be used for the detection of BTV RNA from ruminant whole blood (preferred with anticoagulants, e.g., EDTA-blood) and tissue samples (spleen, lymph nodes) from cattle, sheep and goats.

Due to the high sensitivity of the test, pools of up to 10 individual blood samples may be analyzed. However, the optimum pool size depends on the regional prevalence for BTV.

**Note**: For use in Germany the specifications described in the "Amtliche Methodensammlung" apply.

Prior to real-time PCR, viral RNA must be extracted from the starting material.

INDICAL offers a range of validated kits for the extraction of RNA from animal samples.

## Extraction based on magnetic beads:

- IndiMag<sup>®</sup> Pathogen Kit (SP947457)
- IndiMag Pathogen Kit w/o plastics (SP947257)
- IndiMag Pathogen IM48 Cartridge (SP947654P608, SP947654P224)
- IndiMag Pathogen KF96 Cartridge (SP947855P196)

### Extraction based on spin columns:

- IndiSpin® Pathogen Kit\* (SP54104, SP54106)
- IndiSpin QIAcube® HT Pathogen Kit (SP54161)

If real-time RT-PCR is not performed immediately after extraction, store the RNA at -20°C or at -80°C for longer storage.

For further information on automated and manual extraction of BTV RNA from different sample types, refer to the respective handbook or contact INDICAL Support at support@indical.com.

## 1.8 Important notes

# General precautions

The user should always pay attention to the following:

- Use nuclease-free pipet tips with filters.
- Store and extract positive materials (specimens, positive controls and amplicons) separately from all other reagents and add them to the reaction mix in a spatially separated facility.
- Thaw all components on ice before starting the assay.
- When thawed, mix the components by inverting and centrifuge briefly.
- Do not use components of the test kit past the expiration date.
- Keep samples and controls on ice or in a cooling block during the setup of reactions.

# Negative control

At least one negative control reaction should be included in each PCR run, containing all the components of the reaction except for the pathogen template. This enables assessment of contamination in the reaction.

#### Positive control

When performing PCR on unknown samples, it is recommended to include a positive control reaction in the PCR run, containing a sample that is known to include the targeted viral RNA. A positive control serves to prove the functionality of the pathogen assay, e.g., the correct setup of the reaction mix. Use 5  $\mu$ l of the Positive Control provided with the virotype BTV pan/8 2.0 RT-PCR Kit to test for successful amplification of the target.

# Extraction and amplification control

For increased process safety and convenience, one extraction and amplification control assay is included in the test kit.

An endogenous Internal Control (EC) detects the  $\beta$ -actin gene present within the sample. This allows extraction and amplification to be monitored.

# 2 Procedure

## 2.1 Important points before starting

- Please read "Important notes" before starting.
- Include at least one positive control (Positive Control) and one negative control (Negative Control) per PCR run.
- Before beginning the procedure, read through the protocol and ensure that you are familiar with the operation of the chosen real-time PCR cycler.
- RNA is unstable. Perform the protocol without interruption.

# 2.2 Things to do before starting

- Thaw all reagents on ice and protect from light.
- Before use, spin the reagents briefly.
- Maintain reagents on ice or in a cooling block during PCR setup.

# 2.3 Test procedure

- 1. Pipet 5 μl of RNA samples, Positive Control, and Negative Control into individual reaction tubes. Cover the reaction tubes (e.g., with PCR sealing foil).
  - Include positive and negative control reactions.
  - Positive Control: Use 5 µl of the positive control (Positive Control) instead of sample RNA.
  - Negative Control: Use 5 µl of the negative control (Negative Control) instead of sample RNA.
- 2. Denature the samples and Controls for 5 min at 98°C in a 96-well plate standard cycler with a heated lid.
- 3. Immediately cool down on ice water or liquid nitrogen for at least 20 s. Then store on ice or in a cooling device.
- 4. Before use, mix the Master Mix by inverting 5 times or until mixed thoroughly, then centrifuge briefly to collect the fluids.

5. Pipet 8  $\mu$ l of the Master Mix into each reaction tube. Thus, the final volume of a test is 13  $\mu$ l (Table 1).

Table 1. Preparation of reaction mix

Component	Volume
Master Mix	8 µl
Sample	5 μΙ
Total volume	13 μΙ

- 6. Close the reaction tubes with the corresponding caps.
- 7. Set the filters for the reporter dyes in the software of your thermal cycler according to Table 2.

Table 2. Filter settings for the reporter

Pathogen/ Internal Control	Reporter
Pan BTV	FAM
BTV-8	Cy5
Endogenous Internal Control	HEX/ JOE™¹
Passive reference	Texas Red/ ROX™²

<sup>&</sup>lt;sup>1</sup> Use the option appropriate for your thermal cycler.

8. Run the real-time PCR protocol according to Table 3.

Table 3. Real-time RT-PCR protocol for BTV pan/8 2.0.

_			
Step	Temperature	Time	Number of cycles
Reverse Transcription	50°C	10 min	1
Initial Activation	95°C	2 min	1
2-step cycling			
Denaturation	95°C	5 s	40
Annealing/Extension*	60°C	30 s	40

<sup>\*</sup> Fluorescence data collection, approximate run time 68 min (CFX96, Bio-Rad™)

<sup>&</sup>lt;sup>2</sup> Internal reference for use with Applied Biosystems® ABI PRISM® Sequence Detection Systems

# 3 Data interpretation

#### Interpretation of results

For the assay to be valid the Positive Control must give a signal in the FAM, the Cy5, and HEX/ JOE channels with a  $C_T{}^1 < 35$ . If no FAM and no Cy5 signals of the Positive Controls are measured the denaturation and cooling steps were insufficient and the testing should be repeated. The Negative Control must give no signal.

The following results are possible if working with unknown samples. The possible sample results are also summarized in Table 4.

# The sample is positive for pan BTV and BTV-8, and the assay is valid, if the following criteria are met:

- The sample yields a signal in the FAM, Cy5 and HEX/JOE channel.
- The Positive Control yields a signal in all channels.
- The Negative Control yields no signal in any of the channels.

Note that very high concentrations of BTV RNA or presence of inhibitors in the sample may lead to a reduced HEX/JOE signal or no HEX/JOE signal due to competition with the Internal Control.

# The sample is positive for pan BTV and negative for BTV-8, and the assay is valid, if the following criteria are met:

- The sample yields a signal in the FAM and HEX/JOE channel, but not in the Cy5 channel.
- The Positive Control yields a signal in all channels.
- The Negative Control yields no signal in any of the channels.

Note that very high concentrations of BTV RNA or presence of inhibitors in the sample may lead to a reduced HEX/JOE signal or no HEX/JOE signal due to competition with the Internal Control.

 $<sup>^1</sup>$  Threshold cycle (C<sub>1</sub>) — cycle at which the amplification plot crosses the threshold, i.e., there is the first clearly detectable increase in fluorescence

# The sample is negative for both, pan BTV and BTV-8, and the assay is valid, if the following criteria are met:

- The sample yields a signal only in the HEX/JOE channel.
- The Positive Control yields a signal in all channels.
- The Negative Control yields no signal in any of the channels.

A positive HEX/JOE signal rules out the possibility of PCR inhibition and/ or incorrect RNA extraction as the Internal Control is amplified.

## The sample results are inconclusive, and the assay is invalid, if the following occurs:

• sample yields no signal in any of the fluorescence channels.

If no signal is detected in the FAM (pan BTV), Cy5 (BTV-8) and the HEX/JOE (endogenous Internal Control, EC) channels, the result is inconclusive. The absence of a signal for the housekeeping gene indicates strong PCR inhibition and/or other malfunctions, e.g., during extraction.

To check for inhibition, we recommend 1:5 dilution of the sample RNA in nuclease-free water, to repeat the RNA extraction procedure, or repeat the whole test procedure starting with new sample material.

Check that there is a fluorescence signal in all the channels for the positive control reaction (Positive Control). Absence of a signal for the Positive Control indicates an error, which could be due to incorrect RNA denaturation or incorrect cycling conditions.

Table 4. Results interpretation table\*

Sample recult	FAM	Cy5	HEX
Sample result	(pan BTV)	(BTV-8)	(EC)
pan BTV positive	Χ		(X)
pan BTV and BTV-8 positive	X	X	(X)
BTV negative			Χ
Inconclusive			

<sup>\*</sup> Interpretation of sample results can be determined provided positive and negative control reactions are performed. The Positive Control must yield a signal in the FAM, Cy5 and HEX/JOE channels. The Negative Control must yield no signal in any channel. For a complete explanation of possible sample results please refer to "Data analysis and interpretation"

# 4 Characteristics of the test

# 4.1 Analytical sensitivity

The high analytical sensitivity of the virotype BTV pan/8 2.0 RT-PCR Kit was verified by a titration series of pan BTV and BTV-8 *in vitro* RNA [10<sup>6</sup> – 1 copies/well], performed in triplicates of relevant dilutions using the virotype BTV pan/8 2.0 RT-PCR protocol on the Bio-Rad CFX96, the Thermo Fisher Scientific QuantStudio® 5, the Agilent Technologies Aria Mx, and the Applied Biosystems™ 7500 Fast instrument.

# 4.1.1 Analytical sensitivity of pan BTV and BTV-8 systems using the Bio-Rad CFX96 instrument

### Results / Conclusion

The virotype BTV pan/8 2.0 RT-RT-PCR Kit can detect up to one pan BTV / BTV-8 genome copy per sample (Table 5, Table 6, Figure 1 - Figure 4). There is a high correlation between RNA copy number and amplification results. A correlation coefficient of 0.9990 with an efficiency of 99.6 % for the pan BTV *in vitro* RNA and a correlation coefficient of 0.9970 with an efficiency of 104.7 % for the BTV-8 *in vitro* RNA was calculated when using the virotype BTV pan/8 2.0 RT-PCR Kit on the Bio-Rad CFX96 instrument (Figure 2, Figure 4).

Table 5. Individual and mean  $C_T$  values of **pan BTV** (FAM) *in vitro* RNA titration series in triplicates. The test was performed on the Bio-Rad CFX96 instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

Туре	Copy number	C <sub>T</sub> (FAM)	C <sub>⊤</sub> mean	SD	Result
Standard	106	18.98			+
Standard	106	19.02	19.03	0.05	+
Standard	10 <sup>6</sup>	19.09			+
Standard	105	21.98			+
Standard	105	22.19	22.10	0.11	+
Standard	105	22.14			+
Standard	104	25.47			+
Standard	104	25.43	25.50	0.09	+
Standard	104	25.60			+
Standard	10³	28.76			+
Standard	10 <sup>3</sup>	28.75	28.76	0.02	+
Standard	10 <sup>3</sup>	28.78			+
Standard	100	32.20			+
Standard	100	32.04	32.13	0.08	+
Standard	100	32.14			+
Standard	10	35.91			+
Standard	10	35.91	35.68	0.40	+
Standard	10	35.21			+
Standard	5	35.50			+
Standard	5	35.52	35.53	0.03	+
Standard	5	35.56			+
Standard	1	-			-
Standard	1	38.49	38.49	-	+
Standard	1	-			-

 $SD = standard deviation, - = no C_T$ 

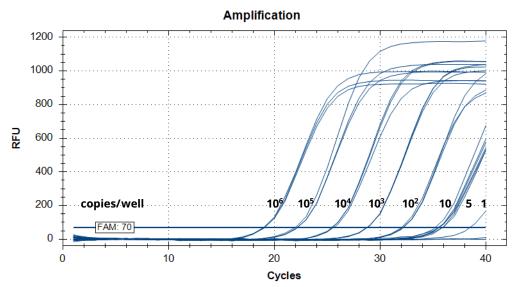


Figure 1. Individual values of a titration series of **pan BTV** (FAM) *in vitro* RNA in triplicates. The test was performed on the Bio-Rad CFX96 instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

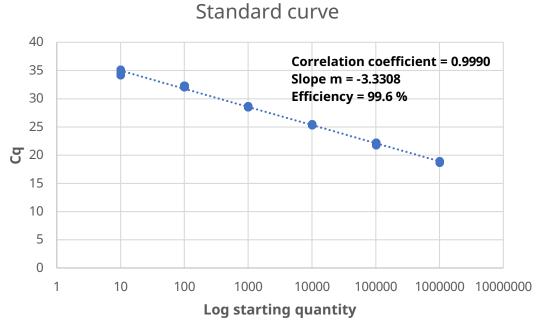


Figure 2. Standard curve of obtained  $C_T$  values for a titration series of **pan BTV** (FAM) *in vitro* RNA. The test was performed on the Bio-Rad CFX96 instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

Table 6. Individual and mean  $C_T$  values of **BTV-8** (Cy5) *in vitro* RNA titration series in triplicates. The test was performed on the Bio-Rad CFX96 instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

Туре	Copy number	С <sub>т</sub> (Су5)	C <sub>⊤</sub> mean	SD	Result
Standard	10 <sup>6</sup>	18.74			+
Standard	10 <sup>6</sup>	18.81	18.82	0.09	+
Standard	106	18.92			+
Standard	10 <sup>5</sup>	22.09			+
Standard	10 <sup>5</sup>	22.24	22.05	0.21	+
Standard	10 <sup>5</sup>	21.83			+
Standard	104	25.32			+
Standard	104	25.45	25.41	0.08	+
Standard	104	25.46			+
Standard	10³	28.56			+
Standard	10 <sup>3</sup>	28.71	28.60	0.10	+
Standard	10 <sup>3</sup>	28.51			+
Standard	100	32.35			+
Standard	100	32.14	32.19	0.14	+
Standard	100	32.08			+
Standard	10	34.23			+
Standard	10	34.47	34.61	0.46	+
Standard	10	35.12			+
Standard	5	35.72			+
Standard	5	35.64	35.60	0.15	+
Standard	5	35.43			+
Standard	1	-			-
Standard	1	-	37.69	-	-
Standard	1	37.69			+

 $<sup>\</sup>overline{SD = standard deviation, - = no C_T}$ 

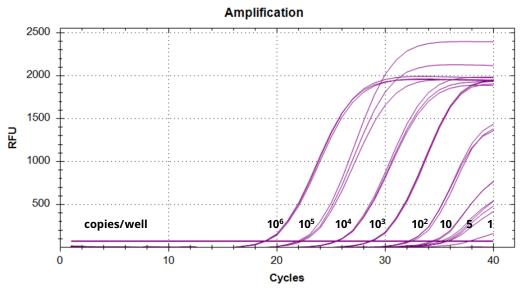


Figure 3. Individual values of a titration series of **BTV-8** (Cy5) *in vitro* RNA in triplicates. The test was performed on the Bio-Rad CFX96 instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

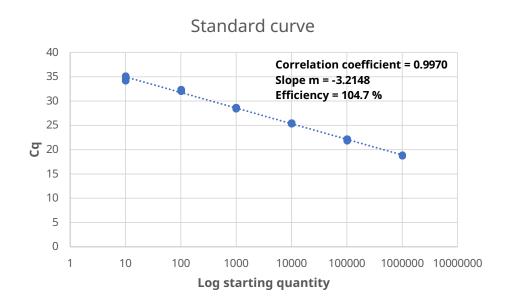


Figure 4. Standard curve of obtained  $C_T$  values for a titration series of **BTV-8** (Cy5) *in vitro* RNA. The test was performed on the Bio-Rad CFX96 instrument using the virotype virotype BTV pan/8 2.0 real-time RT-PCR protocol.

