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Research Article



Evaluation of analytical performance specifications of routine clinical biochemistry tests with biological variation-based total allowable error criteria

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Abstract

Objectives: Total allowable error (TEa) is the analytical quality specification that determines the acceptable limits for a test result. The aim of the present study was to determine whether third-party control data (percentage coefficient of variation (CV), percentage bias, and %TEa) of 26 clinical biochemistry parameters collected over a period of 6 months would meet biological variation (BV)-based analytical quality specifications (minimum, desirable, and optimum) in which BV quality specification might be more appropriate for our laboratory use.

Methods: The study was conducted for 6 months on tests with the TEa values determined according to BV and using third-party controls, Unity Real Time[®] software, and Beckman Coulter AU5800 clinical chemistry analyzer.

Results: The BV minimum specification is the easiest target to be performed by laboratories since it has the widest limits. It is also known that the BV already fulfills the other specifications (minimum and desirable) in tests where the optimum specification targets can be achieved, and the test performance classifications of our work are made according to this information. The minimum specification is only in Level 2 control serum total cholesterol test. The optimum specification with the narrowest limits within the BV criteria and the most difficult to achieve was met in nine tests (alanine aminotransferase, direct bilirubin, total bilirubin, creatine kinase, gamma-glutamyl transferase, ferritin, lipase, triglyceride, and uric acid (UA) for both control levels. Outside of these tests, in general, the desirable specification that each laboratory is aiming to fulfill is met.

Conclusion: It is very valuable to set the boundaries of quality specifications targeted for analytical performance based on BV. These values are determined by the intra-individual and inter-individual variability of the test and, thus, define a test-specific target. In addition to defining the TEa for each specification of BV, the definition of bias and CV boundaries can give a more objective idea of the source of error.

Keywords: Analytical performance, biological variation, total allowable error

Thousands of results are produced in clinical laboratories using assay procedures. These results are used for many purposes, such as group screening, diagnosis, monitoring, and differential diagnosis of diseases. Many variations are encountered in total assay procedures in laboratories, including pre-analytical, analytical, and biological variations (BVs). In order to obtain reliable results, clinical laboratories should minimize all sources of variation and manage all the testing process well. This management is applicable for all types of variation except BV [1, 2].

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BV is defined as the random fluctuations of an analyte concentration around a homeostatic set point in an individual [3]. The application of BV data in the clinical laboratory is very comprehensive and valuable. The components of BV can be used to determine the timing and frequency of sampling in the pre-pre-analytical and pre-analytical phases; quality specifications, such as imprecision, bias, and total error in the analytical phase; and population-based reference intervals, individuality index, and reference change values in the post-analytical and post-post-analytical phases; they are very important during the interpretation and evaluation of test results [2, 4].

The aim of the present study was to determine whether thirdparty control data of 26 clinical biochemistry parameters collected over a period of 6 months would meet BV-based analytical quality specifications (minimum, desirable, and optimum). For this purpose, we decided to use third-party control samples independent of the manufacturers of the devices and reagents present in our laboratory. We procured control samples sufficient for a 1-year period. These samples met the required conditions in terms of stability and analyte concentration at clinical decision level, and so on. We used these control samples for internal quality control in our laboratory, and the results were evaluated by the Unity Real Time[®] program, which was used to compare results on an international scale, and the Westgard Advisor[™] software, which was an integral part of the program.

Using the Unity Real Time[®] program, the mean, standard deviation, and coefficient of variation (CV) values were calculated for the measurands evaluated using the internal quality assurance program.

Bias is defined as "the systematic deviation in the test results indicating the difference between the mean of the serial replicate measurements and the expected true value or the value that is accepted as true" [5]. The programs mentioned above enable us to compare our mean values as calculated for each test using peer-group means, which we accepted as the true value. Unlike the external quality assessment programs, in which a single measurement result is compared with the peer-group mean, these programs allowed us to detect our actual bias.

We investigated whether the CV and bias values of each test and the total analytical error (TAE) derived from these values were within the CV, bias, and total allowable error (TEa) limits that are required by each of the three quality specifications of BV (minimum, desirable, and optimum), and also, which BV quality specification was more appropriate for use in our laboratory.

