# QCMD 2016 Chlamydia trachomatis DNA EQA Programme



Catalogue Code: QAB004101 Ref Code: CTDNA16

Challenge: 1 of 2 Analysis Type:
Qualitative

Dataset: 72268

Report UID: 4254/72268/422

Participant: DE182-01

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

Sample Code	Sample Content Matrix Sample Relationships [1] Detection Frequency [2]	Matrix	Relationships		Sample Status <sup>[3]</sup>	Percentage Correct (AII) <sup>[4]</sup>	
			(%)	(n)			
CTDNA16C1-01	Negative	Simulated Swab		Negative	CORE	97.0	271
CTDNA16C1-02	C. trachomatis (LGV)	Simulated Swab	D1	Frequently Detected	CORE	96.3	271
CTDNA16C1-03	Negative	Urine		Negative	CORE	98.9	271
CTDNA16C1-04	C. trachomatis (LGV)	Simulated Swab	D1	Frequently Detected	CORE	98.2	271
CTDNA16C1-05	C. trachomatis Swedish strain	Urine		Frequently Detected	CORE	98.5	271

- [1] Sample Relationships: Indicates the relationships of the samples within this challenge. Dilution series are indicted by 'DS1' with each panel member in the dilution series represented by a number in order of titre, where DS1\_1 represents the highest titre within that dilution series. Further dilution series are indicated by 'DS2' 'DS3' 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 R	esults
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EQA Assessment Group [1] Commercial

Core Panel Detection (Qualitative) Score [2]

0

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Sample Code	Qualitative Results		Your Quantitative Data (for information only) [3]			
	Percentage Correct (AII) [4]	Your Result <sup>[5]</sup>	Detection Score <sup>[6]</sup>	Reported Value	Unitage	Cycle Threshold
CTDNA16C1-01	97.0	Negative	0		N/A	-
CTDNA16C1-02	96.3	Positive	0		N/A	36.9
CTDNA16C1-03	98.9	Negative	0		N/A	-
CTDNA16C1-04	98.2	Positive	0		N/A	36.9
CTDNA16C1-05	98.5	Positive	0		N/A	32.2

- [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



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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 used to submit your results for this challenge.

Name	CT/NG Workflow double Kit (v2)
Description	Verwendung RealLine Chlamydia trachomatis / Neisseria gonorrhoeae REF VBD0457
Targets	B Chlamydia trachomatis B Neisseria gonorrhoeae
Assays	<ul> <li>Extraction - Manual Extraction Process</li> <li>Commercial</li> <li>Kit Manufacturer: BIORON Diagnostics GmbH</li> <li>Kit Type: RealLine DNA Express REF VBC8899</li> </ul>
	<ul> <li>Amplification - DNA Technology - DTlite Real-Time PCR System</li> <li>Commercial</li> <li>Kit Manufacturer: BIORON Diagnostics GmbH</li> <li>Kit Type: RealLine Chlamydia trachomatis / Neisseria gonorrhoeae REF VBD0457</li> <li>Kit Version: not applicable</li> </ul>

### **Further Programme Details**

Number of Participants	244
Number of Countries	35
Number of Respondents	233
Number of Datasets Submitted	271
Qualitative Results Returned	271 (100.0%)

### **EQA Programme Aims**

To assess the qualitative performance of participants' molecular assays in detecting *Chlamydia trachomatis* (*C. trachomatis*) at various concentrations.

To assess the ability of participants' molecular assays to correctly identify different C. trachomatis strains.

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### 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).

### Additional Information: 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 2016 Chlamydia trachomatis DNA EQA Programme