# 4.1.2 Analytical sensitivity of pan BTV and BTV-8 systems using the Thermo Fisher QuantStudio 5 instrument

# Results / Conclusion

The virotype BTV pan/8 2.0 RT-PCR Kit can detect up to one BTV genome copies per sample (Table 7, Table 8, Figure 5 - Figure 8). There is a high correlation between RNA copy number and amplification results. A correlation coefficient of 0.9998 with an efficiency of 99.6 % for the pan BTV *in vitro* RNA and a correlation coefficient of 0.9946 with an efficiency of 102.3 % for the BTV-8 *in vitro* RNA was calculated when using the virotype BTV pan/8 2.0 RT-PCR Kit on the Thermo Fisher QuantStudio 5 instrument (Figure 6, Figure 8).

Table 7. Individual and mean  $C_T$  values of **pan BTV** (FAM) *in vitro* RNA titration series in triplicates. The test was performed on the Thermo Fisher QuantStudio 5 instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

Туре	Copy number	C <sub>T</sub> (FAM)	C <sub>⊤</sub> mean	SD	Result
Standard	106	18.34			+
Standard	106	18.38	18.35	0.03	+
Standard	10 <sup>6</sup>	18.32			+
Standard	105	21.65			+
Standard	105	21.61	21.64	0.02	+
Standard	105	21.65			+
Standard	104	24.79			+
Standard	104	25.07	24.99	0.18	+
Standard	104	25.12			+
Standard	10³	28.38			+
Standard	10 <sup>3</sup>	28.44	28.42	0.04	+
Standard	10³	28.45			+
Standard	100	31.60			+
Standard	100	31.61	31.61	0.01	+
Standard	100	31.62			+
Standard	10	34.95			+
Standard	10	35.07	35.01	0.06	+
Standard	10	35.00			+
Standard	5	35.35			+
Standard	5	35.19	35.65	0.66	+
Standard	5	36.40			+
Standard	1	37.90			+
Standard	1	36.40	37.15	-	+
Standard	1	-			-

 $SD = standard deviation, - = no C_T$ 

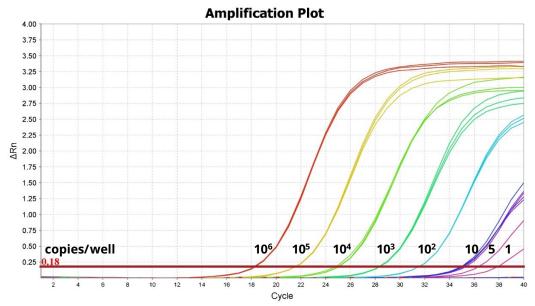


Figure 5. Individual values of a titration series of **pan BTV** (FAM) *in vitro* RNA in triplicates. The test was performed on the Thermo Fisher QuantStudio 5 instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

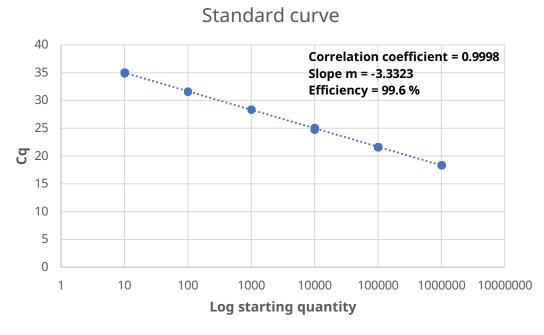


Figure 6. Standard curve of obtained  $C_T$  values for a titration series of **pan BTV** (FAM) *in vitro* RNA. The test was performed on the Thermo Fisher QuantStudio 5 instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

Table 8. Individual and mean  $C_T$  values of **BTV-8** (Cy5) *in vitro* RNA titration series in triplicates. The test was performed on the Thermo Fisher QuantStudio 5 instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

Туре	Copy number	С <sub>т</sub> (Су5)	C <sub>⊤</sub> mean	SD	Result
Standard	10 <sup>6</sup>	17.90			+
Standard	10 <sup>6</sup>	17.72	17.77	0.12	+
Standard	10 <sup>6</sup>	17.68			+
Standard	10 <sup>5</sup>	20.94			+
Standard	10 <sup>5</sup>	20.95	20.93	0.04	+
Standard	10 <sup>5</sup>	20.88			+
Standard	104	24.22			+
Standard	104	24.28	24.25	0.03	+
Standard	104	24.25			+
Standard	10³	27.43			+
Standard	10³	27.51	27.46	0.04	+
Standard	10³	27.46			+
Standard	100	30.83			+
Standard	100	30.63	30.67	0.14	+
Standard	100	30.56			+
Standard	10	35.52			+
Standard	10	33.63	34.16	1.19	+
Standard	10	33.32			+
Standard	5	34.55			+
Standard	5	34.35	34.56	0.21	+
Standard	5	34.77			+
Standard	1	36.44			+
Standard	1	-	36.44	-	-
Standard	1	-			-

 $SD = standard deviation, - = no C_T$ 

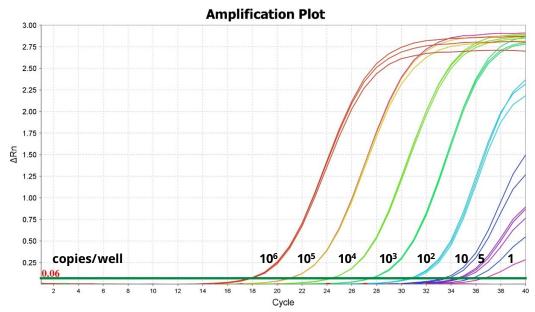


Figure 7. Individual values of a titration series of **BTV-8** (Cy5) *in vitro* RNA in triplicates. The test was performed on the Thermo Fisher QuantStudio 5 instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

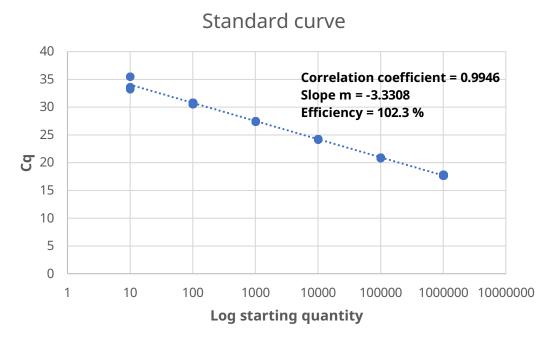


Figure 8. Standard curve of obtained  $C_T$  values for a titration series of **BTV-8** (Cy5) *in vitro* RNA. The test was performed on the Thermo Fisher QuantStudio 5 instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

# 4.1.3 Analytical sensitivity of pan BTV and BTV-8 systems using the Agilent Technologies Aria Mx instrument

# Results / Conclusion

The virotype BTV pan/8 2.0 RT-PCR Kit can detect up to one pan BTV genome copy per sample (Table 9, Figure 9, Figure 10) and up to five BTV-8 genome copies per sample (Table 10, Figure 11, Figure 12). There is a high correlation between RNA copy number and amplification results. A correlation coefficient of 0.9937 with an efficiency of 107.3 % for the pan BTV *in vitro* RNA and a correlation coefficient of 0.9994 with an efficiency of 107.3 % for the BTV-8 *in vitro* RNA was calculated when using the virotype BTV pan/8 2.0 RT-PCR Kit on the Agilent Technologies Aria Mx instrument (Figure 10, Figure 12).

Table 9. Individual and mean  $C_T$  values of **pan BTV** (FAM) *in vitro* RNA titration series in triplicates. The test was performed on the Agilent Technologies Aria Mx instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

Туре	Copy number	C <sub>T</sub> (FAM)	C <sub>⊤</sub> mean	SD	Result
Standard	106	17.84			+
Standard	106	17.97	17.89	0.07	+
Standard	10 <sup>6</sup>	17.85			+
Standard	10 <sup>5</sup>	20.94			+
Standard	10 <sup>5</sup>	21.02	21.02	0.08	+
Standard	10 <sup>5</sup>	21.09			+
Standard	104	24.11			+
Standard	104	24.21	24.14	0.06	+
Standard	104	24.11			+
Standard	10³	27.66			+
Standard	10³	27.34	27.48	0.16	+
Standard	10 <sup>3</sup>	27.44			+
Standard	100	30.23			+
Standard	100	30.45	30.52	0.34	+
Standard	100	30.89			+
Standard	10	32.51			+
Standard	10	34.03	33.63	0.98	+
Standard	10	34.35			+
Standard	5	35.22			+
Standard	5	35.61	35.14	0.51	+
Standard	5	34.59			+
Standard	1	37.62			+
Standard	1	-	37.05	0.81	-
Standard	1	36.48			+

 $SD = standard deviation, - = no C_T$ 

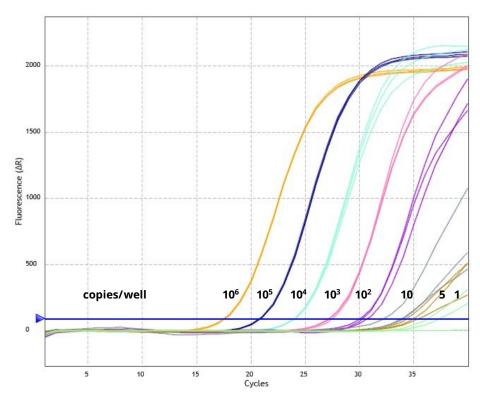


Figure 9. Individual values of a titration series of **pan BTV** (FAM) *in vitro* RNA in triplicates. The test was performed on the Agilent Technologies Aria Mx instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

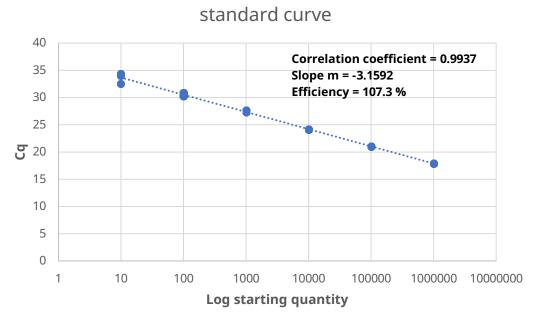


Figure 10. Standard curve of obtained  $C_T$  values for a titration series of **pan BTV** (FAM) *in vitro* RNA. The test was performed on the Agilent Technologies Aria Mx instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

Table 10. Individual and mean  $C_T$  values of **BTV-8** (Cy5) *in vitro* RNA titration series in triplicates. The test was performed on the Agilent Technologies Aria Mx instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

Туре	Copy number	С <sub>т</sub> (Су5)	C <sub>⊤</sub> mean	SD	Result
Standard	10 <sup>6</sup>	17.60			+
Standard	10 <sup>6</sup>	17.92	17.70	0.19	+
Standard	10 <sup>6</sup>	17.59			+
Standard	10⁵	20.78			+
Standard	10 <sup>5</sup>	20.89	20.83	0.06	+
Standard	10 <sup>5</sup>	20.82			+
Standard	104	23.99			+
Standard	104	23.99	23.96	0.06	+
Standard	104	23.89			+
Standard	10³	26.92			+
Standard	10³	27.18	27.04	0.13	+
Standard	10³	27.01			+
Standard	100	30.09			+
Standard	100	30.27	30.20	0.10	+
Standard	100	30.25			+
Standard	10	33.41			+
Standard	10	33.57	33.57	0.17	+
Standard	10	33.74			+
Standard	5	33.71			+
Standard	5	34.67	34.19	0.48	+
Standard	5	34.20			+
Standard	1	-			-
Standard	1	-	-	-	-
Standard	1	-			-

 $SD = standard deviation, - = no C_T$ 

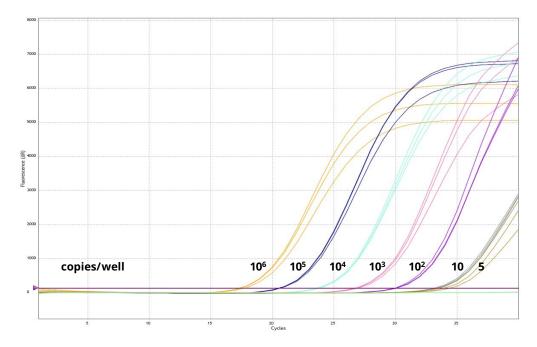


Figure 11. Individual values of a titration series of **BTV-8** (Cy5) *in vitro* RNA in triplicates. The test was performed on the Agilent Technologies Aria Mx instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

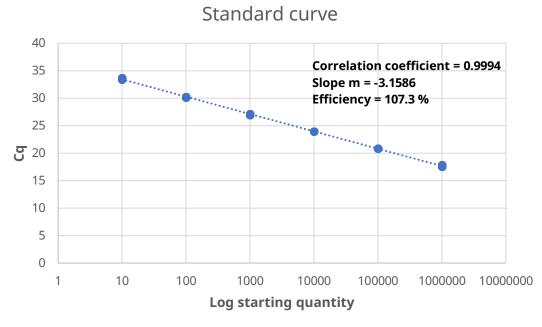


Figure 12. Standard curve of obtained  $C_T$  values for a titration series of **BTV-8** (Cy5) *in vitro* RNA. The test was performed on the Agilent Technologies Aria Mx instrument using the virotype virotype BTV pan/8 2.0 real-time RT-PCR protocol.

