

### Outline of the Talk

- What is biologic variation?
   Do we need to worry about it?
- Setting Goals for Performance from Biologic Variation information
- Reference Change Value
- Current Challenges and Debate on Biologic Variation (Milan Meeting 2014)
- Tools for integrating test results into biologic variation
  - Reference Change Value (RCV)
  - Number of tests required to detect a significant change

## Disclosure: Almost all discussion of Biologic Variation derives from Callum Fraser

**Biologic Variation Data for Setting Quality Specifications** 

https://www.westgard.com/guest12.htm

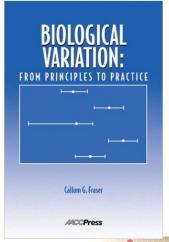
James O. Westgard Foreword:

https://www.westgard.com/guest19.htm

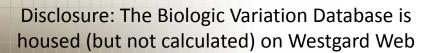
Fraser: Are "Scientific Statements" the

Scientific Truth?

https://www.westgard.com/callumfraser.htm







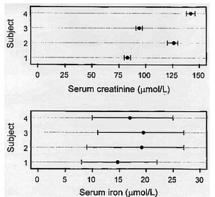
- Biologic Variation Database is compiled and updated by SEQC and Dr. Carmen Ricos
- Sometimes called "Ricos Goals" as a shorthand and tribute to her leadership
- Most popular resource = Desirable specifications for Imprecision, Bias, and Allowable Total Error: <a href="https://www.westgard.com/biodatabase">https://www.westgard.com/biodatabase1.htm</a>
- Minimum Specifications: https://www.westgard.com/minimumbiodatabase1.htm
- Optimal Specifications: https://www.westgard.com/optimal-biodatabase1htm.htm
- Specifications for **patients w/disease**:
  <a href="https://www.westgard.com/biodatabas">https://www.westgard.com/biodatabas</a>
  edisease.htm

	Analytie		Biological Variation		Desirable specification		
		Papers	CV <sub>1</sub>	CVg	1(%)	B(%)	TE(%
5-	11-Desoxycortisol	2	21.3	31.5	10.7	9.5	27.1
S-	17-Hydroxyprogesterone	2	19.6	50.4	9.8	13.5	29.7
U-	4-hydroxy-3-methoximandelate (VMA)	1	22.2	47.0	11.1	13.0	31.3
S-	5' Nucleotidase	2	23.2	19.9	11.6	7.6	26.8
U-	5'-Hydroxyindolacetate, concentration	1	20.3	33.2	10.2	9.7	26.5
8-	a1-Acid Glycoprotein	3	11.3	24.9	5.7	6.8	16.2
S-	a1-Antichymotrypsin	1	13.5	18.3	6.8	5.7	16.8
S-	a1-Antitrypsin	3	5.9	16.3	3.0	4.3	9.2
S-	a1-Globulins	2	11.4	22.6	5.7	6.3	15.7
U-	a1-Microglobulin, concentration, first morning	1	33.0	58.0	16.5	16.7	43.9
P.	«2-Antiplasmin	1	6.2		3.1		-
s-	a2-Globulins	2	10.3	12.7	5.2	4.1	12.6
S-	α2-Macroglobulin	4	3.4	18.7	1.7.	4.75	7.56
U-	a2-Microglobulin output, first morning	1	29.0	32.0	14.5	10.8	34.7
ρ.	a-aminobutryic acid	1	24.7	32.3	12.4	10.2	30.5
S-	a-Amylase	7	8.7	28.3	4.4	7.4	14.6
5-	α-Amylase (pancreatic)	2	11.7	29.9	5.9	8.0	17.7
U-	a-Amylase (pancreatic)	2	69.5	105.0	34.75	31.48	88.82
U-	a-Amylase concentration, random	1	94.0	46.0	47.0	26.2	103.7
P.	a-Carotene	1	24.0	65.0	12.0	17.3	37.1
S-	α-Carotene	1	48.0	65.0	24.0	20.2	59.8
S-	u-Fetoprotein(non hepatic carcinoma)	2	12.2	45.6	6.1	11.8	21.9
S-	a-Tocopherol	3	13.8	15.0	6.9	5.1	16.5
5-	Acid phosphatase	2	8.9	8.0	4.5	3.0	10.3
S-	Acid phosphatase tartrate-resistant (TR-ACP)	2	8.0	13.3	4.0	3.9	10.5