Materials and Methods

In the present study, test performances in Gaziosmanpasa Taksim Education and Research Hospital's Biochemistry Laboratory were evaluated according to the BV acceptance criteria (minimum, desirable, and optimum) as updated by Ricos and colleagues [6]. Furthermore, the quality specifications of BV that could be met by the performance of routinely analyzed biochemical tests were identified. The study was conducted over a 6-month period from February 3 to August 3, 2016 using a Beckman Coulter AU5800 clinical chemistry analyzer. The study duration and the method to be used in data evaluation were both determined according to the Clinical Laboratory and Standards Institute (CLSI) C24-A3 guideline, which states that the mean, standard deviation, and control limits should be determined based on the cumulative data obtained over a period of 3 to 6 months [7].

During the study period, the preliminary data were collected over 20 working days in 1 month, with 40 measurements made for each level, and these data were used to calculate the preliminary mean and standard deviation values (Biorad Lypocheck Clinical Chemistry Control Level 1; lot no: 14461) and Level 2; lot no: 14462). The cumulative mean and standard deviation values were calculated by adding the new data to the initial values. Cumulative control limits are known to be a better indicator of long-term test performance [8].

For each test, the control rules and control measurement numbers were determined using the Westgard Advisor software according to the analytical CV and bias values of the tests. The Unity Real Time[®] software was set to detect control violations automatically. Thus, the control numbers and rules were applied during the process. The detected rule violations were subjected to the necessary intervention, and the interventions

Table 1. Test and methods in our laboratory

Test Name	Test Unit	Method
Albumin	g/dL	Colorimetric BCG
ALT	IU/L	IFCC (and/or PLP Activation)
ALP	IU/L	IFCC AMP Buffer
Amylase	IU/L	CNP-Trioside CNPG3
AST	IU/L	IFCC (and/or PLP Activation)
Direct Bilirubin	mg/dL	Colorimetric Diazo method
Total Bilirubin	mg/dL	Colorimetric Diazo method
Chlorid	mEq/L	ISE indirect
Calcium	mg/dL	Colorimetric Arsenoza III
HDL-Cholesterol	mg/dL	Enzymatic immunoinhibition
Total Cholesterol	mg/dL	CHO-POD
Creatinine kinase	IU/L	Modified IFCC method
Creatinine	mg/dL	Modified Jaffe's Kinetic
GGT	IU/L	G-Glutamyl carboxy nitroanylide
Glucose	mg/dL	Hexokinase
Ferritin	µg/dl	Colorimetric TPTZ
LDH	IU/L	LDH (L-P) IFCC
Lipase	IU/L	Enzymatic 1.2-diglyceride
Magnesium	mEq/L	Xylidyl blue
Potassium	mEq/L	ISE indirect
Phosphor	mg/dL	Phosphomolybdate
Total proteine	g/dL	Biuret
LDL-Cholesterol	mg/dL	Enzymatic immunoinhibition
Sodium	mEq/L	ISE indirect
Triglyceride	mg/dL	GPO-POD
Uric acid	mg/dL	Uricase

