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EXTERNAL QUALITY ASSESSMENT



INTERNAL QUALITY CONTROL



REFERENCE MEASUREMENT SERVICES



EDUCATION & TRAINING

Weqas

GLOBAL PROVIDER OF QUALITY IN DIAGNOSTIC MEDICINE

Getting the most out of EQA

Annette Thomas



Expectations of EQA Provider has changed

Accreditation status – 17043	 If not accredited, labs should justify why 	Scheme designed and overseen by appropriately competent professionals	 Clinical Scientist or medically qualified Independent Scientific or Medical Advisory group.
Clinically relevant Distribution frequency	 Variable across Schemes (EQALM study 2009) * Where EQA used to assess IVDs, minimum of 6 distributions p.a. (BS EN 14136:2004) For core tests - monthly 	Reporting to Professional body / Regulatory body.	• Mechanism for identification and reporting of Persistent Poor performance issues
Clinically relevant material, range and number of samples	 Evidence of reproducibility Cover clinically appropriate range "Blinded" Commutable materials Challenging samples 	Education	TrainingHelplinePre analyticalPost Analytical
Clinically relevant performance criteria	 Based on clinical outcomes Based on biological variation 	Post-marketing surveillance	 Alerts manufacturers Alerts competent authority Alerts laboratories Alerts professional bodies



Objectives of EQA

Provide a measure of the quality of a test To supplement internal quality control procedures Provide a measure of the "state of the art" of a test To obtain consensus values when true values are unknown

- To investigate factors in performance (methods, staff etc)
- To act as an educational stimulus to improvement in performance
- To provide a Post market vigilance service
- To provide evidence and monitoring of harmonisation strategies
- Provide an assessment of the whole testing process



Expectations of EQA Provider





Clinically appropriate Target value

Improvements in the assessment of the analytical phase includes evaluation of trueness using target vales assigned with high order reference methods, utilising performance criteria that are appropriate for the clinical utility of the analyte and the use of clinically challenging samples.



Reference Measurement Service - Weqas EQA programmes

Flame Atomic Absorption/ Emission

Spectrometry

- Sodium, Potassium, Calcium
- Magnesium, Lithium

IFCC Enzymes

• AST, ALT, LDH, GGT

<u>HPLC</u>

• HbA1c **

** Provided by IFCC Ref lab, Netherlands

IDGC-MS & ID-LC-MS/MS

- •17ß-Oestradiol
- Progesterone
- •Testosterone
- Cortisol
- •Bile Acids
- •Creatinine
- •Cholesterol
- •Glucose
- •Urate
- •Triglyceride
- •HDL *

Clinically Relevant Range and number of samples

- Weqas
- Sample numbers for each scheme assessed on an individual basis Appropriate sample matrices, endogenous, commutable, challenging, linear panels to assess method linearity, specificity and sensitivity (to assist with ISO15189).
- Covering pathological and analytical ranges. Careful selection of endogenous material to ensure range is covered, selected sources of patient material
- Cover critical "diagnostic cut points" e.g. high sensitivity Troponin, urine hCG, HbA1c, POCT CRP
- For Qualitative scheme, provide an appropriate number positive and negative pools, underpinned with known quantitative concentrations.

