


Individual Report		QCMD 2022 Respiratory I EQA Programme			 <small>Quality Control for Molecular Diagnostics</small>	
Catalogue Code: QAV164188	Ref Code: RESPI22	Challenge: S	Analysis Type: Multiple Pathogen Qualitative	Dataset: -	Report UID: 0/0/4844	Laboratory

NOTE: Summary information only.


Panel Composition

Sample Code	Sample Content	Matrix	Sample Relationships ^[1]	Detected / Determined ^[2]		Not Detected / Not Determined ^[2]		Not Tested ^[2]	
				(%)	(n)	(%)	(n)	(%)	(n)
RESPI22S-01	Influenza virus A (H3N2)	Transport Medium	DS1_1	95.8	137	3.5	5	0.7	1
RESPI22S-02	Influenza virus A H1N1 (pdm09)	Transport Medium		94.4	135	4.9	7	0.7	1
RESPI22S-03	Influenza virus A (H3N2)	Transport Medium	DS1_2	86.7	124	12.6	18	0.7	1
RESPI22S-04	Negative	Transport Medium		97.2	139	2.8	4	N/A	0
RESPI22S-05	Influenza virus B (Victoria)	Transport Medium	DS2_1	97.9	140	1.4	2	0.7	1
RESPI22S-06	Respiratory Syncytial Virus (type B)	Transport Medium		95.8	137	2.8	4	1.4	2
RESPI22S-07	Influenza virus B (Victoria)	Transport Medium	DS2_2	95.1	136	4.2	6	0.7	1
RESPI22S-08	Influenza (type A) & Respiratory Syncytial Virus (type A)	Transport Medium		90.2	129	7.7	11	2.1	3
RESPI22S-09	Influenza virus B (Yamagata)	Transport Medium		97.2	139	2.1	3	0.7	1
RESPI22S-10	Respiratory Syncytial Virus (type A)	Transport Medium		93	133	5.6	8	1.4	2

[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] **Detected / Determined; Not Detected / Not Determined; Not Tested:** The percentage (%) of datasets reported by all participants in relation to the assigned status of the panel member i.e. 'positive' or 'negative' and the expected pathogen type as defined through pre-testing and the total number of datasets (n) for each panel member.

For further details please refer to the current participant manual.

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EQA Assessment Group ^[1]	N/A (Refer to My Workflow details section below)
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Your Summary Results (Core Samples)

Sample Code	Expected Result ^[2]		Your Final Laboratory Reported Result ^[3]			Sample Status ^[7]	Detection Frequency ^[8]	Detection Score ^[9]
	Qualitative	Pathogen ID	Pathogen included in workflow(s) ^[4] Yes/No	Qualitative ^[5]	Reported Pathogen ID ^[6]			
RESPI22S-01	Positive	Influenza virus	-	-	-	Core	Frequently Detected	-
RESPI22S-02	Positive	Influenza virus	-	-	-	Core	Frequently Detected	-
RESPI22S-03	Positive	Influenza virus	-	-	-	Core	Detected	-
RESPI22S-04	Negative		-	-	-	Core	Negative	-
RESPI22S-05	Positive	Influenza virus	-	-	-	Core	Frequently Detected	-
RESPI22S-06	Positive	Respiratory syncytial virus	-	-	-	Core	Frequently Detected	-
RESPI22S-07	Positive	Influenza virus	-	-	-	Core	Frequently Detected	-
RESPI22S-08	Positive	Influenza virus and Respiratory syncytial virus	-	-	-	Core	Detected	-
RESPI22S-09	Positive	Influenza virus	-	-	-	Core	Frequently Detected	-
RESPI22S-10	Positive	Respiratory syncytial virus	-	-	-	Core	Detected	-

[1] **EQA Assessment Group:** To aid 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] **Expected Result:** positive / negative result and the specific pathogen present within each panel member.

[3] **Your Final Laboratory Reported Result:** the final reported result which may be based on one or more workflows used to test each panel member.

[4] **Pathogen included in workflow(s):** Yes / No answer to whether the expected pathogen was tested for.


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

[6] **Reported Pathogen ID:** The final pathogen(s) identification you reported for each sample within this EQA challenge / distribution.

[7] **Sample Status:** 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.

[8] **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 by participants within the EQA challenge/distribution. Note that the detection frequency is assigned using only datasets submitted using workflows including the target pathogen.

[9] **Detection Score:** Your detection scores are based on the assigned detection frequency of each panel member, where 0 is "highly satisfactory" and 3 (three) is "highly unsatisfactory"

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For further details please refer to the current participant manual.


Multiple Pathogen Programme - Qualitative Assessment of Results

Results are categorised based on the workflow used and the pathogen(s) targeted as shown in the table below.

Expected Qualitative Result	Laboratory Reported Results						Result Category		Sample Weighting			
	Positive	Negative	Not Determined	Expected pathogen(s) included in workflow(s)		Expected pathogen(s) not included in workflow(s)			Frequently Detected (>95% positive)	Detected (Between 65 and 95% positive)	Infrequently Detected (Less than 65% positive)	Negative
				Expected pathogen(s) detected	Expected pathogen(s) not detected							
Positive	✓			✓			Expected Pathogen Reported	Detected / Determined	0	0	0	N/A
Negative		✓					No pathogen reported	Detected / Determined	N/A	N/A	N/A	0
Negative	✓						False Positive	False Positive	3	3	3	N/A
Positive	✓					✓	Reported Pathogen(s) not as expected	False Positive	3	3	3	N/A
Positive	✓					✓	Reported Pathogen(s) not as expected	False Positive	3	3	3	N/A
Positive or Negative			✓				Result reported as not determined	Not Determined	3	2	1	N/A
Positive		✓				✓	No pathogen reported	False Negative	3	2	1	N/A
Positive		✓				✓	Expected pathogen not tested for	Not Tested	Not Scored	Not Scored	Not Scored	N/A

My Workflow Details:

N/A

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Further Programme Details

Number of Participants	130
Number of Countries	31
Number of Respondents	121
Number of Datasets Submitted	143

Comments

Influenza Virus A H1N1 Pdm09: clade 6B.1 strain isolate A/Netherlands/1250/2016, GISAID: EPI_ISL_239751.

Influenza B Virus (Victoria): clade 1A strain isolate B/Netherlands/2518/2016, GISAID: EPI_ISL_239749.

Influenza B Virus (Yamagata): clade 3 strain Isolate B/Netherlands/365/2016, GISAID: EPI_ISL_239748.

Influenza Virus A H3N2: clade 3C.2a strain isolate A/Netherlands/2393/2015, GISAID: EPI_ISL_239750.

EQA Programme Aims


To assess the proficiency of laboratories in the molecular detection and determination of various influenza A & B and respiratory syncytial virus strains.

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.

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

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Individual Panel Member Analysis (Core Samples)

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.

