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EXTERNAL  
QUALITY  
ASSESSMENT



INTERNAL  
QUALITY  
CONTROL



REFERENCE  
MEASUREMENT  
SERVICES



EDUCATION &  
TRAINING

# Weqas

GLOBAL PROVIDER OF QUALITY  
IN DIAGNOSTIC MEDICINE

## POCT Report and Interpretation

Samantha Jones & Nicola Blount

# EQA – key elements in selecting EQA providers

- ✓ Design for clinical need
- ✓ Material & frequency
- ✓ Data analysis – target value
- ✓ Analytical performance specification
- ✓ Reports
- ✓ Education element
- ✓ Troubleshooting support

# Weqas POCT Programmes

## Cuesee

POCT Glucose & Ketones

POCT Urinalysis

## Weqas Connect

POCT INR

POCT Lipids

POCT Bilirubin

POCT Blood Gas

POCT BNP

POCT Cardiac Markers

POCT Co-oximetry

POCT Creatinine

POCT CRP

POCT D-dimer

POCT Haemoglobin

POCT HbA1c

POCT HIV

POCT Influenza & RSV

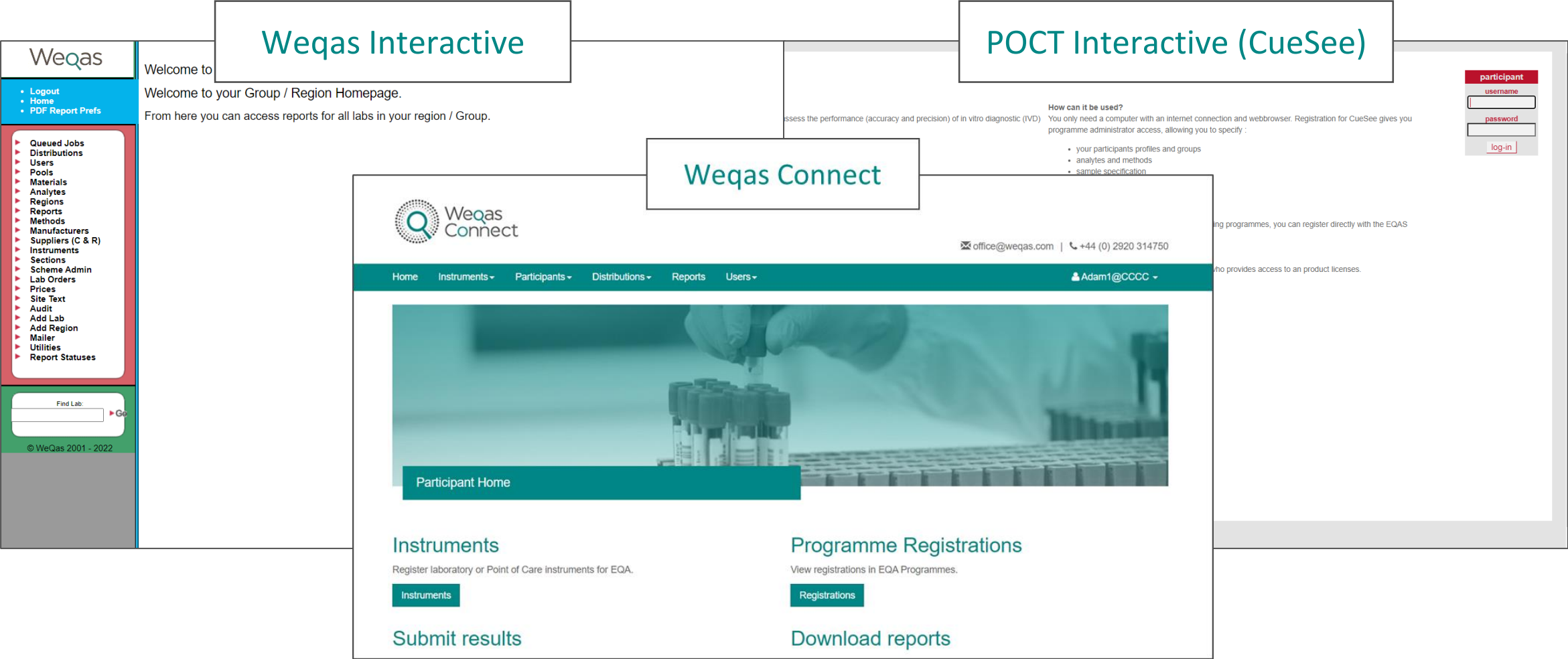
Pre-term Labour Markers

POCT Pregnancy Testing

Urine Drugs of Abuse

Viscoelastic Haemostasis

# Current EQA Web Portals



# Harmonised Performance Scoring

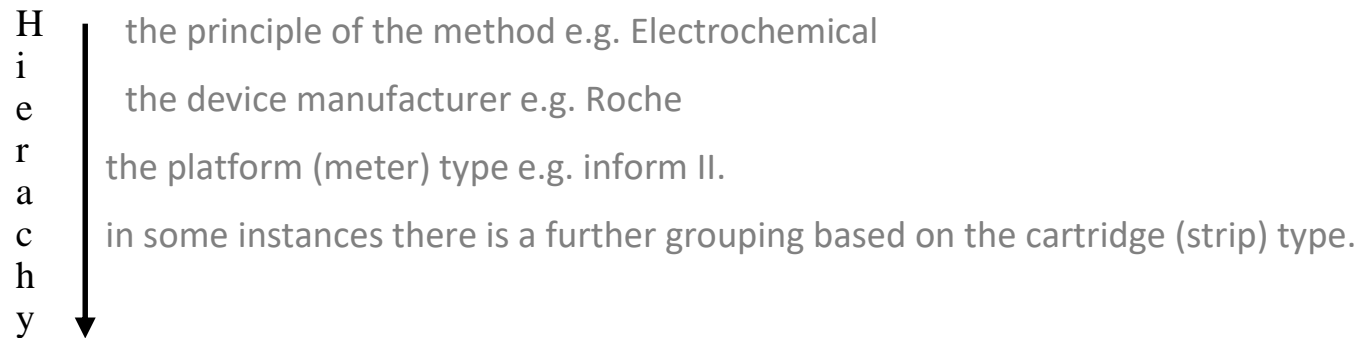
<b>less than 1</b>	Good - all points within $\pm 1$ SD	
<b>1 – 2</b>	Acceptable	
<b>greater than 2</b>	Unacceptable	

# Cuesee Target Value Assignment

Any gross outliers such as transcription, transposition or unit errors are minimised by using a minimum and maximum allowable value (in default unit) that can be entered for a particular sample via the website. This range spans approximately  $\pm 4$  SD from the indicative target value calculated using laboratory methods. Participants will not be allowed to enter results outside this range and will be prompted to contact the Group Administrator for assistance.

For most POCT Programmes assessment is against a peer group target value

Methods are classified into categories based on:



Assessment against the higher order is preferred, however, where there are known differences between the different devices, the most appropriate peer group is selected.

**Target value**

Peer group Median – this is the default target value and is used in the majority of Programmes.

Peer group Mean – this is also calculated and can be selected as the target value.