	Wes	stgard Advisor
Existing QC Rules Configure TE	a Design QC Rules	s Preferences
Analyte	TEa Selection	^
AFP	CLIA	Analyte:
Albumin	CLIA	Insuin
Alkaline Phosphatase	CLIA	
ALT (ALAT/GPT)	CLIA	TEA
Amylase	CLIA	
Anti-Thyroglobulin	BV Des bias/ Des in	BV Des bias/ Des imprecision
Anti-Thyroperoxidase (Anti-TPO)		
AST (ASAT/GOT)	CLIA	BV Des bias/ Des imprecision
Blirubin, Direct/BC (DBIL)	BV Des bias/ Des in	BV Des bias/ Min imprecision
	CLIA	BV Des bias/ Opt imprecision
Blirubin, Total/TBIL		BV Min bias/ Des imprecision
Calcium	CLIA	BV Min bias/ Min imprecision
CEA (Carcinoembryonic Antigen)		mpr BV Min bias/ Opt imprecision
Chloride	CLIA	BV Opt bias/ Des imprecision
Cholesterol, HDL	CLIA	BV Opt bias/ Min imprecision
Cholesterol, LDL	BV Des bias/ Des in	mpr BV Opt bias/ Opt imprecision
Cholesterol, Total	CLIA	GOST
CK (Creatine Kinase)	CLIA	IPH Belgium
CK-MB Activity	BV Des bias/ Des in	
Cortisol	CLIA	QMP-LS - (no bias)
Creatinine	CLIA	QMP-LS (bias)
DHEA-Sulfate	BV Des bias/ Des in	mor RCPA
Estradiol, E2	BV Des bias/ Des in	User Defined
Fentio	BV Des bias/ Des in	Tightest
Folate	BV Des bias/ Des in BV Des bias/ Des in	Loosest
Foliate	BV Des bias/ Des in BV Des bias/ Des in	State of the Att - no bas
GGT (Gamma Glutamyltransfer		
Glucose	CLIA	
hOG	CLIA	
Insulin	BV Des bias/ Des in	mpr
Iron	CLIA	Consensus Group (if applicat
LD (Lactate Dehydrogenase)	CLIA	Const
LH	BV Des bias/ Des in	
Lipase	BV Des bias/ Des in	mpr Peer
Magnesium	CLIA	
Phosphorus	BV Des bias/ Des in	mpr Group Data Range:
Potassium	CLIA	
Progesterone	3SD	Monthly
Prolactin	BV Des bias/ Des in	mor
Protein, Total, Serum	CLIA	For All Analytes
PSA. Free	3SD	- server Analytes
PSA, Total	BV Des bias/ Des in	
PTH, Intact	BV Des bias/ Des in BV Des bias/ Des in	
Sodum	CLIA	Save
	BV Des bias/ Des im	
T3, Free		
T4, Free	CLIA	 Cance
<		>

Figure 1. Determination of the TEa configuration for each test

were recorded. Percentage bias (%B), percentage CV (%CV), and TAE were calculated for 26 tests. We followed the policy that we established as the target for the Clinical Laboratory Improvement Amendments (CLIA) criteria for these tests in our laboratory and then made comparisons with the BV acceptance criteria. During the 6-month study period, performance reports of all tests were obtained separately according to CLIA and BV.

Examined tests and assay methods

All examined tests were analyzed using a Beckman Coulter AU5800 analyzer throughout the 6-month study period in accordance with the planned control procedure. The evaluation comprised 26 tests all analyzed using the routine clinical chemistry analyzer. Table 1 shows these tests, their units, and the assay methods.

Quality control procedure

As shown in Fig. 1, TEa for each test was determined according to BV specifications using the software program. The Westgard Advisor[™] module, which was pre-defined in the program, the calculated %B, %CV, Δ SE (critical systematic error), and sigma (σ) values for the desired parameters for a given time interval. Based on these data, the program automatically calculated the required control rule and control measurement number at the analytical quality assurance percentage, providing the best test performance according to the most appropriate error detection (Ped; probability of error detection) and false rejection (Pfr; probability of false rejection) values with the help of both OPSpecs and critical error graphics.

Allowable error values based on BV specifications

Table 2 shows the %TEa, %B, and %CV limits of the examined tests according to BV minimum, desirable, and optimum specifications.

All %TEa, %B, and %CV values for the desired specifications were derived from the BV data. For minimum and optimum specifications, the %TEa, %B, and %CV limit values for the tests that did not have a specified value in the data were calculated using the formula given below [6]:

For minimum specification:

%CV<0.75CV_w %B<0.375 (CV_w²+CV_b²)^{1/2} %TEa<0.375 (CV_w²+CV_b²)^{1/2}+1.65 (0.75CV_w)

For optimum specification:

%CV<0.25CV

%B<0.125 (CV_w²+CV_b²)^{1/2} %TEa<0.125 (CV_w²+CV_b²)^{1/2}+1.65 (0.25CV_w)

where:

%TEa: total allowable error; a quality requirement for a given test,

%B: deviation from the true value,

%CV: imprecision observed for a given method,

CVw: intra-individual biological variation,

CVb: inter-individual biological variation,

1.65: coefficient for 95% confidence interval.

