

## REFERENCE CHANGE VALUE (RCV)

### A TOOL FOR INTERPRETING SERIAL LABORATORY RESULTS

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### RCV: "WHAT PROBABILITY?"

If either an increase or a decrease in a test result is meaningful (i.e. [serum calcium]), and we want the change to have only a 5% chance of being due to random variation, then  $Z = 1.96$ .

For only a 1% chance,  $Z = 2.58$

If a change in only one direction is meaningful (i.e. [troponin]), then for a 5% chance of being in one tail of the distribution  $Z = 1.65$

$$RCV = (\sqrt{2}) \times (Z) \times \sqrt{(CIVa)^2 + (CVI)^2}$$

### RCV: ANALYTICAL VARIANCE

#### DETERMINATION OF CV<sub>a</sub>

- Initial method validation data for imprecision
- In-house QC data
- Replicate analysis of specimens in the RCV experiment
- Within-run vs. total CV<sub>a</sub>

$$RCV = (\sqrt{2}) \times (Z) \times \sqrt{(CIVa)^2 + (CVI)^2}$$

### RCV: OBJECTIVES

At the end of this presentation, participants will be able to:

1. Explain the statistical theory that underpins the calculation of the relative (reference) change value (RCV).
2. Identify the inputs needed to calculate the RCV.
3. Propose a research design to generate the inputs needed to calculate the RCV.

### RCV: DEFINITION

RCV is the % change in 2 sequential lab results for the same individual that has a specified probability of representing a true change in the state of an individual, as opposed to representing random variation.

$$RCV = (\sqrt{2}) \times (Z) \times \sqrt{(CIVa)^2 + (CVI)^2}$$

CV<sub>a</sub> = analytical coefficient of variation

CV<sub>i</sub> = within-individual coefficient of biological variation

Z = interval of standard deviations (in a Gaussian distribution) between two values that corresponds to a given probability that the difference is due to random variation

### WHY RCV?

#### WHY NOT REFERENCE RANGE?

Many analytes have a low "index of individuality" (Iol).

$$Iol = (CV_i / CV_g)$$

CV<sub>i</sub> is the within-individual biological variation

CV<sub>g</sub> is the between-individual biological variation

If Iol > 1.4, then reference ranges will usually detect biologically significant changes

If Iol < 0.6, then reference ranges will usually fail to detect biologically significant changes

### WHY RCV?

#### WHY NOT REFERENCE RANGE?

Consider the case of serum creatinine:

A 50% increase in serum [creatinine] is associated with a 50% decrease in GFR

Most individuals in the lower half of the reference range can lose 50% of their renal function and still have a "normal" serum [creatinine].

CV<sub>i</sub> for serum [creatinine] = 6%

CV<sub>a</sub> for serum [creatinine] @ 0.74 mg/dL = 3.6%

RCV for serum [creatinine] ~16.3%

### RCV: ANALYTICAL VARIANCE

#### DETERMINATION OF CV<sub>a</sub>

- Initial method validation data for imprecision
  - (+) should have within-run CV<sub>a</sub> (repeatability)
  - (-) data may be out of date
- In-house QC data
  - (+) cumulative s.d. more appropriate for long times between specimens
  - (-) potential matrix effect

### RCV: ANALYTICAL VARIANCE

#### DETERMINATION OF CV<sub>a</sub>

- Replicate analysis of specimens in the RCV experiment (within-run only)
  - (+) permits nested ANOVA when all specimens are analyzed in a single run
  - (-) expensive for some reagents
- Within-run CV<sub>a</sub> permits statistical modeling to extract both analytical and biological variance from a single run
- Total CV<sub>a</sub> better: reflects the imprecision inherent in how sequential specimens are actually run

## RCV: BIOLOGICAL VARIANCE

### DETERMINATION OF CVi

- Look it up vs. measure it
- Population to sample
- Sampling interval
- # of subjects
- # of samples per person

$$RCV = (\sqrt{2}) \times (Z) \times \sqrt{(CVA)^2 + (CVi)^2}$$

## RCV: BIOLOGICAL VARIANCE

### WHY NOT JUST LOOK IT UP?

- Qi et.al. *Clin Chim Acta* 2015; 450:233-236  
vs.  
Matyar et.al. *J Clin Lab Anal* 2016; 30:1081-1085
- RCV for neuron-specific enolase 37% vs. 66%
  - CVi 13.2% vs. 21.5%
  - # subjects 20 vs. 13
  - # samples 7 vs. 4
  - sampling frequency hourly/daily vs. biweekly
  - both used Roche reagents and Cobas analyzers

