# QCMD 2020 Mycobacterium tuberculosis DNA EQA Programme



Catalogue Code: QAB014129

Ref Code: MTBDNA20 Challenge: S **Analysis Type:**Qualitative

Dataset:

Report UID: -/-/2921

Laboratory

NOTE: Summary information only.

### **Intended Results / Panel Composition**

Sample Code	Sample Content	Matrix	Sample Relationships <sup>[1]</sup>	Detection Frequency <sup>[2]</sup>	Sample Status <sup>[3]</sup>	Percentage Correct (All) [4]	
						(%)	(n)
MTBDNA20S-01	M. xenopi	Synthetic Sputum		Negative	CORE	89.7	107
MTBDNA20S-02	Mycobacterium Negative	Synthetic Sputum		Negative	CORE	97.2	107
MTBDNA20S-03	Mycobacterium Negative	Synthetic CSF		Negative	CORE	97.2	107
MTBDNA20S-04	M. bovis (BCG)	Synthetic Sputum	DS2_2	Frequently Detected	CORE	96.3	107
MTBDNA20S-05	M. bovis (BCG)	Synthetic CSF	DS1_3	Frequently Detected	CORE	98.1	107
MTBDNA20S-06	M. bovis (BCG)	Synthetic CSF	DS1_2	Frequently Detected	CORE	98.1	107
MTBDNA20S-07	M. tuberculosis (DR- RIF-R)	Synthetic CSF		Frequently Detected	CORE	99.1	107
MTBDNA20S-08	M. tuberculosis (DR- INH-R)	Synthetic Sputum		Frequently Detected	CORE	99.1	107
MTBDNA20S-09	M. bovis (BCG)	Synthetic Sputum	DS2_1	Frequently Detected	CORE	98.1	107
MTBDNA20S-10	M. bovis (BCG)	Synthetic CSF	DS1_1	Frequently Detected	CORE	95.3	107

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[1] **Sample Relationships:** Indicates the relationships of the samples within this challenge. The highest titre member of dilution series DS1 is indicated by DS1\_1 and further members of the series as DS1\_2, DS1\_3 etc. in order of reducing titre. Additional dilution series are indicated by DS2 (e.g DS2\_1, DS2\_2 etc.), DS3 (e.g. DS3\_1, DS3\_2 etc.). If one duplicate pair is present this is indicated by 'D1'. Further duplicate pairs are indicated by 'D2', 'D3' etc. [2] **Detection Frequency:** To aid qualitative analysis each panel member is assigned a frequency of detection. This is based on the peer group consensus of all qualitative results returned from participants within the EQA challenge / distribution.

[3] **Sample Status:** EQA samples are defined as "CORE" or "EDUCATIONAL". Core proficiency samples are reviewed by the QCMD Scientific Expert(s). This is on the basis of scientific information, clinical relevance, current literature and, where appropriate, professional clinical guidelines. Participating laboratories are expected to report core proficiency samples correctly within the EQA challenge / distribution.

[4] **Percentage Correct (All):** Percentage of datasets (%) reporting the correct qualitative result and the total number of datasets (n) reported for each panel member.

For further details please refer to the current participant manual.

### **Your Summary Results**

EQA Assessment Group [1]	N/A
Core Panel Detection (Qualitative) Score [2]	N/A

#### **Core Panel Members Results**

Sample Code	Qualitative Results		Your Quantitative Data (for information only) [3]			
	Percentage Correct (All) [4]	Your Result	Detection Score	Reported Value	Unitage	Cycle Threshold
MTBDNA20S-01	89.7	-	-	-	-	-
MTBDNA20S-02	97.2	-	-	-	-	-
MTBDNA20S-03	97.2	-	-	-	-	-
MTBDNA20S-04	96.3	-	-	-	-	-
MTBDNA20S-05	98.1	-	-	-	-	-
MTBDNA20S-06	98.1	-	-	-	-	-
MTBDNA20S-07	99.1	-	-	-	-	-
MTBDNA20S-08	99.1	-	-	-	-	-
MTBDNA20S-09	98.1	-	-	-	-	-
MTBDNA20S-10	95.3	-	-	-	-	-