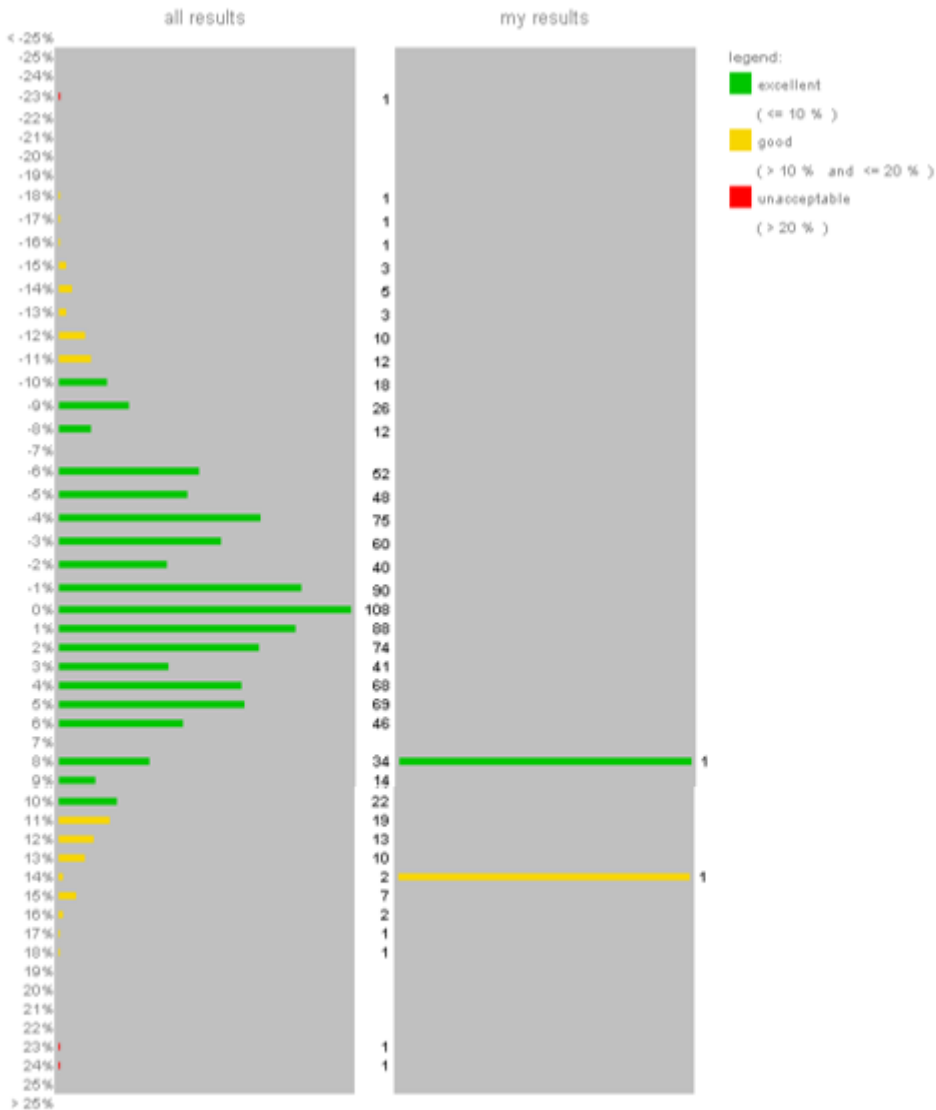
For each analyte for each sample the peer group mean and standard deviation is calculated.

$$SD = \sqrt{\frac{\sum (x - \bar{x})^2}{(n-1)}}$$

Where  $\bar{x}$  = arithmetic mean.

# Cuesee Reports – Standard Report

Report Settings		All Results		My Results	
sample	Weqas 0711	n	1079	n	2
analyte	glu	minimum	7.20	minimum	10.00
reporting in	mmol/l	maximum	11.50	maximum	10.60
deviation	relative (resolution 1%)	average	9.34	average	10.30
reference method	median	median	9.30	median	10.30
reference value	9.30 mmol/l	SD	0.50	SD	0.40
comparison	all results and my results	CV	5.9 %	CV	4.1%



my results

#	instrument	instrument ID	result ID	result
1	Statstrip Glucose (Connectivity)	00002010137	0310270249	10.00
2	Statstrip Glucose (Connectivity)	140002310071	0310270249	10.60

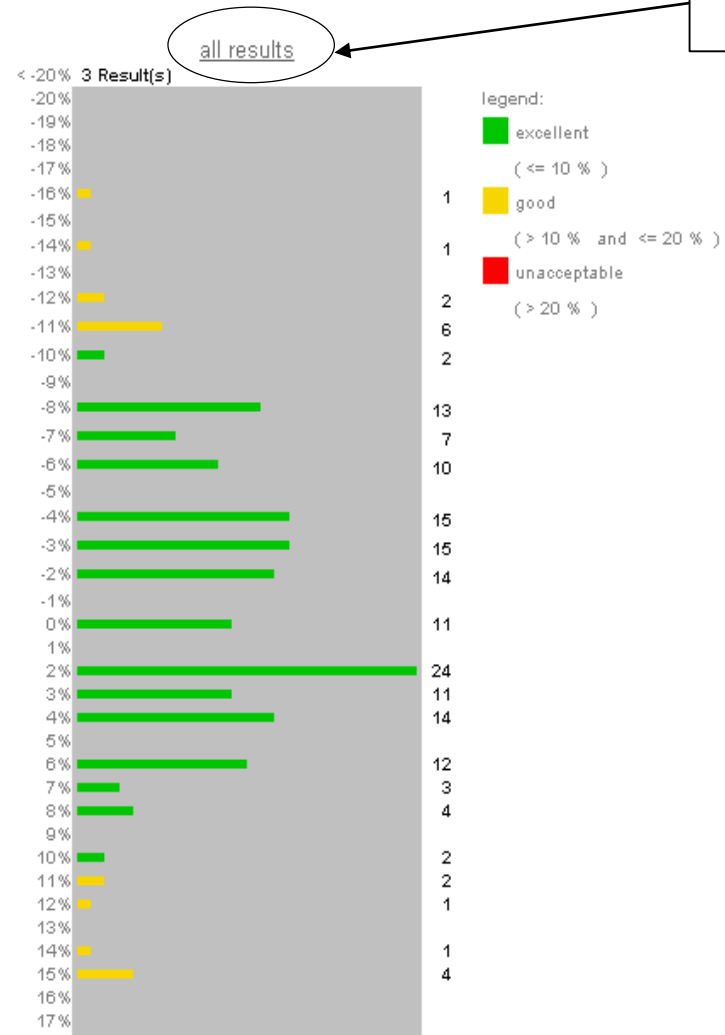


# Cuesee Reports – Managers / Co-ordinators Report

Hyperlinks will take you to further information.

**all results - histogram**  
QUALITY LABORATORY -

report settings		all results	
sample	WEQAS 0110	n	182
analyte	glu	minimum	4.00
reporting in	mmol/l	maximum	11.60
deviation	relative (resolution 1%)	average	7.56
reference method	median	median	7.60
reference value	7.60 mmol/l	SD	0.70
calibration	standard	CV	9.2 %

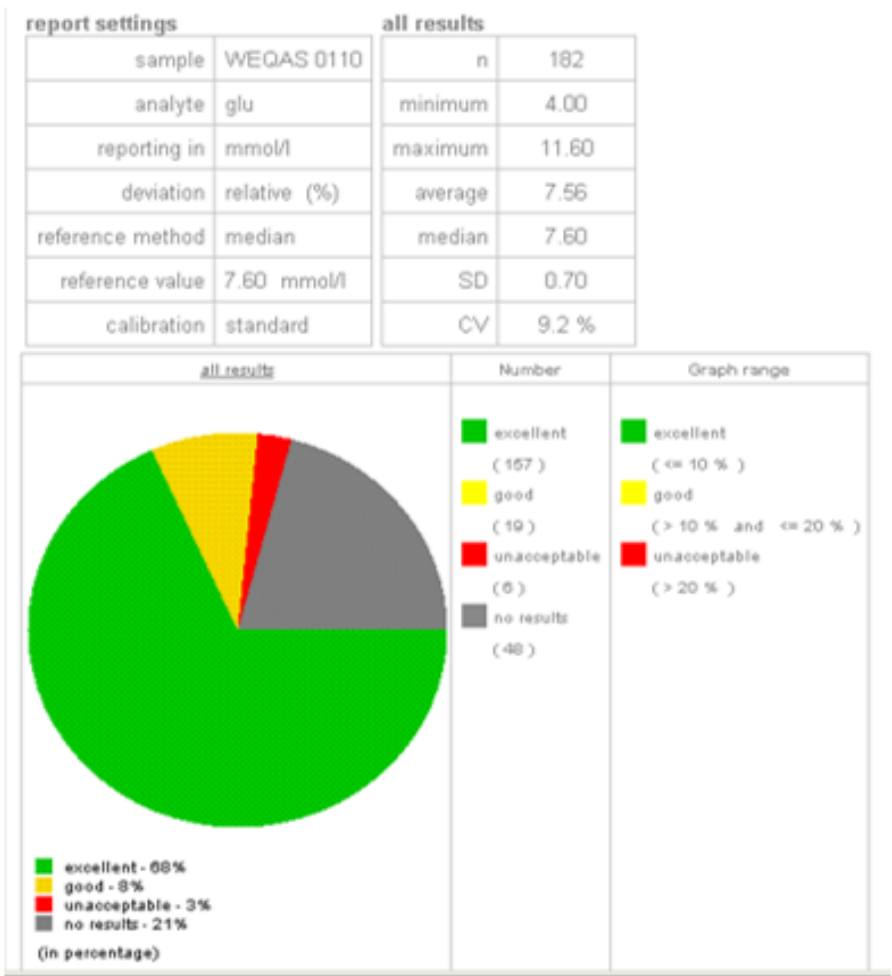


# Tabular data behind hyperlink

instrument	instrument ID	result ID	result	participant contact person	deviation from reference value		ranking
Accu-Check Advantage III	8548307546	0110	4.00	OPD Suite	-48 %	unacceptable	181.0
Accu-Check Advantage III	8547374420	0110	4.80	Paeds Ward	-37 %	unacceptable	180.0
Accu-Check Advantage III	8548421221	0110	4.90	Endoscopy Unit	-36 %	unacceptable	179.0
Accu-Check Advantage III	8541456251	0110	6.40	THEATRE 4	-16 %	good	175.0
Accu-Check Advantage III	8542451290	0110	6.60	A4 South	-14 %	good	169.5
Accu-Check Advantage III	8547261406	0110	6.70	Teenage Cancer Unit	-12 %	good	167.0
Accu-Check Advantage III	8549087265	0110	6.70	Ward 6 Nursing Staff	-12 %	good	167.0
Accu-Check Advantage III	8542487205	0110	6.80	THEATRE	-11 %	good	161.5
Accu-Check Advantage III	8546631983	0110	6.80	WARD	-11 %	good	161.5

Poor Performing sites can be easily reviewed using this table. Unacceptable sites will be shown in red text.