## Biologic Variation: One reason Why we never get the same number twice

- CVG: Within-group variation: people are different from other people
- CVI: Within-subject variation: you are never exactly the same







# Why biologic variation matters: it impacts test results

- The "noise" of the body may obscure the signal of the patient's clinical state
- We want the clinician to treat on signal not noise
- We want to make sure our analytical method variation doesn't add too much additional "noise" and make it even harder for the clinician to determine what's happening with the patient.
- [Which raises the question: how much noise is acceptable?]



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  - Reference Change Value (RCV)
  - Number of tests required to detect a significant change.

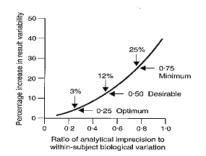
## Westgard QC

# Setting Quality Specifications from Biologic Variation: Imprecision

- Optimal: ONE-QUARTER of CVI
- · Desirable: HALF of CVI
- Minimum: THREE-QUARTERS OF CVI

#### WHY?

- Optimal imprecision will only increase result variability by 3%
- Desirable imprecision will only increase result variability by 12%
- Minimum imprecision will only increase result variability by 25%



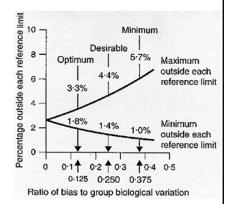


# Setting Quality Specifications from Biologic Variation: Bias

- $desirable < 0.250[CV_1^2 + CV_G^2]^{0.5}$
- optimum <  $0.125[CV_1^2 + CV_G^2]^{0.5}$
- $minimum < 0.375[CV_1^2 + CV_G^2]^{0.5}$

#### WHY?

- Optimal bias increases results outside reference interval by 3.3%
- Desirable bias increases results outside reference interval by 4.4%
- Minimum bias increases results outside reference interval by 5.7%





## Setting Quality Specifications from Biologic Variation: Total Allowable Error

- Total Allowable (Analytical) Error: TEa or ATE = (1.65\*SD) + Bias
- TE < k \* 0.5 CV<sub>I</sub> + 0.25(CV<sub>I</sub><sup>2</sup> + CV<sub>G</sub><sup>2</sup>)<sup>1/2</sup> where k= 1.65 at a=0.05 (limiting chance of a significant error to 5%)

## Disclosure: The Biologic Variation Database is housed (but not calculated) on Westgard Web

- Estimates of CV<sub>I</sub> and CV<sub>G</sub> are compiled through a review of all relevant studies of biologic variation
- · Updated every two years
- As new analytes are introduced, and new studies added, the size and specifications of the database changes
- Unlike all other EQA/PT surveys and government regulations, these are "evidence-driven" quality specifications
- Latest edition covers more than 350 analytes

		Number of	Biological Variation			Desirable specification		
		Papers	CVI	CVg	1(%)	B(%)	TE(%)	
S-	11-Desoxycortisol	2	21.3	31.5	10.7	9.5	27.1	
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### Outline

- Reference Change Value and Number of Tests Required
- Current Challenges and Debate on Biologic Variation (Milan Meeting 2014)
- Tools for integrating test results into biologic variation
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#### Callum Fraser: Reference Change Value (RCV)

Are serial test results clinically different? Or is it just noise?

$$RCV = \sqrt{2} * z * \sqrt{CV_a^2 + CV_i^2}$$

- 2 samples
- need estimates of analytical imprecision and within-subject biologic variation
- Z-value of 1.96 for P < 0.05 or 95% probability (use 2.56 for P < 0.01 or 99% probability)
- No bias included in the calculation



#### What's the deal with the RCV?

- When changes are bigger than the RCV, it is a real difference
- When changes are smaller than the RCV, it may only be noise (imprecision and/or inaccuracy)
- Callum Fraser suggests the following notation on the report
  - \* = significant change;
  - \*\* = highly significant change
- RCV is NOT used to set an analytical goal, but to help interpret test results
   Westgard QC

## Systemic Impact on Clinical Diagnosis: Callum Fraser's Number of tests/samples required

# Tests = 
$$\left(\begin{array}{ccc} z * \sqrt{CV_a^2 + CV_i^2} \\ D \end{array}\right)^2$$

How many tests required to detect a significant change in a patient?

