QCMD 2022 Enterovirus RNA EQA Programme



Catalogue Code: QAV984104

Ref Code: EVRNA22

Challenge:

Analysis Type: Qualitative **Dataset:** 532636

Report UID: 2677/532636/4437

Laboratory CZ023

Intended Results / Panel Composition

Sample Code	Sample Content	Matrix	Sample Relationships ^[1]	Detection Frequency ^[2]	Sample Status ^[3]	Percentage Correct (All) [4]	
						(%)	(n)
EVRNA22C1-01	Coxsackievirus A24	Transport medium	-	Frequently Detected	CORE	95.7	117
EVRNA22C1-02	Negative	Transport medium	-	Negative	CORE	96.6	117
EVRNA22C1-03	Coxsackievirus A9	Transport medium	-	Frequently Detected	CORE	95.7	117
EVRNA22C1-04	Echovirus 6	Transport medium	-	Frequently Detected	CORE	99.1	117
EVRNA22C1-05	Echovirus 25	Transport medium	-	Frequently Detected	CORE	99.1	117

[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.

Y	our	Sum	mary	Res	ults
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EQA Assessment Group [1] Commercial

Core Panel Detection (Qualitative) Score [2]

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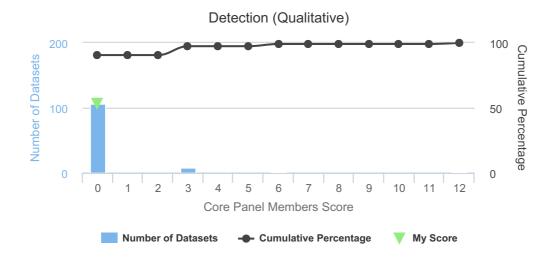
Core Panel Members Results

Sample Code	Qualitative Results	Your Quantitative Data (for information only) [3]				
	Percentage Correct (All) [4]	Your Result	Detection Score	Reported Value	Unitage	Cycle Threshold
EVRNA22C1-01	95.7	Positive	0	9656	Copies/ml	26.5
EVRNA22C1-02	96.6	Negative	0	0	Copies/ml	-
EVRNA22C1-03	95.7	Positive	0	2	Copies/ml	39.68
EVRNA22C1-04	99.1	Positive	0	59706	Copies/ml	23.78
EVRNA22C1-05	99.1	Positive	0	25829	Copies/ml	25.02

- [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.
- $[5] \begin{tabular}{ll} \textbf{Your Result:} The qualitative result you reported for each sample within this EQA challenge / distribution. } \end{tabular}$
- [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(s) used to submit your results for this challenge.

Name	Enterovirus PCR Kit-croBEE (v4)
Description	
Targets	V enterovirus
Assays	Extraction - GeneProof - croBEE NA16 Nucleic Acid Extraction System Commercial Kit Manufacturer: GeneProof Kit Type: croBEE 201A Nucleic Acid Extraction Kit Amplification, GeneProof, croBEE Roal Time BCR System
	 Amplification - GeneProof - croBEE Real-Time PCR System Commercial Kit Manufacturer: GeneProof Kit Type: GeneProof Enterovirus PCR Kit Kit Version: ISEX

Further Programme Details

Number of Participants	105
Number of Countries	25
Number of Respondents	98
Number of Datasets Submitted	117
Qualitative Results Returned	117 (100.0%)

EQA Programme Aims

To assess the ability of laboratories molecular assays to detect different types and concentrations of enterovirus (EV).

To review the performance of laboratories quantitative EV molecular assays.

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

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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.

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Ref Code: EVRNA22

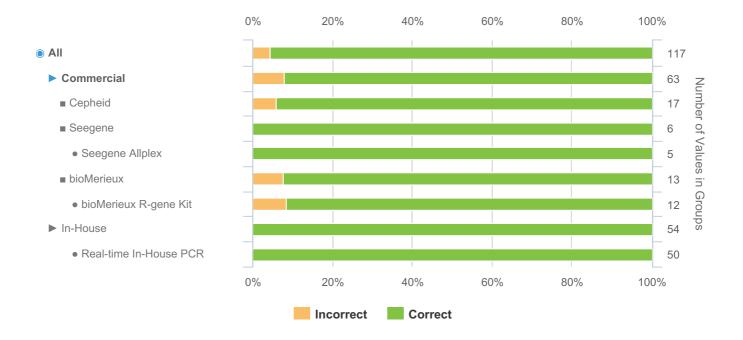
Challenge: C1 Analysis Type: Qualitative **Dataset:** 532636