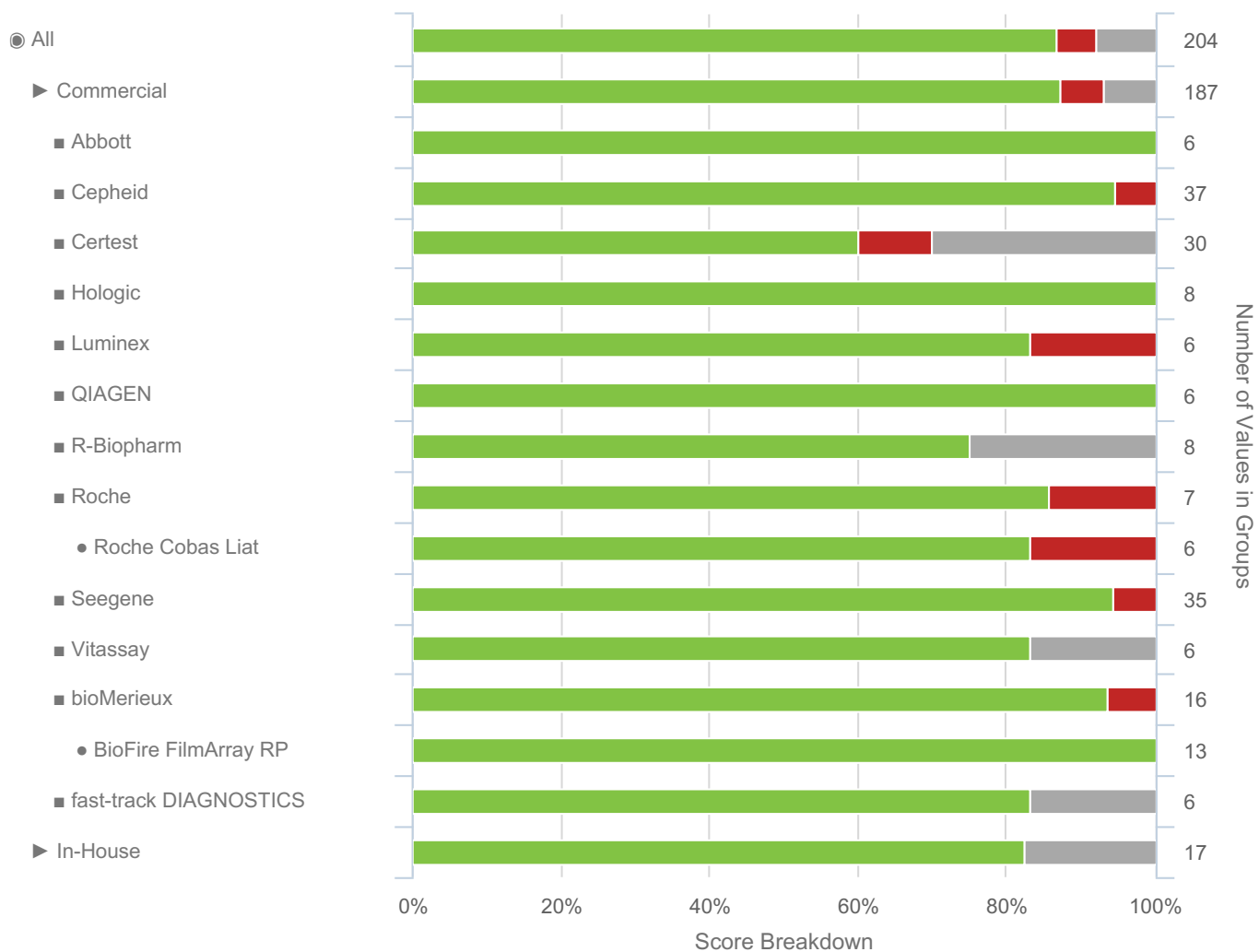
To allow meaningful assessment at the individual method level the EQA assessment group must consist of 5 or more datasets. If there are not sufficient datasets at the individual method level then your results will be included within a higher EQA assessment group based on whether it is a commercial or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

A breakdown of qualitative results reported for all workflows used by participants on each of the panel members within this EQA challenge / distribution is provided below. Note: participants may use multiple workflows for each sample.


The final laboratory result indicates the final reported result which may be based on one or more workflows used to test each panel member.

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Sample Code	Sample Content	Matrix	Sample Relationships	Expected targets	Detected / Determined		Not Detected / Not Determined		Not Tested	
					(%)	(n)	(%)	(n)	(%)	(n)
RESPI22S-01	Influenza virus A (H3N2)	Transport Medium	DS1_1	Influenza virus	95.8	137	3.5	5	0.7	1



■ Detected / Determined ■ Not Detected / Not Determined ■ Not Tested

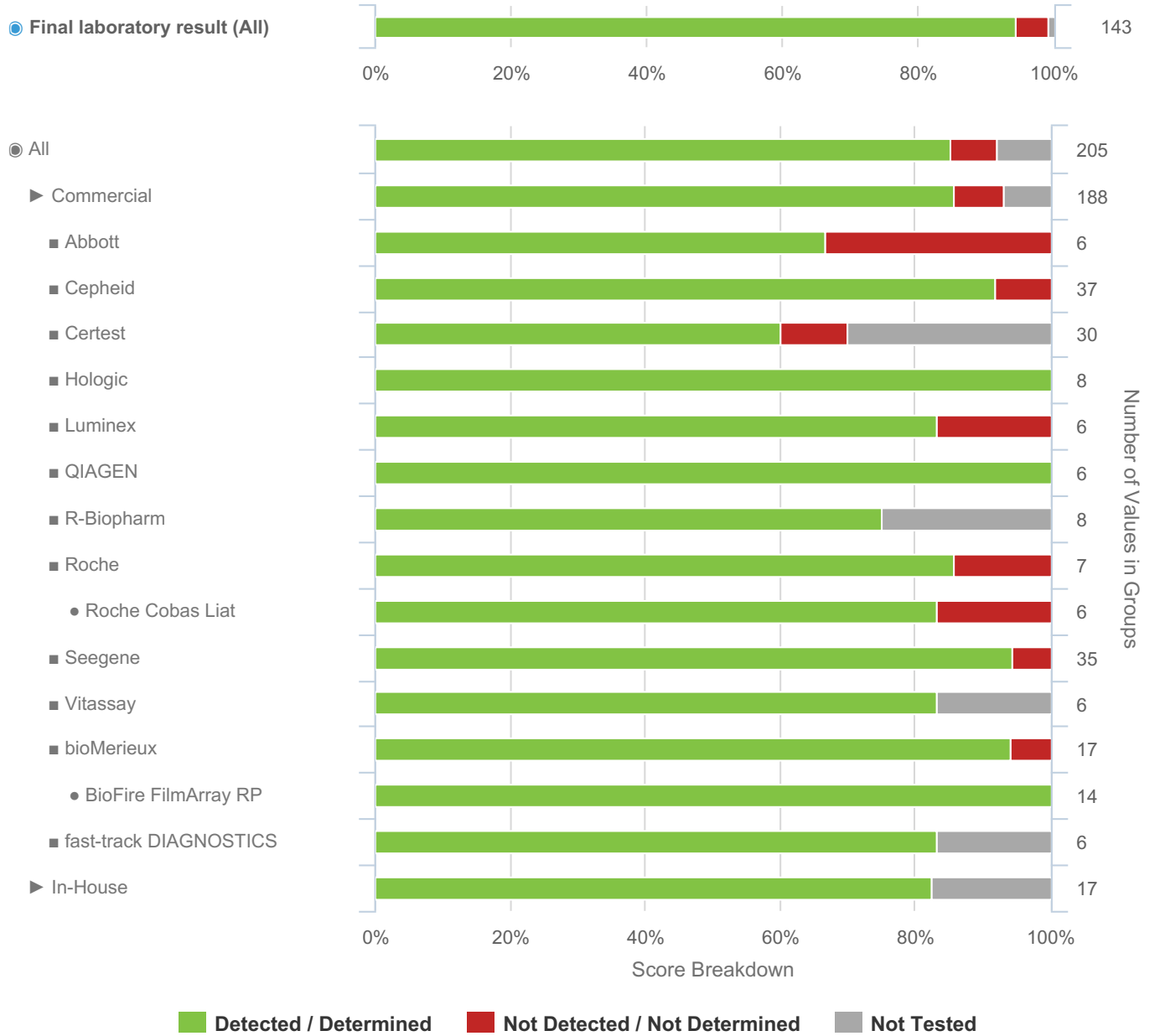
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Groups below n=5: Abbott - Abbott Alinity m (n=3), Abbott - Abbott ID NOW (n=3), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics TandemPlex (n=2), DiaSorin (n=2), DiaSorin - DiaSorin Simplexa (n=2), GenMark Dx (n=3), GenMark Dx - GenMark DX ePlex (n=3), Genematrix (n=1), Genematrix - Genematrix NeoPlex RV-Panel A (n=1), Immundiagnostik (n=1), Immundiagnostik - Immundiagnostik MutaPLEX (n=1), Luminex - Luminex ARIES (n=2), Luminex - Luminex xTAG (n=3), Luminex - Verigene (n=1), PathoFinder (n=4), PathoFinder - PathoFinder Real Time PCR (n=4), QIAGEN - Qiagen NeuMoDx (n=3), QIAGEN - Qiagen QIAstat-Dx (n=3), Roche - Roche Cobas 6800/8800 (n=1), Speedx (n=1), Speedx - Speedx Real Time PCR (n=1), TRUPCR (n=1), TRUPCR - TRUPCR Respiratory (n=1), ThermoFisher (n=1), ThermoFisher - ThermoFisher TaqMan Array (n=1), bioMerieux - BioFire FilmArray PN (n=1), bioMerieux - bioMerieux Primers/Probes (n=1), bioMerieux - bioMerieux R-gene Kit (n=1)