# QCMD 2020 Mycobacterium tuberculosis DNA EQA Programme



Catalogue Code: QAB014129

Ref Code: MTBDNA20 Challenge:

Analysis Type: Qualitative Dataset:

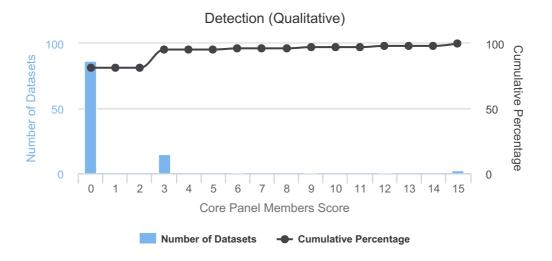
**Report UID:** -/-/2921

Laboratory

- [1] **EQA Assessment Group:** To aid data analysis, participant results are grouped according to the molecular amplification/detection method specified within their molecular workflow for this challenge / distribution. For further details refer to the *Additional Information: Individual Panel Member Analysis* section of this report.
- [2] Core Panel Detection (Qualitative) Score: An overall core panel detection score provided per challenge / distribution.
- [3] **Quantitative Data (for information only):** This is the quantitative value, unitage and cycle threshold you provided when you submitted your results. For qualitative programmes this information is not used as part of your formal EQA assessment.
- [4] Percentage Correct (All): Percentage of datasets (%) reporting the correct qualitative results for each panel member.
- [5] Your Result: The qualitative result you reported for each sample within this EQA challenge / distribution.
- [6] **Detection Score:** Your detection (qualitative) scores are based on the assigned detection frequency of each panel members, where 0 (zero) is "highly satisfactory" and 3 (three) is "highly unsatisfactory". Scores are provided for individual panel members.

For further details please refer to the current participant manual.

### **Core Panel Member Score Breakdown**



Core Panel Member Score Breakdown - Detection: This figure gives you a breakdown of the qualitative detection scores for all qualitative datasets returned within this EQA challenge / distribution independent of the EQA assessment group. Panel detection scores are generated from only those panel members that are defined as "CORE".

For further details please refer to the current participant manual.

#### **Further Programme Details**

Number of Participants	112
Number of Countries	25
Number of Respondents	95
Number of Datasets Submitted	107
Qualitative Results Returned	107 (100.0%)

# QCMD 2020 Mycobacterium tuberculosis DNA EQA Programme



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Ref Code: MTBDNA20 Challenge:

**Analysis Type:**Qualitative

Dataset:

**Report UID:** -/-/2921

Laboratory

### **EQA Programme Aims**

To assess the proficiency of laboratories in the molecular detection of Mycobacterium tuberculosis (M. bovis - BCG).

### Feedback and Enquiries

Participants are encouraged to read the QCMD Participants' Manual, which can be downloaded from the QCMD website.

Any enquiries should be submitted through the 'Contact Us' form that you can find in the 'Help' section of your QCMD (ITEMS) Participant Profile Area.

# QCMD 2020 Mycobacterium tuberculosis DNA EQA Programme



Catalogue Code: QAB014129

Ref Code: MTBDNA20 Challenge:

**Analysis Type:**Qualitative

Dataset:

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Laboratory

Panel member analysis is separated into CORE samples followed by EDUCATIONAL samples.

### **Additional Core Samples Information**

The following section has been categorised as shown below:

Core ► Qualitative

#### **Individual Panel Member Analysis (Qualitative)**

Qualitative analysis for each panel member is provided in relation to your EQA assessment group. EQA assessment groups are established using the molecular workflow information reported by all participants within this EQA challenge / distribution. The principal level of assessment is at the individual method level which is defined based on your reported "amplification/detection method" and other laboratories using the same or similar amplification/detection methods.

To allow meaningful assessment at the individual method level the EQA assessment group must consist of 5 or more datasets. If there are not sufficient datasets at the individual method level then your results will be included within a higher EQA assessment group based on whether it is a commercial or in house technology/method. The highest level assessment grouping is "All" participant reported qualitative results.