# 4.1.4 Analytical sensitivity of pan BTV and BTV-8 systems using the Applied Biosystems™ ABI 7500 Fast instrument

# Results / Conclusion

The virotype BTV pan/8 2.0 RT-PCR Kit can detect up to one BTV genome copy per sample (Table 11, Table 12, Figure 13 -Figure 16). There is a high correlation between RNA copy number and amplification results. A correlation coefficient of 0.9992 with an efficiency of 103.2 % for the pan BTV *in vitro* RNA and a correlation coefficient of 0.9983 with an efficiency of 103.7 % for the BTV-8 *in vitro* RNA was calculated when using the virotype BTV pan/8 2.0 RT-PCR Kit on the ABI 7500 Fast instrument (Figure 14, Figure 16).

Table 11. Individual and mean  $C_T$  values of **pan BTV** (FAM) *in vitro* RNA titration series in triplicates. The test was performed on the ABI 7500 Fast instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

Туре	Copy number	C <sub>T</sub> (FAM)	C <sub>⊤</sub> mean	SD	Result
Standard	106	18.25			+
Standard	106	18.31	18.30	0.04	+
Standard	10 <sup>6</sup>	18.33			+
Standard	105	21.63			+
Standard	105	21.54	21.55	0.07	+
Standard	105	21.48			+
Standard	104	24.78			+
Standard	104	24.88	24.81	0.06	+
Standard	104	24.78			+
Standard	10 <sup>3</sup>	28.00			+
Standard	10³	28.15	28.04	0.10	+
Standard	10³	27.96			+
Standard	100	31.30			+
Standard	100	31.00	31.24	0.21	+
Standard	100	31.40			+
Standard	10	34.93			+
Standard	10	34.19	34.58	0.37	+
Standard	10	34.62			+
Standard	5	35.03			+
Standard	5	34.73	35.13	0.46	+
Standard	5	35.63			+
Standard	1	37.90			+
Standard	1	37.87	37.58	0.53	+
Standard	1	36.96			+

SD = standard deviation

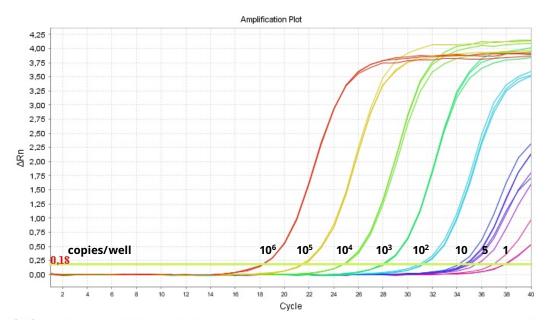


Figure 13. Individual values of a titration series of **pan BTV** (FAM) *in vitro* RNA in triplicates. The test was performed on the ABI 7500 Fast instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

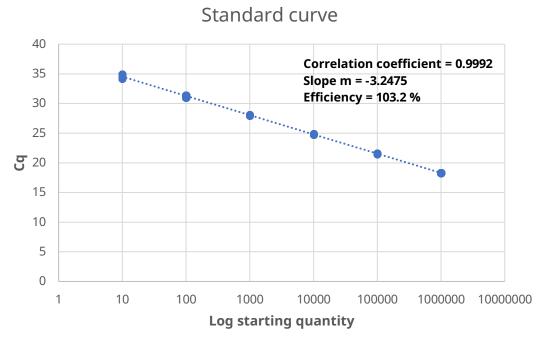


Figure 14. Standard curve of obtained  $C_T$  values for a titration series of **pan BTV** (FAM) *in vitro* RNA. The test was performed on the ABI 7500 Fast instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

Table 12. Individual and mean  $C_T$  values of **BTV-8** (Cy5) *in vitro* RNA titration series in triplicates. The test was performed on the ABI 7500 Fast instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

Туре	Copy number	С <sub>т</sub> (Су5)	C <sub>⊤</sub> mean	SD	Result
Standard	10 <sup>6</sup>	18.43			+
Standard	10 <sup>6</sup>	18.57	18.54	0.10	+
Standard	10 <sup>6</sup>	18.63			+
Standard	10 <sup>5</sup>	21.88			+
Standard	10 <sup>5</sup>	21.97	21.89	0.07	+
Standard	10 <sup>5</sup>	21.82			+
Standard	104	25.29			+
Standard	104	25.16	25.19	0.09	+
Standard	104	25.11			+
Standard	10³	28.57			+
Standard	10³	28.48	28.48	0.09	+
Standard	10 <sup>3</sup>	28.39			+
Standard	100	31.79			+
Standard	100	31.80	31.79	0.01	+
Standard	100	31.77			+
Standard	10	34.85			+
Standard	10	34.02	34.60	0.50	+
Standard	10	34.93			+
Standard	5	35.66			+
Standard	5	35.88	35.50	0.48	+
Standard	5	34.97			+
Standard	1	-			-
Standard	1	-	38.59	-	-
Standard	1	38.59			+

 $SD = standard deviation, - = no C_T$ 

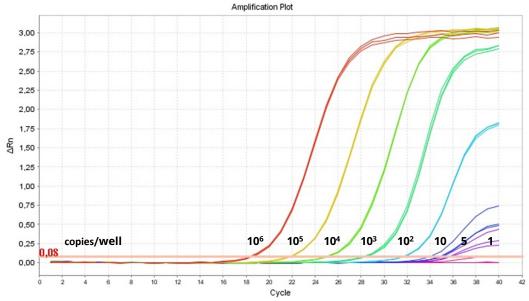


Figure 15. Individual values of a titration series of **BTV-8** (Cy5) *in vitro* RNA in triplicates. The test was performed on the ABI 7500 Fast instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

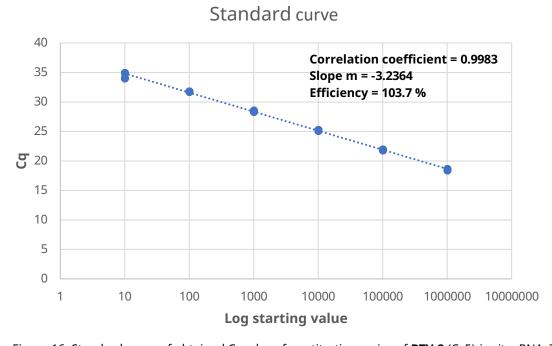


Figure 16. Standard curve of obtained  $C_T$  values for a titration series of **BTV-8** (Cy5) *in vitro* RNA. The test was performed on the ABI 7500 Fast instrument using the virotype virotype BTV pan/8 2.0 real-time RT-PCR protocol.

# 4.1.5 Analytical sensitivity – Limit of detection

The limit of detection (LOD) for the target sequence of the pan BTV and BTV-8 systems was determined by testing individual titration series of *in vitro* RNA of these sequences in octuplicates. The limit of detection with 95 % confidence interval (LOD<sub>95 %</sub>: mean number of copies yielding a probability of detection of 0.95) was determined using the web tool https://quodata.de/content/validation-qualitative-pcr-methods-single-laboratory.

## Results / Conclusion

Results are summarized in Table 13 and Table 14, as well as Figure 17 - Figure 20. Using the virotype pan/8 2.0 RT-PCR Kit, a high correlation between RNA copy number and the amount of amplified product was demonstrated for the BTV targeted sequences. The LOD<sub>95%</sub> for the pan BTV system is 2.9 copies with a 95% confidence interval of [1.307, 6.498] (Figure 17). The LOD<sub>95%</sub> for BTV-8 system is 4.6 copies with a 95% confidence interval of [2.272, 9.317] (Figure 19).

Table 13. Limit of detection for **pan BTV** *in vitro* RNA tested in octuplicates on the Agilent Technologies Aria Mx instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

Copies/test	Total number of replicates	Number of replicates positive	Number of replicates negative
10,000	8	8	0
1,000	8	8	0
100	8	8	0
10	8	8	0
5	8	8	0
1	8	5	3

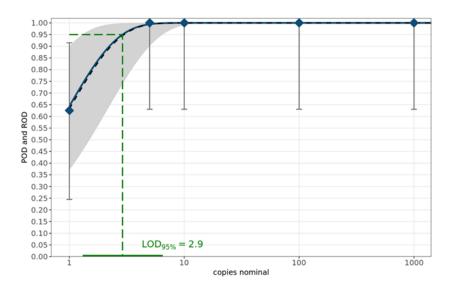


Figure 17. POD (probability of detection) curve and LOD<sub>95 %</sub> for **pan BTV**. The blue diamonds characterize the laboratory-specific rates of detection. The blue curve denotes the mean POD curve along with the corresponding 95 % confidence range highlighted as the grey band. The POD curve under ideal conditions is displayed as the black dashed curve.

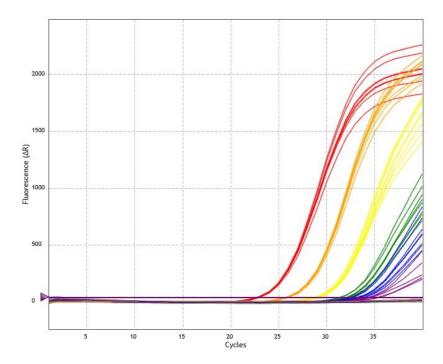


Figure 18. Analytical sensitivity – Limit of detection. Individual values of a titration series of **pan BTV** (FAM) *in vitro* RNA in octuplicates. The test was performed on the Agilent Technologies Aria Mx instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

Table 14. Limit of detection for **BTV-8** *in vitro* RNA tested in octuplicates on the Agilent Technologies Aria Mx instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

Copies/test	Total number of replicates	Number of replicates positive	Number of replicates negative
10,000	8	8	0
1,000	8	8	0
100	8	8	0
10	8	8	0
5	8	8	0
1	8	3	5

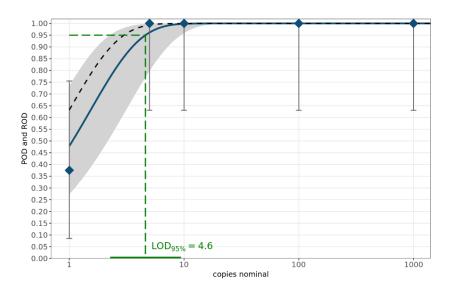


Figure 19. POD (probability of detection) curve and LOD $_{95\,\%}$  for **BTV-8**. The blue diamonds characterize the laboratory-specific rates of detection. The blue curve denotes the mean POD curve along with the corresponding 95 % confidence range highlighted as the grey band. The POD curve under ideal conditions is displayed as the black dashed curve.

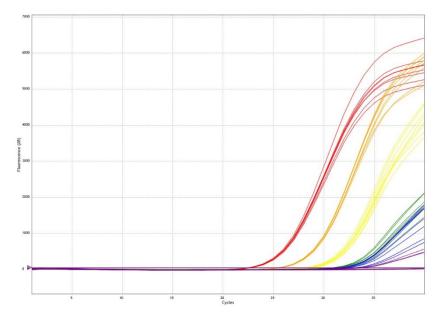


Figure 20. Analytical sensitivity – Limit of detection. Individual values of a titration series of **BTV-8** (Cy5) *in vitro* RNA in octuplicates. The test was performed on the Agilent Technologies Aria Mx instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

# 4.1.6 Analytical sensitivity of pooled samples

Pools were generated by diluting selected BTV-8 positive samples in BTV negative sheep or cattle blood. Samples were extracted with the IndiMag Pathogen kit (INDICAL) on IndiMag 48s following manufacturer's instructions. The resulting pools were tested with the virotype BTV pan/8 2.0 RT-PCR Kit on the Agilent Technologies Aria Mx instrument using the virotype BTV pan/8 2.0 RT-PCR protocol.

#### Results / Conclusion

The  $C_T$  values of the pan BTV (FAM), the BTV-8 (Cy5) and the endogenous Internal Control (JOE) are shown in Table 15. Pooled samples up to at least 20 could be detected with the virotype BTV pan/8 2.0 RT-PCR Kit.

