Individ Report	ual	QCMD 2019 Chlamydia trachomatis and Neisseria gonorrhoeae DNA EQA Programme (Ng)					I for Malecular Diagnostics
Catalogue C						Report UID:	Laboratory
QAB174191						4254/304109/2140	DE182

Intended Results / Panel Composition

Sample Code	Sample Content	Matrix	Sample Relationships ^[1]	Detection Frequency ^[2]	Sample Status ^[3]	Percentage Correct (All) ^[4]	
						(%)	(n)
CTNg19S-01	C. trachomatis (Genovar F)	Urine		Negative	EDUCATIONAL	100.0	71
CTNg19S-02	C. trachomatis (LGV)	Simulated Swab		Negative	EDUCATIONAL	100.0	71
CTNg19S-03	N. gonorrhoeae (St 49226)	Urine	DS2_1	Frequently Detected	CORE	100.0	71
CTNg19S-04	N. gonorrhoeae (St 49226)	Urine	DS2_2	Frequently Detected	CORE	100.0	71
CTNg19S-05	C. trachomatis (LGV)	Urine	DS1_1	Negative	EDUCATIONAL	100.0	71
CTNg19S-06	N. gonorrhoeae (St 49226)	Simulated Swab	D1	Frequently Detected	CORE	97.2	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	Frequently Detected	CORE	100.0	71
CTNg19S-10	C. trachomatis (LGV)	Urine	DS1_2	Negative	EDUCATIONAL	100.0	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.

Individual Report	Neisser		nydia tracho loeae DNA E			
Catalogue Code: QAB174191					Report UID: 4254/304109/2140	Laboratory DE182

Your Summary Results

EQA Assessment Group ^[1]

Core Panel Detection (Qualitative) Score [2]

Commercial

0

Core Panel Members Results

Sample Code	Qualitative Results		Your Quantitative Data (for information only) [3]				
Code	Percentage Correct (All) [4]	Your Result [5]	Detection Score [6]	Reported Value	Unitage	Cycle Threshold	
CTNg19S-03	100.0	Positive	0		N/A	32.3	
CTNg19S-04	100.0	Positive	0		N/A	34.0	
CTNg19S-06	97.2	Positive	0		N/A	30.6	
CTNg19S-07	100.0	Negative	0		N/A	-	
CTNg19S-08	100.0	Negative	0		N/A	-	
CTNg19S-09	100.0	Positive	0		N/A	30.6	

[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 (AII): Percentage of datasets (%) reporting the correct qualitative results for each panel member.

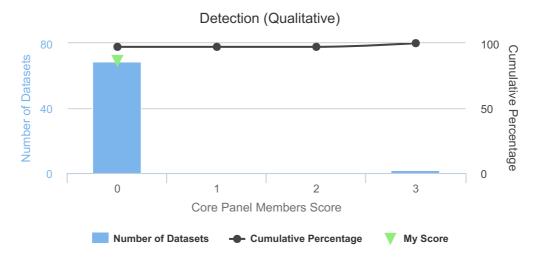
[5] Your Result: The qualitative result you reported for each sample within this EQA challenge / distribution.

[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.

Individual Report		a gonorrh	nydia trachom oeae DNA EQ		I for Molecular Diagnostics
Catalogue Code:	Ref Code:	Challenge:	Analysis Type:	Report UID: 4254/304109/2140	Laboratory
QAB174191	CTNg19	S	Qualitative		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.

Name	CT/NG Workflow double Kit (v3)							
Description	Verwendung RealLine Chlamydia trachomatis / Neisseria gonorrhoeae REF VBD0457							
Targets	B Neisseria gonorrhoeae B Chlamydia trachomatis							
Assays	 <i>Extraction</i> - Manual Extraction Process Commercial Kit Manufacturer: BIORON Diagnostics GmbH Kit Type: RealLine DNA Express REF VBC8899 							
	 Amplification - DNA Technology - DTlite Real-Time PCR System Multiplex Commercial Kit Manufacturer: BIORON Diagnostics GmbH Kit Type: RealLine Chlamydia trachomatis / Neisseria gonorrhoeae REF VBD0457 Kit Version: not applicable 							