A breakdown of qualitative results reported by participants on each of the panel members within this EQA challenge / distribution is provided below. You can compare your results to those within your EQA assessment group and those obtained within other EQA assessment groups or to the overall consensus for each sample within this EQA challenge / distribution.

# QCMD 2020 Mycobacterium tuberculosis DNA EQA Programme



Catalogue Code: QAB014129

Ref Code: MTBDNA20 Challenge: S **Analysis Type:**Qualitative

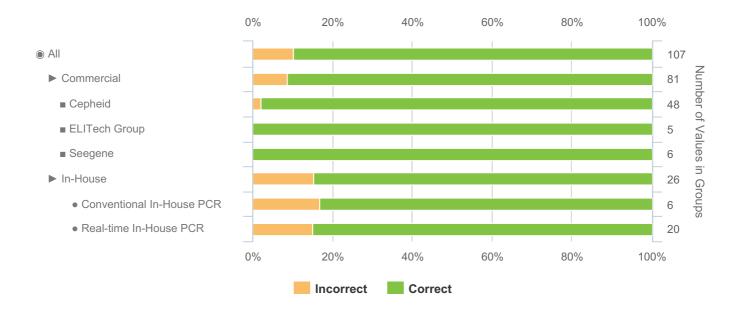
Dataset:

Report UID: -/-/2921

Laboratory

#### MTBDNA20S-01 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (All)		
						(%)	(n)	
MTBDNA20S-01	M. xenopi	Synthetic Sputum		Negative	CORE	89.7	107	



Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AnDiaTec (n=1), AnDiaTec - AnDiaTec Real Time PCR (n=1), BD Molecular Diagnostics (n=1), BD Molecular Diagnostics - BD MAX (n=1), BIORON (n=1), BIORON - BIORON RealLine (n=1), Certest (n=2), Certest - Certest Real Time PCR (n=2), Diagenode (n=2), Diagenode - Diagenode Real Time kit (n=2), Elisabeth Pharmacon (n=1), Elisabeth Pharmacon - Elisabeth Pharmacon EliGene (n=1), GeneProof (n=1), GeneProof - GeneProof Real Time PCR kit (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience GenoType (n=1), PathoFinder (n=3), PathoFinder - PathoFinder Real Time PCR (n=3), Roche (n=4), Roche - Roche Cobas Amplicor (n=1), Roche - Roche Cobas TaqMan (n=3), TIB MOLBIOL (n=2), TIB MOLBIOL - TIB-MolBiol LightMix (n=2), Vitassay (n=2), Vitassay - Vitassay Real-Time PCR (n=2)

# QCMD 2020 Mycobacterium tuberculosis DNA EQA Programme



Catalogue Code: QAB014129

Ref Code: MTBDNA20 Challenge:

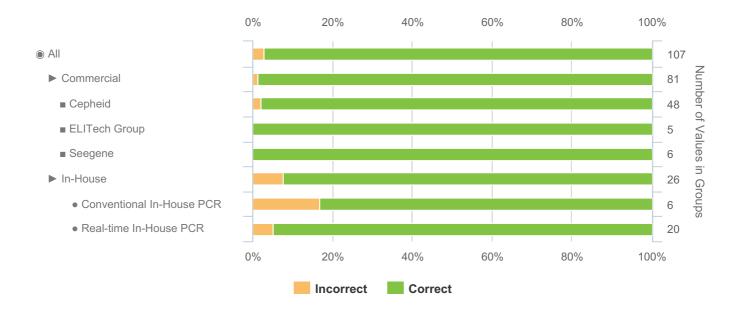
Analysis Type: Qualitative Dataset:

**Report UID:** -/-/2921

Laboratory

#### MTBDNA20S-02 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (All)	
						(%)	(n)
MTBDNA20S-02	Mycobacterium Negative	Synthetic Sputum		Negative	CORE	97.2	107



Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AnDiaTec (n=1), AnDiaTec - AnDiaTec Real Time PCR (n=1), BD Molecular Diagnostics (n=1), BD Molecular Diagnostics - BD MAX (n=1), BIORON (n=1), BIORON - BIORON RealLine (n=1), Certest (n=2), Certest - Certest Real Time PCR (n=2), Diagenode (n=2), Diagenode - Diagenode Real Time kit (n=2), Elisabeth Pharmacon (n=1), Elisabeth Pharmacon - Elisabeth Pharmacon EliGene (n=1), GeneProof (n=1), GeneProof - GeneProof Real Time PCR kit (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience GenoType (n=1), PathoFinder (n=3), PathoFinder - PathoFinder Real Time PCR (n=3), Roche (n=4), Roche - Roche Cobas Amplicor (n=1), Roche - Roche Cobas TaqMan (n=3), TIB MOLBIOL (n=2), TIB MOLBIOL - TIB-MolBiol LightMix (n=2), Vitassay (n=2), Vitassay - Vitassay Real-Time PCR (n=2)

# QCMD 2020 Mycobacterium tuberculosis DNA EQA Programme



Catalogue Code: QAB014129

Ref Code: MTBDNA20 Challenge:

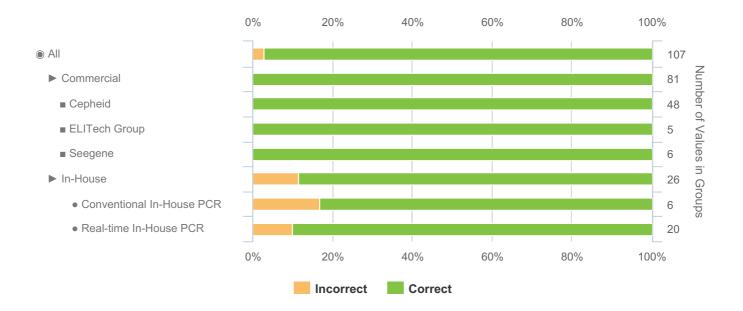
Analysis Type: Qualitative Dataset:

**Report UID:** -/-/2921

Laboratory

#### MTBDNA20S-03 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentag Correct (A	•	
						(%)	(n)	
MTBDNA20S-03	Mycobacterium Negative	Synthetic CSF		Negative	CORE	97.2	107	



Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AnDiaTec (n=1), AnDiaTec - AnDiaTec Real Time PCR (n=1), BD Molecular Diagnostics (n=1), BD Molecular Diagnostics - BD MAX (n=1), BIORON (n=1), BIORON - BIORON RealLine (n=1), Certest (n=2), Certest - Certest Real Time PCR (n=2), Diagenode (n=2), Diagenode - Diagenode Real Time kit (n=2), Elisabeth Pharmacon (n=1), Elisabeth Pharmacon - Elisabeth Pharmacon EliGene (n=1), GeneProof (n=1), GeneProof - GeneProof Real Time PCR kit (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience GenoType (n=1), PathoFinder (n=3), PathoFinder - PathoFinder Real Time PCR (n=3), Roche (n=4), Roche - Roche Cobas Amplicor (n=1), Roche - Roche Cobas TaqMan (n=3), TIB MOLBIOL (n=2), TIB MOLBIOL - TIB-MolBiol LightMix (n=2), Vitassay (n=2), Vitassay - Vitassay Real-Time PCR (n=2)

# QCMD 2020 Mycobacterium tuberculosis DNA EQA Programme



Catalogue Code: QAB014129

Ref Code: MTBDNA20 Challenge:

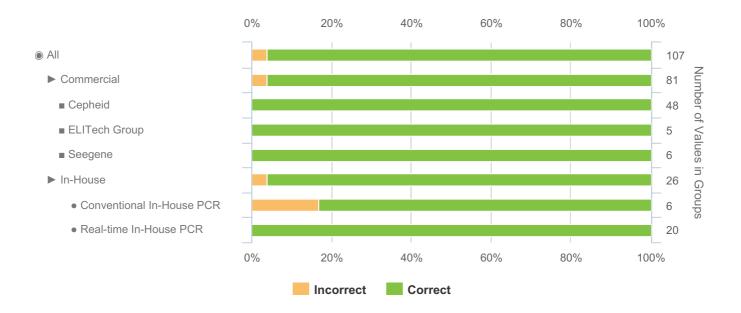
Analysis Type: Qualitative Dataset:

