

# Individual Report

## QCMD 2019 Chlamydia trachomatis and Neisseria gonorrhoeae DNA EQA Programme (CT)



<b>Catalogue Code:</b> QAB174191	<b>Ref Code:</b> CTNg19	<b>Challenge:</b> S	<b>Analysis Type:</b> Qualitative	<b>Dataset:</b> 304097	<b>Report UID:</b> 4254/304097/2141	<b>Laboratory</b> DE182
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### 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) <sup>[4]</sup>	
						(%)	(n)
CTNg19S-01	C. trachomatis (Genovar F)	Urine		Frequently Detected	CORE	95.8	71
CTNg19S-02	C. trachomatis (LGV)	Simulated Swab		Frequently Detected	CORE	98.6	71
CTNg19S-03	N. gonorrhoeae (St 49226)	Urine	DS2_1	Negative	EDUCATIONAL	100.0	71
CTNg19S-04	N. gonorrhoeae (St 49226)	Urine	DS2_2	Negative	EDUCATIONAL	100.0	71
CTNg19S-05	C. trachomatis (LGV)	Urine	DS1_1	Frequently Detected	CORE	98.6	71
CTNg19S-06	N. gonorrhoeae (St 49226)	Simulated Swab	D1	Negative	EDUCATIONAL	98.6	71
CTNg19S-07	Negative	Simulated Swab		Negative	CORE	100.0	71
CTNg19S-08	Negative	Urine		Negative	CORE	100.0	71
CTNg19S-09	N. gonorrhoeae (St 49226)	Simulated Swab	D1	Negative	EDUCATIONAL	100.0	71
CTNg19S-10	C. trachomatis (LGV)	Urine	DS1_2	Frequently Detected	CORE	98.6	71

[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.*

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### Your Summary Results

EQA Assessment Group <sup>[1]</sup>

Commercial

Core Panel Detection (Qualitative) Score <sup>[2]</sup>

0

### Core Panel Members Results

Sample Code	Qualitative Results			Your Quantitative Data (for information only) <sup>[3]</sup>		
	Percentage Correct (All) <sup>[4]</sup>	Your Result <sup>[5]</sup>	Detection Score <sup>[6]</sup>	Reported Value	Unitage	Cycle Threshold
CTNg19S-01	95.8	Positive	0		N/A	40.75
CTNg19S-02	98.6	Positive	0		N/A	37.2
CTNg19S-05	98.6	Positive	0		N/A	35.4
CTNg19S-07	100.0	Negative	0		N/A	-
CTNg19S-08	100.0	Negative	0		N/A	-
CTNg19S-10	98.6	Positive	0		N/A	38.3

**[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.

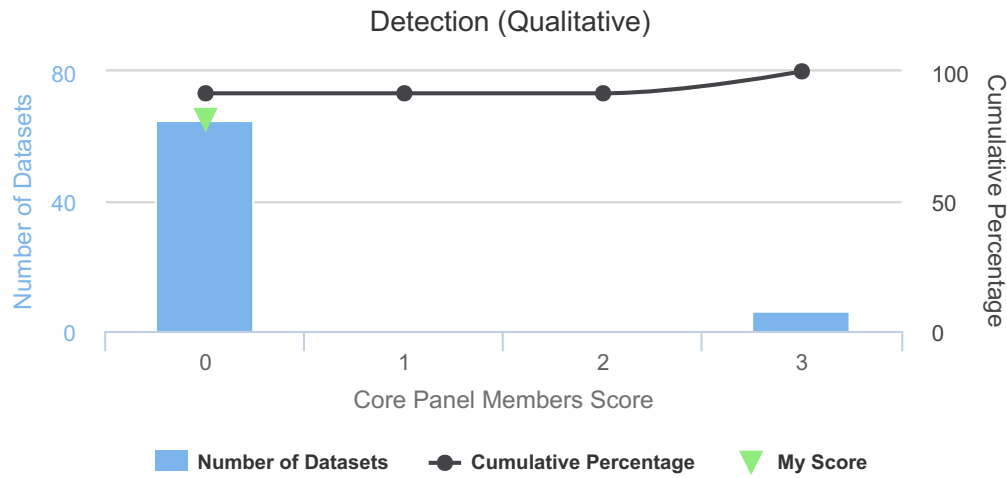
**[5] Your Result:** The qualitative result you reported for each sample within this EQA challenge / distribution.

**[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.**

<b>Individual Report</b>	<b>QCMD 2019 Chlamydia trachomatis and Neisseria gonorrhoeae DNA EQA Programme (CT)</b>				 <b>QCMD</b> <small>Quality Control for Molecular Diagnostics</small>	
<b>Catalogue Code:</b> QAB174191	<b>Ref Code:</b> CTNg19	<b>Challenge:</b> S	<b>Analysis Type:</b> Qualitative	<b>Dataset:</b> 304097	<b>Report UID:</b> 4254/304097/2141	<b>Laboratory</b> DE182

### 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.*

### My Workflow Details

The details of the workflow(s) used to submit your results for this challenge.

<b>Name</b>	CT/NG Workflow double Kit (v3)
<b>Description</b>	Verwendung RealLine Chlamydia trachomatis / Neisseria gonorrhoeae REF VBD0457
<b>Targets</b>	<ul style="list-style-type: none"> <li><span style="color: green; font-weight: bold;">B</span> Neisseria gonorrhoeae</li> <li><span style="color: green; font-weight: bold;">B</span> Chlamydia trachomatis</li> </ul>
<b>Assays</b>	<ul style="list-style-type: none"> <li><span style="color: blue; font-weight: bold;">DNA</span> <i>Extraction</i> - Manual Extraction Process <ul style="list-style-type: none"> <li>● Commercial <ul style="list-style-type: none"> <li>○ Kit Manufacturer: <i>BIORON Diagnostics GmbH</i></li> <li>○ Kit Type: <i>RealLine DNA Express REF VBC8899</i></li> </ul> </li> </ul> </li> <li><span style="color: blue; font-weight: bold;">RNA</span> <i>Amplification</i> - DNA Technology - DTlite Real-Time PCR System <ul style="list-style-type: none"> <li>● Multiplex</li> <li>● Commercial <ul style="list-style-type: none"> <li>○ Kit Manufacturer: <i>BIORON Diagnostics GmbH</i></li> <li>○ Kit Type: <i>RealLine Chlamydia trachomatis / Neisseria gonorrhoeae REF VBD0457</i></li> <li>○ Kit Version: <i>not applicable</i></li> </ul> </li> </ul> </li> </ul>