Groups Rolled Up: Cepheid - Cepheid Xpert kit (n=37), Certest - Certest Real Time PCR (n=30), Hologic - Hologic Panther Fusion (n=8), R-Biopharm - R-Biopharm RIDA Gene (n=8), Seegene - Seegene Allplex (n=35), Vitassay - Vitassay Real-Time PCR (n=6), fast-track DIAGNOSTICS - FTD real time PCR (n=6), In-House - Real-time In-House PCR (n=17)

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Sample Code	Sample Content	Matrix	Sample Relationships	Expected targets	Detected / Determined		Not Detected / Not Determined		Not Tested	
					(%)	(n)	(%)	(n)	(%)	(n)
RESPI22S-02	Influenza virus A H1N1 (pdm09)	Transport Medium		Influenza virus	94.4	135	4.9	7	0.7	1



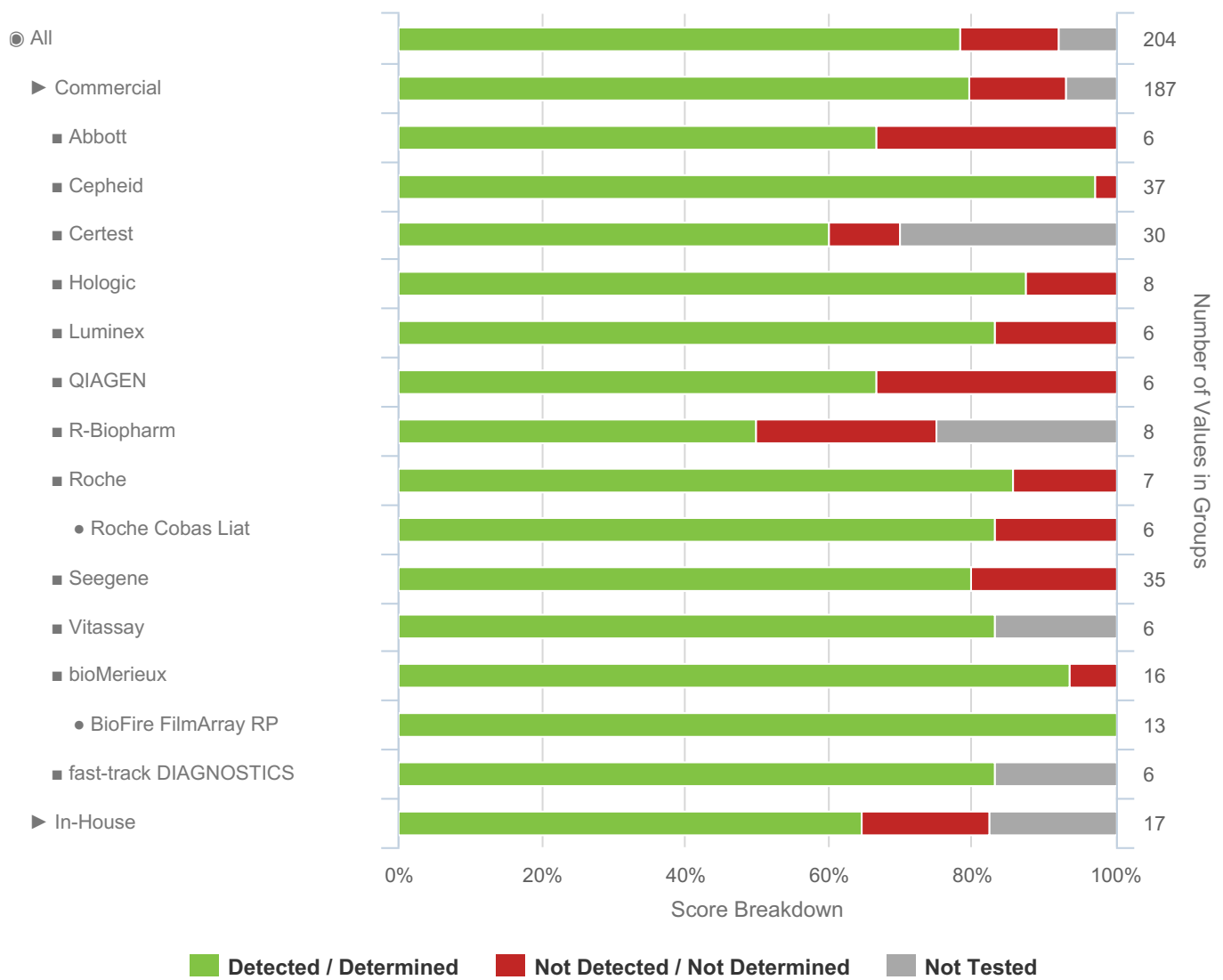
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Groups Rolled Up: Cepheid - Cepheid Xpert kit (n=37), Certest - Certest Real Time PCR (n=30), Hologic - Hologic Panther Fusion (n=8), R-Biopharm - R-Biopharm RIDA Gene (n=8), Seegene - Seegene Allplex (n=35), Vitassay - Vitassay Real-Time PCR (n=6), fast-track DIAGNOSTICS - FTD real time PCR (n=6), In-House - Real-time In-House PCR (n=17)

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Sample Code	Sample Content	Matrix	Sample Relationships	Expected targets	Detected / Determined		Not Detected / Not Determined		Not Tested	
					(%)	(n)	(%)	(n)	(%)	(n)
RESPI22S-03	Influenza virus A (H3N2)	Transport Medium	DS1_2	Influenza virus	86.7	124	12.6	18	0.7	1