**Report UID:** -/-/2921

Laboratory

#### MTBDNA20S-04 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (AII)		
						(%)	(n)	
MTBDNA20S-04	M. bovis (BCG)	Synthetic Sputum	DS2_2	Frequently Detected	CORE	96.3	107	



Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AnDiaTec (n=1), AnDiaTec - AnDiaTec Real Time PCR (n=1), BD Molecular Diagnostics (n=1), BD Molecular Diagnostics - BD MAX (n=1), BIORON (n=1), BIORON - BIORON RealLine (n=1), Certest (n=2), Certest - Certest Real Time PCR (n=2), Diagenode (n=2), Diagenode - Diagenode Real Time kit (n=2), Elisabeth Pharmacon (n=1), Elisabeth Pharmacon - Elisabeth Pharmacon EliGene (n=1), GeneProof (n=1), GeneProof - GeneProof Real Time PCR kit (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience GenoType (n=1), PathoFinder (n=3), PathoFinder - PathoFinder Real Time PCR (n=3), Roche (n=4), Roche - Roche Cobas Amplicor (n=1), Roche - Roche Cobas TaqMan (n=3), TIB MOLBIOL (n=2), TIB MOLBIOL - TIB-MolBiol LightMix (n=2), Vitassay (n=2), Vitassay - Vitassay Real-Time PCR (n=2)

# QCMD 2020 Mycobacterium tuberculosis DNA EQA Programme



Catalogue Code: QAB014129

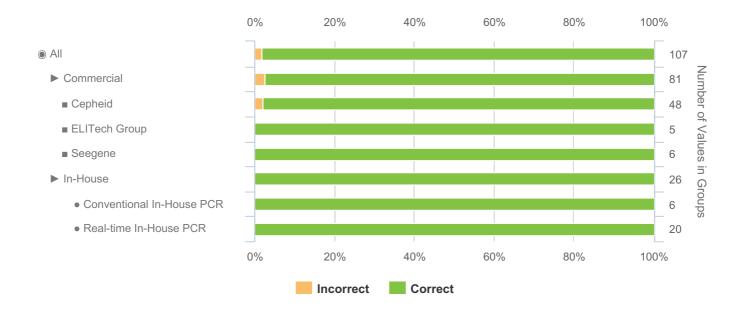
Ref Code: MTBDNA20 Challenge: S Analysis Type: Qualitative Dataset:

**Report UID:** -/-/2921

Laboratory

#### MTBDNA20S-05 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (AII)	
						(%)	(n)
MTBDNA20S-05	M. bovis (BCG)	Synthetic CSF	DS1_3	Frequently Detected	CORE	98.1	107



Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AnDiaTec (n=1), AnDiaTec - AnDiaTec Real Time PCR (n=1), BD Molecular Diagnostics (n=1), BD Molecular Diagnostics - BD MAX (n=1), BIORON (n=1), BIORON - BIORON RealLine (n=1), Certest (n=2), Certest - Certest Real Time PCR (n=2), Diagenode (n=2), Diagenode - Diagenode Real Time kit (n=2), Elisabeth Pharmacon (n=1), Elisabeth Pharmacon - Elisabeth Pharmacon EliGene (n=1), GeneProof (n=1), GeneProof - GeneProof Real Time PCR kit (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience GenoType (n=1), PathoFinder (n=3), PathoFinder - PathoFinder Real Time PCR (n=3), Roche (n=4), Roche - Roche Cobas Amplicor (n=1), Roche - Roche Cobas TaqMan (n=3), TIB MOLBIOL (n=2), TIB MOLBIOL - TIB-MolBiol LightMix (n=2), Vitassay (n=2), Vitassay - Vitassay Real-Time PCR (n=2)