<b>Individual Report</b>	<b>QCMD 2019 Chlamydia trachomatis and Neisseria gonorrhoeae DNA EQA Programme (CT)</b>				 Quality Control for Molecular Diagnostics	
<b>Catalogue Code:</b> QAB174191	<b>Ref Code:</b> CTNg19	<b>Challenge:</b> S	<b>Analysis Type:</b> Qualitative	<b>Dataset:</b> 304097	<b>Report UID:</b> 4254/304097/2141	<b>Laboratory</b> DE182

## Educational Panel Members Results

Sample Code	Qualitative Results			Your Quantitative Data (for information only) <sup>[1]</sup>		
	Percentage Correct (All) <sup>[2]</sup>	Your Result <sup>[3]</sup>	Detection Score <sup>[4]</sup>	Reported Value	Unitage	Cycle Threshold
CTNg19S-03	100.0	Negative	0		N/A	-
CTNg19S-04	100.0	Negative	0		N/A	-
CTNg19S-06	98.6	Negative	0		N/A	-
CTNg19S-09	100.0	Negative	0		N/A	-

[1] **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.

[2] **Percentage Correct (All):** Percentage of datasets (%) reporting the correct qualitative results for each panel member.

[3] **Your Result:** The qualitative result you reported for each sample within this EQA challenge / distribution.

[4] **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.*

## Further Programme Details


Number of Participants	71
Number of Countries	28
Number of Respondents	66
Number of Datasets Submitted	71
Qualitative Results Returned	71 (100.0%)

## Comments

For the analysis of *Chlamydia trachomatis* results, *Neisseria gonorrhoeae* samples were assigned a negative educational status.

## EQA Programme Aims


To assess proficiency of laboratories in the detection of *Chlamydia trachomatis* and *Neisseria gonorrhoeae* using molecular technologies.

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<b>Catalogue Code:</b> QAB174191	<b>Ref Code:</b> CTNg19	<b>Challenge:</b> S	<b>Analysis Type:</b> Qualitative	<b>Dataset:</b> 304097	<b>Report UID:</b> 4254/304097/2141	<b>Laboratory</b> DE182

### Feedback and Enquiries

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

Any queries about this report should be addressed to the QCMD Neutral Office ([neutraloffice@qcmd.org](mailto:neutraloffice@qcmd.org)).

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<b>Catalogue Code:</b> QAB174191	<b>Ref Code:</b> CTNg19	<b>Challenge:</b> S	<b>Analysis Type:</b> Qualitative	<b>Dataset:</b> 304097	<b>Report UID:</b> 4254/304097/2141	<b>Laboratory</b> DE182

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.

# Individual Report

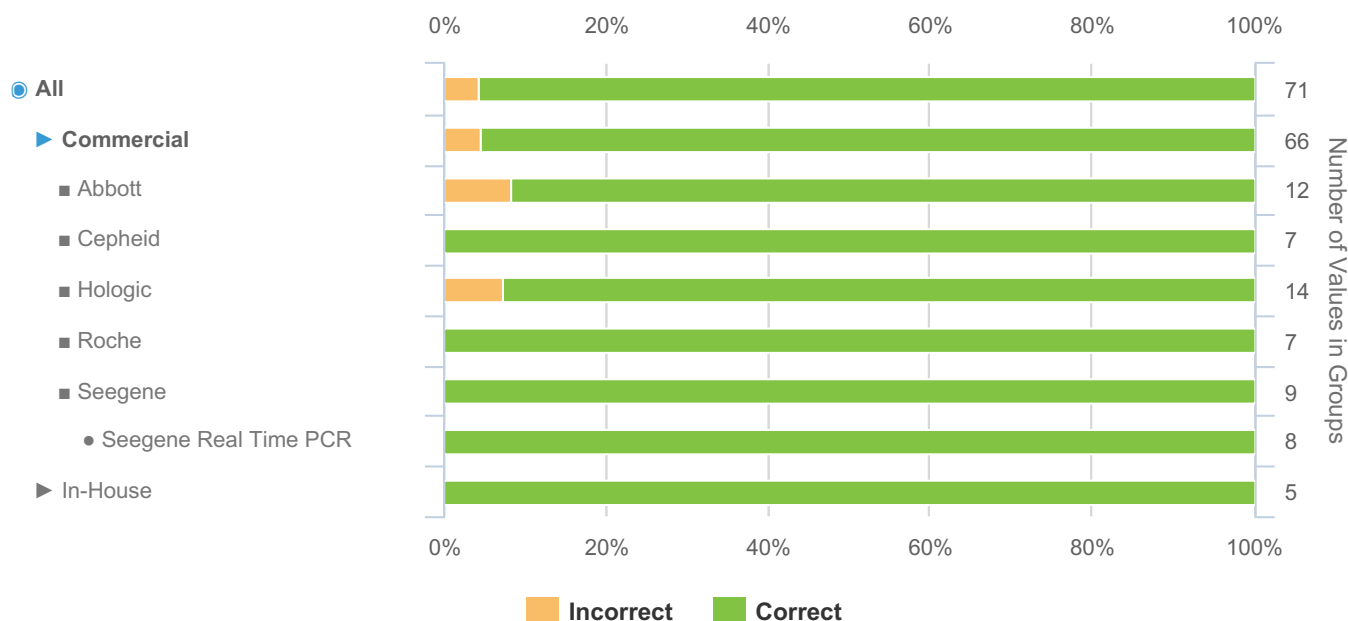
## QCMD 2019 Chlamydia trachomatis and Neisseria gonorrhoeae DNA EQA Programme (CT)



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### CTNg19S-01 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (All)	
						(%)	(n)
CTNg19S-01	C. trachomatis (Genovar F)	Urine		Frequently Detected	CORE	95.8	71



**Groups below n=5:** AmpliSens (n=2), AmpliSens - AmpliSens Real Time PCR (n=2), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=1), BD Molecular Diagnostics - BD ProbeTec (n=2), BIORON (n=2), BIORON - BIORON RealLine (n=2), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), Goffin Molecular Technologies (n=1), Goffin Molecular Technologies - Goffin Presto (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience FlouroType (n=1), Randox (n=1), Randox - Randox Multiplex Array (n=1), Roche - Roche Cobas 4800 (n=4), Roche - Roche Cobas 6800/8800 (n=3), Seegene - Seegene Seeplex (n=1), Siemens (n=1), Siemens - Siemens Versant (n=1), TIB MOLBIOL (n=3), TIB MOLBIOL - TIB MOLBIOL LightMix (n=3)