Report UID: 2677/532636/4437

Laboratory CZ023

EVRNA22C1-01 - Qualitative Results Breakdown

Sample Cod	le Sample Content	Matrix	rix Sample Detection Relationships Frequency		Sample Status	Percentag (All)	ge Correct
						(%)	(n)
EVRNA22C	I-01 Coxsackievirus A24	Transport medium	-	Frequently Detected	CORE	95.7	117



Groups below n=5: AB Analitica (n=2), AB Analitica - AB Analitica REALQUALITY RQ (n=2), Abacus Diagnostica (n=1), Abacus Diagnostica - Abacus GenomeEra (n=1), Altona Diagnostics (n=2), Altona Diagnostics - Altona Diagnostics RealStar (n=2), AusDiagnostics (n=3), AusDiagnostics - AusDiagnostics TademPlex (n=3), Diagenode (n=1), Diagenode - Diagenode Real Time kit (n=1), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), GeneProof (n=4), GeneProof - GeneProof Real Time PCR kit (n=4), Jiangsu Bioperfectus (n=1), Jiangsu Bioperfectus - Jiangsu Bioperfectus Real Time PCR (n=1), Liferiver (n=1), Liferiver - Liferiver Real Time PCR (n=1), Progenie Molecular (n=2), Progenie Molecular - Progenie Molecular RealCycler (n=2), R-Biopharm (n=4), R-Biopharm - R-Biopharm RIDA Gene (n=4), Roche (n=1), Roche - Roche LightCycler (n=1), Seegene - Seegene Anyplex II (n=1), bioMerieux - BioFire FilmArray ME (n=1), fast-track DIAGNOSTICS (n=4), fast-track DIAGNOSTICS - FTD Real Time PCR (n=3), fast-track DIAGNOSTICS - FTD Respiratory Pathogens (n=1), In-House - Conventional In-House PCR (n=3), In-House - Real Time Nested RT PCR (n=1)

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Ref Code: EVRNA22

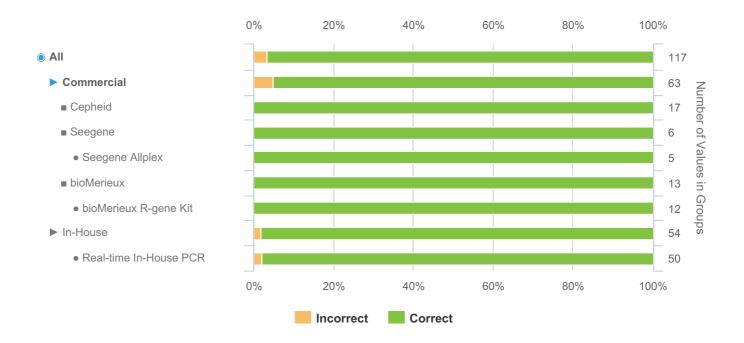
Challenge: C1 Analysis Type: Qualitative **Dataset:** 532636

Report UID: 2677/532636/4437

Laboratory CZ023

EVRNA22C1-02 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix Sample Detection Relationships Frequency		Sample Status	Percentage Correct (AII)		
						(%)	(n)
EVRNA22C1-02	Negative	Transport medium	-	Negative	CORE	96.6	117



Groups below n=5: AB Analitica (n=2), AB Analitica - AB Analitica REALQUALITY RQ (n=2), Abacus Diagnostica (n=1), Abacus Diagnostica - Abacus GenomeEra (n=1), Altona Diagnostics (n=2), Altona Diagnostics - Altona Diagnostics RealStar (n=2), AusDiagnostics (n=3), AusDiagnostics - AusDiagnostics TademPlex (n=3), Diagenode (n=1), Diagenode - Diagenode Real Time kit (n=1), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), GeneProof (n=4), GeneProof - GeneProof Real Time PCR kit (n=4), Jiangsu Bioperfectus (n=1), Jiangsu Bioperfectus - Jiangsu Bioperfectus Real Time PCR (n=1), Liferiver (n=1), Liferiver - Liferiver Real Time PCR (n=1), Progenie Molecular (n=2), Progenie Molecular - Progenie Molecular RealCycler (n=2), R-Biopharm (n=4), R-Biopharm RIDA Gene (n=4), Roche (n=1), Roche - Roche LightCycler (n=1), Seegene - Seegene Anyplex II (n=1), bioMerieux - BioFire FilmArray ME (n=1), fast-track DIAGNOSTICS (n=4), fast-track DIAGNOSTICS - FTD Real Time PCR (n=3), fast-track DIAGNOSTICS - FTD Respiratory Pathogens (n=1), In-House - Conventional In-House PCR (n=3), In-House - Real Time Nested RT PCR (n=1)