Table 15. Analysis of the **pan BTV** (FAM), **BTV-8** (Cy5) and the endogenous **Internal Control** (JOE) signals for simulated pool samples with the virotype BTV pan/8 2.0 RT-PCR Kit tested on the Agilent Technologies Aria Mx instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

Pool size	Sample	Species	virotype I	BTV pan/8 2.0 l	RT-PCR Kit
			pan BTV	BTV-8	Internal Control
			$\mathbf{C}_{T}$	$C_{\tau}$	$C_{T}$
0	WIS 167	Sheep	17.67	16.34	20.99
10			21.48	19.98	18.98
20			22.10	20.96	19.26
0	WIS 168	Sheep	16.31	15.06	18.86
10			20.62	19.45	19.81
20			21.97	20.55	19.75
0	СВ	Cattle	21.87	20.54	25.06
10			24.97	23.15	26.87
100			29.82	27.14	28.59

# 4.2 Specificity

#### 4.2.1 Inclusivity

To confirm that the virotype BTV pan/8 2.0 RT-PCR Kit can detect different serotypes of BTV, a panel n = 32 reference RNA samples comprising of BTV serotypes 1 through 28, 30, 33, 35, and 36 was tested with the kit on the Agilent Technologies Aria Mx instrument. Samples were kindly provided by the Friedrich-Loeffler-Institut (FLI), Greifswald, Germany.

The reference samples were extracted using the IndiMag® Pathogen Kit (INDICAL BIOSCIENCE) following manufacturer's instructions and the RNA samples tested with the virotype BTV pan/8 2.0 RT-PCR Kit.

#### Results / Conclusion

Out of the n = 32 reference samples with different BTV serotypes tested, all were detected using the virotype BTV pan/8 RT-PCR Kit (Table 16).

Table 16. Analysis of RNA samples from the FLI reference panel tested with the virotype BTV pan/8 2.0 RT-PCR on the Agilent Technologies Aria Mx instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

BTV serotypes	virotype BTV pan/	8 2.0 RT-PCR Kit
	pan BTV	BTV-8
	$\mathbf{C}_{T}$	$\mathbf{C}_{T}$
1	24.50	-
2	23.21	-
3	21.80	-
4	23.49	-
5	22.51	-
6	23.15	-
7	21.74	-
8	20.82	19.71
9	21.57	-
10	22.36	-
11	22.02	-
12	21.07	-

13	21.31	-
14	21.82	-
15	23.16	-
16	23.11	-
17	22.60	-
18	22.02	-
19	21.81	-
20	21.41	-
21	24.17	-
22	22.79	-
23	23.34	-
24	22.98	-
25	14.48	-
26	17.87	-
27	21.90	-
28	14.89	-
30	14.76	-
33	13.18	-
35	15.93	-
36	16.15	-

<sup>- =</sup> no C<sub>T</sub>

# 4.2.2 Exclusivity (Discrimination of pathogens for differential diagnosis)

The specificity / exclusivity was tested with nucleic acids from samples positive for other ruminant-related pathogens: *Schmallenberg Virus* (SBV, n = 21), *Bovine Viral Diarrhea Virus* (BVDV, n = 47), *Epizootic Hemorrhagic Disease Virus* (EHDV, n = 15), *Bovine Herpesvirus* 1 (BHV1, n = 9), and *Mycobacterium avium* subsp. *paratuberculosis* (MAP, n = 9). The samples were kindly provided by German Veterinary Laboratories, the FLI (German National Reference Laboratory for BTV) and other reference labs. Specificity testing was performed on the Agilent Technologies Aria Mx or the Thermo Fisher Scientific QuantStudio 5 instrument.

#### Results / Conclusion

No cross-reactivity to other relevant bovine viral and bacterial pathogens was detected using the virotype BTV pan/8 2.0 RT-PCR Kit (Table 17).

Table 17. Cross-reactivity of the virotype BTV pan/8 2.0 RT-PCR Kit to other ruminant-related pathogens on the Agilent Technologies Aria Mx or the Thermo Fisher Scientific QuantStudio 5 instrument using the virotype BTV pan/8 2.0 real-time RT-PCR protocol.

							pe BTV p RT-PCR I	an/8 2.0 (it
	Sample / strain (EHDV)	Species	Dil.	Material	Refe- rence	pan BTV	BTV-8	Int. Control
					$\mathbf{C}_{T}$	C <sub>T</sub>	$\mathbf{C}_{T}$	$\mathbf{C}_{T}$
Schmallenberg	SBV #1	Cattle	1:10	Blood	28.01	-	-	23.87
Virus (SBV)	SBV #2	Cattle	1:10	Blood	25.64	-	-	23.70
	SBV #3	Cattle	1:10	Blood	26.91	-	-	23.38
	SBV #4	Cattle	1:10	Blood	29.59	-	-	23.43
	SBV #5	Cattle	1:10	Blood	31.16	-	-	23.16
	SBV #6	Cattle	1:10	Blood	30.73	-	-	22.83
	SBV #7	Cattle	undil.	Brain	22.20	-	-	21.56
	SBV #8	Cattle	undil.	Brain	30.56	-	-	21.88
	SBV #9	Cattle	undil.	Brain	29.70	-	-	19.78
	SBV #10	Cattle	undil.	Brain	31.92	-	-	22.38
	SBV #10	Cattle	undil.	Brain	27.45	-	-	18.15
	SBV #12	Cattle	undil.	Brain	27.83	-	-	22.19

	SBV #13	Cattle	undil.	Spleen	31.29	-	-	19.24
	SBV #14	Cattle	undil.	Spleen	35.05	-	-	19.99
	SBV #15	Cattle	undil.	Spleen	37.54	-	-	22.23
	SBV #16	Cattle	undil.	Spleen	28.19	-	-	17.10
	SBV #17	Cattle	undil.	Liver	36.55	-	-	19.03
	SBV #18	Cattle	undil.	Liver	37.29	-	-	21.91
	SBV #19	Cattle	undil.	Liver	35.72	-	-	20.51
	SBV #20	Cattle	1:10	Tissue	23.17	-	-	20.61
	SBV #21	Cattle	1:10	Tissue	36.13	-	-	19.26
Bovine Viral	BVDV #1	Cattle	undil.	Blood	19.80	-	-	21.12
Diarrhea Virus (BVDV)	BVDV #2	Cattle	undil.	Blood	21.28	-	-	21.86
	BVDV #3	Cattle	undil.	Blood	21.47	-	-	22.09
	BVDV #4	Cattle	undil.	Blood	37.62	-	-	22.93
	BVDV #5	Cattle	undil.	Blood	34.77	-	-	22.67
	BVDV #6	Cattle	undil.	Blood	20.69	-	-	25.79
	BVDV #7	Cattle	undil.	Blood	20.52	-	-	23.97
	BVDV #8	Cattle	undil.	Blood	20.74	-	-	22.05
	BVDV #9	Cattle	undil.	Blood	22.12	-	-	23.76
	BVDV #10	Cattle	undil.	Blood	20.91	-	-	21.71
	BVDV #10	Cattle	undil.	Blood	20.70	-	-	22.05
	BVDV #12	Cattle	undil.	Blood	21.51	-	-	23.55
	BVDV #13	Cattle	undil.	Blood	18.58	-	-	23.64
	BVDV #14	Cattle	undil.	Blood	20.35	-	-	22.54
	BVDV #15	Cattle	undil.	Blood	21.43	-	-	22.49
	BVDV #16	Cattle	undil.	Blood	19.12	-	-	22.36
	BVDV #17	Cattle	undil.	Blood	21.82	-	-	22.94
	BVDV #18	Cattle	undil.	Blood	20.05	-	-	21.84
	BVDV #19	Cattle	undil.	Blood	20.55	-	-	21.41
	BVDV #20	Cattle	undil.	Blood	20.35	-	-	22.36
	BVDV #21	Cattle	undil.	Blood	20.55	-	-	23.50
	BVDV #22	Cattle	undil.	Blood	20.47	-	-	23.10

	BVDV #23	Cattle	undil.	Blood	23.27	-	-	23.07
	BVDV #24	Cattle	undil.	Blood	20.17	-	-	23.24
	BVDV #25	Cattle	undil.	Blood	19.98	-	-	26.55
	BVDV #26	Cattle	undil.	Blood	21.05	-	-	24.17
	BVDV #27	Cattle	undil.	Blood	20.88	-	-	24.35
	BVDV #28	Cattle	undil.	Blood	22.60	-	-	25.23
	BVDV #29	Cattle	undil.	Blood	26.99	-	-	22.61
	BVDV #30	Cattle	undil.	Blood	21.95	-	-	22.36
	BVDV #31	Cattle	undil.	Blood	20.08	-	-	23.55
	BVDV #32	Cattle	undil.	Blood	37.15	-	-	23.69
	BVDV #33	Cattle	undil.	Blood	18.89	-	-	20.35
	BVDV #34	Cattle	undil.	Blood	30.01	-	-	20.81
	BVDV #35	Cattle	undil.	Blood	19.93	-	-	24.28
	BVDV #36	Cattle	undil.	Blood	18.49	-	-	22.09
	BVDV #37	Cattle	undil.	Blood	19.84	-	-	21.80
	BVDV #38	Cattle	undil.	Blood	19.59	-	-	21.96
	BVDV #39	Cattle	undil.	Blood	18.49	-	-	22.64
	BVDV #40	Cattle	undil.	Blood	20.49	-	-	23.53
	BVDV #41	Cattle	undil.	Liver	24.44	-	-	19.40
	BVDV #42	Cattle	undil.	Liver	27.71	-	-	21.36
	BVDV #43	Cattle	undil.	Liver	27.42	-	-	19.69
	BVDV #44	Cattle	1:50	Tissue	27.12	-	-	24.07
	BVDV #45	Cattle	1:100	Tissue	28.18	-	-	23.67
	BVDV #46	Cattle	1:50	Tissue	26.33	-	-	23.44
	BVDV #47	Cattle	1:100	Tissue	27.47	-	-	23.76
Epizootic	FLI-2422-Alb		undil.	Culture	21.76	-	-	30.08
Hemorrhagic Disease Virus	FLI-2423-3		undil.	Culture	22.98	-	-	28.93
(EHDV)	EHDV-6/CSIR		undil.	Culture	22.89	-	-	-
	EHDV-5/Austra	lia	undil.	Culture	19.10	-	-	-
	EHDV-4/IBAR		undil.	Culture	21.22	-	-	-
	EHDV-3/IBAR		undil.	Culture	21.26	-	-	-