Clinically relevant samples

Scheme: Car Distribution Date	rdiac Mark a: 24/10/17	ker. Distr 7. Final.	ibution Cod Report Issu	le: N207. led: 23/11/17	•							·	0.1	2.07.0					— .		
Troponin T (ng/l	1	1	2	3 4 Analyte SDI									Scheme Distribution D	POCT C	RP. Distr	ibution (Code: R/	4/05/40			
Reported Result	-/	7.0	305.0	21.0 68.0	-								Distribution D	ate: 17/04/	18. Fina	I. Repor	t Issueu:	4/05/18	_		
Method Corrected Result		7.00	305.00 21	1 Scheme: Gly	ated Haem	oglobin. Dis	tributio	n Code: H2	64.				CRP (n	ng/L)		1	2	Analyte SD) s		
Roche High Sensitivity	Mean	7.31	300.39 21	Distribution D	ate: 24/04/1	18. Final. Re	eport Is	sued: 21/05	/18			Rep	orted Result	-		88.00	25.00		1		
,	SD	0.97	9.86 1	HbA1c IFCC (mmol	/mol)	1		2	3	Analyte SDI		Meth	hod Corrected Re	sult		88.000	25.000	1	- r		
	Number	54	57	Reported Result		58	.0	46.0				Quik	Read do		Mean	85,156	23,000	1			
	Uncert.	0.166	1.632 0.	1 Method Corrected Result	Maan	58.0	00	46.00	28.02				and a ge	1	SD	8 054	2 875	1			
Cobas E Module	Mean	7.31	300.98 21	1. Affinity	Mean	59.0	35	4/.51	38.92					ł	Number	0.001	2.010	{			
1	SD	1.00	9.81 1	1.	Number	4.1	17	17	10					-	Number	2 0052	4 2704	4	1		
	Number	50	53		Uncert.	0.81	11	0.501	0.570			0	Dend as		Uncert.	3.8000	1.2704	1			
	Uncert.	0.177	1.684 0.	2 Afinion AS100	Mean	56.8	86	46.26	NNR			QUIK	(Read go	-	Mean	85.150	23.000	1			
Overall	Mean	7.32	300.96 21	1.	SD	1.0	02	1.01						ļ	SD	8.054	2.875				
	SD	0.99	10.34 1	1.	Number	0.40	7	7							Number	7	8				
	Number	56	59	- Overall	Mean	59.9	82 09	47.80	38.82						Uncert.	3.8053	1.2704				
	Uncert.	0.165	1.682 0.	2	SD	2.2	24	2.21	2.89			Over	rall		Mean	78.617	22.000	1			
Reference Values					Number	14	47	145	123					1	SD	6.204	1.574	1			
Ref. Value Uncertainty					Uncert.	0.23	31	0.229	0.326					ł	Number	35	36	1			
Non-scoring Reference values		1 29	24.78 2	Reference Values	T	59.6	0	48.40						ŀ	Uncert.	1.3108	0.3279	1			
SDI		-0.24	0.19 -0), Ref. Value Uncertainty		1.40	0	1.400				Refe	erence Values				0.02.0	1			
Please note: Linear regression .	ISOS CE CO	procted d	lata	Non-scoring Reference Value	s							Ref	Value Uncertaint	v				1			
Flease note. Linear regression t	ises or co	priected d	idid.	WeQas SD		2.9	2	2.50	2.18			Non	-scoring Reference	o Values				1			
				SI	l Cin	-0.4	46	-0.60		0.53		MoC	Dec SD	e values		5 702	1 740	1			
This Distribution N207				Critic	Sig al Level 1: 6	ma metrics						wec	Jas SD			5.703	1.740		_		
				Minimum Acceptable score	1.64	Critical Lev	el 1 Sia	ma score	2.1					SDI		0.50	1.15	0.8	2		
0.0 51.7 103.3 15	5.0 206.7	258.3 31	0.0	MAPS Allowable TE MAPS Allowable bias %	7.7% 3.6%	Lab bias 9	6		3.4%			Pleas	e note: Linear reg	gression us	ses CF co	orrected o	data.				
+ 75.0				MAPS Allowable CV %	2.5%	Lab CV %			2.1%												
+ /5.0			y = 1.01x	-Please note: Linear regression	uses CF co	prrected data	a	0.0	-	0.0	100 010				-						
60.0			IS = 0	0			_	SureScree	en	SureScree	n nCG GHC	GC		Negative	Nega	tive	Negative	Negative	31	10	1.67
45.0			Sy.x = 0.1	71This Distribution H264			_	SureScree	en	SureScree	n nCG GHC	GC		Positive	Posi	ive	Positive	Negative	00	00	0.00
30.0							_	SureScree	en	SureScree	n hCG GHC	GC		Positive	Posi	ive	Positive	Negative	00	00	0.00
15.0		т	X axis = t	ta 28.0 35.0 42.0 49	0 56.0 6	3.0 70.0	_	SureScree	en	SureScree	n hCG GHC	GC		Negative	Nega	tive	Negative	Negative	3 1	10	1.67
15.04			"x" = you	r(₊ 9.8,			Not ca	SureScree	en	SureScree	n hCG GHC	GC		Negative	Posi	tive	Positive	Negative	3 0	0 0	1.00
0.0 🕮			— O = your	n 7.8-			reporte	SureScree	en	SureScree	n hCG GHC	GC		Positive	Posi	ive	Positive	Negative	00	0 0	0.00
15.0			= ±2 V	Ve 59			3 _	SureScree	en	SureScree	n hCG GHC	GC		Positive	Posi	ive	Positive	Negative	00	00	0.00
30.0		-	I = metho	00 39	T		_	SureScree	en	SureScree	n hCG GHC	GC	_	Positive	Posi	tive	Positive	Negative	00	0 0	0.00
			+ = your p	pr 3.8			X axis	SureScree	en	SureScree	n hCG GHC	GC		Positive	Posi	tive	Positive	Negative	0 0	0 0	0.00
45.0				2.0			x =y O = vc	SureScree	en	SureScree	n hCG GHC	GC		Positive	Posi	tive	Positive	Negative	0 0	0 0	0.00
60.0-				0.0			🖬 = yo	SureScree	en	SureScree	n hCG GHC	GC		Positive	Posi	tive	Positive	Negative	0 0	0 0	0.00
- 75.0				2.0	*		= ±: T = me	SureScree	en	SureScree	n hCG GHC	GC		Positive	Posi	tive	Positive	Negative	0 0	0 0	0.00
				3.9	T.		+ = yoi	SureScree	en	SureScreen M	idstream GH	ICGMS		Positive	Posi	tive	Positive	Negative	0 0	0 0	0.00
				5.9-				Veda.Lal	b	Bab	yCheck-1			Positive	Posi	tive	Positive	Negative	0 0	0 0	0.00
				- 9.8			-	Veda.La/	b	Bab	yCheck-1			Positive	Posi	tive	Positive	Negative	0 0	0 0	0.00
							_	Veda.Lal	b	Bab	yCheck-1										-
							_	Veda.Lal	b	Bab	yCheck-1			Positive	Posi	tive	Positive	Negative	0 0	0 0	0.00
							VIS	SITECT Preg	gnancy	VISITED	T Pregnanc	y		Positive	Posi	ive	Positive	Negative	0 0	0 0	0.00
							_							-							
							_		Inter	rpretation				Positive	Equiv furti	ocal, I ner uation in	Equivocal, further	Negative			