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Ref Code: EVRNA22

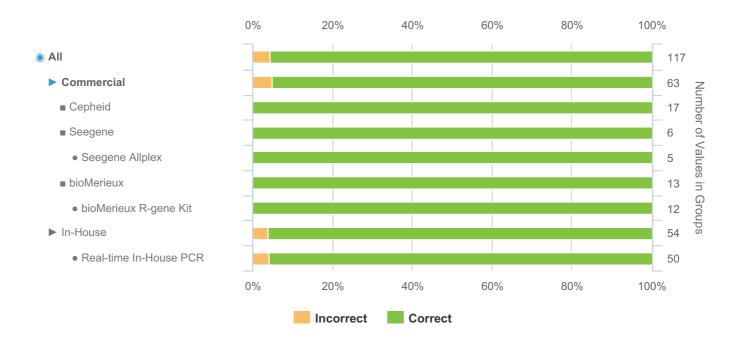
Challenge: C1 Analysis Type: Qualitative **Dataset:** 532636

Report UID: 2677/532636/4437

Laboratory CZ023

EVRNA22C1-03 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Relationships	Detection Frequency	Sample Status	Percentage Correct (AII)	
						(%)	(n)
EVRNA22C1-03	Coxsackievirus A9	Transport medium	-	Frequently Detected	CORE	95.7	117



Groups below n=5: AB Analitica (n=2), AB Analitica - AB Analitica REALQUALITY RQ (n=2), Abacus Diagnostica (n=1), Abacus Diagnostica - Abacus GenomeEra (n=1), Altona Diagnostics (n=2), Altona Diagnostics - Altona Diagnostics RealStar (n=2), AusDiagnostics (n=3), AusDiagnostics - AusDiagnostics TademPlex (n=3), Diagenode (n=1), Diagenode - Diagenode Real Time kit (n=1), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), GeneProof (n=4), GeneProof - GeneProof Real Time PCR kit (n=4), Jiangsu Bioperfectus (n=1), Jiangsu Bioperfectus - Jiangsu Bioperfectus Real Time PCR (n=1), Liferiver (n=1), Liferiver - Liferiver Real Time PCR (n=1), Progenie Molecular (n=2), Progenie Molecular - Progenie Molecular RealCycler (n=2), R-Biopharm (n=4), R-Biopharm - R-Biopharm RIDA Gene (n=4), Roche (n=1), Roche - Roche LightCycler (n=1), Seegene - Seegene Anyplex II (n=1), bioMerieux - BioFire FilmArray ME (n=1), fast-track DIAGNOSTICS (n=4), fast-track DIAGNOSTICS - FTD Real Time PCR (n=3), fast-track DIAGNOSTICS - FTD Respiratory Pathogens (n=1), In-House - Conventional In-House PCR (n=3), In-House - Real Time Nested RT PCR (n=1)

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Catalogue Code: QAV984104

Ref Code: EVRNA22

Challenge: C1 **Analysis Type:**Qualitative

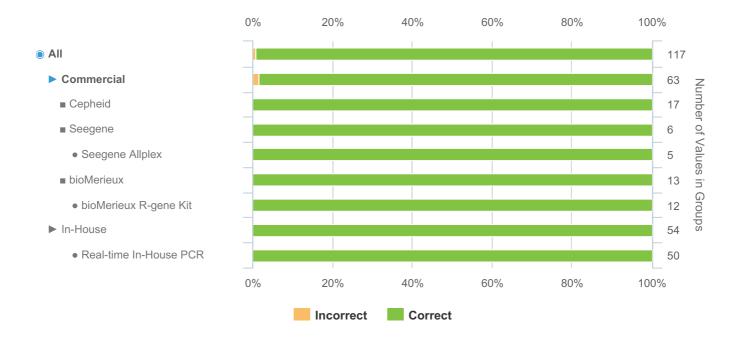
Dataset: 532636

Report UID: 2677/532636/4437

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EVRNA22C1-04 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Matrix Sample Detection Relationships Frequency		Sample Status	Percentage Correct (AII)	
						(%)	(n)
EVRNA22C1-04	Echovirus 6	Transport medium	-	Frequently Detected	CORE	99.1	117