EHDV-2C/Alberta   Undil. Culture   21.39   -   -   -									
EHDV-2A/Ibar		EHDV-2C/Alber	ta	undil.	Culture	21.39	-	-	-
EHDV-1 /NewJer		EHDV-2B/CSIR		undil.	Culture	15.27	-	-	-
EHDV-1/ISR		EHDV-2A/Ibar		undil.	Culture	19.24	-	-	-
EHDV-6/ISR		EHDV-1 /NewJe	er	undil.	Culture	23.55	-	-	-
EHDV-7/ISR		EHDV-1/ISR		undil.	Culture	15.08	-	-	-
EHDV-6   Undil. Culture   21.02   -   -   -		EHDV-6/ISR		undil.	Culture	17.08	-	-	-
Bovine Herpes   BHV #1   Cattle   undil.   Serum   pos   -   -   25.70		EHDV-7/ISR		undil.	Culture	19.68	-	-	28.00
Bovine Herpes   BHV #1   Cattle   undil.   Serum   pos   -   -   25.70		EHDV-6		undil.	Culture	21.02	-	-	-
BHV #2   Cattle   undil. Serum   pos   -   -   23.79		EHDV-8/CSIR		undil.	Culture	16.99	-	-	-
BHV #2 Cattle undil. Serum pos - 23.79 BHV #3 Cattle undil. Serum pos - 26.22 BHV #4 Cattle undil. Serum pos - 27.62 BHV #5 Cattle undil. Serum pos - 26.92 BHV #6 Cattle undil. Serum pos - 26.17 BHV #7 Cattle undil. Serum pos - 25.29 BHV #8 Cattle undil. Serum pos - 25.29 BHV #9 Cattle undil. Serum pos - 25.33 BHV #9 Cattle undil. Serum pos - 25.17  Mycobacterium avium subsp. Paratuber- culosis (MAP) MAP #2 Goat undil. Lymph node 31.81 - 16.47 Culosis (MAP) MAP #3 Goat undil. Lymph node 34.67 - 16.53 MAP #4 Goat undil. Lymph node 32.28 - 16.68 MAP #5 Goat undil. Lymph node 32.28 - 16.68 MAP #6 Goat undil. Liver 28.03 - 19.24 MAP #7 Goat undil. Liver 32.03 - 17.53 MAP #8 Goat undil. Spleen 33.19 - 17.09	-	BHV #1	Cattle	undil.	Serum	pos	-	-	25.70
BHV #4   Cattle   undil. Serum   pos   -   27.62     BHV #5   Cattle   undil. Serum   pos   -   26.92     BHV #6   Cattle   undil. Serum   pos   -   26.17     BHV #7   Cattle   undil. Serum   pos   -   25.29     BHV #8   Cattle   undil. Serum   pos   -   25.33     BHV #9   Cattle   undil. Serum   pos   -   25.17     Mycobacterium avium subsp. Paratuber-culosis (MAP)   MAP #2   Goat   undil. Lymph node   31.81   -   16.47     Culosis (MAP)   MAP #3   Goat   undil. Lymph node   34.67   -   18.06     MAP #4   Goat   undil. Lymph node   34.67   -   16.53     MAP #5   Goat   undil. Lymph node   32.28   -   16.68     MAP #6   Goat   undil. Liver   28.03   -   17.53     MAP #7   Goat   undil. Liver   32.03   -   17.53     MAP #8   Goat   undil. Spleen   33.19   -   17.09	Virus (BHV)	BHV #2	Cattle	undil.	Serum	pos	-	-	23.79
BHV #5   Cattle   undil.   Serum   pos   -   26.92     BHV #6   Cattle   undil.   Serum   pos   -   26.17     BHV #7   Cattle   undil.   Serum   pos   -   25.29     BHV #8   Cattle   undil.   Serum   pos   -   25.33     BHV #9   Cattle   undil.   Serum   pos   -   25.17     Mycobacterium avium subsp. Paratuber-culosis (MAP)   MAP #2   Goat   undil.   Lymph node   33.35   -   16.61     MAP #3   Goat   undil.   Lymph node   31.81   -   16.47     MAP #4   Goat   undil.   Lymph node   34.67   -   16.53     MAP #5   Goat   undil.   Lymph node   32.28   -   16.68     MAP #6   Goat   undil.   Liver   28.03   -   17.53     MAP #7   Goat   undil.   Liver   32.03   -   17.53     MAP #8   Goat   undil.   Spleen   33.19   -   17.09		BHV #3	Cattle	undil.	Serum	pos	-	-	26.22
BHV #6 Cattle undil. Serum pos - 26.17 BHV #7 Cattle undil. Serum pos - 25.29 BHV #8 Cattle undil. Serum pos - 25.33 BHV #9 Cattle undil. Serum pos - 25.17  Mycobacterium avium subsp. Paratuber-culosis (MAP) MAP #3 Goat undil. Lymph node 31.81 - 16.47 MAP #4 Goat undil. Colon 32.21 - 18.06 MAP #5 Goat undil. Lymph node 32.28 - 16.68 MAP #6 Goat undil. Liver 28.03 - 19.24 MAP #7 Goat undil. Liver 32.03 - 17.53 MAP #8 Goat undil. Spleen 33.19 - 17.09		BHV #4	Cattle	undil.	Serum	pos	-	-	27.62
BHV #7 Cattle undil. Serum pos - 25.29  BHV #8 Cattle undil. Serum pos - 25.33  BHV #9 Cattle undil. Serum pos - 25.17  Mycobacterium avium subsp. Paratuber- culosis (MAP)  MAP #2 Goat undil. Lymph node 31.81 - 16.47  MAP #3 Goat undil. Colon 32.21 - 18.06  MAP #4 Goat undil. Lymph node 34.67 - 16.53  MAP #5 Goat undil. Lymph node 32.28 - 16.68  MAP #6 Goat undil. Liver 28.03 - 19.24  MAP #7 Goat undil. Liver 32.03 - 17.53  MAP #8 Goat undil. Spleen 33.19 - 17.09		BHV #5	Cattle	undil.	Serum	pos	-	-	26.92
BHV #8   Cattle   undil.   Serum   pos   -   25.33		BHV #6	Cattle	undil.	Serum	pos	-	-	26.17
Mycobacterium avium subsp. Paratuber-culosis (MAP)         MAP #1 Goat undil. Lymph node 31.81 - 16.47           MAP #4 Goat undil. Lymph node 32.21 - 18.06           MAP #5 Goat undil. Lymph node 32.28 - 16.68           MAP #6 Goat undil. Liver 28.03 - 19.24           MAP #7 Goat undil. Liver 32.03 - 17.53           MAP #8 Goat undil. Spleen 33.19 - 17.09		BHV #7	Cattle	undil.	Serum	pos	-	-	25.29
Mycobacterium avium subsp. Paratuber-culosis (MAP)         MAP #2         Goat undil. Lymph node undil. Colon undil. Lymph node undil. Liver undil. Undil. Liver undil. Undil. Liver undil. Liver undil. Undil. Liver undil. Undil		BHV #8	Cattle	undil.	Serum	pos	-	-	25.33
avium subsp. Paratuber- culosis (MAP)         MAP #2         Goat undil. Lymph node 31.81         - 16.47           MAP #3         Goat undil. Colon 32.21         - 18.06           MAP #4         Goat undil. Lymph node 34.67         - 16.53           MAP #5         Goat undil. Lymph node 32.28         - 16.68           MAP #6         Goat undil. Liver 28.03         - 19.24           MAP #7         Goat undil. Liver 32.03         - 17.53           MAP #8         Goat undil. Spleen 33.19         - 17.09		BHV #9	Cattle	undil.	Serum	pos	-	-	25.17
Paratuber-culosis (MAP)         MAP #2         Goat undil. Lymph node 31.81         - 16.47           MAP #3         Goat undil. Colon 32.21         - 18.06           MAP #4         Goat undil. Lymph node 34.67         - 16.53           MAP #5         Goat undil. Lymph node 32.28         - 16.68           MAP #6         Goat undil. Liver 28.03         - 19.24           MAP #7         Goat undil. Liver 32.03         - 17.53           MAP #8         Goat undil. Spleen 33.19         - 17.09	•	MAP #1	Goat	undil.	Lymph node	33.35	-	-	16.61
MAP #4       Goat       undil. Lymph node       34.67       -       -       16.53         MAP #5       Goat       undil. Lymph node       32.28       -       -       16.68         MAP #6       Goat       undil. Liver       28.03       -       -       19.24         MAP #7       Goat       undil. Liver       32.03       -       -       17.53         MAP #8       Goat       undil. Spleen       33.19       -       -       17.09	•	MAP #2	Goat	undil.	Lymph node	31.81	-	-	16.47
MAP #5 Goat undil. Lymph node 32.28 16.68  MAP #6 Goat undil. Liver 28.03 19.24  MAP #7 Goat undil. Liver 32.03 17.53  MAP #8 Goat undil. Spleen 33.19 17.09	culosis (MAP)	MAP #3	Goat	undil.	Colon	32.21	-	-	18.06
MAP #6 Goat undil. Liver 28.03 19.24  MAP #7 Goat undil. Liver 32.03 17.53  MAP #8 Goat undil. Spleen 33.19 17.09		MAP #4	Goat	undil.	Lymph node	34.67	-	-	16.53
MAP #7 Goat undil. Liver 32.03 17.53  MAP #8 Goat undil. Spleen 33.19 17.09		MAP #5	Goat	undil.	Lymph node	32.28	-	-	16.68
MAP #8 Goat undil. Spleen 33.19 17.09		MAP #6	Goat	undil.	Liver	28.03	-	-	19.24
·		MAP #7	Goat	undil.	Liver	32.03	-	-	17.53
MAP #9 Goat undil. Spleen 33.11 17.24		MAP #8	Goat	undil.	Spleen	33.19	-	-	17.09
		MAP #9	Goat	undil.	Spleen	33.11	-	-	17.24

Dil. = Dilution, Ref. = Reference, undil = undiluted, n.a. = not applicable, pos = positive, - = no C<sub>T</sub>

<sup>\*</sup> Reference assay were as follows: virotype SBV RT-PCR Kit (SBV samples), virotype BVDV RT-PCR Kit BVDV samples), inhouse PCR (EHDV samples), cattletype BHV1 gE Ab (BHV samples), bactotype MAP PCR Kit (MAP samples)

# 4.3 Diagnostic sensitivity, specificity and efficiency

#### 4.3.1 Definition diagnostic sensitivity

Percentage of positive samples in the new test of a population of true positive samples. True positive samples giving negative results in the new test are termed false negative.

Calculation: [true positives / (true positives + false negatives)]\*100

# 4.3.2 Definition diagnostic specificity

Percentage of negative samples in the new test of a population of true negative samples. True negative samples giving positive results in the new test are termed false positive.

Calculation: [true negatives / (false positives + true negatives)]\*100

### 4.3.3 Definition diagnostic efficiency

Diagnostic efficiency refers to the amount of agreement between the results from the new test and those from the reference test. It is expressed as a proportion of correctly identified samples among all samples.

Calculation: [(true positives + true negatives) / (true positives + true negatives + false positives + false negatives)]\*100

#### 4.3.4 Validation of the virotype BTV pan/8 2.0 RT-PCR Kit

For validation of the virotype BTV pan/8 2.0 RT-PCR Kit, n = 348 samples were tested. Samples (n = 182 pan BTV-positive of which were n = 139 BTV-8-positive, n = 166 BTV-negative) were kindly provided by German Veterinary Laboratories, the FLI (German National Reference Laboratory for BTV) and other reference labs.

For validation n = 254 cattle, n = 16 sheep and n = 28 goat samples were tested, of which n = 230 were blood, n = 9 were serum, and n = 59 were tissue samples. In addition, n = 50 culture samples were tested.

The latter samples were processed using the IndiMag Pathogen Kit (INDICAL BIOSCIENCE), the IndiSpin® Pathogen Kit (INDICAL BIOSCIENCE), or the QIAamp® Viral RNA Mini Kit (QIAGEN GmbH, Hilden, Germany), following manufacturer's instructions and tested with the virotype BTV pan/8 2.0 RT-PCR Kit and the virotype BTV pan/8 RT-PCR Kit as reference.

#### Results/ Conclusion

The summary is shown in Table 18. Individual results are shown in Figure 21 (pan BTV) and Figure 22 (BTV-8).

All positive BTV samples were detected correctly with the virotype BTV pan/8 2.0 RT-PCR Kit. The virotype BTV pan/8 2.0 RT-PCR Kit demonstrated a diagnostic sensitivity of 100 %, a diagnostic specificity of 100 % and a diagnostic efficiency of 100 %. In this study the virotype BTV pan/8 2.0 RT-PCR Kit demonstrated an overall higher sensitivity (with often lower  $C_T$  values) compared to the virotype BTV pan/8 RT-PCR Kit.

Table 18. Diagnostic sensitivity, specificity and efficiency of the virotype BTV pan/8 2.0 RT-PCR Kit.

virotype BTV pan/8 2.0	RT-PCR Kit	Re	feren	ce status	
Total	348	Reference positive	182	Reference negative	166
BTV-8-positive	139	true BTV-8-positive	139	false BTV-8-positive	0
pan BTV-positive	182	true pan BTV-positive	182	false pan BTV-positive	0
BTV-negative	166	false-negative	0	true-negative	166

Diagnostic sensitivity: 100 %

Diagnostic specificity: 100 %

Diagnostic efficiency: 100 %

# Comparison virotype BTV pan/8 2.0 RT-PCR Kit vs. virotype BTV pan/8 RT-PCR Kit (BTV pan; FAM; *n* = 182)

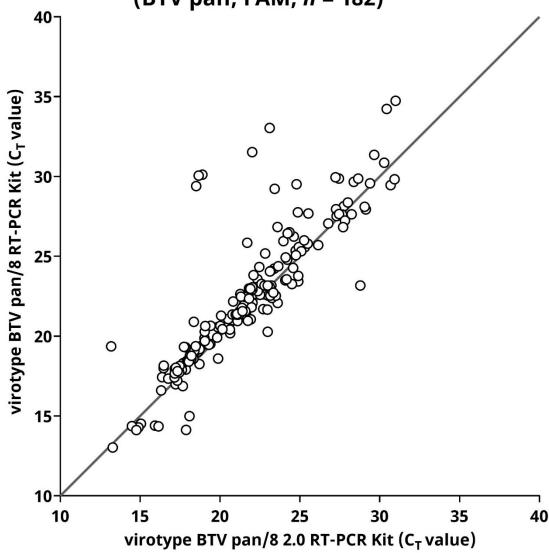


Figure 21. Comparison of FAM  $C_T$  values (**pan BTV**) from n = 182 pan BTV-positive samples tested with the virotype BTV pan/8 2.0 RT-PCR Kit compared to the virotype BTV pan/8 RT-PCR Kit. All samples situated above the black diagonal line showed lower  $C_T$  values with the virotype BTV pan/8 2.0 RT-PCR Kit when compared to the virotype BTV pan/8 RT-PCR Kit.

# Comparison virotype BTV pan/8 2.0 RT-PCR Kit vs. virotype BTV pan/8 RT-PCR Kit

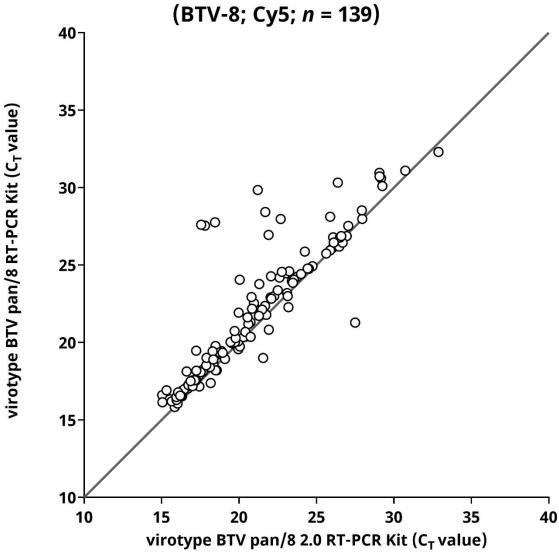


Figure 22. Comparison of Cy5  $C_T$  values (**BTV-8**) from n = 139 BTV-8-positive samples tested with the virotype BTV pan/8 2.0 RT-PCR Kit compared to the virotype BTV pan/8 RT-PCR Kit. All samples situated above the black diagonal line showed lower  $C_T$  values with the virotype BTV pan/8 2.0 RT-PCR Kit when compared to the virotype BTV pan/8 RT-PCR Kit.

# 4.4 Repeatability

The same sample panel comprising four BTV-positive RNA samples (samples 1-4), a negative RNA sample from sheep blood (sample 5), and the Positive Control (sample 5) and was used for assessment of intra-assay variance, inter-assay variance, batch-to-batch variance, and stability, testing. BTV-positive samples 1-4 as well as another negative RNA sample from sheep blood (sample 6) was used for robustness testing for the virotype BTV pan/8 2.0 RT-PCR Kit (Table 19).

Table 19. Sample panel for assessment of intra-assay variance, inter-assay variance, batch-to-batch variance, stability, and robustness testing for the virotype BTV pan/8 2.0 RT-PCR Kit.