Spiked Value

Approximately 35 IU/L 22 IU/L

Approximately 13 IU/L Approximately 4 IU/L

Weqas



Expectations of EQA Provider

Clinically relevant performanc e criteria Based on clinical outcomes

 Based on biological variation Reporting to Professional body / Regulatory body.

 Mechanism for identification and reporting of Persistent Poor performance issues Clin Chem Lab Med 2015; 53(6): 833-835

Sverre Sandberg*, Callum G. Fraser, Andrea Rita Horvath, Rob Jansen, Graham Jones, Wytze Oosterhuis, Per Hyltoft Petersen, Heinz Schimmel, Ken Sikaris and Mauro Panteghini

Defining analytical performance specifications: Consensus Statement from the 1st Strategic Conference of the European Federation of Clinical Chemistry and Laboratory Medicine

DOI 10.1515/cclm-2015-0067

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The Organisers and the Scientific Programme Committee (SPC) of the 1st Strategic Conference of the European Federation of Clinical Chemistry and Laboratory Medicine (EFLM) on 'Defining analytical performance goals 15 years after the Stockholm Conference on Quality Specifications in Laboratory Medicine', held in Milan (IT) on November 24–25, 2014, are pleased to report on the success of the Conference.

The primary aim was to revisit the 'Consensus Agreement' from the Stockholm Conference investigating to what extent the advocated hierarchy is still valid or if it should be changed. A revision of the original hierarchy established by the Stockholm Conference was presented to the meeting with opportunity for discussion and feedback by conference participants. This revision further underwent modification and explanatory additions by the SPC in an attempt to simplify the hierarchy and improve its application by various stakeholders.