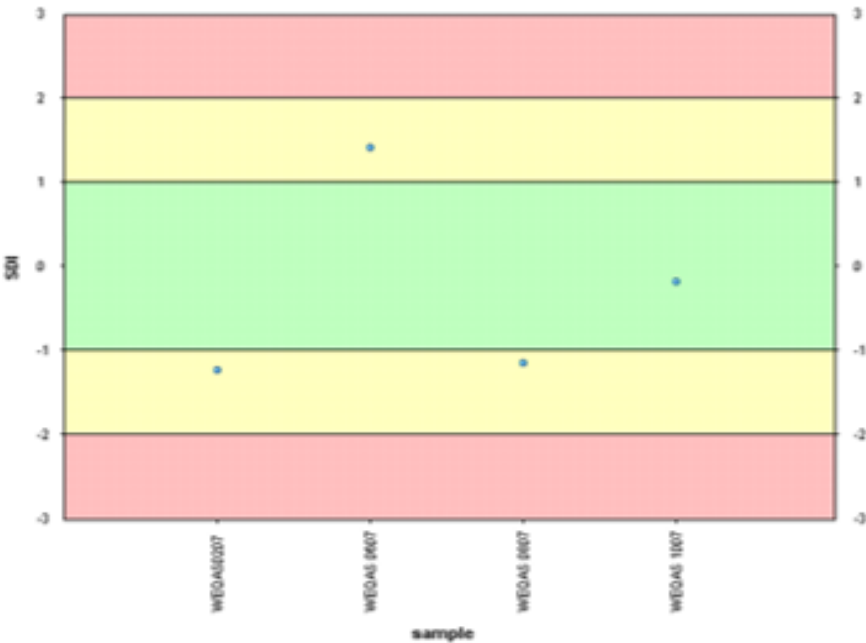
# Cuesee Reports – Managers / Co-ordinators Report



# No Results Booked Summary (Non Compliance Report)

Lipid Programme Dist WEQAS 1216			
contact person	institute	instrument ID	instrument
1 Pharmacist in charge	Store 707	SN920967	BHR - Cardiochek Chol & Glucose
		SN920967	BHR - Cardiochek Chol, HDL & Glucose
2 Pharmacist in charge	Store 323	2014655	BHR - Cardiochek Chol & Glucose
		3019318	BHR - Cardiochek Chol & Glucose
		2014655	BHR - Cardiochek Chol, HDL & Glucose
		3019318	BHR - Cardiochek Chol, HDL & Glucose
3 Pharmacist in charge	Store 61	920503	BHR - Cardiochek Chol & Glucose
		920503	BHR - Cardiochek Chol, HDL & Glucose

# Cumulative graph



report settings  
sample date range 1/1/2007 - 1/1/2008  
your instrument(s) all instruments  
reference value median  
instruments all instruments  
principles all principles  
families all families  
countries all countries  
groups all

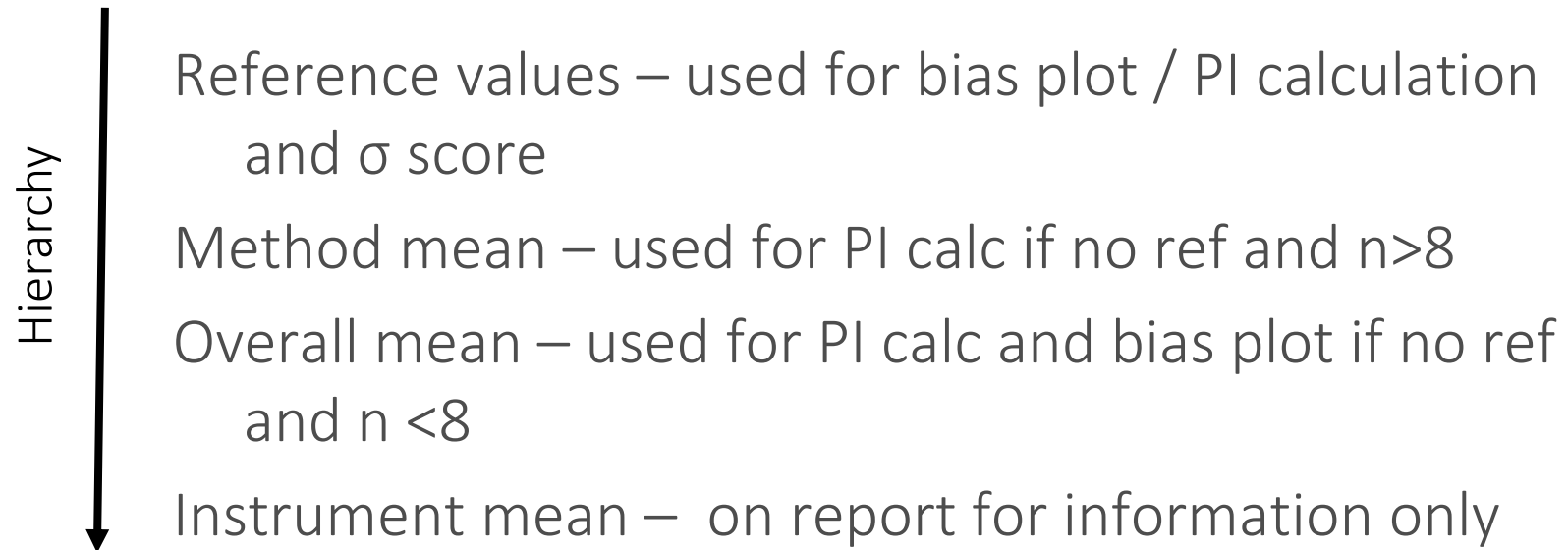
individual results					
	reference	your	bias	SD	SDI
	mmol/l	result	absolute	mmol/l	
WEQAS0207	6.42	5.72	-0.700	0.566	-1.2
WEQAS 0607	3.20	3.60	0.400	0.284	1.4
WEQAS 0807	3.74	3.19	-0.550	0.479	-1.1
WEQAS 1007	4.95	4.81	-0.140	0.776	-0.2

# Design - Why multiple samples are important

- Identifies components of both Imprecision and Inaccuracy
- Identifies systematic errors
- Assesses method linearity – required for ISO 15189
- Powerful error detection tool

# Weqas Interactive

## Target values used in Statistical Analysis



# Harmonised Performance Scoring

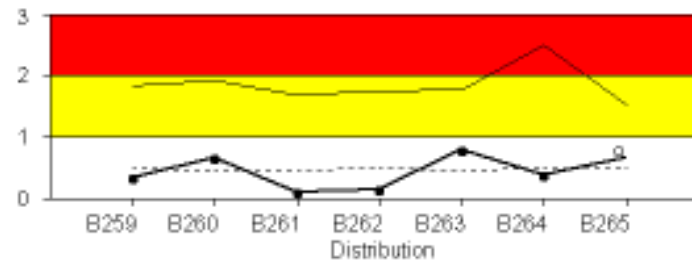
<b>less than 1</b>	Good - all points within $\pm 1$ SD	
<b>1 – 2</b>	Acceptable	
<b>greater than 2</b>	Unacceptable	



# Weqas Interactive Simplified Reports

Distribution Date: 4/01/16. Final. Report Issued: 25/01/16

This Distribution	
Overall Lab SDI:	0.65
Median All Laboratory:	0.49
97.5th centile:	1.52



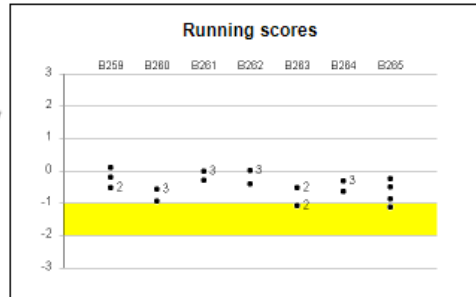
All SDI Ranges	
< 1	Good
1 - 2	Acceptable
> 2	Poor