Catalogue Code: QAB004101

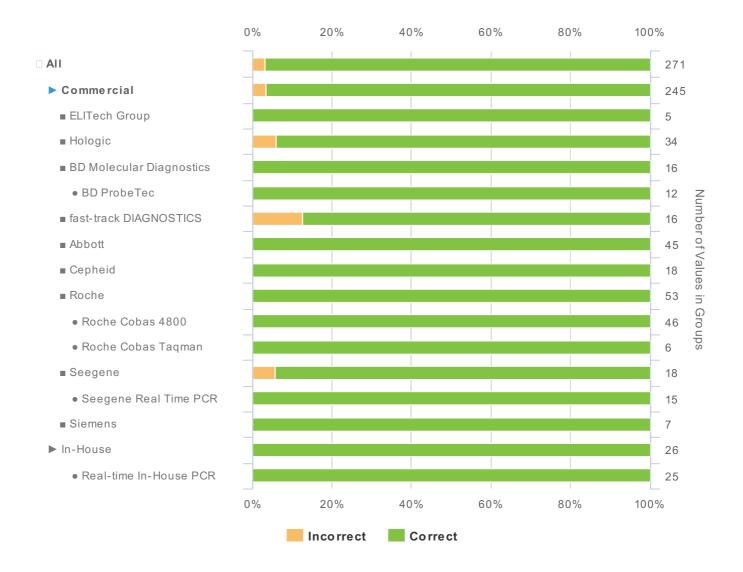
Ref Code: CTDNA16 Challenge: Analysis Type: 1 of 2 Qualitative

Dataset: 72268 Report UID: 4254/72268/422

Participant: DE182-01

#### CTDNA16C1-01 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct All	
						(%)	(n)
CTDNA16C1-01	Negative	Simulated Swab		Negative	CORE	97.0	271



## QCMD 2016 Chlamydia trachomatis DNA EQA Programme



Catalogue Code: QAB004101 Ref Code: CTDNA16 Challenge: 1 of 2

Analysis Type:
Qualitative

Dataset: 72268 Report UID: 4254/72268/422

Participant: DE182-01

Groups below n=5: ELITech Group - Elitech Elite Real time kit (n=1), ELITech Group - Elitech Alert Real Time Q-PCR kit (n=4), BD Molecular Diagnostics - BD MAX (n=4), Roche - Roche Cobas Amplicor (n=1), Seegene - Seegene Seeplex (n=3), BlORON (n=1), BlORON - BlORON RealLine (n=1), AB Analitica (n=4), AB Analitica - AB Analitica REALQUALITY RQ (n=4), EuroClone Diagnostica (n=2), EuroClone Diagnostica - EuroClone Duplica Real-time (n=2), Goffin Molecular Technologies (n=2), Goffin Molecular Technologies - Goffin Molecular Technologies Presto (n=2), Clonit (n=1), Clonit - Clonit PCR Reagents (n=1), Sacace (n=4), Sacace - Sacace Real TM (n=4), InterLabService (n=3), InterLabService - InterLabService AmpliSens (n=3), Hain Lifescience (n=2), Hain Lifescience - Hain Lifescience FlouroType (n=2), Diagenode (n=4), Diagenode - Diagenode Real Time kit (n=4), Randox (n=1), Randox - Randox Multiplex Array (n=1), GeneProof (n=4), GeneProof - GeneProof Real Time PCR kit (n=4), QIAGEN (n=4), QIAGEN - Qiagen Artus Real Time (n=4), Greiner bio one (n=1), Greiner bio one - Greiner bio-one (n=1), In-House - Conventional In-House PCR (n=1)

Groups Rolled Up: Hologic - Hologic Aptima (n=34), fast-track DIAGNOSTICS - FTD Real Time PCR (n=16), Abbott - Abbott Realtime (n=45), Cepheid - Cepheid Xpert kit (n=18), Siemens - Siemens Versant (n=7)