**Groups Rolled Up:** Abbott - Abbott Real Time PCR (n=12), Cepheid - Cepheid Xpert kit (n=7), Hologic - Hologic Aptima (n=14), In-House - Real-time In-House PCR (n=5)

# Individual Report

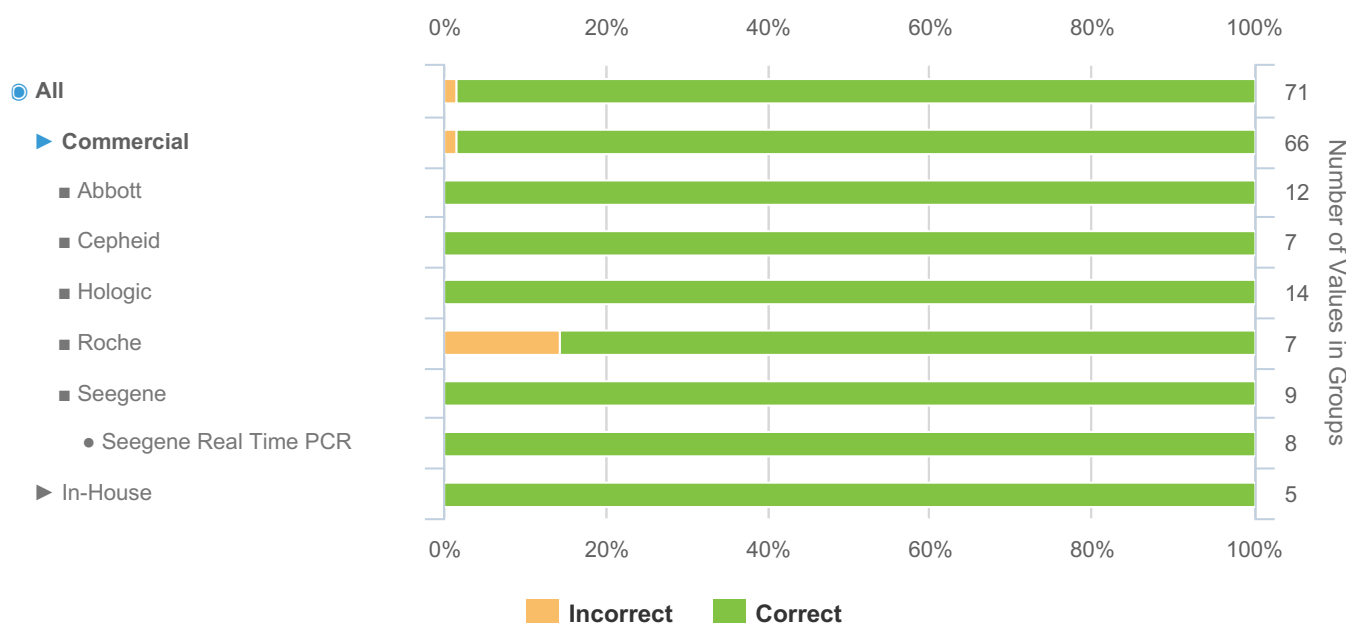
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### CTNg19S-02 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (All)	
						(%)	(n)
CTNg19S-02	C. trachomatis (LGV)	Simulated Swab		Frequently Detected	CORE	98.6	71



**Groups below n=5:** AmpliSens (n=2), AmpliSens - AmpliSens Real Time PCR (n=2), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=1), BD Molecular Diagnostics - BD ProbeTec (n=2), BIORON (n=2), BIORON - BIORON RealLine (n=2), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), Goffin Molecular Technologies (n=1), Goffin Molecular Technologies - Goffin Presto (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience FlouroType (n=1), Randox (n=1), Randox - Randox Multiplex Array (n=1), Roche - Roche Cobas 4800 (n=4), Roche - Roche Cobas 6800/8800 (n=3), Seegene - Seegene Seeplex (n=1), Siemens (n=1), Siemens - Siemens Versant (n=1), TIB MOLBIOL (n=3), TIB MOLBIOL - TIB MOLBIOL LightMix (n=3)

**Groups Rolled Up:** Abbott - Abbott Real Time PCR (n=12), Cepheid - Cepheid Xpert kit (n=7), Hologic - Hologic Aptima (n=14), In-House - Real-time In-House PCR (n=5)



# Individual Report

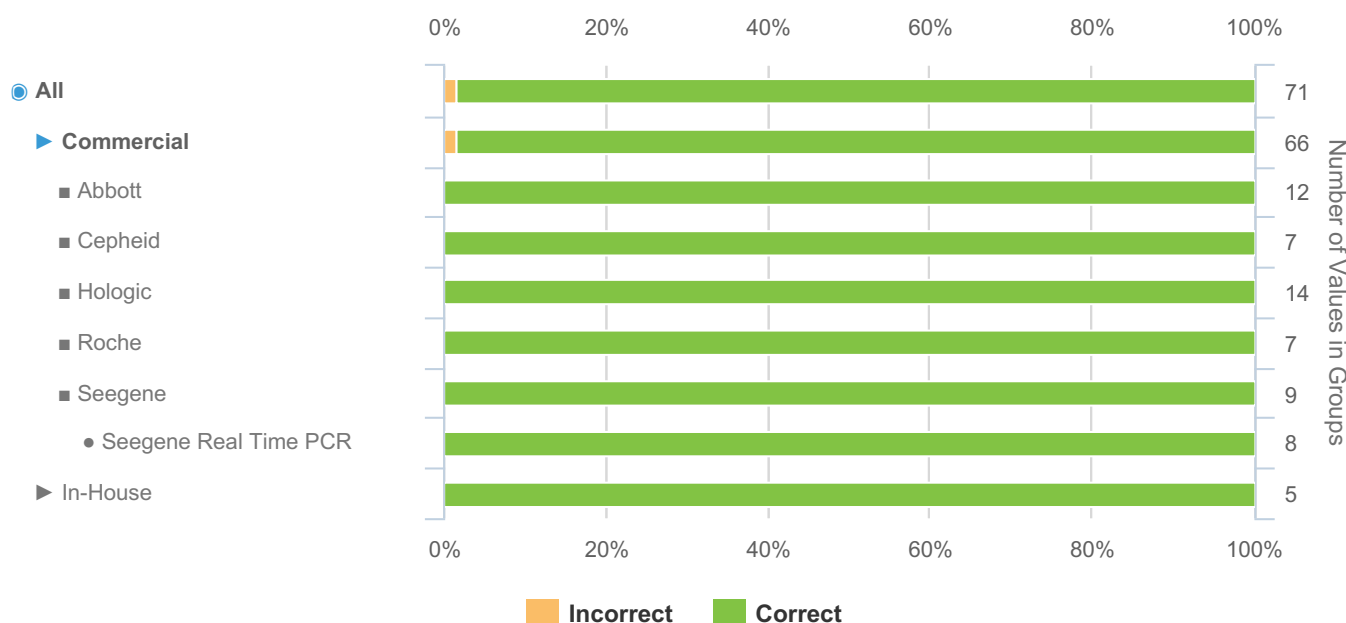
## QCMD 2019 Chlamydia trachomatis and Neisseria gonorrhoeae DNA EQA Programme (CT)