## Section SDI scores for this distribution

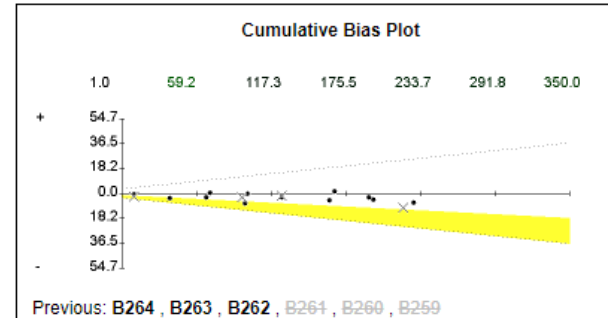
Section	Biochemistry Cobas
Overall	0.65
Total Bilirubin	0.65

Scheme: Bilirubin. Distribution Code: B265. Distribution Date: 4/01/16. Final Report Issued: 25/01/16				
Total Bilirubin (µmol/L)	Sample 1	Sample 2	Sample 3	Sample 4
Reported Result	124.0	8.0	210.0	92.0
Overall Mean	133.40	12.06	234.71	101.19
Method Mean: Roche Diazo	125.67	10.33	220.74	95.04
Instrument Mean: Cobas C Module	125.46	10.23	220.45	94.87
Your results are compared against	125.67	10.33	220.74	95.04

Good
Acceptable
Unacceptable Discuss with QC officer



Colour coded graphical representation of Individual SDI scores over 6 month time frame



6 month time frame of bias plot across concentration range.

Cumulative Submitted results

	B259	B260	B261	B262	B263	B264	B265
Sample 1	148.0	168.0	39.0	70.0	158.0	64.0	124.0
Sample 2	231.0	72.0	235.0	168.0	90.0	123.0	8.0
Sample 3	48.0	261.0	11.0	99.0	35.0	10.0	210.0
Sample 4	189.0	39.0	130.0	193.0	222.0	191.0	92.0

Key	
?	Analyte enrolled but no results returned
N/A	Not enrolled for this analyte

### ⚠ Performance Alert

Your results show a consistent bias over an extended period. Please discuss your results with your EQA Officer.

Performance alert based on Westgard rules

The following rules are used for the Performance alert.



At least 1 result with  $SDI > 3$



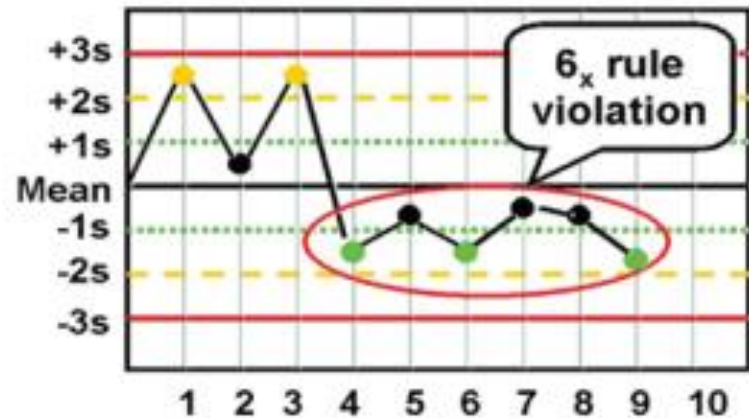
At least 2 results with  $SDI > 2$

If number of samples in current distribution  $\geq 2$  then applies to this dist only.

If number of samples in current = 1 then applies to this and at least one in previous distribution.

### Consistent Bias in one direction.

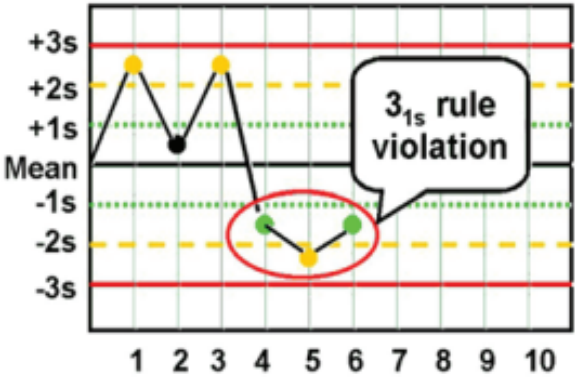
This is calculated over several distributions and will depend on the number of samples distributed per round e.g. if 3 samples are distributed per round, then the rule will be calculated over 2 distributions.



# samples in current with analyte	Number of samples overall to consider	trigger
1	6 samples 6 distributions	6 <sub>x</sub> : 6 samples with SDI > 0.2 <i>or</i> 6 samples with SDI < -0.2
2	6 samples 3 distributions	6 <sub>x</sub> : 6 samples with SDI > 0.2 <i>or</i> 6 samples with SDI < -0.2
3	6 samples 2 distributions	6 <sub>x</sub> : 6 samples with SDI > 0.2 <i>or</i> 6 samples with SDI < -0.2
4	8 samples 2 distributions	8 <sub>x</sub> : 8 samples with SDI > 0.2 <i>or</i> 8 samples with SDI < -0.2
5	10 samples 2 distributions	10 <sub>x</sub> : 10 samples with SDI > 0.2 <i>or</i> 10 samples with SDI < -0.2
6	6 samples 1 distribution	6 <sub>x</sub> : 6 samples with SDI > 0.2 <i>or</i> 6 samples with SDI < -0.2

At least 3 results > 1 SDI.

This is calculated over several distributions and will depend on the number of samples distributed per round e.g. if only 1 sample is distributed per round, then the rule will be calculated over 3 distributions.

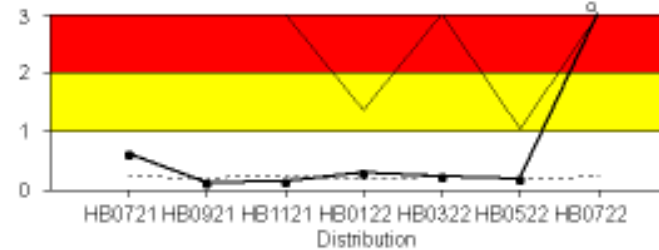


# samples in current with analyte	Number of samples overall to consider	trigger
1	3 samples 3 distributions	3 <sub>1s</sub> : 3 samples with SDI > 1 <i>or</i> 3 samples with SDI < -1
2	4 samples 2 distributions	4 <sub>1s</sub> : 4 samples with SDI > 1 <i>or</i> 4 samples with SDI < -1
3	3 samples 1 distributions	3 <sub>1s</sub> : 3 samples with SDI > 1 <i>or</i> 3 samples with SDI < -1
4	4 samples 1 distributions	4 <sub>1s</sub> : 4 samples with SDI > 1 <i>or</i> 4 samples with SDI < -1
5 or more	All in current 1 distribution	4 <sub>1s</sub> : 4 samples with SDI > 1 <i>or</i> 4 samples with SDI < -1

# Simplified Reports – poor performance example

Lab: Scheme: POCT Haemoglobin. Distribution Code: HB0722.  
Distribution Date: 5/07/22. Final. Report Issued: 29/07/22

This Distribution	
Overall Lab SDI:	7.06
Median All Laboratory:	0.21
97.5th centile:	7.06



All SDI Ranges	
< 1	Good
1 - 2	Acceptable
> 2	Poor

---- Median —●— Lab SDI — 97.5th

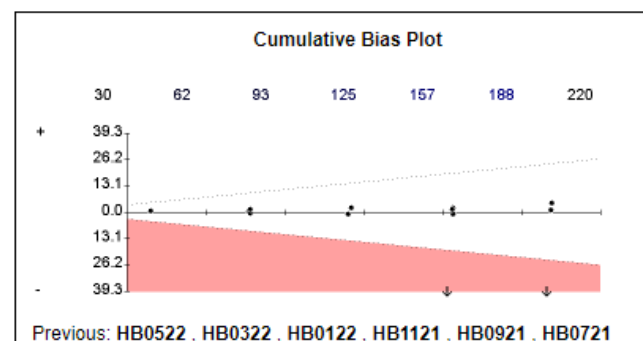
## Section SDI scores for this distribution

Section		
Overall	** 7.06	** 7.06
Haemoglobin	** 7.06	** 7.06

Scheme: POCT Haemoglobin. Distribution Code: HB0722.  
Distribution Date: 5/07/22. Final Report Issued: 29/07/22