Clin Chem Lab Med 2017; 55(7): 949-955

Opinion Paper

Graham R.D. Jones*, Stephanie Albarede, Dagmar Kesseler, Finlay MacKenzie, Joy Mammen, Morten Pedersen, Anne Stavelin, Marc Thelen, Annette Thomas, Patrick J. Twomey, Emma Ventura and Mauro Panteghini, for the EFLM Task Finish Group – Analytical Performance Specifications for EQAS (TFG-APSEQA)

Analytical performance specifications for external quality assessment – definitions and descriptions

DOI 10.1515/cclm-2017-0151 Received February 21, 2017; accepted April 18, 2017; previously published online May 23, 2017

Abstract: External Quality Assurance (EQA) is vital to ensure acceptable analytical quality in medical laboratories. A key component of an EQA scheme is an analytical performance specification (APS) for each measurand that a laboratory can use to assess the extent of deviation of the obtained results from the target value. A consensus conference held in Milan in 2014 has proposed three models to set APS and these can be applied to setting APS for EQA. A goal arising from this conference is the harmonisation of EQA APS between different schemes to deliver consistent quality messages to laboratories irrespective

Laboratory Medicine (EFLM) Task and Finish Group on Performance Specifications for External Quality Assurance Schemes (TFG-APSEQA) and on clear terminology for EQA APS. The recommended terminology covers six elements required to understand APS: 1) a statement on the EQA material matrix and its commutability; 2) the method used to assign the target value; 3) the data set to which APS are applied; 4) the applicable analytical property being assessed (i.e. total error, bias, imprecision, uncertainty); 5) the rationale for the selection of the APS; and 6) the type of the Milan model(s) used to set the APS. The terminology is required for EQA participants and other interested parties to understand the meaning of meeting or not meeting APS.



Clinically Relevant Performance Specification





"State of the art" v Biology

Creatinine Precision Profile (CV %)



Glucose Precision Profile (CV%)



Biological goal is only achievable down to 4 mmol/l Glucose

Table 1: Examples of current variation in models used to assign analytical performance specifications (APS) to External Quality Assurance(EQA) schemes.

EQA Program	Models
CSCQ Switzerland	Governmental regulations (combination of BV and state of the art) and Combination of limits given by scientific societies and Z-score
CTCB France	Z-score/state of the art/limits given by scientific societies or other/limits based on clinical impact
DEKS Denmark	Combination of BV, state of the art and expert opinion
NOKLUS Norway	Fixed percentage limits and based on a combination of BV, state of the art and expert opinion
RCPAQAP Australia	Combination of BV and state of the art
SEHH Spain	Statistical/state of the art/BV
SEQC Spain	Combination of BV and statistical results
SKML The Netherlands	Combination of BV and state of the art
WEQAS UK	Combination of BV and state of the art
CMCEQAS	Combination of state of the art and statistical considerations



Expectations of EQA Provider



Weqas Pre / Post Analytical / Interpretive Programmes

- Serum Indices Programme, report semi-quantitative and qualitative results
- Questionnaires sent out as part of Programme repertoire to assess current practice and use of National guidelines e.g. pre analytical sample handling, Adjusted Calcium equations, Porphyrin Standards
- Interference Studies e.g. HIL samples various programmes, Bilirubin
 effect on Salicylate & Paracetamol
- Post analytical cases provided with Programmes e.g. Porphyrin interpretation cases, Macroprolactin samples
- EQA for calculated parameters.
- Foetal Fibronectin, Pre-eclampsia, D-dimer, participants can report quantitative, qualitative (post-analytical interpretation)

Weoas



Interference Reports – bilirubin effect on creatinine



The reference value (ID-GCMS) was 184.3 µmol/L for sample 2 and 184.4 µmol/L for sample 3

Specificity and Sensitivity Studies

Assessment of analytical sensitivity of Pregnancy testing kits.



Sure Screen hCG GHCGC



Percentage of negative results (red), equivocal results (green), positive results (purple) and positive plus equivocal (light blue) for each pregnancy testing kit / device at each conc.