<b>Catalogue Code:</b> QAB174191	<b>Ref Code:</b> CTNg19	<b>Challenge:</b> S	<b>Analysis Type:</b> Qualitative	<b>Dataset:</b> 304097	<b>Report UID:</b> 4254/304097/2141	<b>Laboratory:</b> DE182
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### CTNg19S-05 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (All)	
						(%)	(n)
CTNg19S-05	C. trachomatis (LGV)	Urine	DS1_1	Frequently Detected	CORE	98.6	71



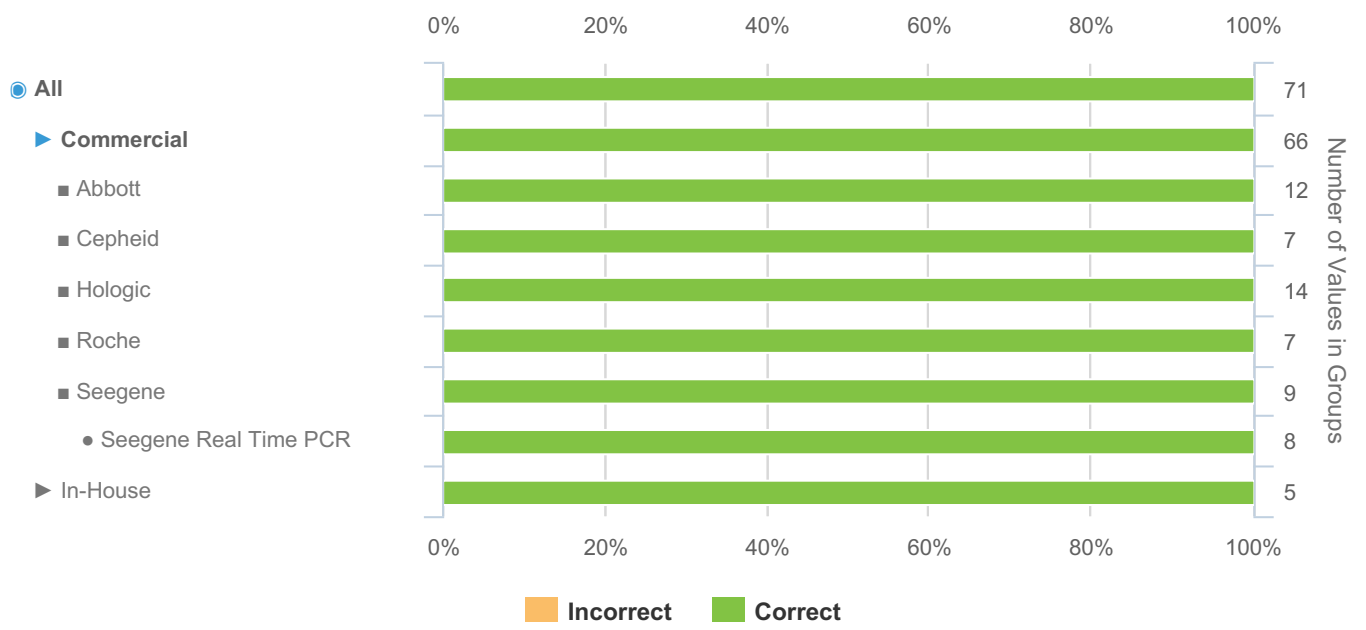
**Groups below n=5:** AmpliSens (n=2), AmpliSens - AmpliSens Real Time PCR (n=2), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=1), BD Molecular Diagnostics - BD ProbeTec (n=2), BIORON (n=2), BIORON - BIORON RealLine (n=2), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), Goffin Molecular Technologies (n=1), Goffin Molecular Technologies - Goffin Presto (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience FlouroType (n=1), Randox (n=1), Randox - Randox Multiplex Array (n=1), Roche - Roche Cobas 4800 (n=4), Roche - Roche Cobas 6800/8800 (n=3), Seegene - Seegene Seeplex (n=1), Siemens (n=1), Siemens - Siemens Versant (n=1), TIB MOLBIOL (n=3), TIB MOLBIOL - TIB MOLBIOL LightMix (n=3)

**Groups Rolled Up:** Abbott - Abbott Real Time PCR (n=12), Cepheid - Cepheid Xpert kit (n=7), Hologic - Hologic Aptima (n=14), In-House - Real-time In-House PCR (n=5)

Catalogue Code: QAB174191	Ref Code: CTNg19	Challenge: S	Analysis Type: Qualitative	Dataset: 304097	Report UID: 4254/304097/2141	Laboratory DE182
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**CTNg19S-07 - Qualitative Results Breakdown**

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (All)	
						(%)	(n)
CTNg19S-07	Negative	Simulated Swab		Negative	CORE	100.0	71



**Groups below n=5:** AmpliSens (n=2), AmpliSens - AmpliSens Real Time PCR (n=2), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=1), BD Molecular Diagnostics - BD ProbeTec (n=2), BIORON (n=2), BIORON - BIORON RealLine (n=2), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), Goffin Molecular Technologies (n=1), Goffin Molecular Technologies - Goffin Presto (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience FlouroType (n=1), Randox (n=1), Randox - Randox Multiplex Array (n=1), Roche - Roche Cobas 4800 (n=4), Roche - Roche Cobas 6800/8800 (n=3), Seegene - Seegene Seeplex (n=1), Siemens (n=1), Siemens - Siemens Versant (n=1), TIB MOLBIOL (n=3), TIB MOLBIOL - TIB MOLBIOL LightMix (n=3)