Haemoglobin (g/L)	Sample 1	Sample 2
Reported Result	80	119
Overall Mean	198.5	158.8
Method Mean: HemoCue 201+ / DM	198.5	158.8
Instrument Mean: HemoCue 201+	198.5	158.8
Your results are compared against	198.5	158.8

Good
Acceptable
Unacceptable Discuss with QC officer



#### Cumulative Submitted results

	HB0721	HB0921	HB1121	HB0122	HB0322	HB0522	HB0722
Sample 1	81	201	40	122	163	79	80
Sample 2	162	79	160	205	40	118	119

Key	
?	Analyte enrolled but no results returned
N/A	Not enrolled for this analyte

#### ⚠ Performance Alert

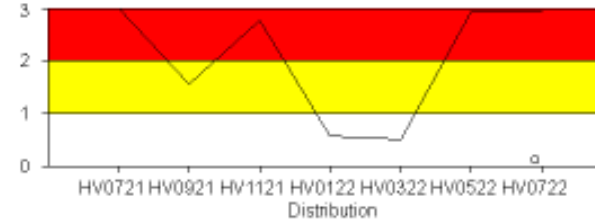
At least 1 sample in the current distribution with a very poor score -  $|SDI| > 3$

At least 2 samples in current/prior distribution with poor score -  $|SDI| > 2$

# Weqas Interactive Qualitative Reports

Distribution Date: 26/07/22. Final. Report Issued: 26/08/22

This Distribution	
Overall Lab SDI:	0.00
Median All Laboratory:	0.00
97.5th centile:	2.92



All SDI Ranges	
< 1	Good
1 - 2	Acceptable
> 2	Poor

## Section SDI scores for this distribution

Section	Determine Combo
Overall	0.00
p24 Ag	0.00
HIV Ab	0.00

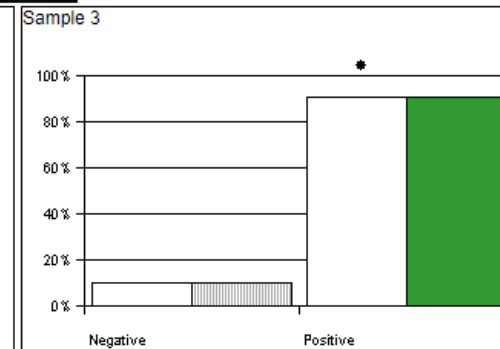
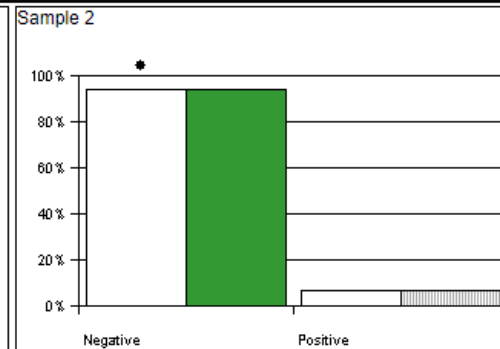
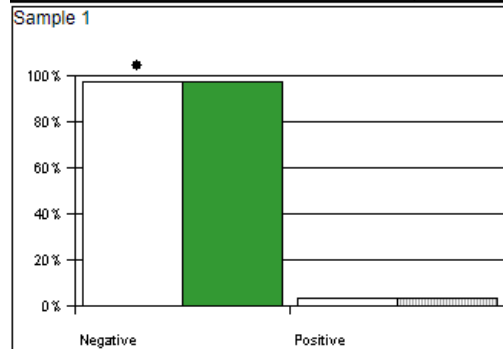


## Qualitative Report

Distribution HV0722

p24 Ag Results

Lab Code	Section	Method	Instrument	Sample Number			Sample Score			Average Score
				1	2	3	1	2	3	(Average)
AAE	Determine Combo	Alere	Determine	Negative	Negative	Positive	0	0	0	0.00
Interpretation				Negative	Negative	Positive				
Spiked Value				Negative	Negative	p24 Ag spiked to give a positive response				



Legend	
	All
	Alert
	Correct Level (all)
	Your result

Number of Responses	Sample 1		Sample 2		Sample 3	
	Method	Overall	Method	Overall	Method	Overall
Negative	30	30	29	29	3	3
Positive	1	1	2	2	28	28
TOTAL	31	31	31	31	31	31

### Individual Sample Score Interpretation

Score	Interpretation
0	Good
1	Acceptable
2	Warning
> 2	Unacceptable

# Summary data

Analyte: HIV Ab

Lab Code	Instrument	Sample Number			Sample Score			Average Score
		1	2	3	1	2	3	
AAE	Determine	Negative	Positive	Positive	0	0	0	0
ACN	Determine							
ACS	Determine							
AKT	Determine	Negative	Negative	Positive	0	4	0	2
AKT	Determine	Negative	Positive	Positive	0	0	0	0
AKT	Determine	Negative	Positive	Positive	0	0	0	0
AKT	Determine							
AY	Determine	Negative	Positive	Positive	0	0	0	0
AZK	Determine	Negative	Positive	Positive	0	0	0	0
BBE	Determine							
HP	Determine	Negative	Positive	Positive	0	0	0	0
HP	Determine	Negative	Positive	Positive	0	0	0	0
KT	Determine	Negative	Positive	Positive	0	0	0	0
MI	Determine	Negative	Positive	Positive	0	0	0	0
MI	Determine	Negative	Positive	Positive	0	0	0	0
MI	Determine	Negative	Positive	Positive	0	0	0	0
MO	Determine	Negative	Positive	Positive	0	0	0	0
MP	Determine	Negative	Positive	Positive	0	0	0	0
MP	Determine	Negative	Positive	Positive	0	0	0	0
MP	Determine	Negative	Positive	Positive	0	0	0	0
AGF	HIV Combo	Negative	Positive	Positive	0	0	0	0
AGF	HIV Combo	Negative	Positive	Positive	0	0	0	0
AGF	HIV Combo	Negative	Positive	Negative	0	0	4	2
AHX	HIV Combo							
AHX	HIV Combo							
AJD	HIV Combo	Negative	Positive	Positive	0	0	0	0
DJ	HIV Combo							
FP	HIV Combo	Negative	Positive	Positive	0	0	0	0
GC	HIV Combo	Positive	Positive	Negative	3	0	4	2.33
ME	HIV Combo	Negative	Positive	Positive	0	0	0	0
ME	HIV Combo	Negative	Positive	Positive	0	0	0	0
ME	HIV Combo	Negative	Positive	Positive	0	0	0	0
MQ	HIV Combo							
MQ	HIV Combo							
MQ	HIV Combo							
MQ	HIV Combo							
RJ	HIV Combo	Negative	Positive	Positive	0	0	0	0
RJ	HIV Combo	Negative	Positive	Positive	0	0	0	0
RJ	HIV Combo	Negative	Positive	Positive	0	0	0	0
TW	HIV Combo	Negative	Positive	Positive	0	0	0	0
TW	HIV Combo	Negative	Positive	Negative	0	0	4	2
YS	HIV Combo	Negative	Positive	Positive	0	0	0	0

Interpretation	Negative	Positive	Positive
Spiked Value	Negative	HIV-1 spiked to give a positive response	HIV-1 spiked to give a positive response

[www.weqas.com](http://www.weqas.com)

# Simplified Reports

Reports are very interactive.

You can hover over various parts of any of the graphical displays within the reports to focus in on the element you wish to see.