- D is the % deviation allowed from homeostatic set point (input the quality requirement here)
- value of <1 means only 1 test is needed to detect a significant change in patient status.



## Example Using Validation, Six Sigma and Biologic Variation calculations: Homocysteine

#### Performance Characteristics of Six Homocysteine Assays

Sonia L. La'ulu, <sup>1</sup> Mindy L. Rawlins, <sup>1</sup> Christine M. Pfeiffer, PhD, <sup>2</sup> Mindy Zhang, MD, <sup>2</sup> and William L. Roberts, MD, PhD<sup>5</sup>

Key Words: Homocysteine; Method comparison; Imprecision

DOI: 10.1309/AJCP64BJIPNPSQDJ

American Journal of Clinical Pathology 2008; 130:969-975 Comparison of Six Homocysteine methods on 5 instruments



## Homocysteine:

Determining the Size and Shape of the Target

- · Sigma-metrics as an assessment tool
- Find the quality requirement:
  - Non-regulated analyte by CLIA
  - Ricos et al database gives 17.7%
- · Pick critical level of performance: 15 umol/L



## Homocysteine:

Measuring the Method Performance (arrow)

- CV: total imprecision study performed
  - Method A at mean of 17 umol/L, 2.1% CV



## Homocysteine:

Measuring the Method Performance (arrow)

- How to calculate Bias? (comparison study with HPLC reference method, Deming Regression used)
  - Use the Regression equation:

```
NewMethod= (slope * OldMethod) + Y-intercept
```

Bias (in units) = (NewMethod – OldMethod)

Bias% = |Bias| / OldMethod



## Homocysteine:

Measuring the Method Performance (arrow)

- Bias: comparison study with HPLC reference method, Deming Regression used
  - Method A: slope = 0.93, Y-Intercept = 0.64
  - Bias = NewMethod OldMethod = (((15\*0.93)+0.64) - 15) = (13.95 + 0.64) - 15 = 14.59 - 15 = - 0.41
  - Bias % = abs(-0.41) / 15 = 2.73%



## Homocysteine:

Measuring the Method Performance (arrow)

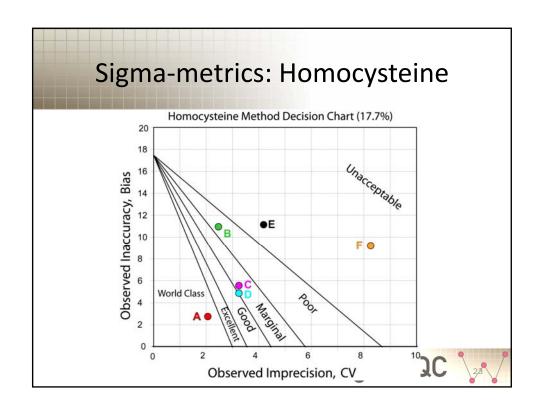
- Sigma-metric: (TEa Bias ) / CV
  - -(17.7-2.73)/2.1
  - -14.97/2.1=7.1
  - Method A Sigma-metric: 7.1