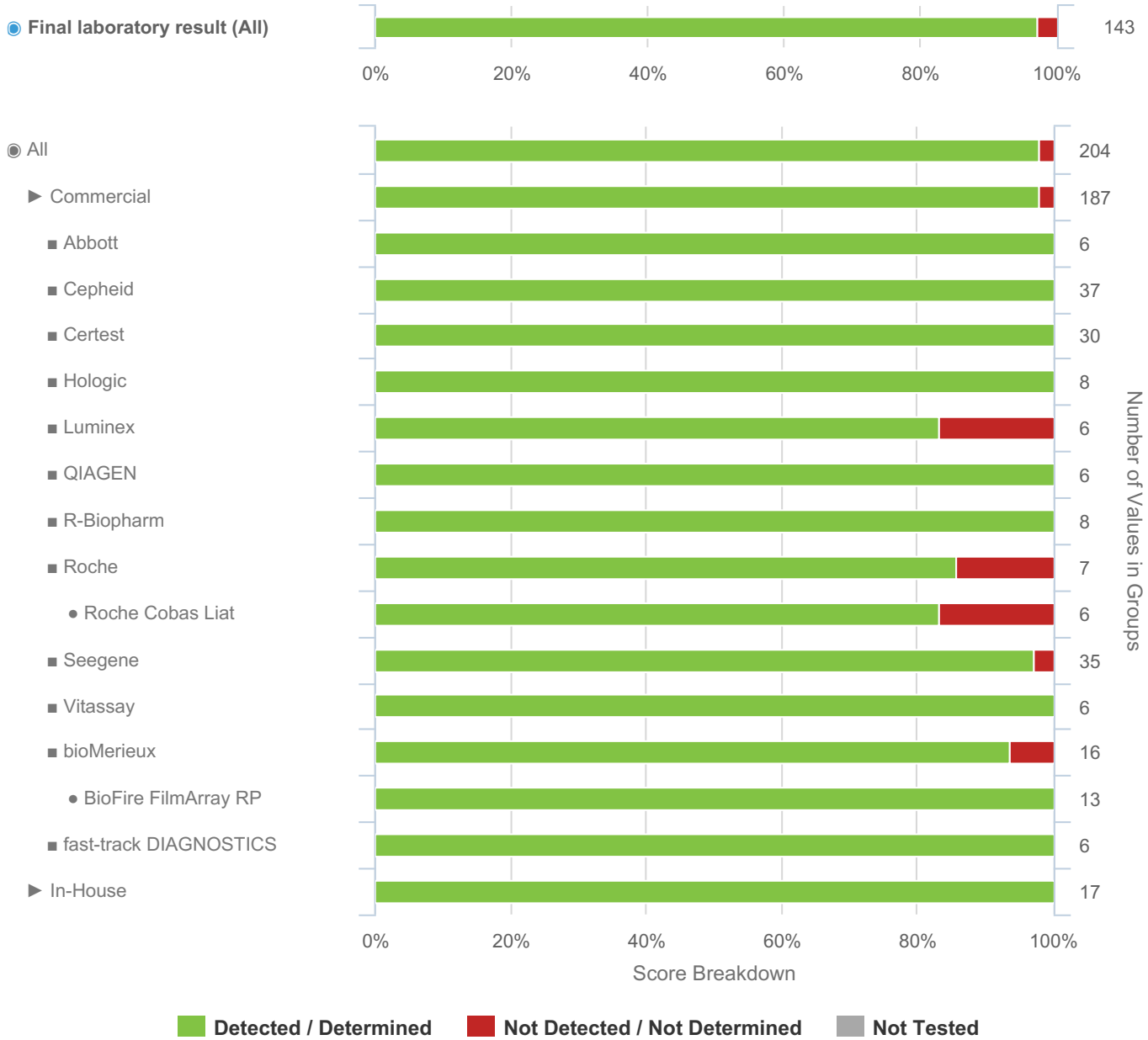



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Groups Rolled Up: Cepheid - Cepheid Xpert kit (n=37), Certest - Certest Real Time PCR (n=30), Hologic - Hologic Panther Fusion (n=8), R-Biopharm - R-Biopharm RIDA Gene (n=8), Seegene - Seegene Allplex (n=35), Vitassay - Vitassay Real-Time PCR (n=6), fast-track DIAGNOSTICS - FTD real time PCR (n=6), In-House - Real-time In-House PCR (n=17)

Sample Code	Sample Content	Matrix	Sample Relationships	Expected targets	Detected / Determined		Not Detected / Not Determined		Not Tested	
					(%)	(n)	(%)	(n)	(%)	(n)
RESPI22S-04	Negative	Transport Medium			97.2	139	2.8	4	N/A	0



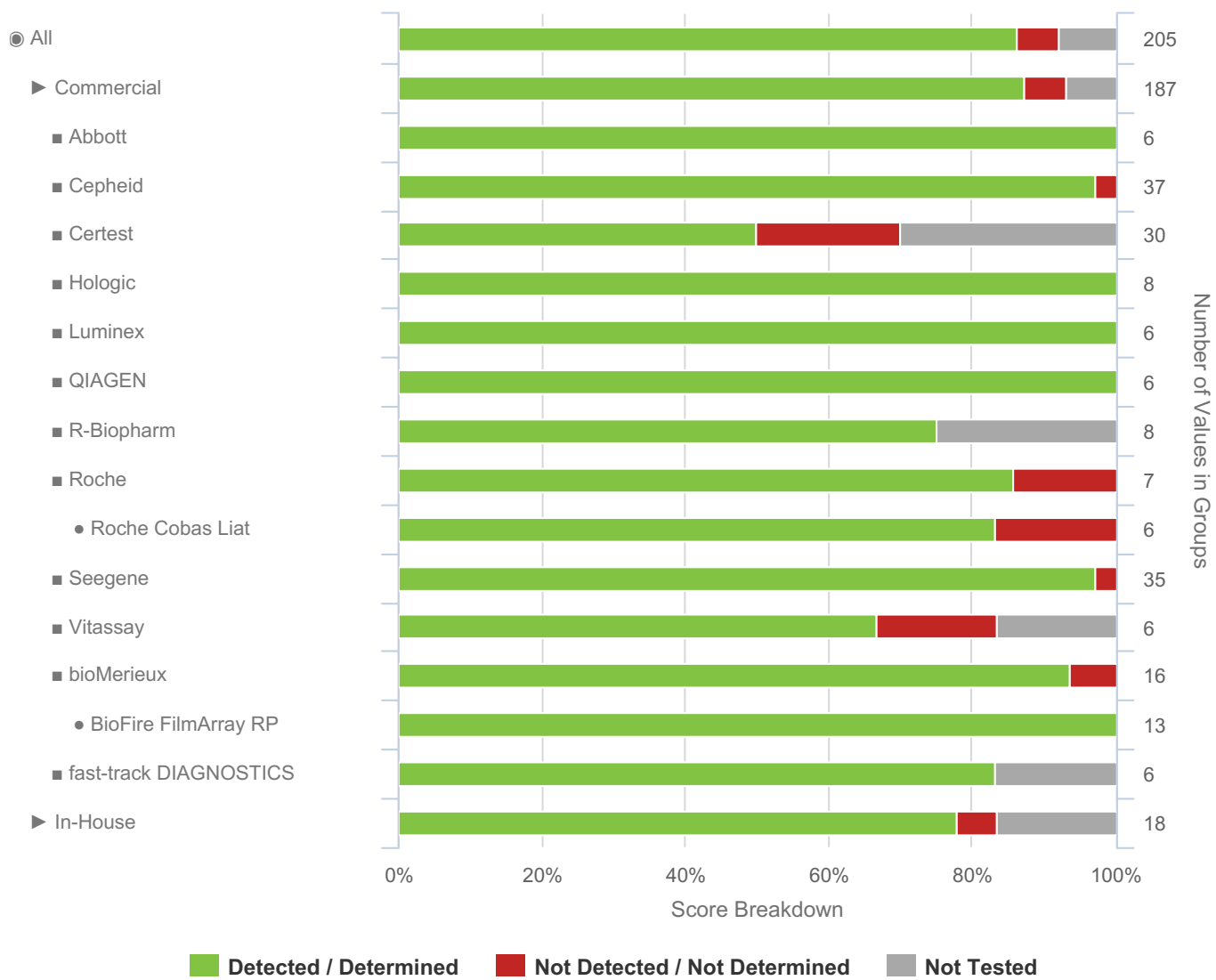
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					(%)	(n)	(%)	(n)	(%)	(n)
RESPI22S-05	Influenza virus B (Victoria)	Transport Medium	DS2_1	Influenza virus	97.9	140	1.4	2	0.7	1