Groups below n=5: AB Analitica (n=2), AB Analitica - AB Analitica REALQUALITY RQ (n=2), Abacus Diagnostica (n=1), Abacus Diagnostica - Abacus GenomeEra (n=1), Altona Diagnostics (n=2), Altona Diagnostics - Altona Diagnostics RealStar (n=2), AusDiagnostics (n=3), AusDiagnostics - AusDiagnostics TademPlex (n=3), Diagenode (n=1), Diagenode - Diagenode Real Time kit (n=1), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), GeneProof (n=4), GeneProof - GeneProof Real Time PCR kit (n=4), Jiangsu Bioperfectus (n=1), Jiangsu Bioperfectus - Jiangsu Bioperfectus Real Time PCR (n=1), Liferiver (n=1), Liferiver - Liferiver Real Time PCR (n=1), Progenie Molecular (n=2), Progenie Molecular - Progenie Molecular RealCycler (n=2), R-Biopharm (n=4), R-Biopharm - R-Biopharm RIDA Gene (n=4), Roche (n=1), Roche - Roche LightCycler (n=1), Seegene - Seegene Anyplex II (n=1), bioMerieux - BioFire FilmArray ME (n=1), fast-track DIAGNOSTICS (n=4), fast-track DIAGNOSTICS - FTD Real Time PCR (n=3), fast-track DIAGNOSTICS - FTD Respiratory Pathogens (n=1), In-House - Conventional In-House PCR (n=3), In-House - Real Time Nested RT PCR (n=1)

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Ref Code: EVRNA22

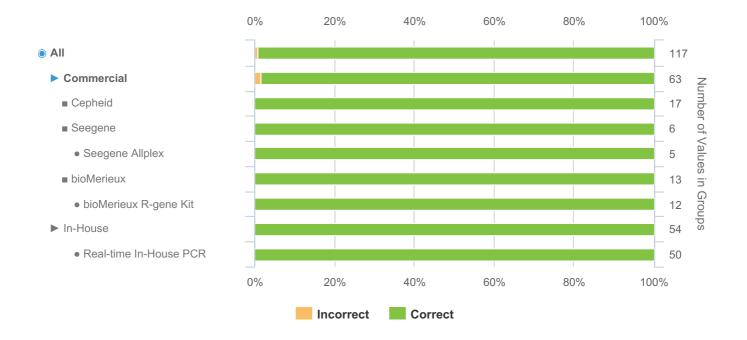
Challenge: C1 Analysis Type: Qualitative **Dataset:** 532636

Report UID: 2677/532636/4437

Laboratory CZ023

EVRNA22C1-05 - Qualitative Results Breakdown

Sample Code	Sample Content	Matrix	Sample Detection Relationships Frequency		Sample Status	Percentag (All)	je Correct
						(%)	(n)
EVRNA22C1-05	Echovirus 25	Transport medium	-	Frequently Detected	CORE	99.1	117



Groups below n=5: AB Analitica (n=2), AB Analitica - AB Analitica REALQUALITY RQ (n=2), Abacus Diagnostica (n=1), Abacus Diagnostica - Abacus GenomeEra (n=1), Altona Diagnostics (n=2), Altona Diagnostics - Altona Diagnostics RealStar (n=2), AusDiagnostics (n=3), AusDiagnostics - AusDiagnostics TademPlex (n=3), Diagenode (n=1), Diagenode - Diagenode Real Time kit (n=1), ELITech Group (n=1), ELITech Group - Elitech Elite Real Time kit (n=1), GeneProof (n=4), GeneProof - GeneProof Real Time PCR kit (n=4), Jiangsu Bioperfectus (n=1), Jiangsu Bioperfectus - Jiangsu Bioperfectus Real Time PCR (n=1), Liferiver (n=1), Liferiver - Liferiver Real Time PCR (n=1), Progenie Molecular (n=2), Progenie Molecular - Progenie Molecular RealCycler (n=2), R-Biopharm (n=4), R-Biopharm - R-Biopharm RIDA Gene (n=4), Roche (n=1), Roche - Roche LightCycler (n=1), Seegene - Seegene Anyplex II (n=1), bioMerieux - BioFire FilmArray ME (n=1), fast-track DIAGNOSTICS (n=4), fast-track DIAGNOSTICS - FTD Real Time PCR (n=3), fast-track DIAGNOSTICS - FTD Respiratory Pathogens (n=1), In-House - Conventional In-House PCR (n=3), In-House - Real Time Nested RT PCR (n=1)

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Ref Code: EVRNA22

Challenge: Analysis Type:
C1 Qualitative

Dataset: 532636

Report UID: 2677/532636/4437

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