Individual Report	Neisseri		nydia trachor loeae DNA EC		
Catalogue Code:	Ref Code:	Challenge:	Analysis Type:	Report UID: 4254/304109/2140	Laboratory
QAB174191	CTNg19	S	Qualitative		DE182

Educational Panel Members Results

Sample Code	Qualitative Results		Your Quantitative Data (for information only) ^[1]			
Coue	Percentage Correct (All) ^[2]	Your Result ^[3]	Detection Score ^[4]	Reported Value	Unitage	Cycle Threshold
CTNg19S-01	100.0	Negative	0		N/A	-
CTNg19S-02	100.0	Negative	0		N/A	-
CTNg19S-05	100.0	Negative	0		N/A	-
CTNg19S-10	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	65
Number of Datasets Submitted	71
Qualitative Results Returned	71 (100.0%)

Comments

For the analysis of Neisseria gonorrhoeae results, Chlamydia trachomatis 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.

Individual Report	Neisseri		nydia trachor loeae DNA EC		QUALITY Control for Molecular Diagnostics			
Catalogue Code: QAB174191	Ref Code: CTNg19	Challenge: S	Analysis Type: Qualitative	Dataset: 304109	Report UID: 4254/304109/2140	Laboratory 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).

Individual Report		a gonorrh	nydia trachom oeae DNA EQ		
Catalogue Code:	Ref Code:	Challenge:	Analysis Type:	Report UID: 4254/304109/2140	Laboratory
QAB174191	CTNg19	S	Qualitative		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.

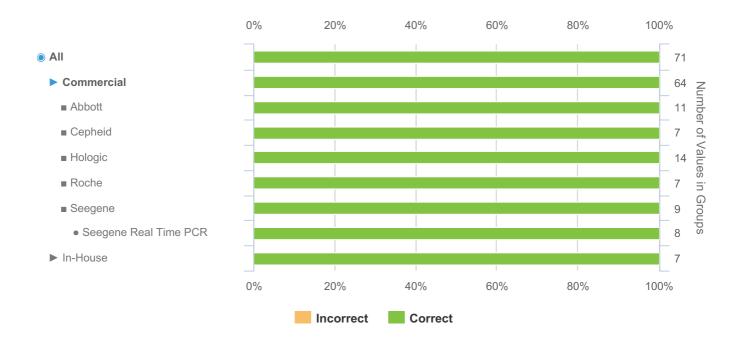
To allow meaningful assessment at the individual method level the EQA assessment group must consist of 5 or more datasets. If there are not sufficient datasets at the individual method level then your results will be included within a higher EQA assessment group based on whether it is a commercial or 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.

		a gonorrh	nydia trachom oeae DNA EQ		
Catalogue Code:	Ref Code:	Challenge:	Analysis Type:	Report UID:	Laboratory
QAB174191	CTNg19	S	Qualitative	4254/304109/2140	DE182

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	Frequently Detected	CORE	100.0	71	

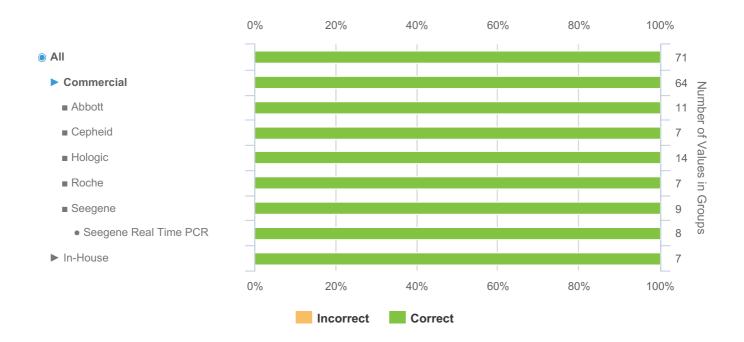


Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=2), BD Molecular Diagnostics - BD ProbeTec (n=1), BIORON (n=2), BIORON - BIORON RealLine (n=2), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), Goffin Molecular Technologies (n=1), Goffin Molecular Technologies (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)

		019 Chlam a gonorrho me (Ng)				
talogue Code:Ref Code:Challenge:Analysis Type:Dataset:B174191CTNg19SQualitative304109					Report UID: 4254/304109/2140	Laboratory DE182

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	Frequently Detected	CORE	100.0	71	