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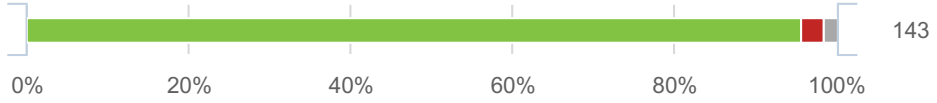
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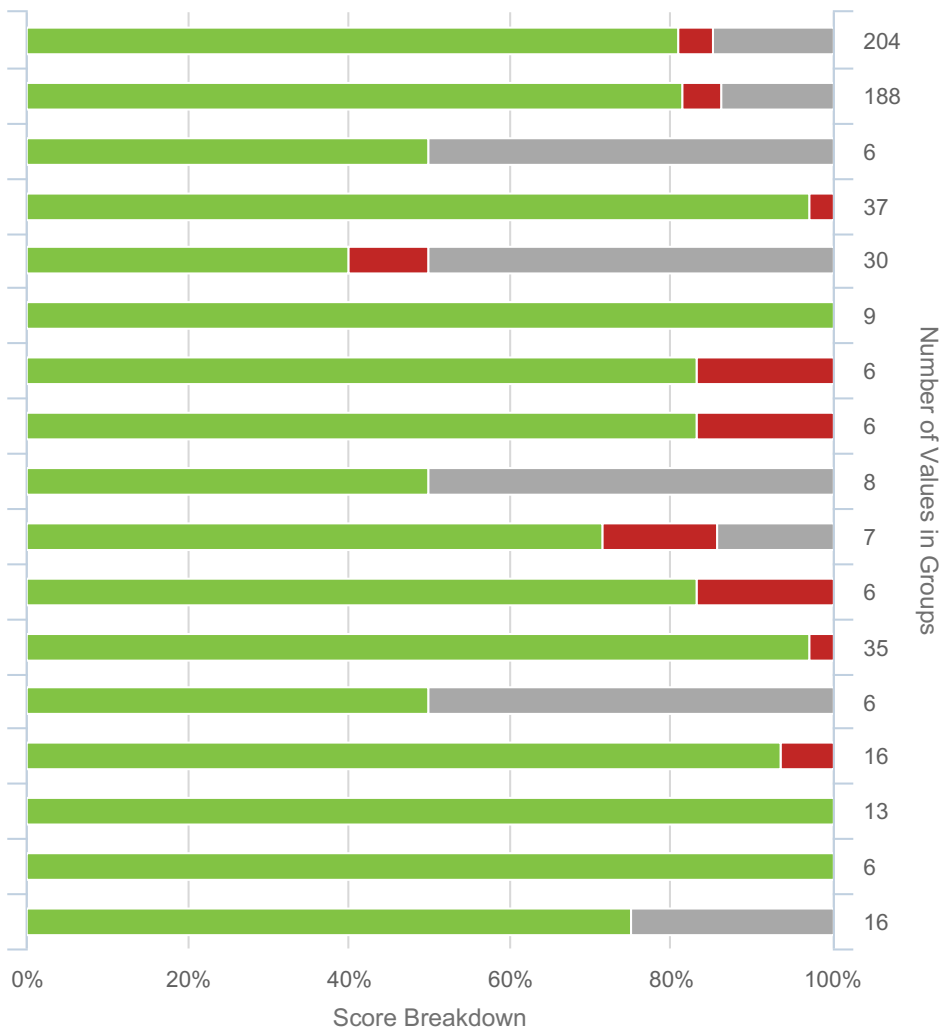
Sample Code	Sample Content	Matrix	Sample Relationships	Expected targets	Detected / Determined		Not Detected / Not Determined		Not Tested	
					(%)	(n)	(%)	(n)	(%)	(n)
RESPI22S-06	Respiratory Syncytial Virus (type B)	Transport Medium		Respiratory syncytial virus	95.8	137	2.8	4	1.4	2

Final laboratory result (All)




All

- ▶ Commercial
 - Abbott
 - Cepheid
 - Certest
 - Hologic
 - Luminex
 - QIAGEN
 - R-Biopharm
 - Roche
 - Roche Cobas Liat
 - Seegene
 - Vitassay
 - bioMerieux
 - BioFire FilmArray RP
 - fast-track DIAGNOSTICS
- ▶ In-House



■ Detected / Determined ■ Not Detected / Not Determined ■ Not Tested

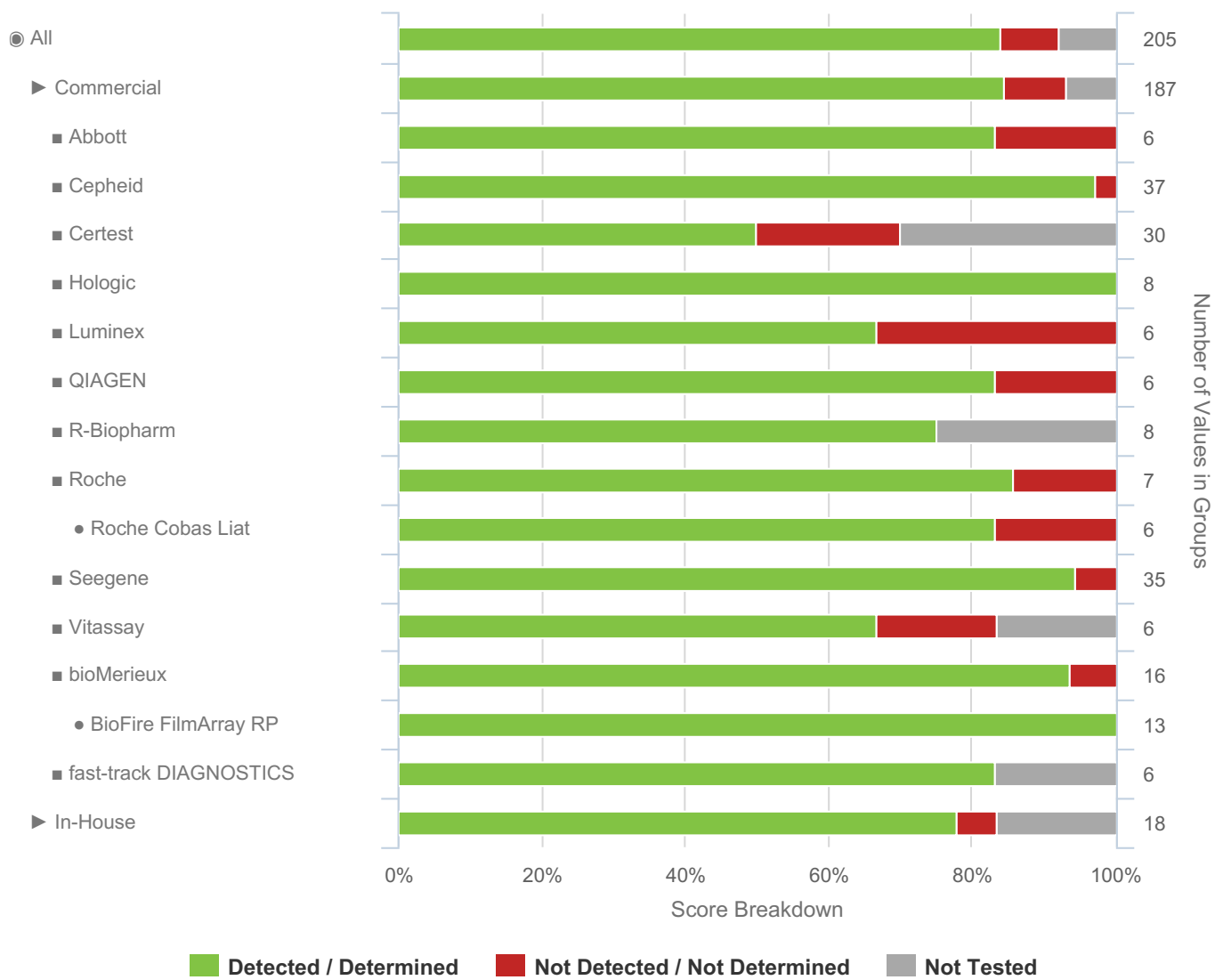
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Catalogue Code: QAV164188	Ref Code: RESPI22	Challenge: S	Analysis Type: Multiple Pathogen Qualitative	Dataset: -	Report UID: 0/0/4844	Laboratory