## RCV: BIOLOGICAL VARIANCE

### SAMPLING INTERVAL

- Hilderink et.al. *Clin Chem Lab Med* 2017; 55:1013-1024
- Measured CBC parameters in older adults
  - Samples drawn hourly for 24 hours from venous catheter
- Fraser et.al. *Am J Clin Pathol* 1989; 92:465-470
- Measured CBC parameters in older adults
  - Samples drawn biweekly for 18-20 weeks
- CVi similar in both studies except for WBC and platelets

## RCV: BIOLOGICAL VARIANCE

### DETERMINATION OF CVi

<https://www.westgard.com/biodatabase1.htm>

Database originally published by Ricos et.al. *Scand J Clin Lab Invest* 1999;59:491-500

Last updated in 2014...now ~370 analytes

$$RCV = (\sqrt{2}) \times (Z) \times \sqrt{(CVA)^2 + (CVi)^2}$$

## RCV: BIOLOGICAL VARIANCE

### POPULATION TO SAMPLE FOR CVi HEALTHY vs. STABLE DISEASE

- Braga et.al. *Clin Chim Acta* 2010; 411:1606-1610
- meta analysis for biological variation in HgbA1c
  - 4 studies using healthy individuals: CVi all <2%
  - 10 studies using diabetics: CVi from 2.4-9.8%
- Aakre et.al. *Clin Chem* 2014; 60:838-847
- CVi for cTnI=15.6%, cTnT=8.3% for healthy
  - CVi for cTnI=14.3%, cTnT=8.3% for hemodialysis

## RCV: BIOLOGICAL VARIANCE

### # of SUBJECTS, # of SAMPLES

- Roraa et.al. *Clin Chem* 2012; 58:1306-1313
- Used computer simulations (10,000 data sets) and nested ANOVA to study the effect of varying:
- # of subjects (10, 15, 20, 40, 80, 160)
  - # of samples per subject (2, 4, 6, 8, 10)
  - # of replicates measured per sample (2, 3, 4)
  - (CVA/Cvi) (0.25, 0.5, 0.75, 1.0, 1.5, 2.0, 3.0)
- on the confidence interval for the estimate of CVi, and on the power to estimate CVi

## RCV: BIOLOGICAL VARIANCE

### WHY NOT JUST LOOK IT UP?

Carobene et.al. *Clin Chem Lab Med* 2013; 51:1997-2007

- meta analysis for biological variation in AST,ALT,GGT
- CVi from 3% - 58%
- # subjects from 10-274
- # samples from 2-16
- sampling frequency from daily to monthly

## RCV: BIOLOGICAL VARIANCE

### SAMPLING INTERVAL

- Wu et.al. *Clin Chem* 2009; 55:52-58  
Wu et.al. *Clin Biochem* 2012; 45:714-716
- CVi for cTnI
- |                     |                        |                       |
|---------------------|------------------------|-----------------------|
| <u>Hourly (4hr)</u> | <u>biweekly (8wks)</u> | <u>9 months (avg)</u> |
| 9.7%                | 14%                    | 28%                   |

## RCV: BIOLOGICAL VARIANCE

### # of SUBJECTS, # of SAMPLES

- Roraa et.al. *Clin Chem* 2012; 58:1306-1313
- The more subjects, the smaller the  $CI_{95\%}$  for CVi
  - Minimal decrease in  $CI_{95\%}$  with more than 8 samples per individual
  - If  $(CVA/Cvi) < 1$ , minimal decrease in  $CI_{95\%}$  with more than 2 replicates per sample
  - If  $(CVA/Cvi) \leq 1$ , power to detect  $CVi > 0$  can be optimized with 10 subjects, 8 samples each, analyzed in duplicate
  - Fewer samples per subject required if  $> 10$  subjects

## RCV: EXPERIMENTAL DESIGN

### ALL IN ONE RUN vs. REAL TIME ANALYSIS

#### ALL IN ONE RUN

- Freeze samples as they are collected
- Thaw all samples when study complete
- Analyze all samples in duplicate at the same time
- (+) no between-run or between-day CVa
- (+) data can be analyzed with nested ANOVA

#### REAL TIME ANALYSIS

- Analyze samples as they are collected
- (+) avoids freeze-thaw effect
- (+) reflects how sequential clinical samples are run

## RCV: NESTED ANOVA

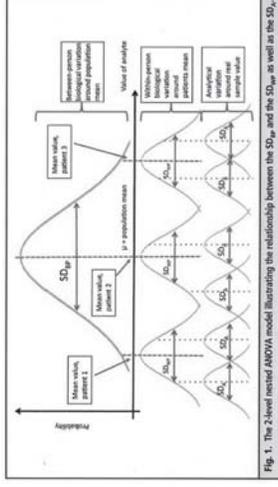


Fig. 1. The 2-level nested ANOVA model illustrating the relationship between the  $SD_p$  and the  $SD_{sp}$  as well as the  $SD_{sp}$ .