Assessment of analytical specificity of total Bile Acid methods

Four pools of human serum were prepared, and spiked with approximately 100µmol/L of each of the major bile acids and target value assigned using an ID-GCMS method

	CHOLIC	ACID	ID-GC	MS Target	DEOXYCHOLIC ID-GCMS Target					
	103.2 µn	nol/L			108.8 µı	mol/L				
Returned results	mean	SD	n	% recovery	mean	SD	n	% recovery		
overall	101.18	7.54	111	98.06	137.80	15.87	110	126.68		
Enz-Thio-NADH	99.89	6.59	95	96.81	141.27	15.64	94	129.87		
Enz-Formazan	89.5	1.50	5	86.74	137.00	15.00	2	125.94		
Enz-Formazan (Sentinel)	112.41	4.90	15	108.95	119.42	5.08	15	109.78		
POOL ID	URSOD	EOXYC	HOLI	C Gavimetric	CHENO	DEOXY	CHOL	IC ID-GCMS		
	target				Target					
	100 µmc	ol/L			77.1 μmol/L					
Returned results	mean	SD	n	% recovery	mean	SD	n	% recovery		
overall	57.81	8.44	107	57.81	56.05	7.30	107	72.66		
Enz-Thio-NADH	56.00	4.44	98	56.00	54.25	4.61	95	70.32		
Enz-Formazan	51.50	0.5	2	51.50	51.00	2.00	2	66.11		
Enz-Formazan (Sentinel)	90.47	3.33	15	90.47	77.05	2.88	12	99.88		

The study highlights the importance of using reference methods to assign target values rather than consensus mean and presents strong evidence on the variability in specificities of the methods for the different bile.

Weqas Post Analytical / Interpretive Programmes – Porphyrin Programme

include monitoring PBG/Creatinine, urine porphyrin HPLC, plasma

offered following identifying type of acute porphyria. Family studies/family history of porphyria and appropriate cascade

porphyrin scan, faecal porphyrin HPLC, plasma porphyrin scan, faecal porphyrin HPLC (ensure samples light protected). Genetic analysis

ramme						
PBG Case 1						
Comment	Elevated PBG	Acute Porphyria / Acute attack	Result telephoned	Advise discuss with NAPS	Samples for further investigation / ID	Total Out of 5
Increased Urine Porphobilinogen, consistent with a diagnosis of acute porphyria. Contact metabolic medicine or the National Acute Porphyria Service for advice on management	1	1		1		3
Result suggest acute porphyria. Please send 5-10 ml EDTA blood and a small portion (5-10g wet weight) of fresh faeces in a universal container. All samples should be protected from light.		1			1	2
Significantly raised urine PBG, consistent with active acute porphyria. Advise collect further urine, blood and stool sample for further characterisation. Seek advice from national acute porphyria service regarding patient management.	1	1		1	1	4
Significantly raised PBG + PBG/Creatinine ratio consistent with acute porphyria attack. Result telephoned to ITU + discussed with medical staff. Advise discuss with NAPS for appropriate haem arginate treatment, current medication to determine any porphyrinogenic drugs + appropriate management of hyponatraemia (NAPS emergency protocol), follow up to determine specific form of acute porphyria to						

1

		screening, patient hospital records + emergency pr porphyria) alert added. When well consult patient re card/other.	otocol (acute garding medi-alert
www.wegas.com			

Result

(umol/mmol

creatinine)

52.6

62.6

51.5

58.2

Result

(umol/l)

644.3

789

627

664

Lab Code

AXP

ΒZ

EN

FB

5



Educational role (quality improvement)

Pre-analytical effects Performance of methods – state of the art accuracy precision limits of detection linearity Susceptibility of methods to interference including other analytes and matrix Interpretation of results – standard units, global cut off Undertaking audit of clinical services – identify good practice Understanding how to use Quality tools – IQC, EQA, audit

Method performance – hs Tnl



Weqas

Glucose (mmol/l)		1	2	3	4	Analyte SD
Reported Result		11.4	8.1	20.7	1.8	
Method Corrected Result		11.40	8.10	20.70	1.80	
Hexokinase	Mean	11.42	8.21	21.11	1.85	
	SD	0.20	0.15	0.41	0.05	
	Number	170	172	169	168	
	Uncert.	0.015	0.012	0.032	0.004	
Cobas C Module	Mean	11.45	8.26	21.13	1.88	
	SD	0.17	0.13	0.31	0.04	1
	Number	91	95	92	91	1
	Uncert.	0.018	0.013	0.033	0.004	
Overall	Mean	11.39	8.21	21.05	1.86	1
	SD	0.22	0.15	0.46	0.06	
	Number	191	188	188	186	
	Uncert.	0.010	0.011	0.033	0.004	1
Reference Values ID-GCMS		11.40	8.15	20.95	1.76	
Ret. value Uncertainty		0.100	0.070	0 190	0.020	
Non-scoring Reference Values						
WeQas SD		0.34	0.25	0.65	0.12	1
SDI		0.00	-0.20	-0.38	0.34	0.23
	Sign	na Metric	s			
	Critical Le	evel 1: 7	mmol/l			
Minimum Acceptable score	1.62	Critical I	Level 1 S	igma sco	ore	7.4
MAPS Allowable TE	6.9%					
MAPS Allowable bias %	2.20%	Lab bia	0.2%			
MAPS Allowable CV %	2.90%	Lab CV	%			0.9%