Sample	BTV status	Species	Description	Dilution
1	positive	Cattle	Extracted nucleic acids from blood of a BTV-8 infected cow	
2	positive	Cattle	sample 1 diluted in RNA from neg cow blood	1:10
3	positive	Cattle	sample 1 diluted in RNA from neg cow blood	1:100
4	positive	Cattle	sample 1 diluted in RNA from neg cow blood	1:1,000
5	negative	Sheep	Extracted nucleic acids from blood	
6	negative	Sheep	Extracted nucleic acids from blood	
PC	PC			

PC = Positive Control

# 4.4.1 Intra-assay variance

The sample panel listed in Table 19 (sample 1-5, PC) was tested in a sevenfold setup in the same PCR run with the virotype BTV pan/8 2.0 RT-PCR Kit on the Agilent Technologies Aria Mx instrument.

#### Results / Conclusion

The intra-assay variance is on average 0.75 % for pan BTV (FAM; Table 20), 0.53 % for BTV-8 (Cy5; Table 21), and 0.84 % for the endogenous Internal Control (JOE; Table 22).

These results show an excellent reproducibility in the same RT-PCR run for the virotype BTV pan/8 2.0 RT-PCR Kit.

Table 20. Intra-assay variance for **pan BTV** (FAM) for the virotype BTV pan/8 2.0 RT-PCR Kit using the Agilent Technologies Aria Mx instrument.

		]	intra-as	ssay va	riance	for par	n BTV (F	AM)			
Sample	BTV			Reactio	ns (C <sub>T</sub>	values)	)		Mean	SD	CV%
	status	1	2	3	4	5	6	7			
1	pos	20.69	20.56	20.56	20.50	20.64	20.47	20.90	20.62	0.146	0.71
2	pos	23.82	23.77	23.76	23.50	23.58	23.78	23.97	23.74	0.156	0.66
3	pos	26.93	26.87	27.32	27.11	27.39	26.92	26.92	27.07	0.213	0.79
4	pos	30.38	30.32	30.44	30.07	30.16	30.18	29.79	30.19	0.220	0.73
5	neg	-	-	-	-	-	-		-	-	-
PC	pos	27.53	27.24	27.97	27.72	27.36	27.48	27.50	27.54	0.240	0.87
Mean											0.75

neg = negative, pos = positive, SD = standard deviation, CV = coefficient of variation, - = no C<sub>T</sub>

Table 21. Intra-assay variance for **BTV-8** (Cy5) for the virotype BTV pan/8 2.0 RT-PCR Kit using the Agilent Technologies Aria Mx instrument.

			Intr	a-assay	variar	ce for	BTV-8 (	Cy5)			
Sample	BTV			Reactio	ns (C <sub>T</sub>	values)	)		Mean	SD	CV%
	status	1	2	3	4	5	6	7	•		
1	pos	19.50	19.42	19.36	19.32	19.40	19.43	19.69	19.45	0.122	0.63
2	pos	22.67	22.50	22.56	22.44	22.51	22.40	22.48	22.51	0.088	0.39
3	pos	25.89	25.54	25.68	25.69	26.00	25.70	25.65	25.74	0.156	0.61
4	pos	29.16	28.60	28.84	28.96	28.78	28.73	28.89	28.85	0.179	0.62
5	neg	-	-	-	-	-	-		-	-	-
PC	pos	26.51	26.29	26.49	26.41	26.25	26.28	26.30	26.36	0.107	0.41
Mean										'	0.53

neg = negative, pos = positive, SD = standard deviation, CV = coefficient of variation,  $- = no C_T$ 

Table 22. Intra-assay variance for the endogenous **Internal Control** (JOE) for the virotype BTV pan/8 2.0 RT-PCR Kit using the Agilent Technologies Aria Mx instrument.

		Intra	a-assay	varian	ce for t	he Inte	ernal C	ontrol	(JOE)		
Sample	BTV			Reactio	ns (C <sub>T</sub>	values)			Mean	SD	CV%
	status	1	2	3	4	5	6	7	-		
1	pos	24.30	24.65	24.27	24.23	24.26	24.50	24.71	24.42	0.201	0.82
2	pos	24.56	24.22	24.50	24.15	24.44	24.40	24.54	24.40	0.159	0.65
3	pos	24.65	24.52	24.51	24.39	24.44	24.51	24.50	24.50	0.080	0.33
4	pos	25.33	24.33	24.44	23.89	24.34	24.43	24.83	24.51	0.453	1.85
5	neg	24.61	24.52	24.35	24.56	24.72	24.40	24.50	24.52	0.125	0.51
PC	pos	27.01	26.37	26.47	26.61	26.54	26.30	26.60	26.56	0.230	0.87
Mean											0.84

neg = negative, pos = positive, SD = standard deviation, CV = coefficient of variation

# Intra-assay variance virotype BTV pan/8 2.0 RT-PCR Kit

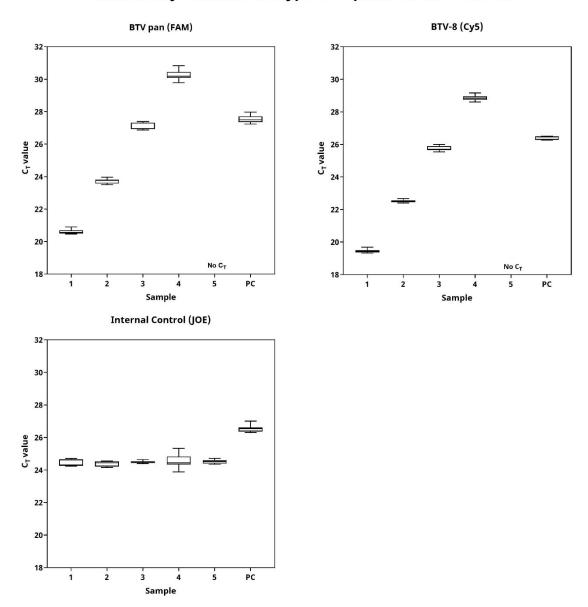


Figure 23. Boxplots of intra-assay variance for pan BTV (FAM), BTV-8 (Cy5), and the endogenous Internal Control (JOE) for the virotype BTV 2.0 pan/8 RT-PCR Kit tested using the Agilent Technologies Aria Mx instrument.

# 4.4.2 Inter-assay variance

The sample panel listed in Table 19 (sample 1-5, PC) was tested in six different PCR runs using the virotype BTV pan/8 2.0 RT-PCR Kit on the Agilent Technologies Aria Mx instrument.

#### Results / Conclusion

The inter-assay variance is on average 1.08 % for pan BTV (FAM; Table 23), 0.85 % for BTV-8 (Cy5; Table 24), and 0.96 % for the endogenous Internal Control (JOE; Table 25). These results show an excellent reproducibility in different RT-PCR runs for the virotype BTV pan/8 2.0 RT-PCR Kit.

Table 23. Inter-assay variance for **pan BTV** (FAM) for the virotype BTV pan/8 2.0 RT-PCR Kit using Agilent Technologies Aria Mx instrument.

Inter-assay variance for pan BTV (FAM)										
Sample	BTV		Rea	ctions	(C <sub>T</sub> valu	ıes)		Mean	SD	CV%
	status	1	2	3	4	5	6			
1	pos	21.21	20.69	21.55	21.59	21.11	20.82	21.16	0.368	1.74
2	pos	24.16	23.82	23.90	24.17	24.00	24.23	24.05	0.165	0.69
3	pos	27.24	27.11	27.22	27.24	27.13	27.23	27.2	0.059	0.22
4	pos	30.22	30.32	29.38	30.23	30.55	30.43	30.19	0.415	1.38
5	neg	-	-	-	-	-		-	-	-
PC	pos	27.29	27.24	26.92	27.64	27.52	28.00	27.44	0.372	1.36
Mean										1.08

neg = negative, pos = positive, SD = standard deviation, CV = coefficient of variation, - = no C<sub>T</sub>

Table 24. Inter-assay variance for **BTV-8** (Cy5) for the virotype BTV pan/8 2.0 RT-PCR Kit using the using the Agilent Technologies Aria Mx instrument.

		In	ter-ass	ay vari	ance fo	r BTV-8	3 (Cy5)			
Sample	BTV		Rea	ctions	(C <sub>T</sub> valu	ues)		Mean	SD	CV%
	status	1	2	3	4	5	6			
1	pos	19.94	19.5	20.02	20.02	19.70	19.71	19.82	0.211	1.07
2	pos	22.74	22.67	22.72	22.89	22.58	22.76	22.73	0.103	0.45
3	pos	25.70	25.69	25.68	25.83	25.69	25.88	25.75	0.087	0.34
4	pos	29.00	28.6	28.56	29.52	29.56	29.07	29.05	0.43	1.48
5	neg	-	-	-	-	-		-	-	-
PC	pos	26.42	26.29	26.10	26.56	26.69	26.73	26.47	0.243	0.92
Mean										0.85

neg = negative, pos = positive, SD = standard deviation, CV = coefficient of variation, - = no  $C_T$ 

Table 25. Inter-assay variance for endogenous **Internal Control** (JOE) for the virotype BTV pan/8 2.0 RT-PCR Kit using the Agilent Technologies Aria Mx instrument.

	Inter-ass	ay vari	ance fo	r the e	ndogei	nous In	ternal	Control	(JOE)	
Sample	BTV		Rea	ctions	(C <sub>⊤</sub> valı	ues)		Mean	SD	CV%
	status	1	2	3	4	5	6			
1	pos	24.77	24.3	24.83	24.74	24.64	24.64	24.65	0.188	0.76
2	pos	24.51	24.56	24.13	24.41	24.32	24.40	24.39	0.153	0.63
3	pos	24.43	24.39	24.31	24.73	24.50	24.31	24.45	0.157	0.64
4	pos	24.27	24.33	24.56	24.30	24.78	24.58	24.47	0.202	0.83
5	neg	24.26	24.52	24.19	24.41	24.65	24.61	24.44	0.187	0.77
PC	pos	26.13	26.37	26.39	27.29	27.15	27.47	26.80	0.568	2.12
Mean										0.96

neg = negative, pos = positive, SD = standard deviation, CV = coefficient of variation

# Inter-assay variance virotype BTV pan/8 2.0 RT-PCR Kit

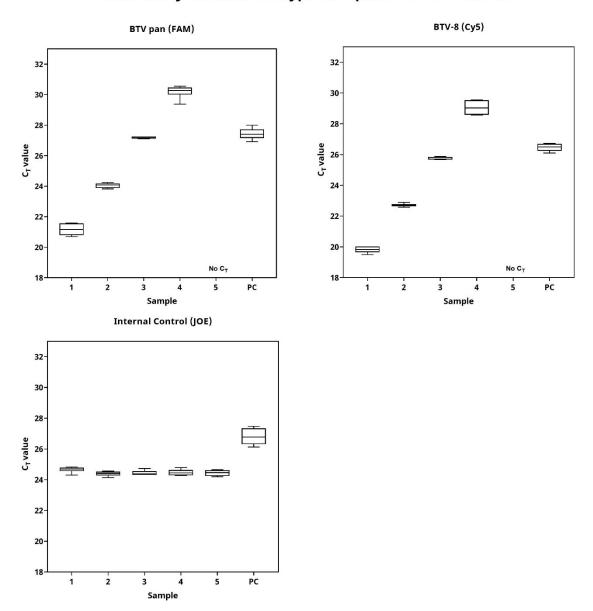


Figure 24. Boxplots of inter-assay variance for pan BTV (FAM), BTV-8 (Cy5), and the endogenous Internal Control (JOE) for the virotype BTV 2.0 pan/8 RT-PCR Kit tested using the Agilent Technologies Aria Mx instrument.

# 4.4.3 Batch-to-batch comparison

The sample panel listed in Table 19 (sample 1-5, PC) was tested in in the same PCR run using three different batches (1, 2, 3) of the virotype BTV pan/8 2.0 RT-PCR Kit on the Agilent Technologies Aria Mx instrument.

#### Results / Conclusion

The batch-to-batch performance showed on average variance of 1.36 % for pan BTV (FAM), 0.94 % for BTV-8 (Cy5), 0.78 % for the endogenous Internal Control (JOE).

These results show an excellent reproducibility of different batches for the virotype BTV pan/8 2.0 RT-PCR Kit (Table 26 - Table 28, Figure 25).

Table 26. Batch-to-batch variance for **pan BTV** (FAM) for the virotype BTV pan/8 2.0 RT-PCR Kit using the Agilent Technologies Aria Mx instrument.

		Batch-to-bat	ch variance f	or pan BTV (	FAM)		
Sample	BTV	Batch	number (C <sub>T</sub> \	Mean	SD	CV%	
	status	1	2	3	_		
1	pos	21.59	21.61	21.62	21.61	0.015	0.07
2	pos	24.17	24.36	23.81	24.11	0.279	1.16
3	pos	27.81	26.82	27.24	27.29	0.497	1.82
4	pos	29.90	30.23	30.63	30.25	0.366	1.21
5	neg	-	-	-	-	-	-
PC	pos	28.52	27.64	27.13	27.76	0.703	2.53
Mean							1.36

neg = negative, pos = positive, SD = standard deviation, CV = coefficient of variation, - = no C<sub>T</sub>

Table 27. Batch-to-batch variance for **BTV-8** (Cy5) for the virotype BTV pan/8 2.0 RT-PCR Kit using the Agilent Technologies Aria Mx instrument.

		Batch-to-b	atch variance	for BTV-8 (C	y5)		
Sample	BTV	Batch	number (C <sub>T</sub> v	Mean	SD	CV%	
	status	1	2	3	_		
1	pos	20.02	19.98	20.02	20.01	0.023	0.12
2	pos	22.89	22.67	22.61	22.72	0.147	0.65
3	pos	26.35	25.67	25.83	25.95	0.356	1.37
4	pos	29.00	29.52	29.43	29.32	0.278	0.95
5	neg	-	-	-	-	-	-
PC	pos	27.10	26.56	26.24	26.63	0.435	1.63
Mean							0.94

neg = negative, pos = positive, SD = standard deviation, CV = coefficient of variation,  $- = no C_T$ 

Table 28. Batch-to-batch variance for the endogenous **Internal Control** (JOE) for the virotype BTV pan/8 2.0 RT-PCR Kit using the Agilent Technologies Aria Mx instrument.