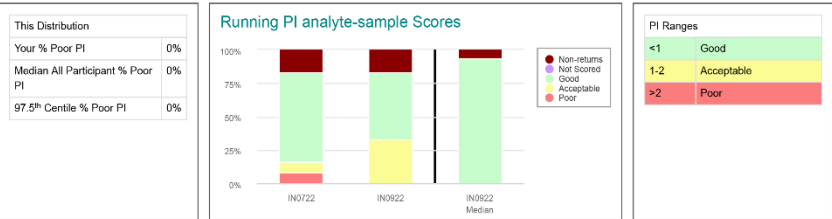
[Sign in - Weqas Connect](#)

Participant Code: WQ00000  
#####  
#####  
#####  
#####

Weqas

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Tel: 02920 314750  
Email: office@weqas.com  
Programme Organiser: Mrs Annette Thomas

Programme: POCT INR • Distribution Code: IN0922  
Distribution Start: 13-Sep-2022 • Distribution End: 11-Oct-2022 • Report Issued: n/a • Report Status: n/a  
Requested By: #####



Location	Instrument Serial #	Instrument Name	INR	Overall % Poor PI	Overall % Non-Return
Anti-Coag Clinic	TZ1000001		Good	0%	0%
	TZ1000002		Acceptable	0%	0%
Day Surgery Unit	TZ1000003		Good	0%	0%
Endoscopy Unit	TZ1000004		Good	0%	0%
Ward 1	TZ1000005		Acceptable	0%	0%
Ward 2	UQ0000001		Acceptable	0%	0%
Pathology 1	UQ0000002		Good	0%	0%
Ward 3	TZ1000006		Good	0%	0%
Ward 4	TZ1000007		Acceptable	0%	0%
Ward 5	UQ0000003		NRR	0%	100%
West Building	TZ1000008		NRR	0%	100%
West Building Ward 1	UQ0000004		Good	0%	0%

PI Code	Meaning
N/A	Not enrolled for this analyte
NRR	Analyte enrolled but no results returned
N/S	This quantitative analyte not scored for this distribution

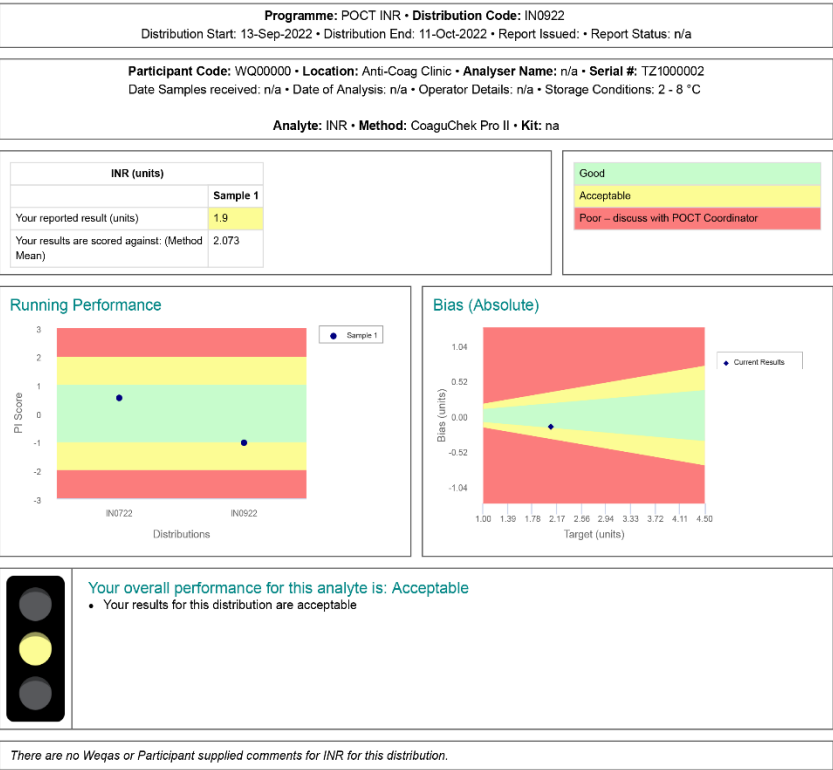
\*\*\*\*\* Please note\*\*\*\*\*

The NQAAP expects 100% compliance on EQA returns. If you are enrolled for an analyte and have not returned a result (denoted by NRR in your PI table), this will be treated as a poor performance score. Please let us know as soon as possible if you are no longer providing this analyte as part of your diagnostic service.

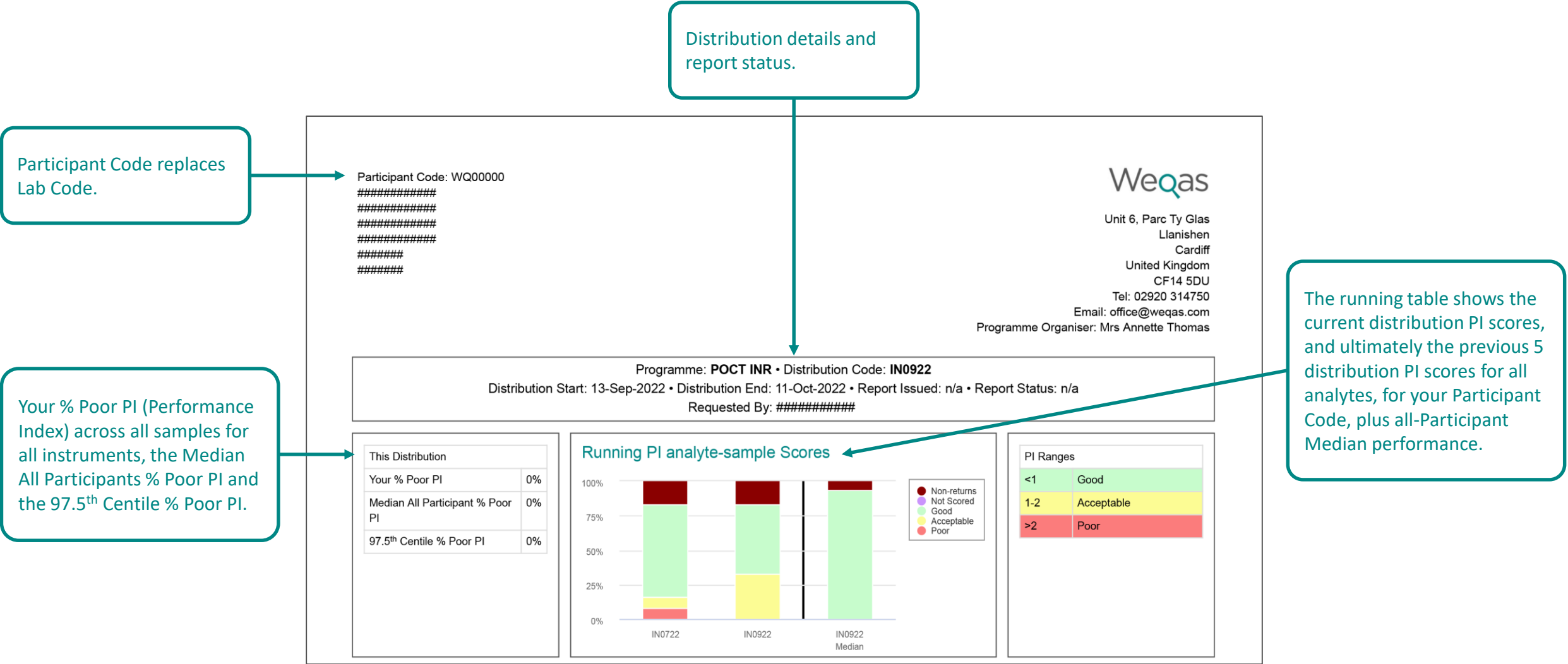
Distribution Specific Comment  
There is no specific comment for this distribution.

Participant submitted comments

Location	Instrument Name / Serial #	Distribution specific comment from the participant
West Building	TZ1000008	ANALYSER NOT IN CLINICAL USE - AWAITING TRAINING FROM ROCHE



# Simplified Reports



The PI Scores table shows performance for all analytes and all instruments.

PI Scores are converted into performance categories and colour coded.

Location	Instrument Serial #	Instrument Name	INR	Overall % Poor PI	Overall % Non-Return
Anti-Coag Clinic	TZ1000001		Good	0%	0%
	TZ1000002		Acceptable	0%	0%
Day Surgery Unit	TZ1000003		Good	0%	0%
Endoscopy Unit	TZ1000004		Good	0%	0%
Ward 1	TZ1000005		Acceptable	0%	0%
Ward 2	UQ0000001		Acceptable	0%	0%
Pathology 1	UQ0000002		Good	0%	0%
Ward 3	TZ1000006		Good	0%	0%
Ward 4	TZ1000007		Acceptable	0%	0%
Ward 5	UQ0000003		NRR	0%	100%
West Building	TZ1000008		NRR	0%	100%
West Building Ward 1	U00000004		Good	0%	0%

Overall % Poor PI across all analytes for each instrument, and Overall % non return for each instrument.

Key for PI Codes.

PI Code	Meaning
N/A	Not enrolled for this analyte
NRR	Analyte enrolled but no results returned
N/S	This quantitative analyte not scored for this distribution

\*\*\*\*\* Please note\*\*\*\*\*

The NQAAP expects 100% compliance on EQA returns. If you are enrolled for an analyte and have not returned a result (denoted by NRR in your PI table), this will be treated as a poor performance score. Please let us know as soon as possible if you are no longer providing this analyte as part of your diagnostic service.

Distribution Specific Comment

There is no specific comment for this distribution.

Participant submitted comments

Location	Instrument Name / Serial #	Distribution specific comment from the participant
West Building	TZ1000008	ANALYSER NOT IN CLINICAL USE - AWAITING TRAINING FROM ROCHE

Distribution Specific Comments from Weqas or participant.