This Distribution RH





ISO 15180 tools from EQA Weqas reports

traceability to higher order method

Linearity assessment



Troubleshooting Support & post market vigilance

Now part of the EQA providers role

To provide **help** with

Participant Performance queries

report interpretation

Provide additional material for problem solving

To alert manufacturers of potential issue To assist in issue resolution

To alert regulatory authority

Weqas

DCA 2000/ Vantage HbA1c Bias Plot





Distributions (June 2016 - June 2019)



Post Market vigilance – INR thromboplastin



INR results classified into pre and post recalibration.

participants using strips calibrated to WHO reference thromboplastin rTF/09

The pre calibration strips compared well with the results from Distribution 0517 (Median 2.8) however much higher results and a wider distribution of results was observed for the post calibration strips. Wegas immediately contacted the manufacturer and sent them the data.

Aug 2018 – Urgent field safety notice issued to inform users that the manufacturer was reverting back to previous WHO reference standard.



Web portals

A wealth of additional information can be provided to participants with direct links to the Weqas databases providing useful troubleshooting tools.

Can also provide tools for laboratories to achieve ISO 15189 accreditation



ISO 15189 clause 5.3.1.4 - Traceability

- Equipment calibration and metrological traceability
- The laboratory shall have a documented procedure for the calibration of equipment that directly or indirectly affects examination results. This procedure includes:
- b) recording the metrological traceability of the calibration standard and the traceable calibration of the item of equipment;
- c) verifying the required measurement accuracy and the functioning of the measuring system at defined intervals.
- Metrological traceability shall be to a reference material or reference procedure of the higher metrological order available.
- NOTE Documentation of calibration traceability to a higher order reference material or reference procedure may be provided by an examination system manufacturer. Such documentation is acceptable as long as the manufacturer's examination system and calibration procedures are used without modification.
- Where this is not possible or relevant, other means for providing confidence in the results shall be applied, including but not limited to the following:
- — use of certified reference materials;.....





Traceability – Weqas QC-RM

Testosterone and Cortisol Tandem MS standards are assayed quality control material for verification of "in house" prepared calibrators
Standards prepared and value assigned using the Weqas Reference Measurement Laboratory using traceable material of the highest metrological order.

•Assists with ISO 15189:2012 compliance.



estosterone l'argeted Calibrators											
Sample ID	Target Value (nmol/L)	SD	%CV	Expanded Uncertainty							
Level 0	0	-	-	-							
Level 1*	0.50	-	-	-							
Level 2	1.04	0.02	2.00	0.03							
Level 3	2.88	0.05	1.85	0.09							
Level 4	7.63	0.16	2.09	0.24							
Level 5	15.1	0.31	2.06	0.48							
Level 6	23.48	0.29	1.24	0.74							
Level 7	38.32	0.84	2.20	1.21							