**Groups Rolled Up:** Abbott - Abbott Real Time PCR (n=12), Cepheid - Cepheid Xpert kit (n=7), Hologic - Hologic Aptima (n=14), In-House - Real-time In-House PCR (n=5)

# Individual Report

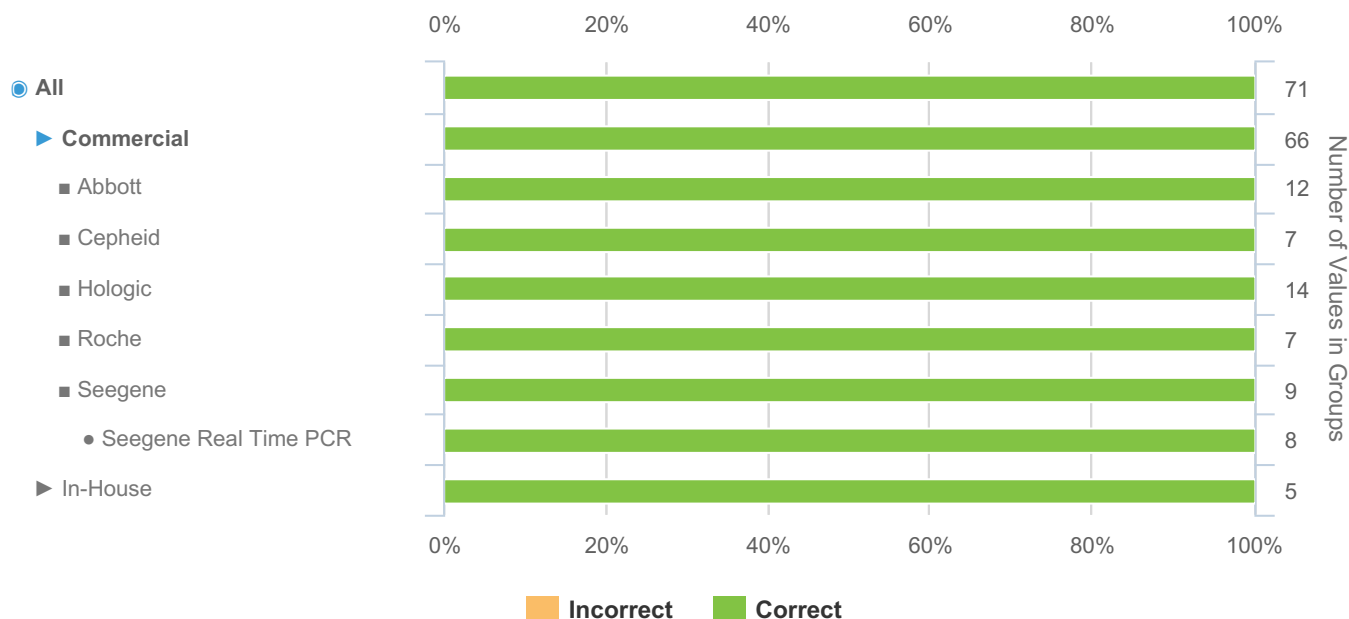
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### CTNg19S-08 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (All)	
						(%)	(n)
CTNg19S-08	Negative	Urine		Negative	CORE	100.0	71



**Groups below n=5:** AmpliSens (n=2), AmpliSens - AmpliSens Real Time PCR (n=2), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=1), BD Molecular Diagnostics - BD ProbeTec (n=2), BIORON (n=2), BIORON - BIORON RealLine (n=2), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), Goffin Molecular Technologies (n=1), Goffin Molecular Technologies - Goffin Presto (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience FlouroType (n=1), Randox (n=1), Randox - Randox Multiplex Array (n=1), Roche - Roche Cobas 4800 (n=4), Roche - Roche Cobas 6800/8800 (n=3), Seegene - Seegene Seeplex (n=1), Siemens (n=1), Siemens - Siemens Versant (n=1), TIB MOLBIOL (n=3), TIB MOLBIOL - TIB MOLBIOL LightMix (n=3)

**Groups Rolled Up:** Abbott - Abbott Real Time PCR (n=12), Cepheid - Cepheid Xpert kit (n=7), Hologic - Hologic Aptima (n=14), In-House - Real-time In-House PCR (n=5)

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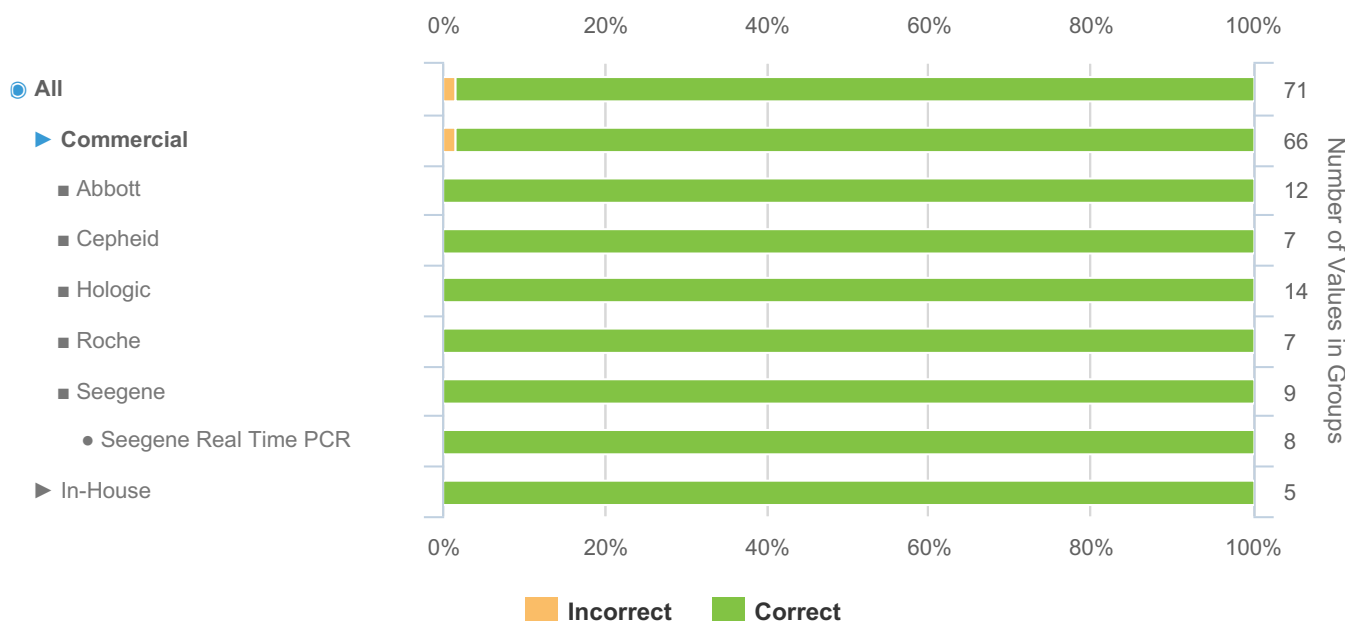
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### CTNg19S-10 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (All)	
						(%)	(n)
CTNg19S-10	C. trachomatis (LGV)	Urine	DS1_2	Frequently Detected	CORE	98.6	71