Roraas et al. *Clin Chem* 2012; 58:1306-1313

## RCV: ASSUMPTIONS (so far)

- Distributions of measured results are Gaussian
- Differences between measures are random with no autocorrelation between sequential results
- Variances are homogeneous
- No pre-analytical variation

## RCV: NON-GAUSSIAN DISTRIBUTION

- Calculate the ln of each data point
- Calculate CVa and CVi using ln-transformed data
- $RCV_{pos} = (\exp[1.96(2)^{1/2}(CVa^2 + CVi^2)^{1/2}] - 1) \times 100\%$
- $RCV_{neg} = (\exp[-1.96(2)^{1/2}(CVa^2 + CVi^2)^{1/2}] - 1) \times 100\%$
- $RCV_{pos}$  may not =  $RCV_{neg}$  when underlying distribution is skewed

## RCV: CASE STUDY FRUCTOSAMINE vs. FRUCTOSAMINE/ALBUMIN

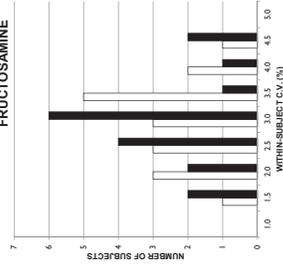
"Biological Variation of Serum Fructosamine in Healthy Subjects"

Supported by a Member Grant from the ASCLS E&R Fund and by CTSA #UL1TR000058 from NIH

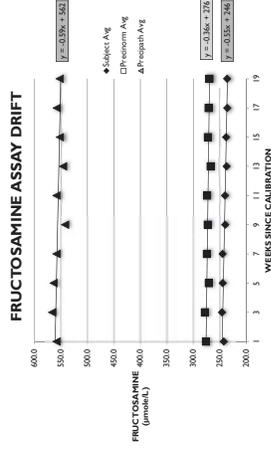
Does correcting the serum fructosamine result for the serum [albumin] in the same specimen reduce the CVi?

## RCV: CASE STUDY FRUCTOSAMINE vs. FRUCTOSAMINE/ALBUMIN

DISTRIBUTION OF WITHIN SUBJECT C.V.s FOR FRUCTOSAMINE

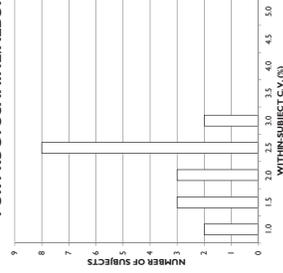


## RCV: CASE STUDY FRUCTOSAMINE vs. FRUCTOSAMINE/ALBUMIN



## RCV: CASE STUDY FRUCTOSAMINE vs. FRUCTOSAMINE/ALBUMIN

DISTRIBUTION OF WITHIN-SUBJECT C.V.s FOR FRUCTOSAMINE/ALBUMIN RATIO



### RCV: CASE STUDY FRUCTOSAMINE vs. FRUCTOSAMINE/ALBUMIN

CVi	MEAN/MEDIAN	RANGE
Fructosamine (real time)	3.2% / 3.4%	1.8% - 4.8%
Fructosamine (all in one run)	3.0% / 3.1%	1.6% - 5.0%
Fructosamine/albumin (real time)	2.4% / 2.6%	1.2% - 3.2%

Using total CVa for the low control (1.3%) and Z= 1.96:  
RCV for fructosamine = 10.1%

### RCV: ALTERNATIVE USE

$$RCV = (\sqrt{2}) \times (Z) \times \sqrt{(CVa)^2 + (CVi)^2}$$

$$Z = \frac{RCV}{(\sqrt{2}) \sqrt{(CVa)^2 + (CVi)^2}}$$

### RCV: SUMMARY

RCV is a more sensitive indicator of a change in the biological / physiological state of an individual than whether or not sequential results fall within or outside the reference range for an analyte, when the within-individual biological CV is less than the between-individual CV.

How the RCV is determined must be customized for each analyte, with respect to:

- the population to which it will be applied
- how often the test is normally ordered
- the achievable CVa and (CVa/CVi)

### RCV: QUESTIONS