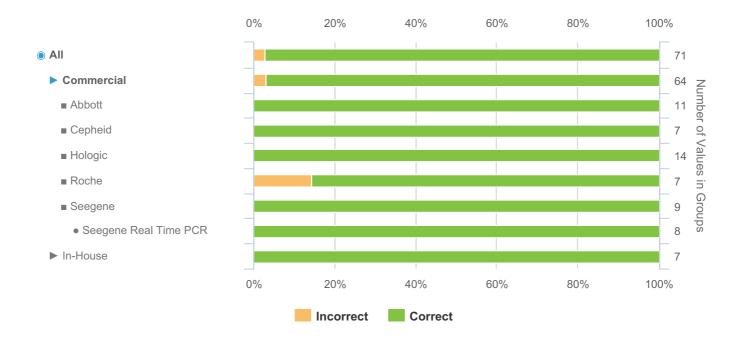


Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=2), BD Molecular Diagnostics - BD ProbeTec (n=1), BIORON (n=2), BIORON - BIORON RealLine (n=2), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), Goffin Molecular Technologies (n=1), Goffin Molecular Technologies (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)

		019 Chlam a gonorrho me (Ng)				
talogue Code:Ref Code:Challenge:Analysis Type:Dataset:B174191CTNg19SQualitative304109					Report UID: 4254/304109/2140	Laboratory DE182

CTNg19S-06 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage (All)	e Correct
						(%)	(n)
CTNg19S-06	N. gonorrhoeae (St 49226)	Simulated Swab	D1	Frequently Detected	CORE	97.2	71

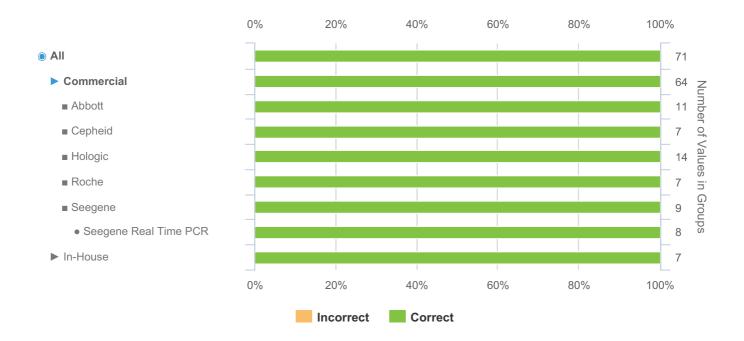


Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=2), BD Molecular Diagnostics - BD ProbeTec (n=1), BIORON (n=2), BIORON - BIORON RealLine (n=2), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), Goffin Molecular Technologies (n=1), Goffin Molecular Technologies (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)

Individual Report	QCMD 2 Neisser Progran	matis and QA				
Catalogue Code: QAB174191	Ref Code: CTNg19	Challenge: S	Analysis Type: Qualitative	Dataset: 304109	Report UID: 4254/304109/2140	Laboratory DE182

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

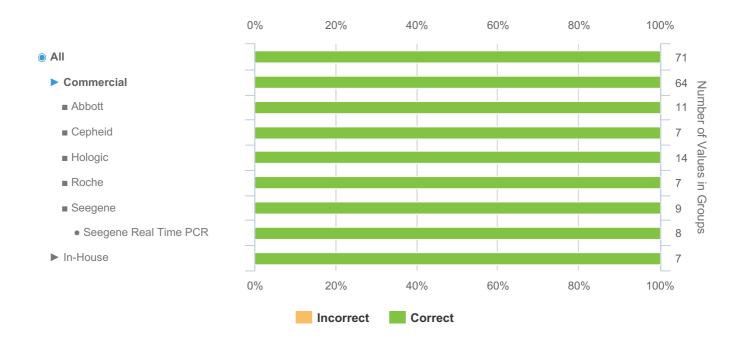


Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=2), BD Molecular Diagnostics - BD ProbeTec (n=1), BIORON (n=2), BIORON - BIORON RealLine (n=2), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), Goffin Molecular Technologies (n=1), Goffin Molecular Technologies (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)

)19 Chlam a gonorrho me (Ng)				
Catalogue Code: QAB174191	Ref Code: CTNg19	Challenge: S	Analysis Type: Qualitative	Dataset: 304109	Report UID: 4254/304109/2140	Laboratory DE182