# QCMD 2016 Chlamydia trachomatis DNA EQA Programme



Catalogue Code: QAB004101

Ref Code: CTDNA16 Challenge: Analysis Type: 1 of 2 Qualitative

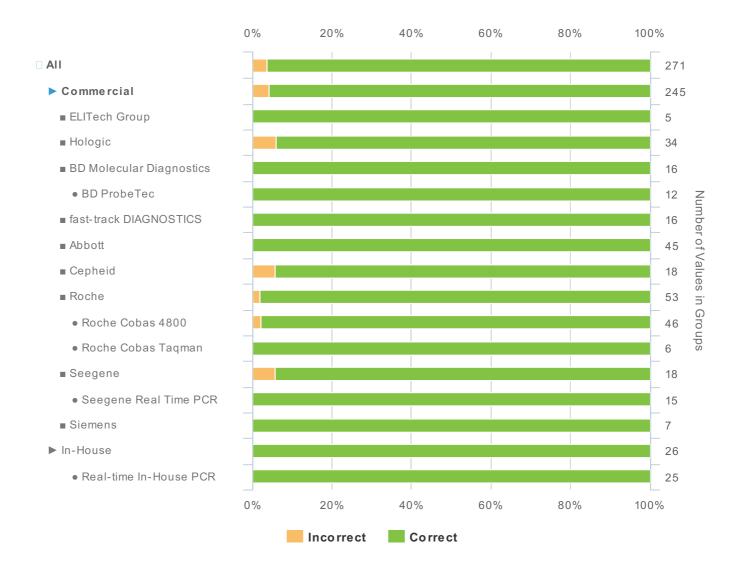
Dataset: 72268

Report UID: 4254/72268/422

Participant: DE182-01

#### CTDNA16C1-02 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percenta Correct	•
						(%)	(n)
CTDNA16C1-02	C. trachomatis (LGV)	Simulated Swab	D1	Frequently Detected	CORE	96.3	271



## QCMD 2016 Chlamydia trachomatis DNA EQA Programme



Catalogue Code: QAB004101 Ref Code: CTDNA16 Challenge: 1 of 2 Analysis Type:
Qualitative

Dataset: 72268 Report UID: 4254/72268/422 Participant: DE182-01

Groups below n=5: ELITech Group - Elitech Elite Real time kit (n=1), ELITech Group - Elitech Alert Real Time Q-PCR kit (n=4), BD Molecular Diagnostics - BD MAX (n=4), Roche - Roche Cobas Amplicor (n=1), Seegene - Seegene Seeplex (n=3), BlORON (n=1), BlORON - BlORON RealLine (n=1), AB Analitica (n=4), AB Analitica - AB Analitica REALQUALITY RQ (n=4), EuroClone Diagnostica (n=2), EuroClone Diagnostica - EuroClone Duplica Real-time (n=2), Goffin Molecular Technologies (n=2), Goffin Molecular Technologies - Goffin Molecular Technologies Presto (n=2), Clonit (n=1), Clonit - Clonit PCR Reagents (n=1), Sacace (n=4), Sacace - Sacace Real TM (n=4), InterLabService (n=3), InterLabService - InterLabService AmpliSens (n=3), Hain Lifescience (n=2), Hain Lifescience - Hain Lifescience FlouroType (n=2), Diagenode (n=4), Diagenode - Diagenode Real Time kit (n=4), Randox (n=1), Randox - Randox Multiplex Array (n=1), GeneProof (n=4), GeneProof - GeneProof Real Time PCR kit (n=4), QIAGEN (n=4), QIAGEN - Qiagen Artus Real Time (n=4), Greiner bio one (n=1), Greiner bio one - Greiner bio-one (n=1), In-House - Conventional In-House PCR (n=1)

Groups Rolled Up: Hologic - Hologic Aptima (n=34), fast-track DIAGNOSTICS - FTD Real Time PCR (n=16), Abbott - Abbott Realtime (n=45), Cepheid - Cepheid Xpert kit (n=18), Siemens - Siemens Versant (n=7)

# QCMD 2016 Chlamydia trachomatis DNA EQA Programme



Catalogue Code: QAB004101

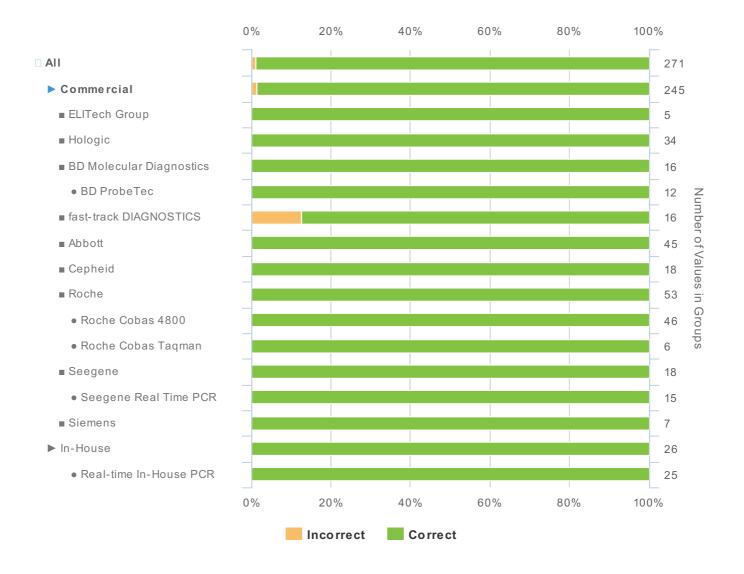
Ref Code: CTDNA16 Challenge: Analysis Type: 1 of 2 Qualitative