Groups below n=5: Abbott - Abbott Alinity m (n=3), Abbott - Abbott ID NOW (n=3), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics TandemPlex (n=2), DiaSorin (n=2), DiaSorin - DiaSorin Simplexa (n=2), GenMark Dx (n=3), GenMark Dx - GenMark DX ePlex (n=3), Genematrix (n=1), Genematrix - Genematrix NeoPlex RV-Panel A (n=1), Immundiagnostik (n=1), Immundiagnostik - Immundiagnostik MutaPLEX (n=1), Luminex - Luminex ARIES (n=2), Luminex - Luminex xTAG (n=3), Luminex - Verigene (n=1), PathoFinder (n=4), PathoFinder - PathoFinder Real Time PCR (n=4), QIAGEN - Qiagen NeuMoDx (n=3), QIAGEN - Qiagen QIAstat-Dx (n=3), Roche - Roche Cobas 6800/8800 (n=1), Speedx (n=1), Speedx - Speedx Real Time PCR (n=1), TRUPCR (n=1), TRUPCR - TRUPCR Respiratory (n=1), ThermoFisher (n=1), ThermoFisher - ThermoFisher TaqMan Array (n=1), bioMerieux - BioFire FilmArray PN (n=1), bioMerieux - bioMerieux Primers/Probes (n=1), bioMerieux - bioMerieux R-gene Kit (n=1)

Groups Rolled Up: Cepheid - Cepheid Xpert kit (n=37), Certest - Certest Real Time PCR (n=30), Hologic - Hologic Panther Fusion (n=9), R-Biopharm - R-Biopharm RIDA Gene (n=8), Seegene - Seegene Allplex (n=35), Vitassay - Vitassay Real-Time PCR (n=6), fast-track DIAGNOSTICS - FTD real time PCR (n=6), In-House - Real-time In-House PCR (n=16)

Catalogue Code: QAV164188	Ref Code: RESPI22	Challenge: S	Analysis Type: Multiple Pathogen Qualitative	Dataset: -	Report UID: 0/0/4844	Laboratory
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Sample Code	Sample Content	Matrix	Sample Relationships	Expected targets	Detected / Determined		Not Detected / Not Determined		Not Tested	
					(%)	(n)	(%)	(n)	(%)	(n)
RESPI22S-07	Influenza virus B (Victoria)	Transport Medium	DS2_2	Influenza virus	95.1	136	4.2	6	0.7	1



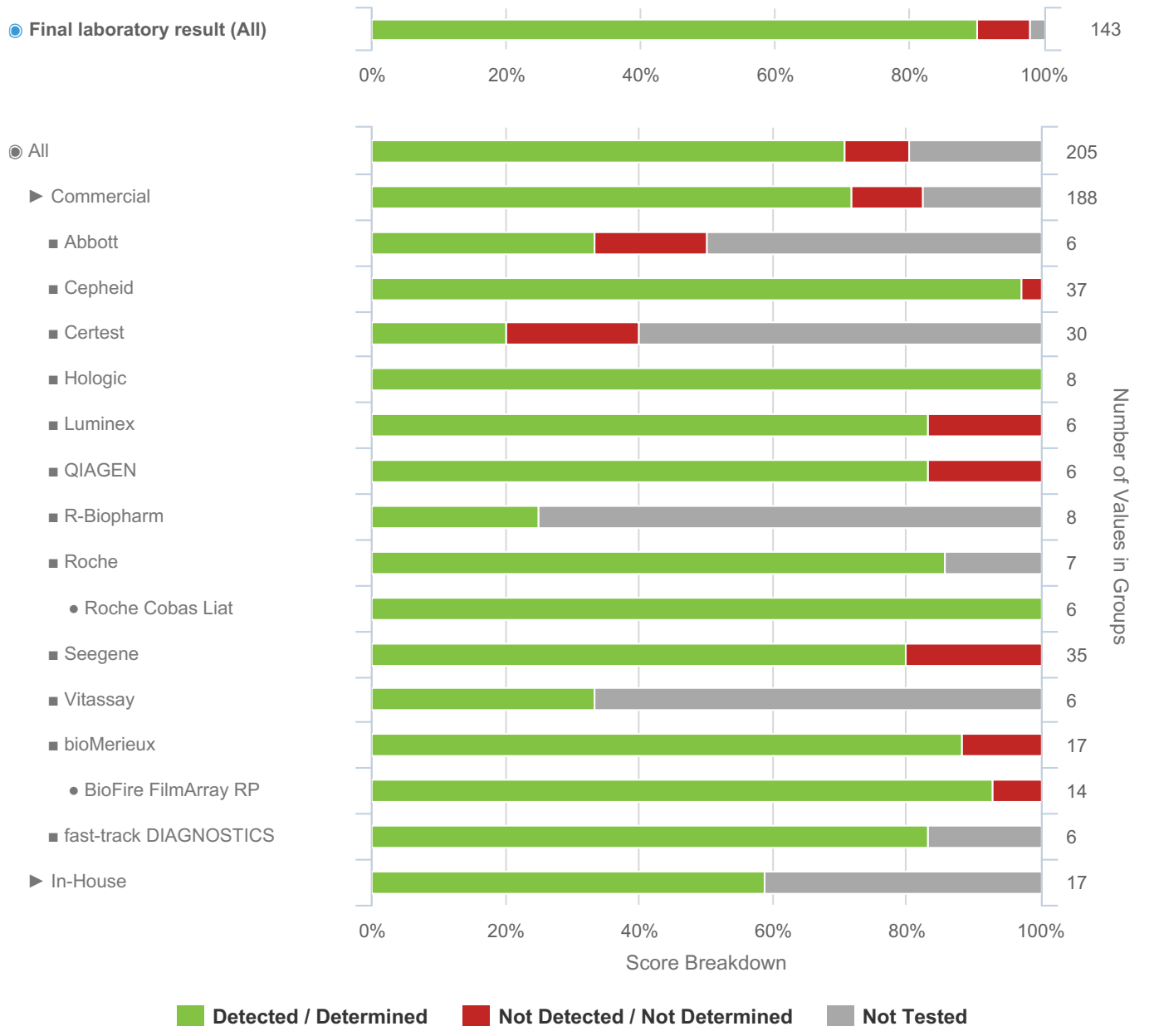
Individual Report		QCMD 2022 Respiratory I EQA Programme				
Catalogue Code: QAV164188	Ref Code: RESPI22	Challenge: S	Analysis Type: Multiple Pathogen Qualitative	Dataset: -	Report UID: 0/0/4844	Laboratory


Groups below n=5: Abbott - Abbott Alinity m (n=3), Abbott - Abbott ID NOW (n=3), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics TandemPlex (n=2), DiaSorin (n=2), DiaSorin - DiaSorin Simplexa (n=2), GenMark Dx (n=3), GenMark Dx - GenMark DX ePlex (n=3), Genematrix (n=1), Genematrix - Genematrix NeoPlex RV-Panel A (n=1), Immundiagnostik (n=1), Immundiagnostik - Immundiagnostik MutaPLEX (n=1), Luminex - Luminex ARIES (n=2), Luminex - Luminex xTAG (n=3), Luminex - Verigene (n=1), PathoFinder (n=4), PathoFinder - PathoFinder Real Time PCR (n=4), QIAGEN - Qiagen NeuMoDx (n=3), QIAGEN - Qiagen QIAstat-Dx (n=3), Roche - Roche Cobas 6800/8800 (n=1), Speedx (n=1), Speedx - Speedx Real Time PCR (n=1), TRUPCR (n=1), TRUPCR - TRUPCR Respiratory (n=1), ThermoFisher (n=1), ThermoFisher - ThermoFisher TaqMan Array (n=1), bioMerieux - BioFire FilmArray PN (n=1), bioMerieux - bioMerieux Primers/Probes (n=1), bioMerieux - bioMerieux R-gene Kit (n=1)