Scheme: Serum Chemistry. Distribution Code: RH. Distribution Date: 2/01/18. Final. Report Issued: 24/01/18							
Glucose (mmol/l)		1	2	3	4	Analyte SDI	
Reported Result		11.4	8.1	20.7	1.8		
Method Corrected Result		11.40	8.10	20.70	1.80]	
Hexokinase	Mean	11.42	8.21	21.11	1.85]	
	SD	0.20	0.15	0.41	0.05	1	
	Number	170	172	169	168]	
	Uncert.	0.015	0.012	0.032	0.004		
Cobas C Module	Mean	11.45	8.26	21.13	1.88]	
	SD	0.17	0.13	0.31	0.04	1	
	Number	91	95	92	91	1	
	Uncert.	0.018	0.013	0.033	0.004	1	
Overall	Mean	11.39	8.21	21.05	1.86	1	
	SD	0.22	0.15	0.46	0.06	1	
Nur		191	188	188	186	1	
	Uncert.	0.016	0.011	0.000	0.004	1	
Reference Values ID-GCMS		11.40	8.15	20.95	1.76		
Ref. Value Uncontainty		0 100	0.070	0.100	U.020		
Non-scoring Reference Values]	
WeQas SD		0.34	0.25	0.65	0.12		
SDI		0.00	-0.20	-0.38	0.34	0.23	
	Sign	na Metrio	s				
	Critical Le	evel 1: 7	mmol/l				
Minimum Acceptable score	1.62	Critical I	Level 1 S	igma sco	ore	7.4	
MAPS Allowable TE	6.9%						
MAPS Allowable bias %	2.20%	Lab bia	s %			0.2%	
MAPS Allowable CV %	2.90%	Lab CV	%			0.9%	

Please note: Linear regression uses CF corrected d

This Distribution RH



Prev

Traceability From Weqas Reports

- Reference measurement values shown on report (and reference value uncertainty)
- Full traceability chain to SI units available.
- Lab results compared directly to reference values
- SDI scores, Sigma scores and bias plot based on reference values

Scheme: Ser Distribution Dat	um Chem e: 2/01/18	istry. Dis . Final. F	tribution Report Is:	Code: R sued: 24	H. /01/18	
Glucose (mmol/l)	1	2	3	4	Analyte SDI
Reported Result		11.4	8.1	20.7	1.8	
Method Corrected Result		11.40	8.10	20.70	1.80	
Hexokinase	Mean	11.42	8.21	21.11	1.85	
	SD	0.20	0.15	0.41	0.05	
	Number	170	172	169	168	
	Uncert.	0.015	0.012	0.032	0.004	
Cobas C Module	Mean	11.45	8.26	21.13	1.88	
	SD	0.17	0.13	0.31	0.04	
	Number	91	95	92	91	
	Uncert.	0.018	0.013	0.033	0.004	
Overall	Mean	11.39	8.21	21.05	1.86	
	SD	0.22	0.15	0.46	0.06	
	Number	191	188	188	186	
	Uncert.	0.016	0.011	0.033	0.004	
Reference Values ID-GCMS		11.40	8.15	20.95	1.76	
Ref. Value Uncertainty		0.100	0.070	0.190	0.020	
Non-scoring Reference Values						
WeQas SD		0.34	0.25	0.65	0.12	
SDI		0.00	-0.20	-0.38	0.34	0.23
	Sign	na Metric	s			
	Critical Le	evel 1: 7 i	mmol/l			
Minimum Acceptable score MAPS Allowable TE	1.62 6.9%	Critical I	Level 1 S	igma sco	ore	7.4
MAPS Allowable bias %	2.20%	Lab bia	s %			0.2%
MAPS Allowable CV %	2.90%	Lab CV	%			0.9%

Please note: Linear regression uses CF corrected data.

This Distribution RH



2.4

Prev

+

-

Uncertainty

Laboratory within run Imprecision: Sy.x = 0.06 mmol/LCV% = (Sy.x/x)*100 = 0.06/7*100 = 0.86%





Method Uncertainty From Weqas End of Batch Reports

Analyte: Creatinine (µmol/L)

Method: Jaffe - IDMS	M891a	M892	M893	M894	M895	M896	M897	M898
Section Stats								
Mean reported results	64.5	133.6	206.8	276.7	346.9	420.1	490.6	558.4
SD reported results	2.9	3.4	7.3	8.6	7.8	11.0	14.7	12.7
CV(%) reported results	4.51	2.52	3.52	3.10	2.25	2.61	2.99	2.27
Number of results	5	5	4	3	5	5	5	6
Method Result Stats								
Mean method mean	67.7	139.3	213.3	286.6	357.4	428.9	498.4	570.0
Median CV	3.08	2.52	1.91	2.00	2.14	2.14	1.88	2.11
Overall Result Stats								
Median CV	2.44	2.19	1.69	1.81	1.92	1.97	1.70	1.85