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

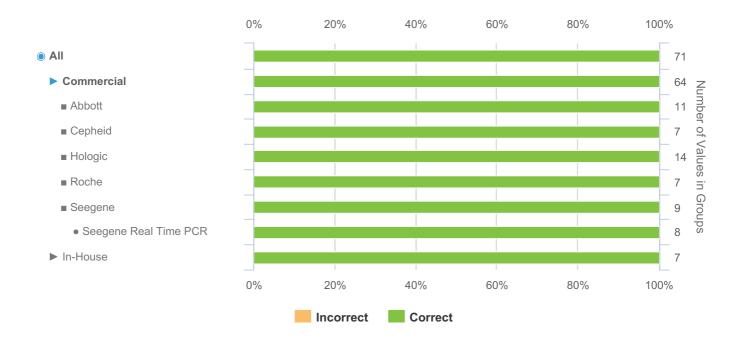


Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=2), BD Molecular Diagnostics - BD ProbeTec (n=1), BIORON (n=2), BIORON - BIORON RealLine (n=2), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), Goffin Molecular Technologies (n=1), 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)

		019 Chlam a gonorrho me (Ng)				
talogue Code:Ref Code:Challenge:Analysis Type:Dataset:B174191CTNg19SQualitative304109					Report UID: 4254/304109/2140	Laboratory DE182

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	Frequently Detected	CORE	100.0	71



Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=2), BD Molecular Diagnostics - BD ProbeTec (n=1), 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 (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)

Individual Report		a gonorrh	nydia trachon oeae DNA EC			
Catalogue Code:	Ref Code:	Challenge:	Analysis Type:	Dataset: 304109	Report UID:	Laboratory
QAB174191	CTNg19	S	Qualitative		4254/304109/2140	DE182

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.

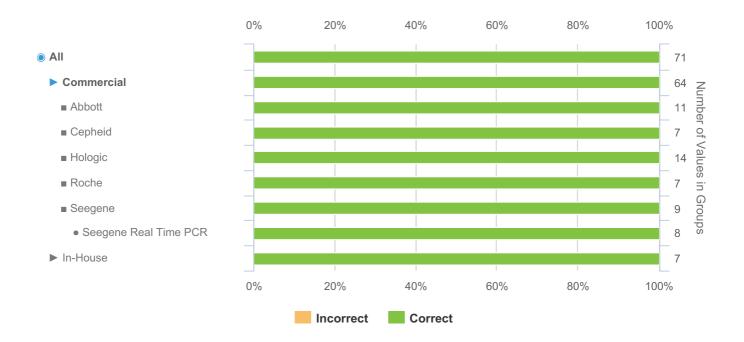
To allow meaningful assessment at the individual method level the EQA assessment group must consist of 5 or more datasets. If there are not sufficient datasets at the individual method level then your results will be included within a higher EQA assessment group based on whether it is a commercial or 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	Neisser	2019 Chlar ia gonorrh nme (Ng)				
Catalogue Code: QAB174191	5		Analysis Type: Qualitative	Dataset: 304109	Report UID: 4254/304109/2140	Laboratory DE182

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		Negative	EDUCATIONAL	100.0	71

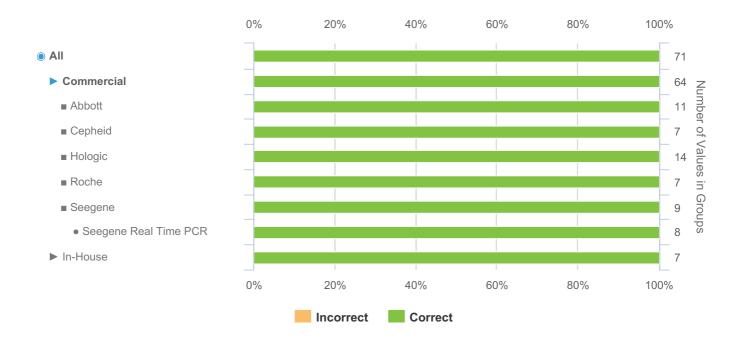


Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=2), BD Molecular Diagnostics - BD ProbeTec (n=1), BIORON (n=2), BIORON - BIORON RealLine (n=2), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), Goffin Molecular Technologies (n=1), Goffin Molecular Technologies (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)