Groups Rolled Up: Cepheid - Cepheid Xpert kit (n=37), Certest - Certest Real Time PCR (n=30), Hologic - Hologic Panther Fusion (n=8), R-Biopharm - R-Biopharm RIDA Gene (n=8), Seegene - Seegene Allplex (n=35), Vitassay - Vitassay Real-Time PCR (n=6), fast-track DIAGNOSTICS - FTD real time PCR (n=6), In-House - Real-time In-House PCR (n=18)

Catalogue Code: QAV164188	Ref Code: RESPI22	Challenge: S	Analysis Type: Multiple Pathogen Qualitative	Dataset: -	Report UID: 0/0/4844	Laboratory
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Sample Code	Sample Content	Matrix	Sample Relationships	Expected targets	Detected / Determined		Not Detected / Not Determined		Not Tested	
					(%)	(n)	(%)	(n)	(%)	(n)
RESPI22S-08	Influenza (type A) & Respiratory Syncytial Virus (type A)	Transport Medium		Influenza virus and Respiratory syncytial virus	90.2	129	7.7	11	2.1	3



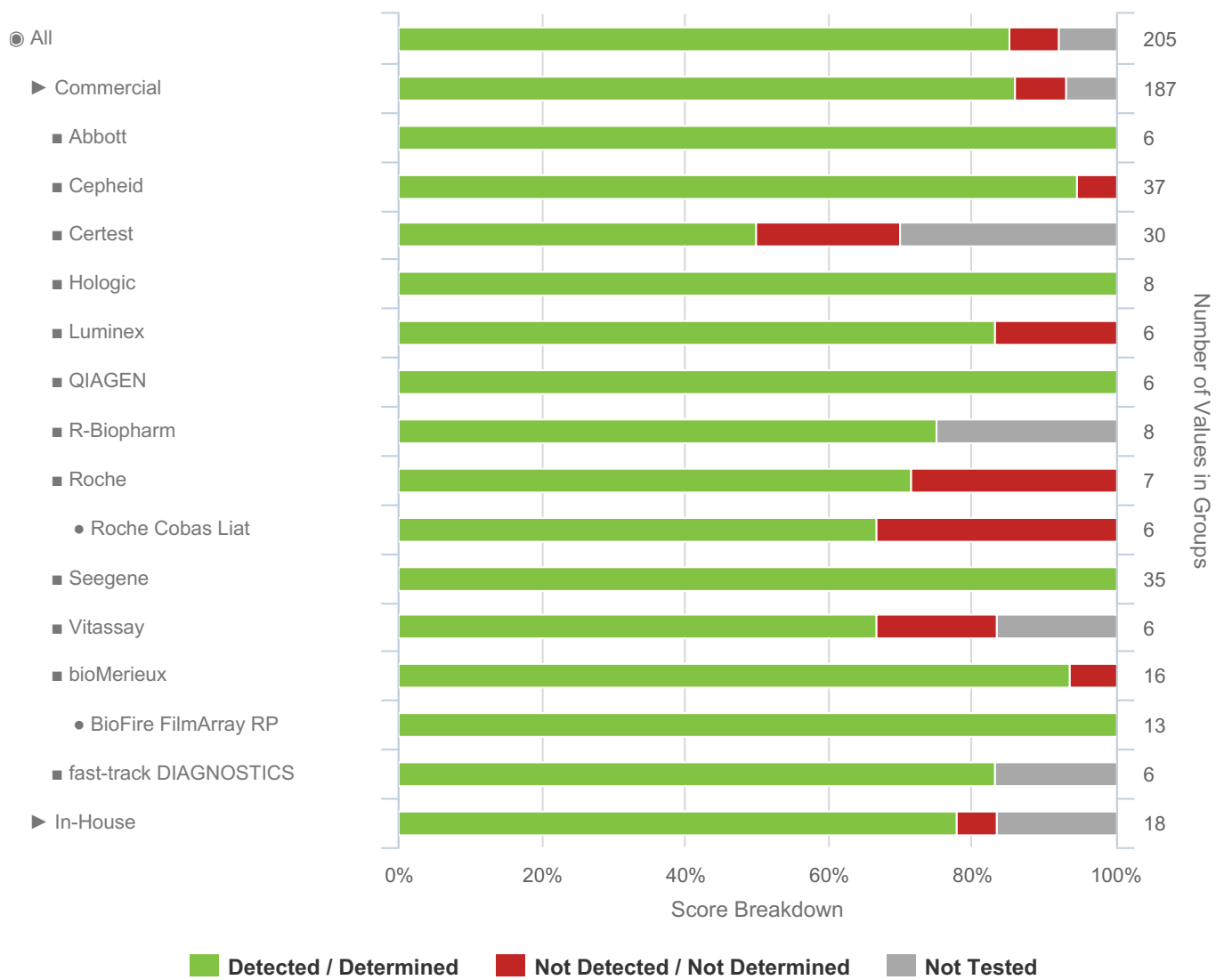
Individual Report		QCMD 2022 Respiratory I EQA Programme			 <small>Quality Control for Molecular Diagnostics</small>	
Catalogue Code: QAV164188	Ref Code: RESPI22	Challenge: S	Analysis Type: Multiple Pathogen Qualitative	Dataset: -	Report UID: 0/0/4844	Laboratory


Groups below n=5: Abbott - Abbott Alinity m (n=3), Abbott - Abbott ID NOW (n=3), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics TandemPlex (n=2), DiaSorin (n=2), DiaSorin - DiaSorin Simplexa (n=2), GenMark Dx (n=3), GenMark Dx - GenMark DX ePlex (n=3), Genematrix (n=1), Genematrix - Genematrix NeoPlex RV-Panel A (n=1), Immundiagnostik (n=1), Immundiagnostik - Immundiagnostik MutaPLEX (n=1), Luminex - Luminex ARIES (n=2), Luminex - Luminex xTAG (n=3), Luminex - Verigene (n=1), PathoFinder (n=4), PathoFinder - PathoFinder Real Time PCR (n=4), QIAGEN - Qiagen NeuMoDx (n=3), QIAGEN - Qiagen QIAstat-Dx (n=3), Roche - Roche Cobas 6800/8800 (n=1), Speedx (n=1), Speedx - Speedx Real Time PCR (n=1), TRUPCR (n=1), TRUPCR - TRUPCR Respiratory (n=1), ThermoFisher (n=1), ThermoFisher - ThermoFisher TaqMan Array (n=1), bioMerieux - BioFire FilmArray PN (n=1), bioMerieux - bioMerieux Primers/Probes (n=1), bioMerieux - bioMerieux R-gene Kit (n=1)

Groups Rolled Up: Cepheid - Cepheid Xpert kit (n=37), Certest - Certest Real Time PCR (n=30), Hologic - Hologic Panther Fusion (n=8), R-Biopharm - R-Biopharm RIDA Gene (n=8), Seegene - Seegene Allplex (n=35), Vitassay - Vitassay Real-Time PCR (n=6), fast-track DIAGNOSTICS - FTD real time PCR (n=6), In-House - Real-time In-House PCR (n=17)

Catalogue Code: QAV164188	Ref Code: RESPI22	Challenge: S	Analysis Type: Multiple Pathogen Qualitative	Dataset: -	Report UID: 0/0/4844	Laboratory
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Sample Code	Sample Content	Matrix	Sample Relationships	Expected targets	Detected / Determined		Not Detected / Not Determined		Not Tested	
					(%)	(n)	(%)	(n)	(%)	(n)
RESPI22S-09	Influenza virus B (Yamagata)	Transport Medium		Influenza virus	97.2	139	2.1	3	0.7	1



Individual Report		QCMD 2022 Respiratory I EQA Programme			 <small>Quality Control for Molecular Diagnostics</small>	
Catalogue Code: QAV164188	Ref Code: RESPI22	Challenge: S	Analysis Type: Multiple Pathogen Qualitative	Dataset: -	Report UID: 0/0/4844	Laboratory