**Groups below n=5:** AmpliSens (n=2), AmpliSens - AmpliSens Real Time PCR (n=2), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=1), BD Molecular Diagnostics - BD ProbeTec (n=2), BIORON (n=2), BIORON - BIORON RealLine (n=2), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), Goffin Molecular Technologies (n=1), Goffin Molecular Technologies - Goffin Presto (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience FlouroType (n=1), Randox (n=1), Randox - Randox Multiplex Array (n=1), Roche - Roche Cobas 4800 (n=4), Roche - Roche Cobas 6800/8800 (n=3), Seegene - Seegene Seeplex (n=1), Siemens (n=1), Siemens - Siemens Versant (n=1), TIB MOLBIOL (n=3), TIB MOLBIOL - TIB MOLBIOL LightMix (n=3)

**Groups Rolled Up:** Abbott - Abbott Real Time PCR (n=12), Cepheid - Cepheid Xpert kit (n=7), Hologic - Hologic Aptima (n=14), In-House - Real-time In-House PCR (n=5)

<b>Individual Report</b>	<b>QCMD 2019 Chlamydia trachomatis and Neisseria gonorrhoeae DNA EQA Programme (CT)</b>					
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### Additional Educational Samples Information

The following section has been categorised as shown below:

Educational ► 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.

# Individual Report

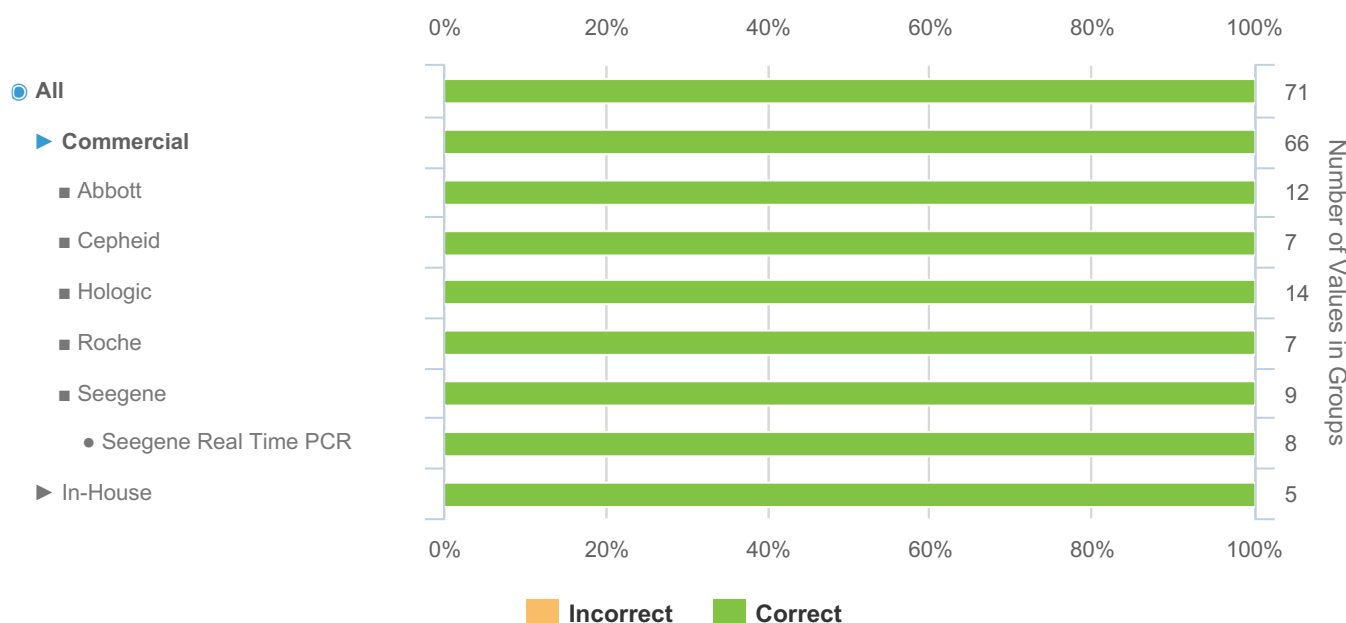
## QCMD 2019 Chlamydia trachomatis and Neisseria gonorrhoeae DNA EQA Programme (CT)



<b>Catalogue Code:</b> QAB174191	<b>Ref Code:</b> CTNg19	<b>Challenge:</b> S	<b>Analysis Type:</b> Qualitative	<b>Dataset:</b> 304097	<b>Report UID:</b> 4254/304097/2141	<b>Laboratory</b> DE182
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### CTNg19S-03 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (All)	
						(%)	(n)
CTNg19S-03	N. gonorrhoeae (St 49226)	Urine	DS2_1	Negative	EDUCATIONAL	100.0	71



**Groups below n=5:** AmpliSens (n=2), AmpliSens - AmpliSens Real Time PCR (n=2), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=1), BD Molecular Diagnostics - BD ProbeTec (n=2), BIORON (n=2), BIORON - BIORON RealLine (n=2), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), Goffin Molecular Technologies (n=1), Goffin Molecular Technologies - Goffin Presto (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience FlouroType (n=1), Randox (n=1), Randox - Randox Multiplex Array (n=1), Roche - Roche Cobas 4800 (n=4), Roche - Roche Cobas 6800/8800 (n=3), Seegene - Seegene Seeplex (n=1), Siemens (n=1), Siemens - Siemens Versant (n=1), TIB MOLBIOL (n=3), TIB MOLBIOL - TIB MOLBIOL LightMix (n=3)