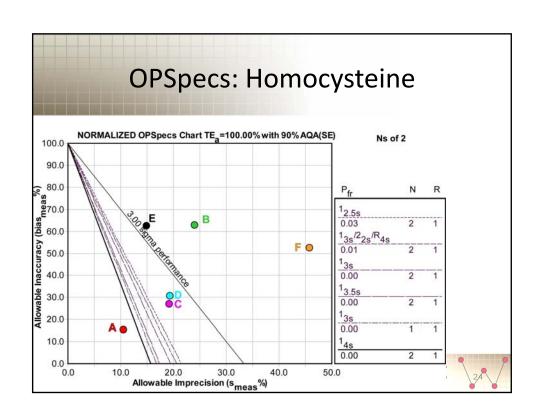


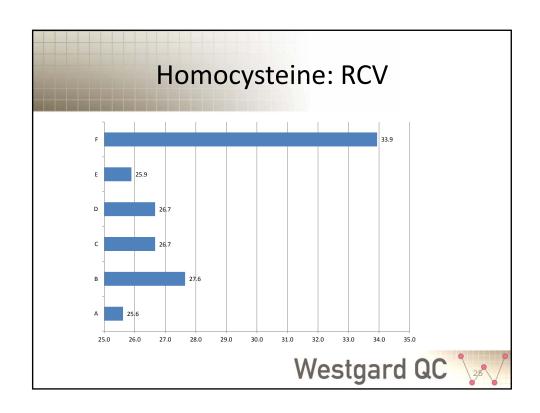
# Homocysteine: Data table

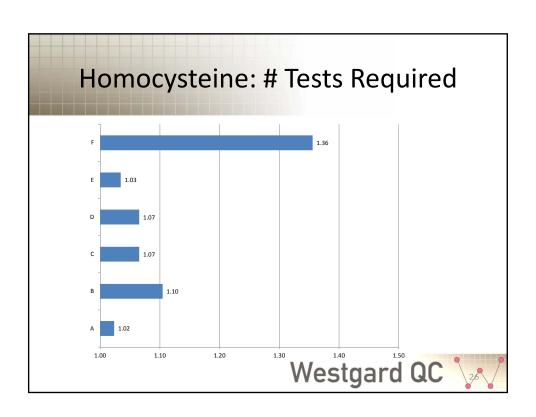
Method	Imprecision	Bias	Sigma-metric
Α	2.1	2.73	7.1
В	4.3	11.3	1.5
С	3.4	4.93	3.8
D	3.4	5.33	3.6
E	2.5	11.2	2.6
F	8.3	9.1	1.0











### Outline



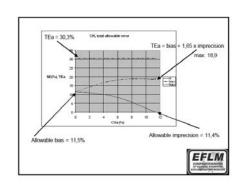
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## Milan 2014, Oosterhuis Outcry

Gross Overestimation of Total Allowable Error Based on Biologic Variation, Wytze P. Oosterhuis, Clin Chem 2011; 57:1334

- Biologic I and B specifications are maximums that should not be combined in traditional Tea format
- Example CK, biodatabase estimates 30.3% TEa while Oosterhuis model projects maximum of 18.9% Tea
- Essentially, Biodatabase goals are too big





## Milan 2014, Carobene Correction

- Around 40-80% of analytes have their quality requirements specified on the basis of just one paper = TOO FEW
- Only 25% of the papers have been published in last 14 years (after 2000) = TOO OLD
- It is unknown whether all of the papers in the database adhered to the proper study protocol = TOO UNRELIABLE
- Impossible to calculate confidence intervals Westgard QC

#### Can labs hit the current Ricos Goals?

#### **UK MAPS project estimates:**

- < 25% labs achieve HbA1c goals (7.0%)</li>
- < 25% labs achieve glucose goals (7.0%)</li>
- NO labs achieve creatinine goals (8.2%)
- Half of labs achieve cholesterol goals (8.5%)
- Less than half of labs achieve HDL goals (11.1%)

Opinion Paper

Nuthar Jassam\*, John Yundt-Pacheco, Rob Jansen, Annette Thomas and Julian H. Barth

Can current analytical quality performance of

UK clinical laboratories support evidence-based
guidelines for diabetes and ischaemic heart
disease? — A pilot study and a proposal

Keywords: analytical performance; internal quality control; σ metric.

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ohn Yunder Pacheco, Bio Sald Labostories (2 quality system Obvision,
riano, Fairiew, TK, USA
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## Sodium goals

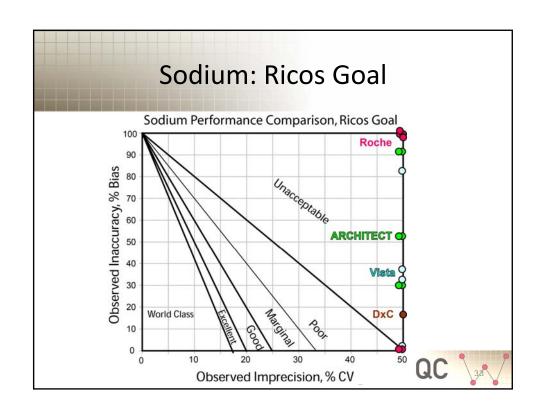
Analyte	Quality Goal				
	CLIA	Desirable Biologic Goal	RCPA	Rilibak	Spanish Minimum Consensus
Sodium	± 4 mmol/L	± 0.9%	± 3 mmol/L ≤ 150 mmol/L; ± 2% > 150 mmol/L	± 5.0%	± 5.0%