# Simplified Reports

‘Your reported result’ will be shown in the units entered, in the Weqas standard units (if units entered are different from the Weqas standard units), and method corrected result (if appropriate for the analyte).

This will show whether you have been scored against Reference Value, Method Mean or Overall Mean and the actual target value will be shown here.

**Programme:** POCT INR • **Distribution Code:** IN0922  
Distribution Start: 13-Sep-2022 • Distribution End: 11-Oct-2022 • Report Issued: • Report Status: n/a

**Participant Code:** WQ00000 • **Location:** Anti-Coag Clinic • **Analyser Name:** n/a • **Serial #:** TZ1000002  
Date Samples received: n/a • Date of Analysis: n/a • Operator Details: n/a • Storage Conditions: 2 - 8 °C  
**Analyte:** INR • **Method:** CoaguChek Pro II • **Kit:** na

INR (units)	
	Sample 1
Your reported result (units)	1.9
Your results are scored against: (Method Mean)	2.073

Good

Acceptable

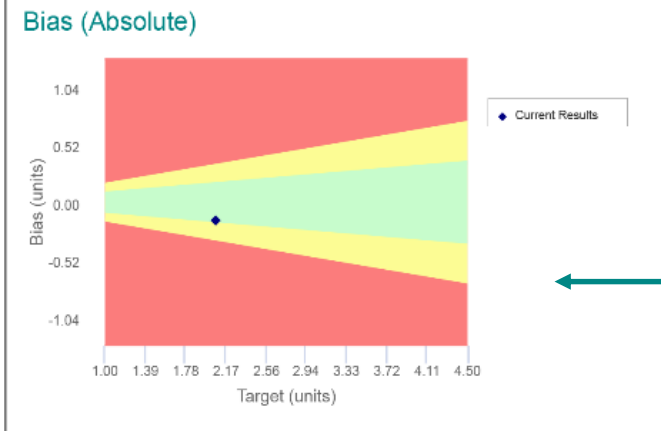
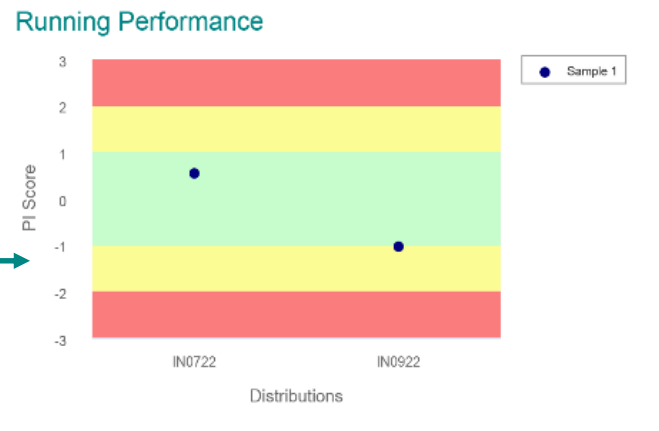
Poor – discuss with POCT Coordinator

The individual Location and instrument details will be shown here.



# Simplified Reports

Running Performance chart shows the PI (Performance Index) score for each sample for the current distribution and the last 5 distributions, with colour coding for each performance category.



The Bias (Absolute) Plot shows absolute bias in Weqas standard units for each sample for the current distribution. The x-axis is the target value; if the target value is a Reference Value or the Overall Mean, your Method Means will be plotted.

# Simplified Reports

Overall Performance is shown as a 'Traffic light', with performance colour coded to easily identify the performance category.



Your overall performance for this analyte is: Acceptable

- Your results for this distribution are acceptable

Overall Performance is stated here along with a basic interpretation of your performance and potential error identified.

*There are no Weqas or Participant supplied comments for INR for this distribution.*

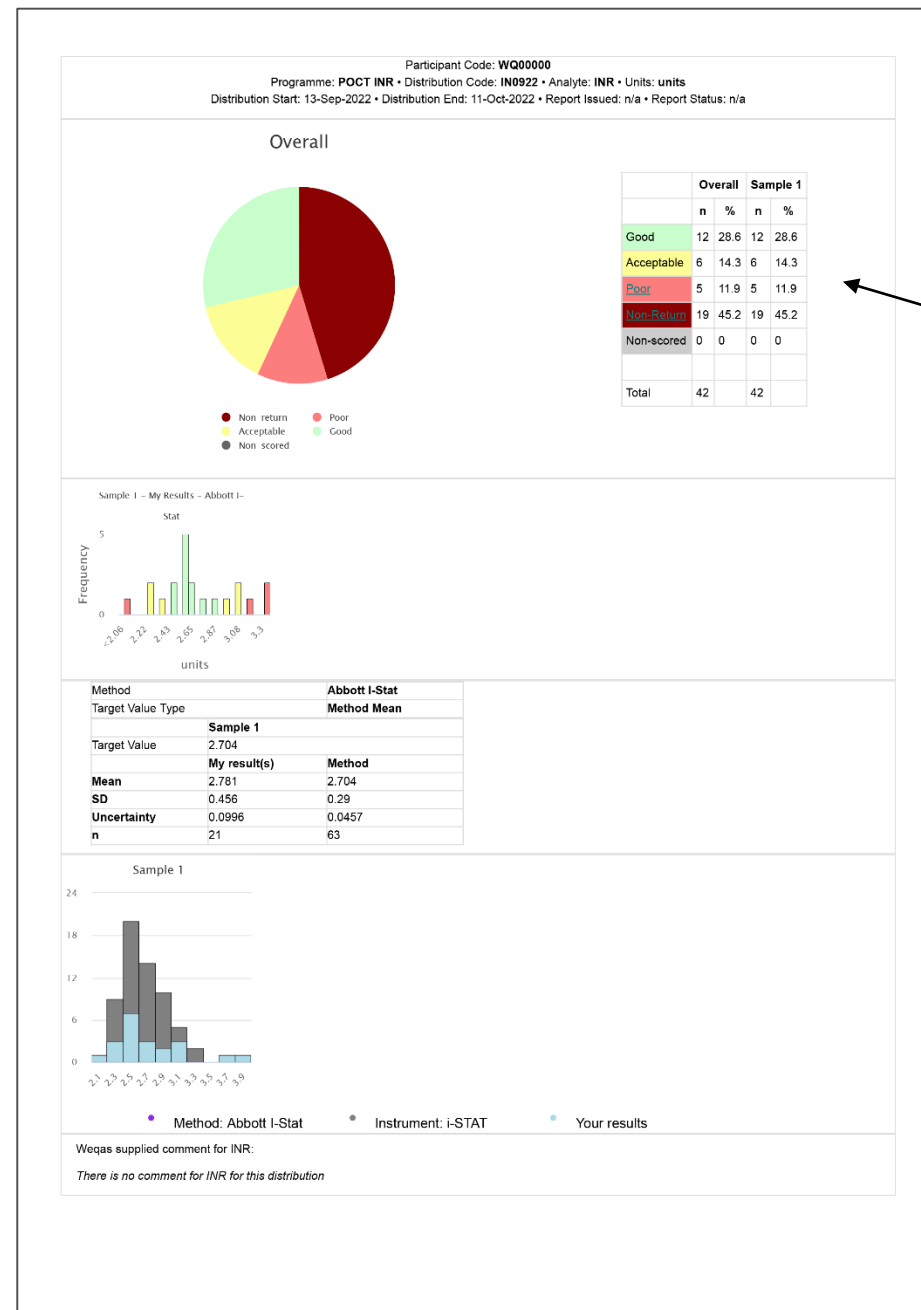
Analyte Specific Comments from Weqas or participant.

# POCT Co-ordinators Report

Reports are very interactive.

You can hover over various parts of any of the graphical displays within the reports to focus in on the element you wish to see.

[Sign in - Weqas Connect](#)



Hyperlinks on 'Poor' and 'Non Returns' to take you to further information.

Participant Code: **WQ00**

Programme: **POCT INR** • Distribution Code: **IN0922** • Analyte: **INR** • Units: **units**

Distribution Start: **13-Sep-2022** • Distribution End: **11-Oct-2022** • Report Issued: **26-Oct-2022** • Report Status: **Final**

Location	Serial #	Name	Sample 1 result	Sample 1 score	Overall performance
	338	n/a	3.3	2.76	Poor
	338	n/a	3.2	2.29	Poor
	338	n/a	3.3	2.76	Poor
	34	n/a	3.2	2.29	Poor

Request PDF of Simplified Analyte reports for all poor performing instruments for this analyte

[Request PDF](#)

PDF will be created only for the sites that have poor performance to allow full review of the individual reports.