Between batch CV% provided on End of Batch reports (6-12 month review)



Pool M891a - CV% of reported results: 4.51%

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Method Linearity from Weqas

Weq	as
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Scheme: S Distribution	erum Che Date: 3/04	emistry. D /18. Final	istributior . Report Is	Code: Ri sued: 8/0	K. 5/18	
Potassium (mmol	1	2	3	4	Analyte SDI	
Reported Result	6.50	4.10	5.70	1.90		
Method Corrected Result		6.500	4.100	5.700	1.900]
Indirect ISE	Mean	6.379	4.073	5.618	1.841]
	SD	0.088	0.065	0.077	0.048]
	Number	179	176	174	177]
	Uncert.	0.0066	0.0049	0.0058	0.0036]
Cobas C Module	Mean	6.437	4.119	5.666	1.857]
	SD	0.049	0.036	0.048	0.046	1
	Number	97	95	99	99	1
	Uncert.	0.0050	0.0037	0.0049	0.0047	1
Overall	Mean	6.380	4.075	5.618	1.837]
	SD	0.088	0.065	0.078	0.052	1
	Number	184	183	181	183	1
	Uncert.	0.0065	0.0048	0.0058	0.0038]
Reference Values FAAS / FAES		6.330	4.050	5.560	1.810	
Ref. Value Uncertainty		0.0440	0.0280	0.0390	0.0130	1
Non-scoring Reference Values]
WeQas SD		0.127	0.076	0.104	0.077	L
SDI		1.34	0.66	1.34	1.17	Γ

Please note: Linear regression uses CF corrected data.



Linear series of 8 pools distributed for most routine chemistry programmes

Linear series cover wide analytical and pathological range

Chemistry QCRM linearity panel available as a range of up to 8 samples and are suitable for ISO 15189 method verification

Previou		u D.	Weq	Serum Chemistry QCRM					
	c).), S	Serum Chemistry Qua	lity Control Reference Mate	erial	Creatinine			
t value ent results rod	0.	o. S	Size: 3.5mL Lot No: 080118/93	Exp.: Jan-2020		Pool	ID-GCMS Target Value (μmol/L)	Expanded Uncertainty	
od specific instrument s SD	C	D.				932	110.98	1.38	
SD ous results	c	D. D.				934	262.15	3.25	
	c	D.				936	414.37	5.14	
	- c	J.				938	559.71	6.94	

Weqas Regional EQA Reports

5. EOB Absolute Deviation Report

For the labs in the region, for the selected batch, deviation reports give mean reported results and look at absolute and % deviation from the regional mean on a per analyte basis.

		B291	B297	B300	B296	B301	B292	B293	B294	B295	B298	B299	B302
FH	Advia 2400 1	-0.59	-9.31	-10.89	-5.39	-14.48	-2.56	-3.67	-5.22	-7.38	-6.15	-10.04	-11.82
FH	Advia 2400 2	-1.92	-11.48	-18.52	-9.36	-13.35	-2.92	-4.20	-5.86	-7.84	-5.65	-14.51	-14.05
GF	Beta	1.21	5.82	9.58	9.14	8.08	2.24	1.13	4.21	0.06	8.95	0.72	6.72
GF	Alpha (4)	1.21	3.49	11.58	7.81	16.42	3.24	2.47	6.88	3.72	11.62	11.72	10.72
KJ	ARCHITECT 2	0.88	11.49	8.25	3.48	2.42	1.91 *	1.13	4.21 *	3.39	-3.88	2.06	-0.62
KJ	ARCHITECT 1	-0.79	6.49 *	1.25 *	-5.69	0.92	0.91 *	3.13	1.21 *	8.06	-4.88	10.06	9.05
KJ	ARCHITECT 1	3.21 *	16.49 *	9.25 *	16.81 *	-	-	-	-	-	-	-	-
	Wegas SD	2.31	8.61	12.09	7.39	12.90	3.24	4.27	5.34	6.41	9.47	10.93	14.24



Weqas



EQA Challenges – the patient test workflow

How do we assess the full patient testing pathway ? How can we mimic the laboratory and POCT test workflow with greater use of automation? How can we assess the integrity of the data?

Weqas Laboratory test workflow









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Thank you

Any Questions?