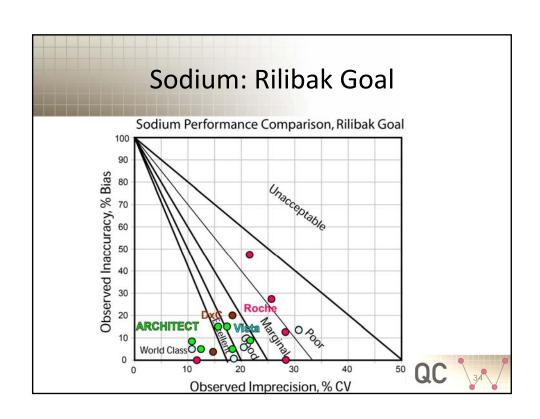


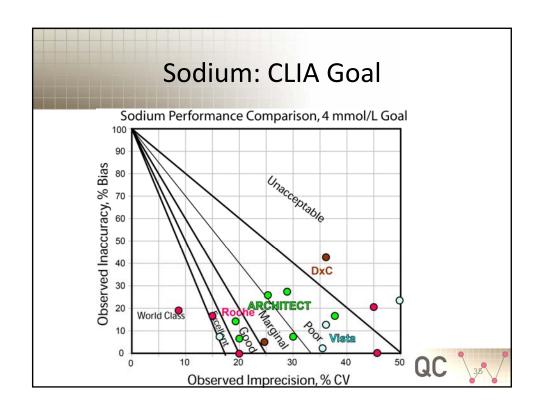
### **Data Sources**

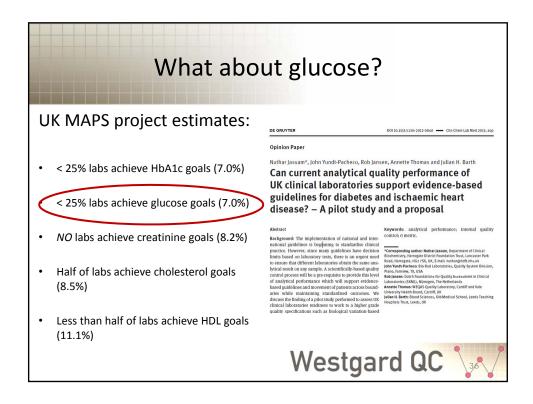
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- Sigma metrics used to access analytical quality of clinical chemistry assays: importance of the allowable total error (TEa) target. Hens K, Berth M, Armbruster D, Westgard S. Clin Chem Lab Med 2014 (July issue)