# QCMD 2020 Mycobacterium tuberculosis DNA EQA Programme



Catalogue Code: QAB014129

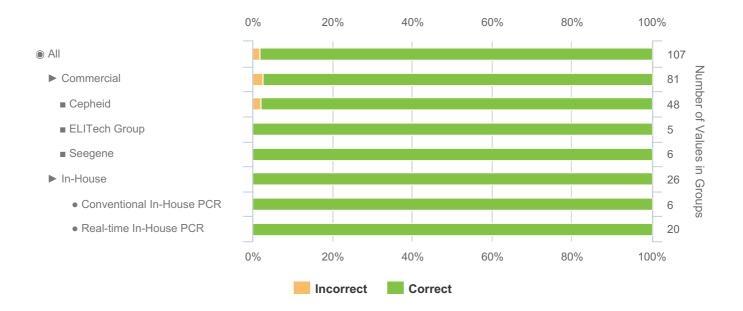
Ref Code: MTBDNA20 Challenge: S Analysis Type: Qualitative Dataset:

Report UID: -/-/2921

Laboratory

#### MTBDNA20S-06 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (AII)		
						(%)	(n)	
MTBDNA20S-06	M. bovis (BCG)	Synthetic CSF	DS1_2	Frequently Detected	CORE	98.1	107	



Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AnDiaTec (n=1), AnDiaTec - AnDiaTec Real Time PCR (n=1), BD Molecular Diagnostics (n=1), BD Molecular Diagnostics - BD MAX (n=1), BIORON (n=1), BIORON - BIORON RealLine (n=1), Certest (n=2), Certest - Certest Real Time PCR (n=2), Diagenode (n=2), Diagenode - Diagenode Real Time kit (n=2), Elisabeth Pharmacon (n=1), Elisabeth Pharmacon - Elisabeth Pharmacon EliGene (n=1), GeneProof (n=1), GeneProof - GeneProof Real Time PCR kit (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience GenoType (n=1), PathoFinder (n=3), PathoFinder - PathoFinder Real Time PCR (n=3), Roche (n=4), Roche - Roche Cobas Amplicor (n=1), Roche - Roche Cobas TaqMan (n=3), TIB MOLBIOL (n=2), TIB MOLBIOL - TIB-MolBiol LightMix (n=2), Vitassay (n=2), Vitassay - Vitassay Real-Time PCR (n=2)

# QCMD 2020 Mycobacterium tuberculosis DNA EQA Programme



Catalogue Code: QAB014129

Ref Code: MTBDNA20 Challenge:

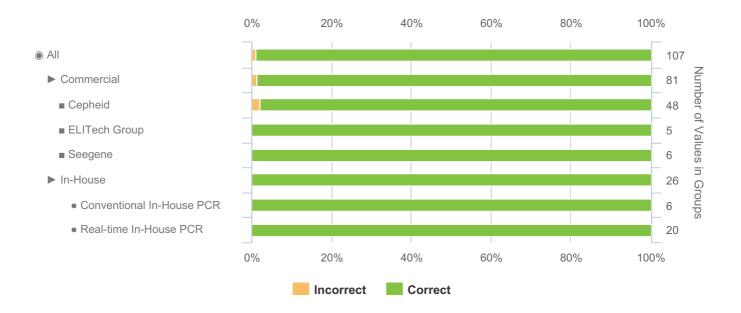
Analysis Type: Qualitative Dataset:

**Report UID:** -/-/2921

Laboratory

#### MTBDNA20S-07 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (All)	
						(%)	(n)
MTBDNA20S-07	M. tuberculosis (DR-RIF-R)	Synthetic CSF		Frequently Detected	CORE	99.1	107



Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AnDiaTec (n=1), AnDiaTec - AnDiaTec Real Time PCR (n=1), BD Molecular Diagnostics (n=1), BD Molecular Diagnostics - BD MAX (n=1), BIORON (n=1), BIORON - BIORON RealLine (n=1), Certest (n=2), Certest - Certest Real Time PCR (n=2), Diagenode (n=2), Diagenode - Diagenode Real Time kit (n=2), Elisabeth Pharmacon (n=1), Elisabeth Pharmacon - Elisabeth Pharmacon EliGene (n=1), GeneProof (n=1), GeneProof - GeneProof Real Time PCR kit (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience GenoType (n=1), PathoFinder (n=3), PathoFinder - PathoFinder Real Time PCR (n=3), Roche (n=4), Roche - Roche Cobas Amplicor (n=1), Roche - Roche Cobas TaqMan (n=3), TIB MOLBIOL (n=2), TIB MOLBIOL - TIB-MolBiol LightMix (n=2), Vitassay (n=2), Vitassay - Vitassay Real-Time PCR (n=2)

# QCMD 2020 Mycobacterium tuberculosis DNA EQA Programme



Catalogue Code: QAB014129

Ref Code: MTBDNA20 Challenge:

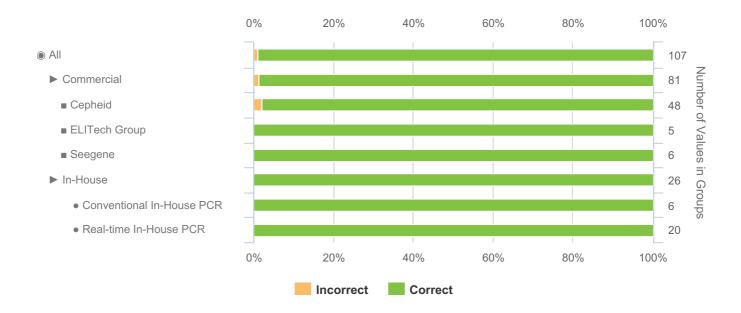
Analysis Type: Qualitative Dataset:

**Report UID:** -/-/2921

Laboratory

#### MTBDNA20S-08 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (All)	
						(%)	(n)
MTBDNA20S-08	M. tuberculosis (DR-INH-R)	Synthetic Sputum		Frequently Detected	CORE	99.1	107



Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AnDiaTec (n=1), AnDiaTec - AnDiaTec Real Time PCR (n=1), BD Molecular Diagnostics (n=1), BD Molecular Diagnostics - BD MAX (n=1), BIORON (n=1), BIORON - BIORON RealLine (n=1), Certest (n=2), Certest - Certest Real Time PCR (n=2), Diagenode (n=2), Diagenode - Diagenode Real Time kit (n=2), Elisabeth Pharmacon (n=1), Elisabeth Pharmacon - Elisabeth Pharmacon EliGene (n=1), GeneProof (n=1), GeneProof - GeneProof Real Time PCR kit (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience GenoType (n=1), PathoFinder (n=3), PathoFinder - PathoFinder Real Time PCR (n=3), Roche (n=4), Roche - Roche Cobas Amplicor (n=1), Roche - Roche Cobas TaqMan (n=3), TIB MOLBIOL (n=2), TIB MOLBIOL - TIB-MolBiol LightMix (n=2), Vitassay (n=2), Vitassay - Vitassay Real-Time PCR (n=2)

# QCMD 2020 Mycobacterium tuberculosis DNA EQA Programme



Catalogue Code: QAB014129

Ref Code: MTBDNA20 Challenge:

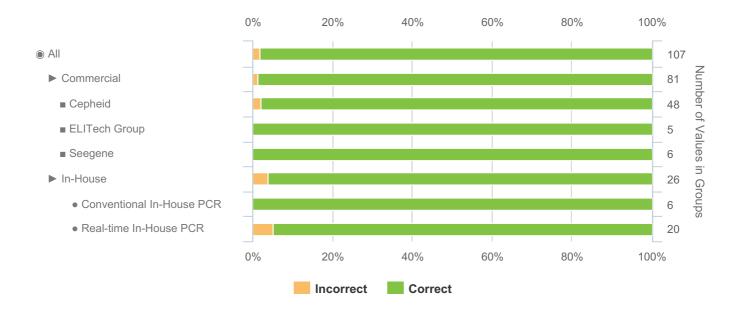
Analysis Type: Qualitative Dataset:

Report UID: -/-/2921

Laboratory

#### MTBDNA20S-09 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (All)	
						(%)	(n)
MTBDNA20S-09	M. bovis (BCG)	Synthetic Sputum	DS2_1	Frequently Detected	CORE	98.1	107



Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AnDiaTec (n=1), AnDiaTec - AnDiaTec Real Time PCR (n=1), BD Molecular Diagnostics (n=1), BD Molecular Diagnostics - BD MAX (n=1), BIORON (n=1), BIORON - BIORON RealLine (n=1), Certest (n=2), Certest - Certest Real Time PCR (n=2), Diagenode (n=2), Diagenode - Diagenode Real Time kit (n=2), Elisabeth Pharmacon (n=1), Elisabeth Pharmacon - Elisabeth Pharmacon EliGene (n=1), GeneProof (n=1), GeneProof - GeneProof Real Time PCR kit (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience GenoType (n=1), PathoFinder (n=3), PathoFinder - PathoFinder Real Time PCR (n=3), Roche (n=4), Roche - Roche Cobas Amplicor (n=1), Roche - Roche Cobas TaqMan (n=3), TIB MOLBIOL (n=2), TIB MOLBIOL - TIB-MolBiol LightMix (n=2), Vitassay (n=2), Vitassay - Vitassay Real-Time PCR (n=2)

# QCMD 2020 Mycobacterium tuberculosis DNA EQA Programme



Catalogue Code: QAB014129

Ref Code: MTBDNA20 Challenge:

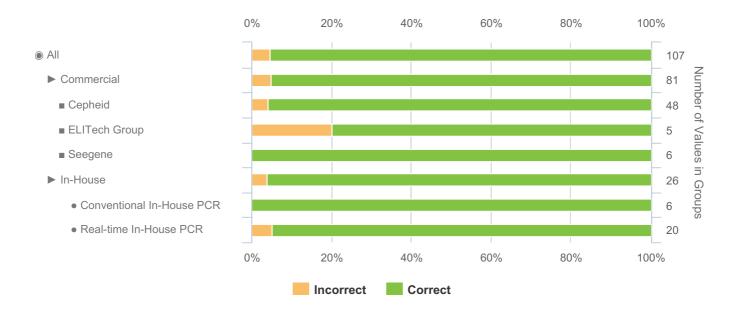
Analysis Type: Qualitative Dataset:

Report UID: -/-/2921

Laboratory

#### MTBDNA20S-10 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (All)	
						(%)	(n)
MTBDNA20S-10	M. bovis (BCG)	Synthetic CSF	DS1_1	Frequently Detected	CORE	95.3	107



Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AnDiaTec (n=1), AnDiaTec - AnDiaTec Real Time PCR (n=1), BD Molecular Diagnostics (n=1), BD Molecular Diagnostics - BD MAX (n=1), BIORON (n=1), BIORON - BIORON RealLine (n=1), Certest (n=2), Certest - Certest Real Time PCR (n=2), Diagenode (n=2), Diagenode - Diagenode Real Time kit (n=2), Elisabeth Pharmacon (n=1), Elisabeth Pharmacon - Elisabeth Pharmacon EliGene (n=1), GeneProof (n=1), GeneProof - GeneProof Real Time PCR kit (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience GenoType (n=1), PathoFinder (n=3), PathoFinder - PathoFinder Real Time PCR (n=3), Roche (n=4), Roche - Roche Cobas Amplicor (n=1), Roche - Roche Cobas TaqMan (n=3), TIB MOLBIOL (n=2), TIB MOLBIOL - TIB-MolBiol LightMix (n=2), Vitassay (n=2), Vitassay - Vitassay Real-Time PCR (n=2)

# QCMD 2020 Mycobacterium tuberculosis DNA EQA Programme



Catalogue Code: QAB014129

Ref Code: MTBDNA20 Challenge:

**Analysis Type:**Qualitative

Dataset:

**Report UID:** -/-/2921

Laboratory

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