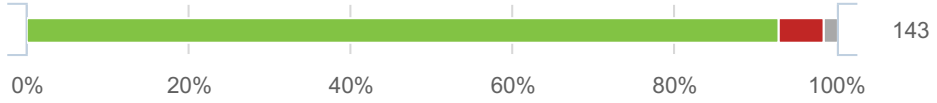
Groups below n=5: Abbott - Abbott Alinity m (n=3), Abbott - Abbott ID NOW (n=3), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics TandemPlex (n=2), DiaSorin (n=2), DiaSorin - DiaSorin Simplexa (n=2), GenMark Dx (n=3), GenMark Dx - GenMark DX ePlex (n=3), Genematrix (n=1), Genematrix - Genematrix NeoPlex RV-Panel A (n=1), Immundiagnostik (n=1), Immundiagnostik - Immundiagnostik MutaPLEX (n=1), Luminex - Luminex ARIES (n=2), Luminex - Luminex xTAG (n=3), Luminex - Verigene (n=1), PathoFinder (n=4), PathoFinder - PathoFinder Real Time PCR (n=4), QIAGEN - Qiagen NeuMoDx (n=3), QIAGEN - Qiagen QIAstat-Dx (n=3), Roche - Roche Cobas 6800/8800 (n=1), Speedx (n=1), Speedx - Speedx Real Time PCR (n=1), TRUPCR (n=1), TRUPCR - TRUPCR Respiratory (n=1), ThermoFisher (n=1), ThermoFisher - ThermoFisher TaqMan Array (n=1), bioMerieux - BioFire FilmArray PN (n=1), bioMerieux - bioMerieux Primers/Probes (n=1), bioMerieux - bioMerieux R-gene Kit (n=1)

Groups Rolled Up: Cepheid - Cepheid Xpert kit (n=37), Certest - Certest Real Time PCR (n=30), Hologic - Hologic Panther Fusion (n=8), R-Biopharm - R-Biopharm RIDA Gene (n=8), Seegene - Seegene Allplex (n=35), Vitassay - Vitassay Real-Time PCR (n=6), fast-track DIAGNOSTICS - FTD real time PCR (n=6), In-House - Real-time In-House PCR (n=18)

Catalogue Code: QAV164188	Ref Code: RESPI22	Challenge: S	Analysis Type: Multiple Pathogen Qualitative	Dataset: -	Report UID: 0/0/4844	Laboratory
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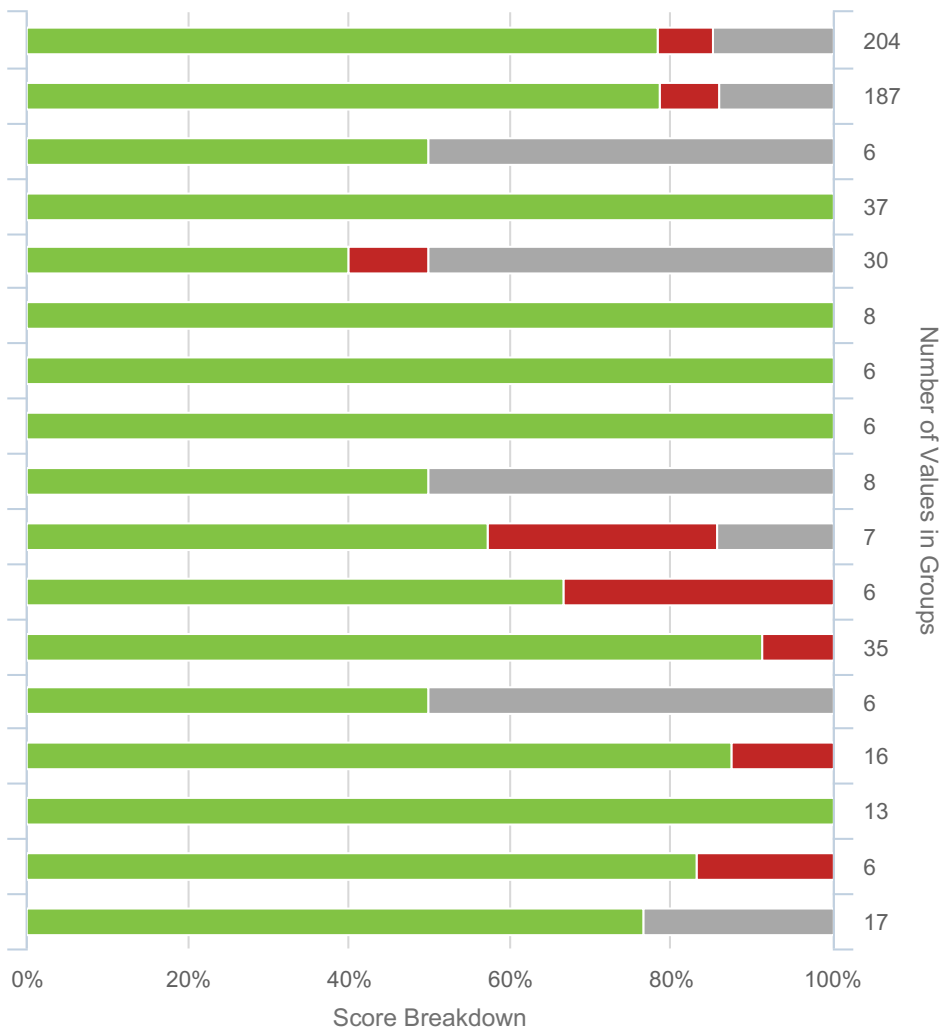
Sample Code	Sample Content	Matrix	Sample Relationships	Expected targets	Detected / Determined		Not Detected / Not Determined		Not Tested	
					(%)	(n)	(%)	(n)	(%)	(n)
RESPI22S-10	Respiratory Syncytial Virus (type A)	Transport Medium		Respiratory syncytial virus	93	133	5.6	8	1.4	2

Final laboratory result (All)




All

- ▶ Commercial
- Abbott
- Cepheid
- Certest
- Hologic
- Luminex
- QIAGEN
- R-Biopharm
- Roche
- Roche Cobas Liat
- Seegene
- Vitassay
- bioMerieux
- BioFire FilmArray RP
- fast-track DIAGNOSTICS
- ▶ In-House



■ Detected / Determined
 ■ Not Detected / Not Determined
 ■ Not Tested

Individual Report		QCMD 2022 Respiratory I EQA Programme			 <small>Quality Control for Molecular Diagnostics</small>	
Catalogue Code: QAV164188	Ref Code: RESPI22	Challenge: S	Analysis Type: Multiple Pathogen Qualitative	Dataset: -	Report UID: 0/0/4844	Laboratory

Groups below n=5: Abbott - Abbott Alinity m (n=3), Abbott - Abbott ID NOW (n=3), AusDiagnostics (n=2), AusDiagnostics - AusDiagnostics TandemPlex (n=2), DiaSorin (n=2), DiaSorin - DiaSorin Simplexa (n=2), GenMark Dx (n=3), GenMark Dx - GenMark DX ePlex (n=3), Genematrix (n=1), Genematrix - Genematrix NeoPlex RV-Panel A (n=1), Immundiagnostik (n=1), Immundiagnostik - Immundiagnostik MutaPLEX (n=1), Luminex - Luminex ARIES (n=2), Luminex - Luminex xTAG (n=3), Luminex - Verigene (n=1), PathoFinder (n=4), PathoFinder - PathoFinder Real Time PCR (n=4), QIAGEN - Qiagen NeuMoDx (n=3), QIAGEN - Qiagen QIAstat-Dx (n=3), Roche - Roche Cobas 6800/8800 (n=1), Speedx (n=1), Speedx - Speedx Real Time PCR (n=1), TRUPCR (n=1), TRUPCR - TRUPCR Respiratory (n=1), ThermoFisher (n=1), ThermoFisher - ThermoFisher TaqMan Array (n=1), bioMerieux - BioFire FilmArray PN (n=1), bioMerieux - bioMerieux Primers/Probes (n=1), bioMerieux - bioMerieux R-gene Kit (n=1)

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