Individual Report	Neisser	QCMD 2019 Chlamydia trachomatis and Neisseria gonorrhoeae DNA EQA Programme (Ng)						
Catalogue Code: QAB174191			Analysis Type: Qualitative	Dataset: 304109	Report UID: 4254/304109/2140	Laboratory DE182		

CTNg19S-02 - Qualitative Results Breakdown

Sample Code	e	Sample Content	Matrix	Sample Relationships	Detection Sample Stat Frequency		Percentag Correct (A	
							(%)	(n)
CTNg1	9S-02	C. trachomatis (LGV)	Simulated Swab		Negative	EDUCATIONAL	100.0	71

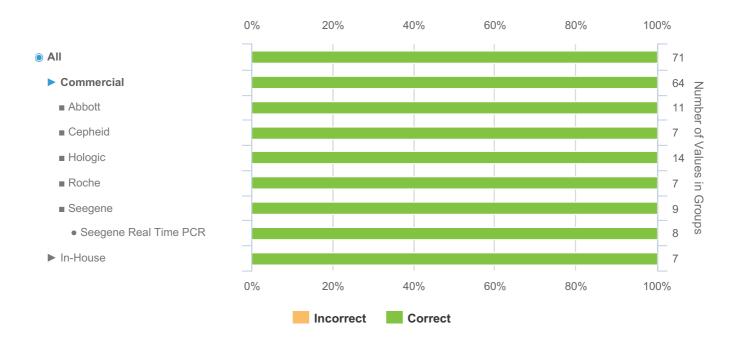


Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=2), BD Molecular Diagnostics - BD ProbeTec (n=1), BIORON (n=2), BIORON - BIORON RealLine (n=2), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), Goffin Molecular Technologies (n=1), Goffin Molecular Technologies (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)

)19 Chlam a gonorrho me (Ng)				
Catalogue Code: QAB174191	Ref Code: CTNg19	Challenge: S	Analysis Type: Qualitative	Dataset: 304109	Report UID: 4254/304109/2140	Laboratory DE182

CTNg19S-05 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage (All)	Correct
						(%)	(n)
CTNg19S-05	C. trachomatis (LGV)	Urine	DS1_1	Negative	EDUCATIONAL	100.0	71

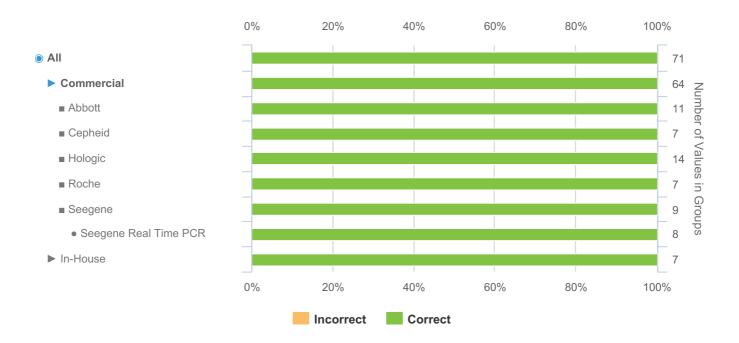


Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=2), BD Molecular Diagnostics - BD ProbeTec (n=1), BIORON (n=2), BIORON - BIORON RealLine (n=2), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), Goffin Molecular Technologies (n=1), Goffin Molecular Technologies (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)

)19 Chlam a gonorrho me (Ng)				
Catalogue Code: QAB174191	Ref Code: CTNg19	Challenge: S	Analysis Type: Qualitative	Dataset: 304109	Report UID: 4254/304109/2140	Laboratory DE182

CTNg19S-10 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage (All)	Correct
						(%)	(n)
CTNg19S-10	C. trachomatis (LGV)	Urine	DS1_2	Negative	EDUCATIONAL	100.0	71



Groups below n=5: AmpliSens (n=1), AmpliSens - AmpliSens Real Time PCR (n=1), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics Easy-Plex (n=2), BD Molecular Diagnostics (n=3), BD Molecular Diagnostics - BD MAX (n=2), BD Molecular Diagnostics - BD ProbeTec (n=1), 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 (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)

Individual Report	Neisseri		nydia trachon oeae DNA EG			
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