Dataset: 72268 Report UID: 4254/72268/422

Participant: DE182-01

#### CTDNA16C1-03 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	·		Percenta Correct A	•	
						(%)	(n)
CTDNA16C1-03	Negative	Urine		Negative	CORE	98.9	271



### **QCMD 2016 Chlamydia trachomatis DNA EQA Programme**



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Ref Code: CTDNA16

Challenge: 1 of 2

Analysis Type: Qualitative

Dataset: 72268

Report UID: 4254/72268/422 Participant: DE182-01

Groups below n=5: ELITech Group - Elitech Elite Real time kit (n=1), ELITech Group - Elitech Alert Real Time Q-PCR kit (n=4), BD Molecular Diagnostics - BD MAX (n=4), Roche - Roche Cobas Amplicor (n=1), Seegene - Seegene Seeplex (n=3), BIORON (n=1), BIORON - BIORON RealLine (n=1), AB Analitica (n=4), AB Analitica - AB Analitica REALQUALITY RQ (n=4), Euro Clone Diagnostica (n=2), EuroClone Diagnostica - EuroClone Duplica Real-time (n=2), Goffin Molecular Technologies (n=2), Goffin Molecular Technologies - Goffin Molecular Technologies Presto (n=2), Clonit (n=1), Clonit - Clonit PCR Reagents (n=1), Sacace (n=4), Sacace - Sacace Real TM (n=4), InterLab Service (n=3), InterLab Service - InterLab Service AmpliSens (n=3), Hain Lifescience (n=2), Hain Lifescience - Hain Lifescience Flouro Type (n=2), Diagenode (n=4), Diagenode - Diagenode Real Time kit (n=4), Randox (n=1), Randox - Randox Multiplex Array (n=1), GeneProof (n=4), GeneProof - GeneProof Real Time PCR kit (n=4), QIAGEN (n=4), QIAGEN -Qiagen Artus Real Time (n=4), Greiner bio one (n=1), Greiner bio one - Greiner bio-one (n=1), In-House - Conventional In-House

Groups Rolled Up: Hologic - Hologic Aptima (n=34), fast-track DIAGNOSTICS - FTD Real Time PCR (n=16), Abbott - Abbott Realtime (n=45), Cepheid - Cepheid Xpert kit (n=18), Siemens - Siemens Versant (n=7)

# QCMD 2016 Chlamydia trachomatis DNA EQA Programme



Catalogue Code: QAB004101

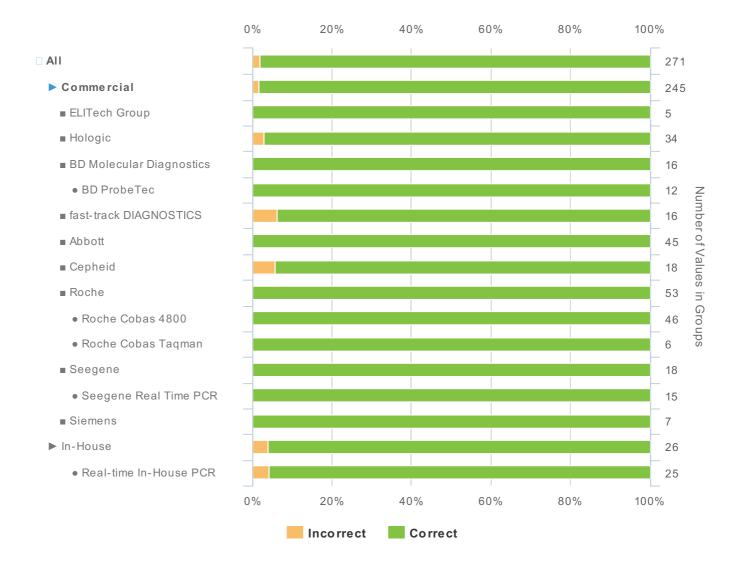
Ref Code: CTDNA16 Challenge: Analysis Type: 1 of 2 Qualitative

Dataset: 72268 Report UID: 4254/72268/422

Participant: DE182-01

#### CTDNA16C1-04 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percenta Correct	•
						(%)	(n)
CTDNA16C1-04	C. trachomatis (LGV)	Simulated Swab	D1	Frequently Detected	CORE	98.2	271