	Batch-to-bat	tch variance	for the endo	genous Inter	nal Contro	l (JOE)	
Sample	BTV	Batch	number (C <sub>T</sub> v	alues)	Mean	SD	CV%
	status	1	2	3	_		
1	pos	24.74	24.72	24.81	24.76	0.047	0.19
2	pos	24.41	24.19	24.04	24.21	0.186	0.77
3	pos	24.60	24.60	24.73	24.64	0.075	0.30
4	pos	24.26	24.30	24.40	24.32	0.072	0.230
5	neg	24.41	24.65	24.78	24.61	0.188	0.76
PC	pos	28.33	27.29	27.13	27.58	0.652	2.36
Mean					1	1	0.78

neg = negative, pos = positive, SD = standard deviation, CV = coefficient of variation

# Batch-to-batch variance virotype BTV pan/8 2.0 RT-PCR Kit

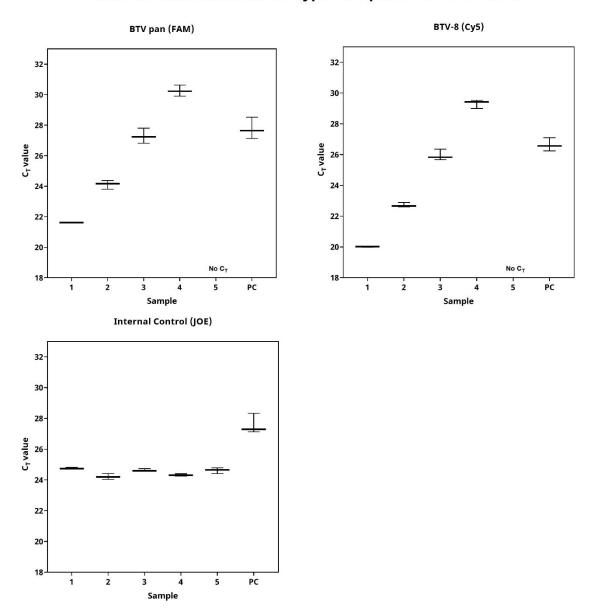


Figure 25. Boxplots of batch-to-batch variance for pan BTV (FAM), BTV-8 (Cy5), and the endogenous Internal Control (JOE) for the virotype BTV 2.0 pan/8 RT-PCR Kit tested using the Agilent Technologies Aria Mx instrument.

# 4.4.4 Comparison of real-time PCR thermocyclers

The virotype BTV pan/8 2.0 RT-PCR Kit can be used on different standard real-time PCR cyclers. Table 29 gives an overview of selected PCR cyclers and their approximate run times, using the virotype BTV pan/8 2.0 RT-PCR protocol.

Note: The use of the RT-PCR Kit is not limited to the mentioned instruments.

Table 29. Selected overview of real-time thermocyclers and their approximate run times for the virotype BTV pan/8 2.0 RT-PCR protocol.

	Thermod	cycler		
	Manufacturer	Model	Filters	Run time [minutes]
Α	Agilent Technologies, Santa Clara, California, USA	AriaMx	FAM, HEX, Cy5	68
В	Bio-Rad Laboratories, Inc., Hercules, California, USA	CFX96	FAM, HEX, Cy5	67
С	Thermo Fisher Scientific Inc., Waltham, Massachusetts, USA	Applied Biosystems <sup>™</sup> 7500 Fast <sup>1</sup>	FAM, JOE, Cy5	52
D	Thermo Fisher Scientific Inc., Waltham, Massachusetts, USA	QuantStudio 5	FAM, VIC/JOE, Cy5	68

<sup>&</sup>lt;sup>1</sup> Fast Mode setting

A titration series of pan BTV and BTV-8 *in vitro* RNA (10<sup>6</sup> - 1 copies/reaction) were analyzed in the same PCR run using the virotype BTV pan/8 2.0 RT-PCR Kit when determining the variance in FAM (pan BTV) and Cy5 channels (BTV-8). Furthermore, six bovine blood samples were analyzed in the same PCR run using the virotype BTV pan/8 2.0 RT-PCR Kit when determining the variance in the JOE channel (endogenous Internal Control). For the Thermo Fisher ABI 7500 Fast instrument, the Standard Mode with ROX as passive reference dye was performed.

#### Results / Conclusion

The results are summarized in Table 30 (pan BTV / FAM channel), Table 31 (BTV-8, Cy5 channel), and Table 32 (endogenous Internal Control/ JOE channel). All samples tested on different real-time PCR thermocycler instruments showed comparable results.

Based on this data set, INDICAL can recommend using the virotype BTV pan/8 2.0 RT-PCR Kit on the Bio-Rad CFX96, Agilent Technologies AriaMx, Thermo Fisher ABI 7500 Fast, and Thermo Fisher QuantStudio 5 instruments.

Table 30. Inter-thermocycler variance for **pan BTV** (FAM).

	Inte	r-thermocycl	er variance f	or pan BTV (F	AM)			
Сору	Material	BTV	Thermocycler (C <sub>T</sub> values)					
number		status	Α	В	С	D		
10 <sup>6</sup>	in-vitro RNA	pos	17.89	19.03	18.30	18.35		
10 <sup>5</sup>	<i>in-vitro</i> RNA	pos	21.02	22.10	21.55	21.64		
104	<i>in-vitro</i> RNA	pos	24.14	25.50	24.81	24.99		
10³	<i>in-vitro</i> RNA	pos	27.48	28.76	28.04	28.42		
10 <sup>2</sup>	<i>in-vitro</i> RNA	pos	30.52	32.13	31.24	31.61		
10	in-vitro RNA	pos	33.63	35.68	34.58	35.01		
5	in-vitro RNA	pos	35.14	35.53	35.13	35.65		
1	in-vitro RNA	pos	37.05	38.49	37.58	37.15		

pos = positive

Table 31. Inter-thermocycler variance for **BTV-8** (Cy5).

		Inter-cycler	variance for	BTV-8 (Cy5)		
Сору	Material	BTV		Thermocycle	er (C <sub>T</sub> values)	
number		status	Α	В	С	D
10 <sup>6</sup>	in-vitro RNA	pos	17.70	18.82	18.54	17.77
10 <sup>5</sup>	<i>in-vitro</i> RNA	pos	20.83	22.05	21.89	20.93
10 <sup>4</sup>	<i>in-vitro</i> RNA	pos	23.96	25.41	25.19	24.25
10³	<i>in-vitro</i> RNA	pos	27.04	28.60	28.48	27.46
10 <sup>2</sup>	<i>in-vitro</i> RNA	pos	30.20	32.19	31.79	30.67
10	in-vitro RNA	pos	33.57	34.61	34.60	34.16
5	in-vitro RNA	pos	34.19	35.60	35.50	34.56
1	in-vitro RNA	pos	-	37.69	38.59	36.44

pos = positive, - = no  $C_T$ 

Table 32. Inter-thermocycler variance for the endogenous **Internal Control** (JOE).

Inter-cycler variance for the endogenous Internal Control (JOE)									
Sample	Material	BTV	Thermocycler (C <sub>T</sub> values)						
		status	Α	В	С	D			
S1	Blood	pos	24.97	23.05	24.27	24.97			
S2	Blood	pos	22.22	23.34	22.17	22.22			
S3	Blood	pos	24.37	25.36	23.40	24.37			
S4	Blood	pos	21.51	22.97	21.97	21.51			
S5	Blood	pos	22.00	22.61	22.61	22.00			
S6	Blood	pos	17.51	19.42	18.43	17.51			

pos = positive

# 4.5 Stability testing

#### 4.5.1 Freeze-thaw-cycles

The sample panel listed in Table 19 (sample 1-5, PC) was tested at the time of production and after ten freeze/thaw cycles. The mean value (Mean), standard deviation (SD) and coefficient of variation (CV) were calculated.

# Results / Conclusion

The virotype BTV pan/8 2.0 RT-PCR Kit shows excellent stability after 10 freeze/thaw cycles of the virotype BTV pan/8 2.0 Master Mix. The average variance was 0.70 % for pan BTV (FAM), 0.53 % for BTV-8 (Cy5), and 0.99 % for the endogenous Internal Control (JOE) (Table 33 - Table 35).

Table 33. Stability testing for **pan BTV** (FAM) of the virotype BTV pan/8 2.0 RT-PCR Kit using the Agilent Technologies Aria Mx instrument.

		Stability fo	r pan BTV (FAM)			
Sample	BTV	Freeze-thaw-c	ycle (C <sub>⊤</sub> values)	Mean	SD	CV%
	status	1	10	_		
1	pos	21.21	20.82	21.02	0.276	1.31
2	pos	24.16	24.23	24.20	0.049	0.21
3	pos	27.17	27.23	27.20	0.042	0.16
4	pos	30.44	30.43	30.44	0.007	0.02
5	neg	-	-	-	-	-
PC	pos	27.29	28.00	27.65	0.502	1.82
Mean						0.70

PC = Positive Control, neg = negative, pos = positive, SD = standard deviation, CV = coefficient of variation, -= no  $C_T$ 

Table 34. Stability testing for **BTV-8** (Cy5) of the virotype BTV pan/8 2.0 RT-PCR Kit using the Agilent Technologies Aria Mx instrument.

		Stability f	or BTV-8 (Cy5)			
Sample	BTV	ΓV Freeze-thaw-cycle (C <sub>τ</sub> values)			SD	CV%
	status	1	10	_		
1	pos	19.94	19.71	19.83	0.163	0.82
2	pos	22.74	22.76	22.75	0.014	0.06
3	pos	25.75	25.88	25.82	0.092	0.36
4	pos	28.84	29.07	28.96	0.163	0.56
5	neg	-	-	-	-	-
PC	pos	26.42	26.73	26.58	0.219	0.83
Mean						0.53

PC = Positive Control, neg = negative, pos = positive, SD = standard deviation, CV = coefficient of variation, -= no  $C_T$ 

Table 35. Stability testing for the endogenous **Internal Control** (JOE) of the virotype BTV pan/8 2.0 RT-PCR Kit using the Agilent Technologies Aria Mx instrument.

	Stabil	ity for the endoge	enous Internal Co	ntrol (JOE)		
Sample	BTV	Freeze-thaw-c	ycle (C⊤ values)	Mean	SD	CV%
	status	1	10	_		
1	pos	24.77	24.64	24.71	0.092	0.37
2	pos	24.51	24.40	24.46	0.078	0.32
3	pos	24.31	24.31	24.31	0.000	0.00
4	pos	24.44	24.58	24.51	0.099	0.40
5	neg	26.13	27.47	26.8	0.948	3.54
PC	pos	18.56	18.22	18.39	0.240	1.31
Mean				1		0.99

PC = Positive Control, neg = negative, pos = positive, SD = standard deviation, CV = coefficient of variation

# 4.5.2 Heparin inhibition

To test the stability of the virotype BTV pan/8 2.0 RT-PCR Kit, sample inhibition was simulated by treating a BTV-positive sample with an increasing concentration of heparin (0.025 – 1.00  $\mu$ g/reaction). The sample was tested in duplicate using the virotype BTV pan/8 2.0 RT-PCR Kit.

#### Results / Conclusion

The inhibition study using heparin showed that the endogenous Internal Control signal will be affected first in case of sample inhibition. The increased  $C_T$  value will help to identify partially and fully inhibited samples (Table 36).

Table 36. Stability testing of the virotype BTV pan/8 2.0 RT-PCR Kit for inhibition by heparin.

# Stability (heparin inhibition)

# virotype BTV pan/8 2.0 RT-PCR

		, ,	
Heparin	pan BTV	BTV-8	Internal Control
[µg / reaction]	$\mathbf{C}_{T}$	$\mathbf{C}_{T}$	$\mathbf{C}_{T}$
0.00	25.14	24.16	20.19
0.00	24.94	23.66	20.22
0.025	24.79	23.89	19.61
0.023	25.06	23.7	20.09
0.05	24.88	23.62	20.16
0.03	24.78	23.56	20.03
0.25	26.10	24.43	21.85
0.23	26.30	24.54	22.00
0.50	34.65	31.64	-
0.30	32.83	29.45	-
0.75	36.47	35.04	-
0.75	36.98	33.36	-
1.00	-	-	-
1.00	-	-	-

- = no C<sub>T</sub>

#### 4.5.3 EDTA inhibition

To test the stability of the virotype BTV pan/8 2.0 RT-PCR Kit, sample inhibition was simulated by treating a positive sample with an increasing concentration of Ethylenediaminetetraacetic acid (0.1 – 0.6 mM EDTA final concentration in reaction mix). The sample was tested in duplicate using the virotype BTV pan/8 2.0 RT-PCR Kit.

#### **Results / Conclusion**

The inhibition study using EDTA showed that the endogenous Internal Control signal will be affected first in case of sample inhibition. The increased  $C_T$  value will help to identify partially and fully inhibited samples (Table 37).

Table 37. Stability testing of the virotype BTV pan/8 2.0 RT-PCR Kit for inhibition by EDTA.