**Groups Rolled Up:** Abbott - Abbott Real Time PCR (n=12), Cepheid - Cepheid Xpert kit (n=7), Hologic - Hologic Aptima (n=14), In-House - Real-time In-House PCR (n=5)

# Individual Report

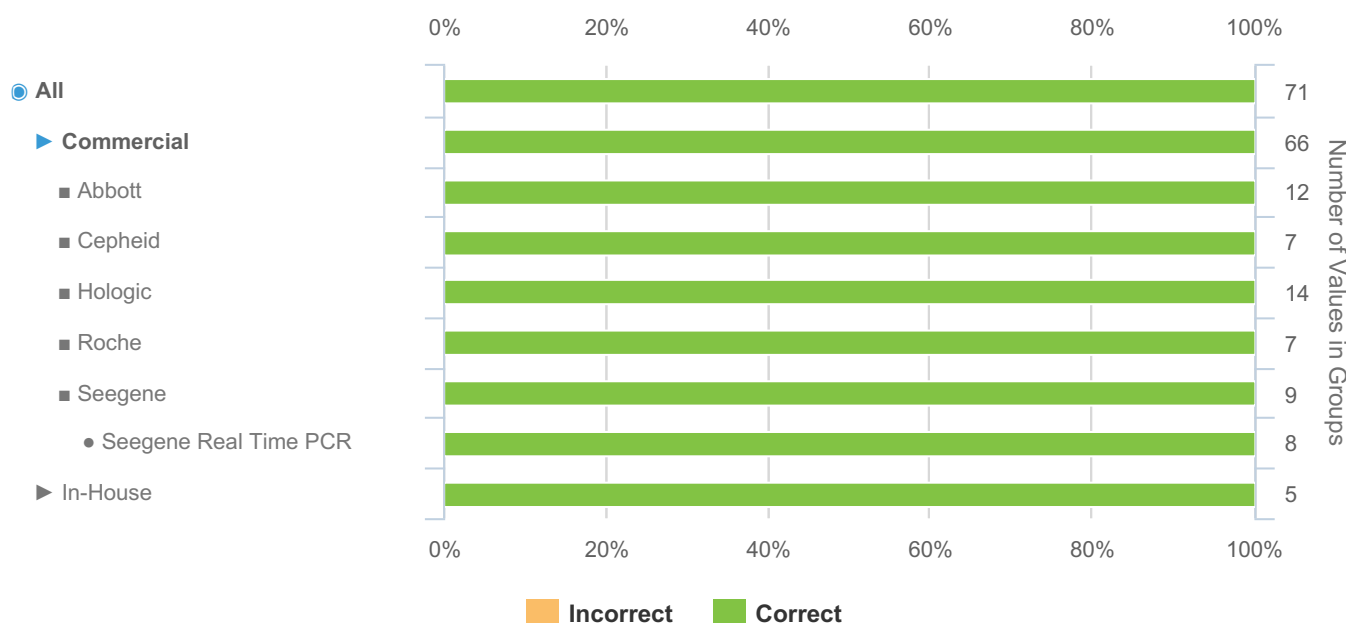
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### CTNg19S-04 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (All)	
						(%)	(n)
CTNg19S-04	N. gonorrhoeae (St 49226)	Urine	DS2_2	Negative	EDUCATIONAL	100.0	71



**Groups below n=5:** AmpliSens (n=2), AmpliSens - AmpliSens Real Time PCR (n=2), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=1), BD Molecular Diagnostics - BD ProbeTec (n=2), BIORON (n=2), BIORON - BIORON RealLine (n=2), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), Goffin Molecular Technologies (n=1), Goffin Molecular Technologies - Goffin Presto (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience FlouroType (n=1), Randox (n=1), Randox - Randox Multiplex Array (n=1), Roche - Roche Cobas 4800 (n=4), Roche - Roche Cobas 6800/8800 (n=3), Seegene - Seegene Seeplex (n=1), Siemens (n=1), Siemens - Siemens Versant (n=1), TIB MOLBIOL (n=3), TIB MOLBIOL - TIB MOLBIOL LightMix (n=3)

**Groups Rolled Up:** Abbott - Abbott Real Time PCR (n=12), Cepheid - Cepheid Xpert kit (n=7), Hologic - Hologic Aptima (n=14), In-House - Real-time In-House PCR (n=5)

# Individual Report

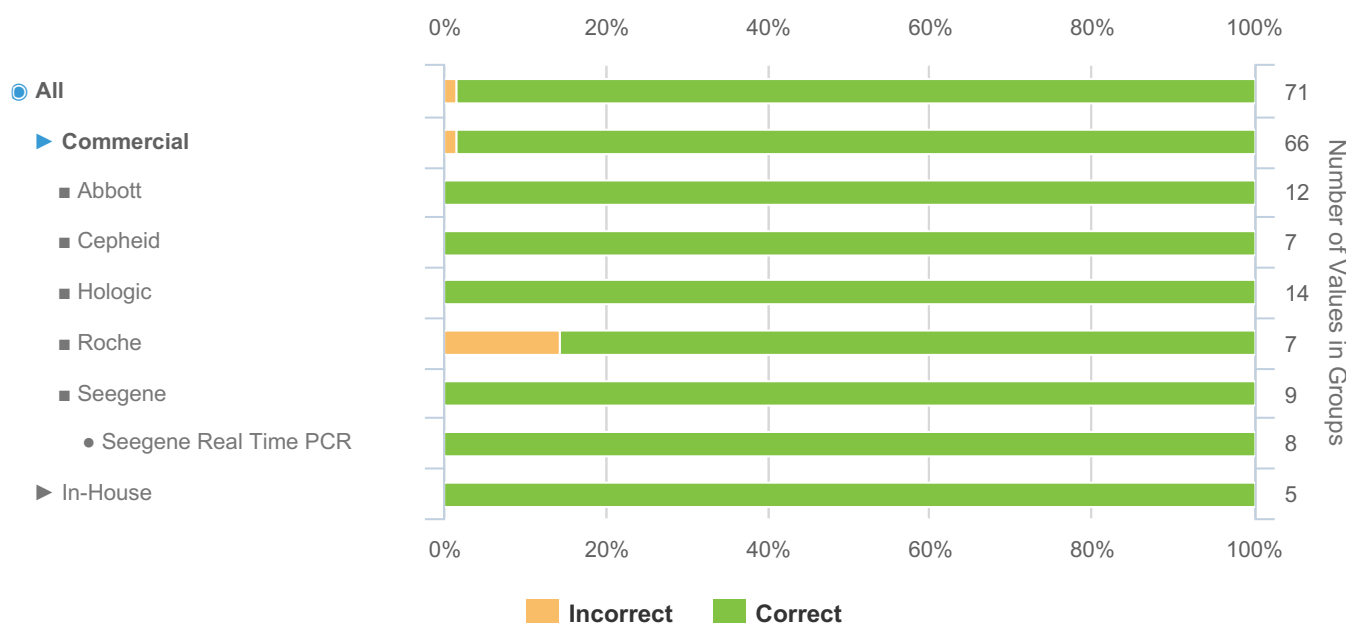
## QCMD 2019 Chlamydia trachomatis and Neisseria gonorrhoeae DNA EQA Programme (CT)