## Non>Returns Report

Participant: **WQ00001**

Programme: **POCT INR** • Distribution Code: **IN0922** • Analyte: **INR**

Distribution start: **13-Sep-2022** • Distribution End: **11-Oct-2022** • Report Issued: **26-Oct-2022** • Report Status: **n/a**

Participant Code	Location	Serial #	Name
WQ00001	POCT Lab		n/a
WQ00001	POCT Lab		n/a

# POCT Co-ordinators Report

Participant Code,  
Distribution details and  
report status.

Analyte specific breakdown  
of performance categories  
for all samples.

Participant Code: **WQ00000**  
Programme: **POCT INR** • Distribution Code: **IN0922** • Analyte: **INR** • Units: **units**  
Distribution Start: 13-Sep-2022 • Distribution End: 11-Oct-2022 • Report Issued: n/a • Report Status: n/a

Overall



● Non-return ● Poor  
● Acceptable ● Good  
● Non-scored

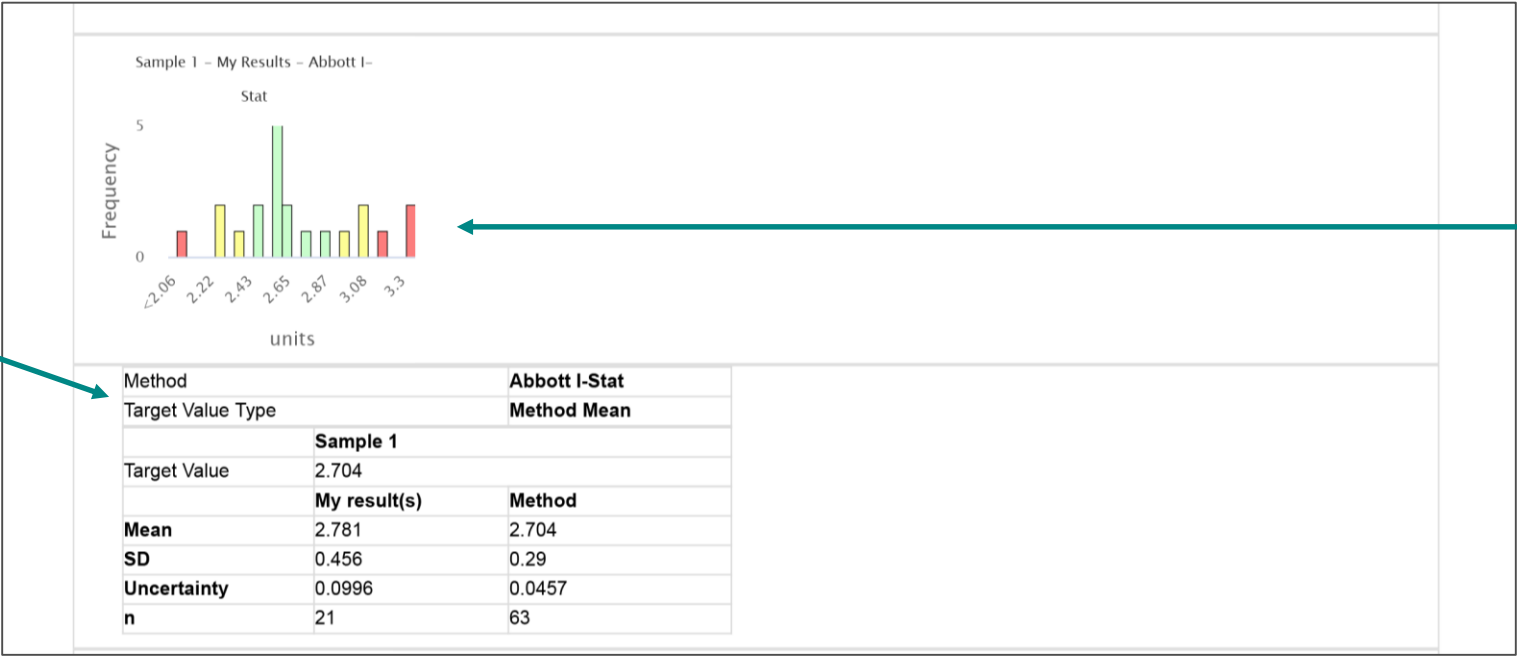
	Overall		Sample 1	
	n	%	n	%
Good	12	28.6	12	28.6
Acceptable	6	14.3	6	14.3
Poor	5	11.9	5	11.9
Non-Return	19	45.2	19	45.2
Non-scored	0	0	0	0
Total	42		42	

Numerical analyte specific  
breakdown of performance  
categories for each sample  
and overall for your  
Participant, Group or Super  
Group dependent on access  
level and report chosen.

Method specific data table showing:

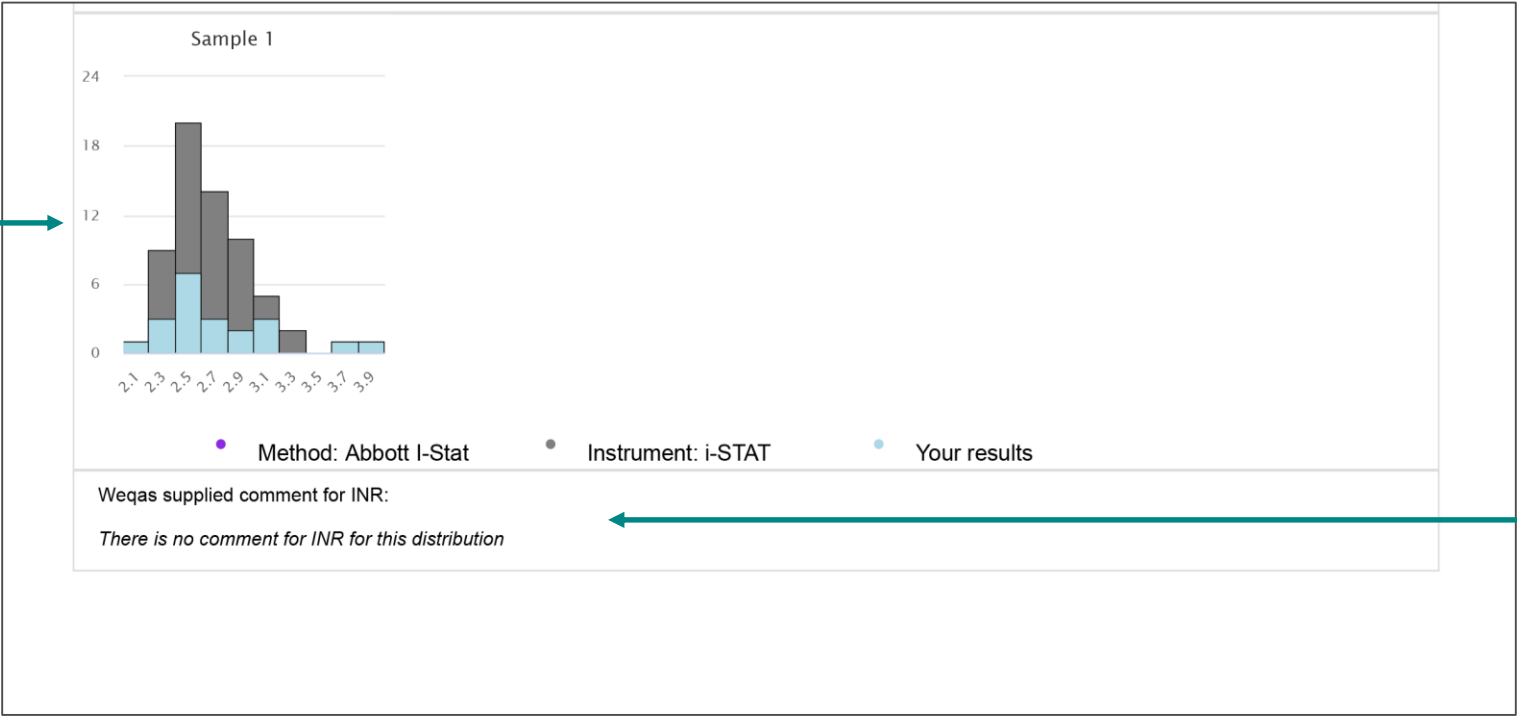
- Target Value to which the submitted result is compared against and Target Value type e.g. Method Mean.
- Mean and SD of your submitted results for this method / instrument.
- Method data (all participants within this method) is also supplied in the right-hand column.

If you have more than one meter type, this data will be shown in a separate table.



Histogram showing the spread of your results. Bars are colour coded to show performance categories across the range.

Histogram showing breakdown of results for each sample, showing the spread of results for your method, your instrument and your submitted results.



Distribution Specific Comments from Weqas or participant.