	Stability (ED	TA inhibition)	
	viro	type BTV pan/8 2.0 F	RT-PCR
EDTA	pan BTV	BTV-8	Internal Control
[mM]	$\mathbf{C}_{T}$	$\mathbf{C}_{T}$	$\mathbf{C}_{T}$
0.0	24.54	23.43	19.75
0.0	24.52	23.50	19.64
0.1	24.01	23.15	19.28
0.1	23.95	23.09	19.46
0.2	24.04	23.18	19.22
0.2	24.19	23.16	18.96
0.3	23.80	22.77	19.63
0.5	24.63	23.08	20.28
0.4	24.68	23.59	21.4
0.4	24.22	23.03	20.9
0.5	31.39	36.2	-
0.5	31.42	36.91	-
0.6	-	-	-
0.0	-	-	-

 $<sup>- =</sup> no C_T$ 

#### 4.6 Robustness

#### 4.6.1 Robustness: Variation of sample volume

To test the robustness of the virotype BTV pan/8 2.0 RT-PCR Kit, the sample volume was varied by 10 % to simulate errors in the reaction mix preparation. Therefore, the sample panel listed in Table 19 (sample 1-4, 6, PC) was tested in triplicates in one PCR run on the Agilent Technologies Aria Mx instrument.

#### Results / Conclusion

Results are shown in Table 38 - Table 40. The virotype BTV pan/8 2.0 RT-PCR Kit shows excellent robustness for 10 % sample volume variation with an average variance of 1.20 % for pan BTV (FAM), 1.01 % for BTV-8 (Cy5), and 0.86 % for the endogenous Internal Control (JOE).

Table 38. Robustness testing of the virotype BTV pan/8 2.0 RT-PCR Kit (**pan BTV** / FAM) for different sample volumes used in the RT-qPCR. The tests were performed on the Agilent Technologies Aria Mx instrument.

	pan BTV (FAM) C <sub>τ</sub>					
Sample	S	ample volum	е	Mean	SD	CV%
-	4.5 µl	5.0 μl	5.5 µl	-		
	22.48	21.76	21.18			
1	21.60	21.11	21.31	21.49	0.470	2.19
	21.80	21.17	21.01			
	24.19	24.39	24.52			
2	24.45	24.00	24.16	24.29	0.174	0.72
	24.25	24.18	24.46			
	27.25	26.78	26.95			
3	27.58	27.13	27.26	27.08	0.346	1.28
	27.24	26.37	27.16			
	29.99	30.95	30.45			
4	30.18	30.85	30.58	30.54	0.302	0.99
	30.62	30.55	30.67			
	-	-	-			
5	-	-	-	-	-	-
	-	-	-			
	27.61	28.00	27.69			
PC	27.83	27.52	27.70	27.77	0.234	0.84
	28.18	27.91	27.46			
Mean						1.20

 $<sup>\</sup>overline{SD}$  = standard deviation;  $\overline{CV}$  = coefficient of variation, - = no  $\overline{C_T}$ 

Table 39. Robustness testing of the virotype BTV pan/8 2.0 RT-PCR Kit (**BTV-8** / Cy5) for different sample volumes used in the RT-qPCR. The tests were performed on the Agilent Technologies Aria Mx instrument.

			BTV-8 (Cy5) C	т		
Sample	S	ample volum	е	Mean	SD	CV%
-	4.5 µl	5.0 µl	5.5 µl	-		
	20.91	20.16	19.64			
1	20.07	19.70	19.70	20.00	0.415	2.07
	20.28	19.79	19.71			
	22.88	22.95	22.92			
2	23.02	22.58	22.81	22.85	0.147	0.64
	22.88	22.64	22.94			
	26.23	25.76	25.68			
3	26.30	25.69	25.83	25.93	0.247	0.95
	26.22	25.81	25.87			
	29.22	29.27	29.18			
4	29.28	29.40	28.89	29.29	0.196	0.67
	29.50	29.56	29.30			
	-	-	-			
5	-	-	-	-	-	-
	-	-	-			
	26.55	26.71	26.63			
PC	26.75	26.69	26.72	26.69	0.196	0.73
	26.91	26.97	26.30			
Mean						1.01

SD = standard deviation;  $CV = coefficient of variation, - = no C_T$ 

Table 40. Robustness testing of the virotype BTV pan/8 2.0 RT-PCR Kit (endogenous **Internal Control** / JOE) for different sample volumes used in the RT-qPCR. The tests were performed on the Agilent Technologies Aria Mx instrument.

		Endogenous	Internal Co	ntrol (JOE) C <sub>T</sub>		
Sample	S	ample volum	e	Mean	SD	CV%
-	4.5 µl	5.0 μl	5.5 µl	-		
	25.26	24.92	24.59			
1	25.15	24.64	24.56	24.86	0.362	1.46
	25.48	24.71	24.43			
	24.68	24.73	24.65			
2	24.76	24.32	24.43	24.60	0.156	0.63
	24.48	24.64	24.74			
	24.80	24.37	24.38			
3	24.95	24.50	24.53	24.57	0.201	0.82
	24.51	24.40	24.68			
	24.40	24.50	24.52			
4	24.70	24.73	24.48	24.61	0.155	0.63
	24.85	24.78	24.57			
	18.39	18.56	18.31			
5	18.59	18.39	18.38	18.44	0.135	0.73
	18.27	18.67	18.41			
	27.31	27.23	26.91			
PC	27.03	27.15	26.84	27.06	0.235	0.87
	27.30	27.16	26.61			
Mean						0.86

SD = standard deviation; CV = coefficient of variation

# 4.6.2 Robustness: Variation of annealing time

To test the robustness of the virotype BTV pan/8 2.0 RT-PCR Kit, the annealing time during RT-qPCR reaction was varied by 10 % to simulate cycling errors. Therefore, the sample panel listed in Table 19 (sample 1-4, 6, PC) was tested in triplicates in one PCR run on the Agilent Technologies Aria Mx instrument.

#### Results / Conclusion

Results are shown in Table 41 - Table 43. The virotype BTV pan/8 2.0 RT-PCR Kit shows excellent robustness for 10 % annealing time variation with an average variance of 1.13 % for pan BTV (FAM), 0.86 % for BTV-8 (Cy5), and 0.80 % for the endogenous Internal Control (JOE).

Table 41. Robustness testing of the virotype BTV pan/8 2.0 RT-PCR Kit (**pan BTV** / FAM) for different annealing times used in the RT-qPCR. The tests were performed on the Agilent Technologies Aria Mx instrument.

		pa	an BTV (FAM)	Ст		
Sample	A	nnealing tim	e	Mean	SD	CV%
	27 sec	30 sec	33 sec	_		
	21.38	21.76	21.59			
1	21.67	21.11	21.70	21.46	0.253	1.18
	21.17	21.17	21.55			
	24.33	24.39	24.39			
2	24.20	24.00	24.34	24.28	0.146	0.60
	24.21	24.18	24.48			
	27.33	26.78	27.32			
3	27.23	27.13	27.05	27.01	0.317	1.18
	26.74	26.37	27.10			
	30.26	30.95	30.63			
4	30.00	30.85	30.08	30.28	0.509	1.68
	29.72	30.55	29.46			
	-	-	-			
5	-	-	-	-	-	-
	-	-	-			
	27.91	28.00	27.27			
PC	27.52	27.52	27.58	27.61	0.285	1.03
	27.18	27.91	27.64			
Mean						1.13

SD = standard deviation; CV = coefficient of variation, - = no  $C_T$ 

Table 42. Robustness testing of the virotype BTV pan/8 2.0 RT-PCR Kit (**BTV-8** / Cy5) for different annealing times used in the RT-qPCR. The tests were performed on the Agilent Technologies Aria Mx instrument.

			BTV-8 (Cy5) C	т		
Sample	P	nnealing tim	e	Mean	SD	CV%
	27 sec	30 sec	33 sec	-		
	20.00	20.16	20.15			
1	20.15	19.70	20.27	20.03	0.202	1.01
	19.87	19.79	20.20			
	22.84	22.95	22.98			
2	22.83	22.58	23.04	22.92	0.245	1.07
	22.96	22.64	23.42			
	25.91	25.76	26.21			
3	25.87	25.69	26.09	25.90	0.167	0.65
	25.79	25.81	25.98			
	29.22	29.27	29.71			
4	29.04	29.40	28.96	29.25	0.263	0.90
	29.06	29.56	29.01			
	-	-	-			
5	-	-	-	-	-	-
	-	-	-			
	26.69	26.71	26.51			
PC	26.44	26.69	26.76	26.67	0.178	0.67
	26.45	26.97	26.83			
Mean						0.86

SD = standard deviation;  $CV = coefficient of variation, - = no C_T$ 

Table 43. Robustness testing of the virotype BTV pan/8 2.0 RT-PCR Kit (endogenous **Internal Control** / JOE) for different annealing times used in the RT-qPCR. The tests were performed on the Agilent Technologies Aria Mx instrument.

	Endogenous Internal Control (JOE) C <sub>⊤</sub>					
Sample	P	nnealing tim	е	Mean	SD	CV%
	27 sec	30 sec	33 sec	-		
	24.89	24.92	24.88			
1	25.30	24.64	24.99	24.91	0.186	0.75
	24.91	24.71	24.98			
	24.79	24.73	24.58			
2	24.48	24.32	24.49	24.58	0.141	0.57
	24.60	24.64	24.63			
	24.67	24.37	24.44			
3	24.72	24.50	24.40	24.49	0.124	0.50
	24.50	24.40	24.43			
	24.36	24.50	24.67			
4	24.65	24.73	24.39	24.55	0.173	0.70
	24.60	24.78	24.30			
	18.25	18.56	18.10			
5	18.31	18.39	18.38	18.44	0.224	1.22
	18.84	18.67	18.44			
	27.92	27.23	27.04			
PC	27.16	27.15	27.11	27.19	0.287	1.05
	26.94	27.16	27.02			
Mean						0.80

SD = standard deviation; CV = coefficient of variation

# 4.6.3 Robustness: Variation of annealing temperature

To test the robustness of the virotype BTV pan/8 2.0 RT-PCR Kit, the annealing temperature during RT-qPCR reaction was varied by 1°C to simulate cycling errors. Therefore, the sample panel listed in Table 19 (sample 1-4, 6, PC) was tested in duplicates in one PCR run on the Agilent Technologies Aria Mx instrument.

#### Results / Conclusion

Results are shown in Table 44 - Table 46. The virotype BTV pan/8 2.0 RT-PCR Kit shows excellent robustness for 1°C annealing temperature variation with an average variance of 0.95 % for pan BTV (FAM), 0.69 % for BTV-8 (Cy5), and 1.27 % for the endogenous Internal Control (JOE).

Table 44. Robustness testing of the virotype BTV pan/8 2.0 RT-PCR Kit (**pan BTV** / FAM) for different annealing temperatures used in the RT-qPCR. The tests were performed on the Agilent Technologies Aria Mx instrument.

		pa	an BTV (FAM)	C <sub>T</sub>		
Sample	Anne	aling temper	ature	Mean	SD	CV%
-	59°C	60°C	61°C			
1	21.49	21.11	21.20	21.24	0.216	1.01
ı	21.42	21.17	21.66	21.34	0.216	1.01
2	24.36	24.00	24.73	24.40	0.274	1.12
	24.49	24.18	24.62	24.40	0.274	1.12
3	27.43	26.78	27.76	27.25	0.357	1.31
3	27.41	27.13	26.96	27.25	0.557	1.51
4	30.72	30.85	30.66	30.62	0.172	0.56
4	30.60	30.55	30.34	30.02	0.172	0.50
5	-	-	-			
J	-	-	-	-	-	-
PC	27.98	27.52	28.00	27.04	0.205	0.73
PC	27.94	27.91	28.12	27.91	0.205	0.75
Mean						0.95

SD = standard deviation; CV = coefficient of variation, - = no C<sub>T</sub>

Table 45. Robustness testing of the virotype BTV pan/8 2.0 RT-PCR Kit (**BTV-8** / Cy5) for different annealing temperatures used in the RT-qPCR. The tests were performed on the Agilent Technologies Aria

			BTV-8 (Cy5) C	т		
Sample	Anne	aling temper	ature	Mean	SD	CV%
-	59°C	60°C	61°C	_		
1	19.89	20.16	19.92	19.99	0.151	0.76
1	20.03	19.79	20.16	19.99	0.131	0.70
2	22.96	22.95	23.15	22.96	0.193	0.84
	22.90	22.64	23.17	22.96	0.195	0.04
2	26.04	25.76	26.14	25.99	0.169	0.65
3	26.17	25.81	26.02	23.99	0.109	0.03
4	29.16	29.27	29.26	29.31	0.169	0.58
4	29.14	29.56	29.47	29.51	0.109	0.58
5	-	-	-	_		
J	-	-	-	_	-	-
PC	26.51	26.71	26.88	26.77	0.162	0.61
	26.70	26.97	26.83	20.77	0.102	0.01
Mean				1		0.69

SD = standard deviation;  $CV = coefficient of variation, - = no C_T$ 

Table 46. Robustness testing of the virotype BTV pan/8 2.0 RT-PCR Kit (**endogenous Internal Control** / JOE) for different annealing temperatures used in the RT-qPCR. The tests were performed on the Agilent Technologies Aria Mx instrument.

		Endogenous	s Internal Cor	ntrol (JOE) C <sub>T</sub>		
Sample	Anne	aling temper	ature	Mean	SD	CV%
-	59°C	60°C	61°C	-		
1	24.33	24.92	24.79	24.67	0.200	0.81
'	24.60	24.71	24.68	24.67	0.200	0.61
2	24.33	24.73	24.79	24.63	0.161	0.65
2	24.60	24.64	24.68	24.03	0.101	0.05
3	24.53	24.37	24.86	24.53	0.175	0.71
	24.50	24.40	24.52	24.33	0.175	0.71
4	24.52	24.50	24.62	24.64	0.181	0.73
4	24.94	24.78	24.50	24.04	0.101	0.73
5	18.66	18.56	18.34	18.47	0.195	1.06
5	18.40	18.67	18.18	10.47	0.155	1.00
PC	27.23	27.23	28.95	27.79	1.009	3.63
FC	26.95	27.16	29.21	27.73	1.009	3.03
Mean					· '	1.27

SD = standard deviation; CV = coefficient of variation