# QCMD 2016 Chlamydia trachomatis DNA EQA Programme



Catalogue Code: QAB004101 Ref Code: CTDNA16 Challenge: Ai

Analysis Type:
Qualitative

Dataset: 72268 Report UID: 4254/72268/422

Participant: DE182-01

Groups below n=5: ELITech Group - Elitech Elite Real time kit (n=1), ELITech Group - Elitech Alert Real Time Q-PCR kit (n=4), BD Molecular Diagnostics - BD MAX (n=4), Roche - Roche Cobas Amplicor (n=1), Seegene - Seegene Seeplex (n=3), BlORON (n=1), BlORON - BlORON RealLine (n=1), AB Analitica (n=4), AB Analitica - AB Analitica REALQUALITY RQ (n=4), EuroClone Diagnostica (n=2), EuroClone Diagnostica - EuroClone Duplica Real-time (n=2), Goffin Molecular Technologies (n=2), Goffin Molecular Technologies - Goffin Molecular Technologies Presto (n=2), Clonit (n=1), Clonit - Clonit PCR Reagents (n=1), Sacace (n=4), Sacace - Sacace Real TM (n=4), InterLabService (n=3), InterLabService - InterLabService AmpliSens (n=3), Hain Lifescience (n=2), Hain Lifescience - Hain Lifescience FlouroType (n=2), Diagenode (n=4), Diagenode - Diagenode Real Time kit (n=4), Randox (n=1), Randox - Randox Multiplex Array (n=1), GeneProof (n=4), GeneProof - GeneProof Real Time PCR kit (n=4), QIAGEN (n=4), QIAGEN - Qiagen Artus Real Time (n=4), Greiner bio one (n=1), Greiner bio one - Greiner bio-one (n=1), In-House - Conventional In-House PCR (n=1)

Groups Rolled Up: Hologic - Hologic Aptima (n=34), fast-track DIAGNOSTICS - FTD Real Time PCR (n=16), Abbott - Abbott Realtime (n=45), Cepheid - Cepheid Xpert kit (n=18), Siemens - Siemens Versant (n=7)

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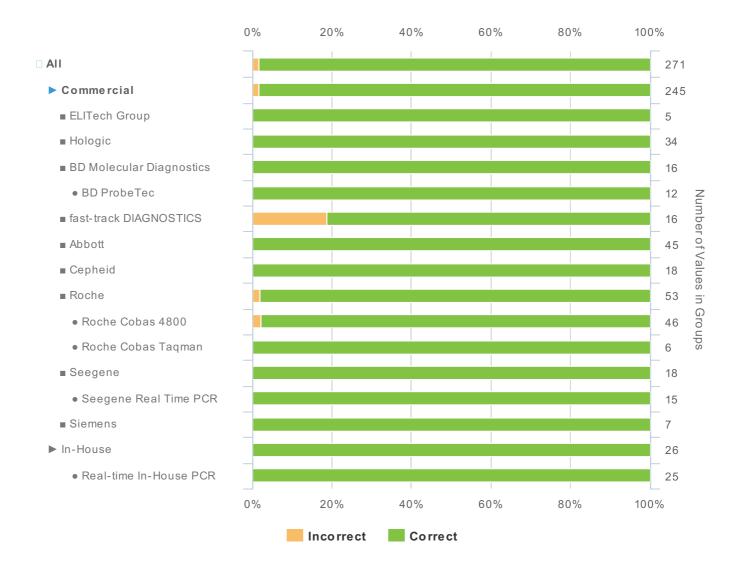
Ref Code: CTDNA16 Challenge: Analysis Type: 1 of 2 Qualitative

Dataset: 72268 Report UID: 4254/72268/422

Participant: DE182-01

#### CTDNA16C1-05 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percent Correct	•
						(%)	(n)
CTDNA16C1-05	C. trachomatis Swedish strain	Urine		Frequently Detected	CORE	98.5	271



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Dataset: 72268 Report UID: 4254/72268/422 Participant: DE182-01

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Groups Rolled Up: Hologic - Hologic Aptima (n=34), fast-track DIAGNOSTICS - FTD Real Time PCR (n=16), Abbott - Abbott Realtime (n=45), Cepheid - Cepheid Xpert kit (n=18), Siemens - Siemens Versant (n=7)

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