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### CTNg19S-06 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (All)	
						(%)	(n)
CTNg19S-06	N. gonorrhoeae (St 49226)	Simulated Swab	D1	Negative	EDUCATIONAL	98.6	71



**Groups below n=5:** AmpliSens (n=2), AmpliSens - AmpliSens Real Time PCR (n=2), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=1), BD Molecular Diagnostics - BD ProbeTec (n=2), BIORON (n=2), BIORON - BIORON RealLine (n=2), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), Goffin Molecular Technologies (n=1), Goffin Molecular Technologies - Goffin Presto (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience FlouroType (n=1), Randox (n=1), Randox - Randox Multiplex Array (n=1), Roche - Roche Cobas 4800 (n=4), Roche - Roche Cobas 6800/8800 (n=3), Seegene - Seegene Seeplex (n=1), Siemens (n=1), Siemens - Siemens Versant (n=1), TIB MOLBIOL (n=3), TIB MOLBIOL - TIB MOLBIOL LightMix (n=3)

**Groups Rolled Up:** Abbott - Abbott Real Time PCR (n=12), Cepheid - Cepheid Xpert kit (n=7), Hologic - Hologic Aptima (n=14), In-House - Real-time In-House PCR (n=5)



# Individual Report

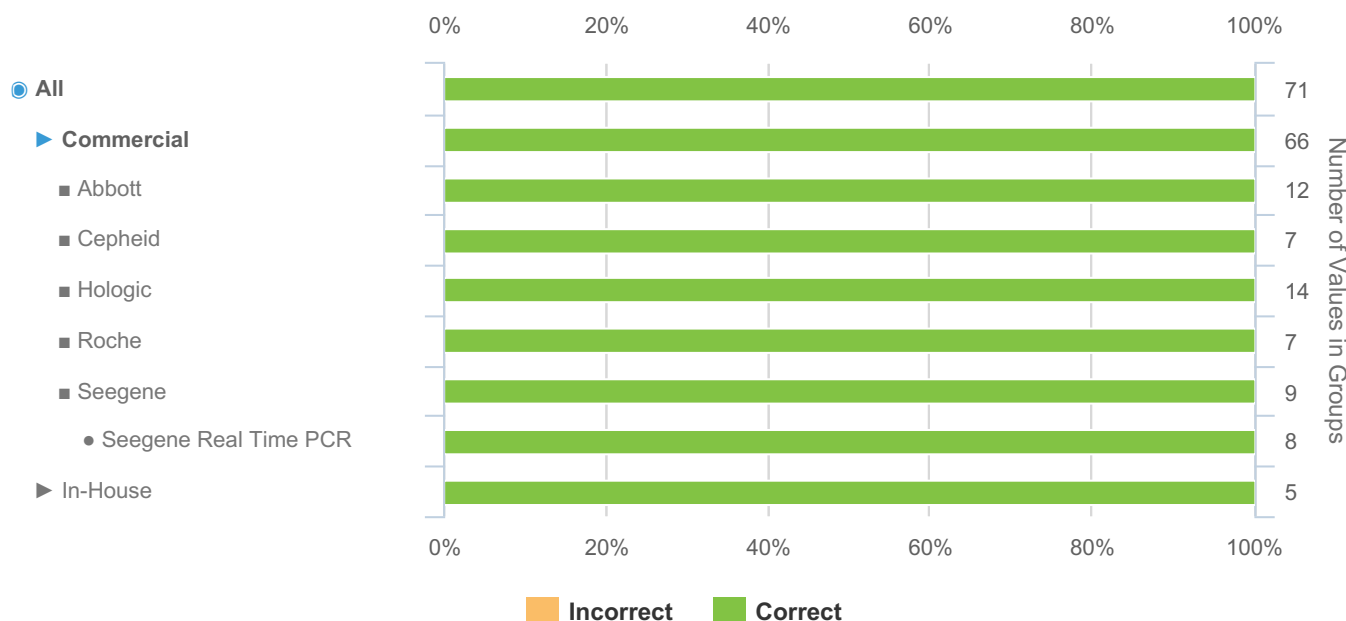
## QCMD 2019 Chlamydia trachomatis and Neisseria gonorrhoeae DNA EQA Programme (CT)



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### CTNg19S-09 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (All)	
						(%)	(n)
CTNg19S-09	N. gonorrhoeae (St 49226)	Simulated Swab	D1	Negative	EDUCATIONAL	100.0	71



**Groups below n=5:** AmpliSens (n=2), AmpliSens - AmpliSens Real Time PCR (n=2), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=1), BD Molecular Diagnostics - BD ProbeTec (n=2), BIORON (n=2), BIORON - BIORON RealLine (n=2), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), Goffin Molecular Technologies (n=1), Goffin Molecular Technologies - Goffin Presto (n=1), Hain Lifescience (n=1), Hain Lifescience - Hain Lifescience FlouroType (n=1), Randox (n=1), Randox - Randox Multiplex Array (n=1), Roche - Roche Cobas 4800 (n=4), Roche - Roche Cobas 6800/8800 (n=3), Seegene - Seegene Seeplex (n=1), Siemens (n=1), Siemens - Siemens Versant (n=1), TIB MOLBIOL (n=3), TIB MOLBIOL - TIB MOLBIOL LightMix (n=3)

**Groups Rolled Up:** Abbott - Abbott Real Time PCR (n=12), Cepheid - Cepheid Xpert kit (n=7), Hologic - Hologic Aptima (n=14), In-House - Real-time In-House PCR (n=5)

**Individual Report****QCMD 2019 Chlamydia trachomatis and Neisseria gonorrhoeae DNA EQA Programme (CT)**

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