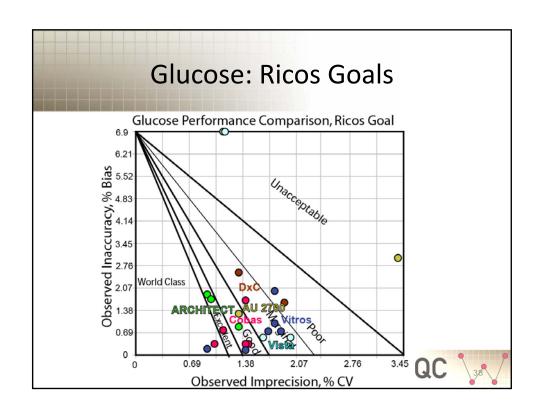


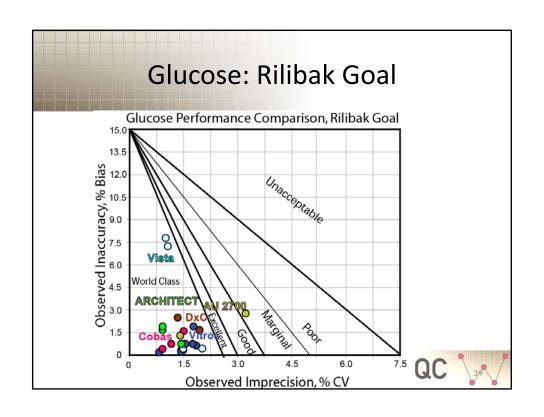


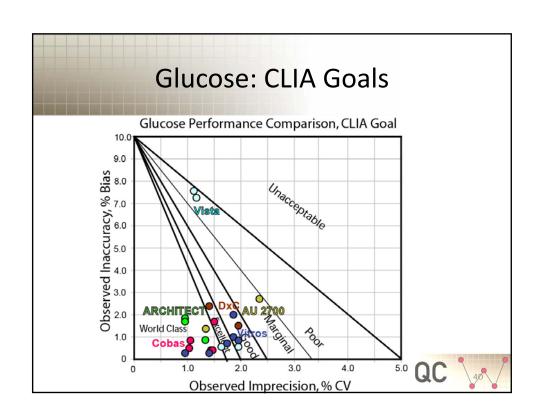




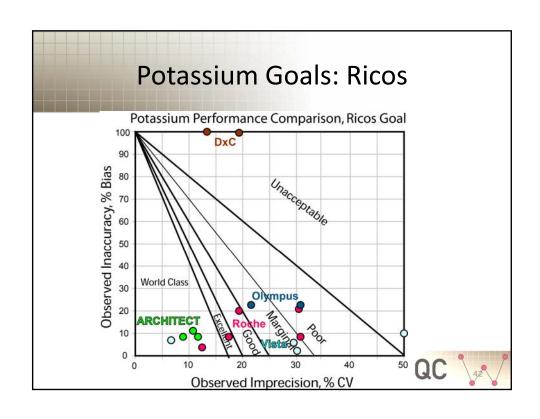
Glucose goals								
Analyte	Quality Goal							
	CLIA	Desirable Biologic Goal	RCPA	Rilibak	Spanish Minimum Consensus			
Glucose	± 10%	± 6.9%	± 0.4 mmol/L ≤ 5.0 mmol/L; ± 8% > 5.0 mmol/L	± 15.0%	± 11%			
			West	gard	QC 3			

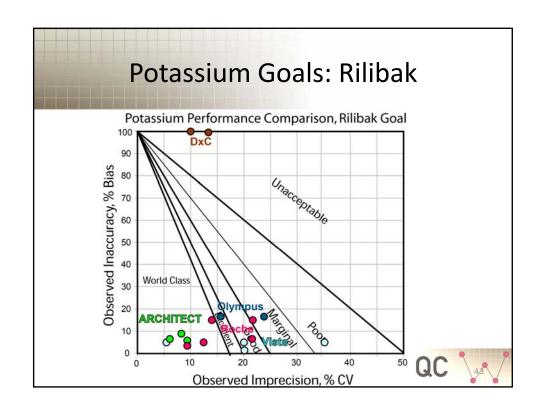


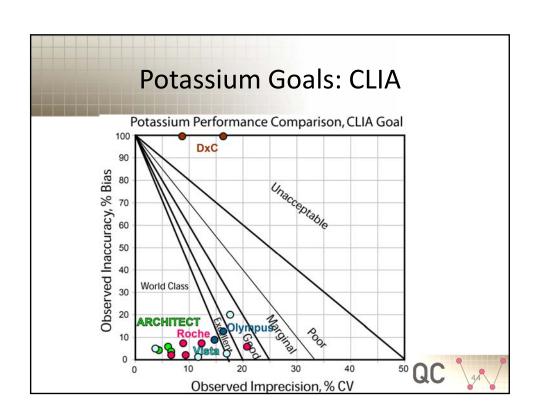




	Pc	otassiu	m Goal	S	
Analyte	Quality Goal				
	CLIA	Desirable Biologic Goal	RCPA	Rilibak	Spanish Minimum Consensus
Potassium	± 0.5 mmol/L	± 5.8%	± 0.2 mmol/L ≤ 4.0 mmol/L; ± 5% > 4.0 mmol/L	± 8.0%	± 8.0%
			West	nard	QC \48







	C	hlorid	e Goals		
Analyte	<b>Quality Goal</b>				
	CLIA	Desirable Biologic Goal	RCPA	Rilibak	Spanish Minimum Consensus
Chloride	± 5.0%	± 1.5%	± 3.0 mmol/L	± 8.0%	± 9.0%
			Westo	gard	QC \45