For all tests, TAE, %CV, and %B were calculated, and the BV acceptance criteria (minimum, desirable, and optimum) were used for comparison. TAE for each level of each test was calculated using the TAE=%B+1.65×CV formula in the Excel program. The %B value was accepted as its absolute value. 1.65 is the coefficient used for 95% confidence interval. The performance reports include the observed %B, the %CV values for the given level, and the recommended control rule and numbers based on the BV minimum, desirable, and optimum acceptance criteria.

Results

The tests using the Level 1 and Level 2 control materials that we assayed in the clinical biochemistry laboratory following

Test Name	Minir	num specific	ations	Desir	able specific	ations	Optimum specifications		
	CV(%)	B(%)	TEa(%)	CV(%)	B(%)	TEa(%)	CV(%)	B(%)	TEa(%)
Albumin	2.4	2.1	6.1	1.6	1.43	4.07	0.8	0.7	2.0
ALT	14.6	17.2	41.2	9.7	11.48	27.48	4.9	5.7	13.7
ALP	4.8	10.1	18.1	3.23	6.72	12.04	1.6	3.4	6.0
Amylase	6.5	11.1	21.9	4.4	7.4	14.6	2.2	3.7	7.3
AST	9.2	9.8	25.0	6.15	6.54	16.69	3.1	3.3	8.3
Direct Bilirubin	27.6	21.3	66.8	18.4	14.2	44.5	9.2	7.1	22.3
Total Bilirubin	16.4	13.4	40.4	10.9	8.95	26.94	5.5	4.5	13.5
Chlorid	0.9	0.7	2.2	0.6	0.5	1.5	0.3	0.3	0.75
Calcium	1.4	1.3	3.6	1.05	0.82	2.55	0.5	0.4	1.3
HDL-Cholesterol	5.5	8.4	17.4	3.65	5.61	11.63	1.8	2.8	5.8
Total Cholesterol	4.5	6.2	13.5	2.98	4.1	9.01	1.5	2.1	4.5
Creatinine kinase	17.1	17.3	45.5	11.4	11.5	30.3	5.7	5.8	15.2
Creatinine	4.5	5.9	13.4	2.98	3.96	8.87	2.8	3.2	7.7
GGT	10.1	16.6	33.2	6.7	11.06	22.11	3.4	5.5	11.1
Glucose	4.2	3.5	10.4	2.8	2.34	6.96	1.4	1.2	3.5
Ferritin	20.0	13.2	46	13.3	8.8	30.7	6.7	4.4	15.3
LDH	6.5	6.5	17	4.3	4.3	11.4	2.2	2.2	5.7
Lipase	24.2	17.0	56.8	16.1	11.31	37.88	8.1	5.7	19
Magnesium	2.7	2.8	7.2	1.8	1.8	4.8	0.9	0.9	2.4
Potassium	3.5	2.7	8.4	2.3	1.81	5.61	1.2	0.9	2.8
Phosphor	6.1	5.1	15.2	4.08	3.38	10.11	2.0	1.7	5.1
Total proteine	2.1	2.0	5.4	1.38	1.36	3.63	0.7	0.7	1.8
LDL-Cholesterol	5.9	8.2	17.8	3.9	5.46	11.9	2.0	2.7	6.0
Sodium	0.5	0.3	1.1	0.3	0.23	0.73	0.2	0.1	0.37
Triglyceride	14.9	14.4	39	9.95	9.57	25.99	5.0	4.8	13
Uric acid	6.5	7.3	18	4.3	4.87	11.97	2.2	2.4	6.0

Table 2. %TEa, %Bias (%B), and %CV limits of the examined tests according to BV minimum, desirable, and optimum specifications

Table 3. Classification of tests in Level 1 and Level 2 control materials that met the BV TEa acceptance criteria (minimum, desirable, and optimum)

Biologic Variation Acceptance	Criteria Level 1	Level 2
Optimum Specification TEa	ALT, Direct Bilirubin, Total Bilirubin,	ALT, Amylase, AST, Direct Bilirubin, Total Bilirubin,
	HDL-Cholesterol, GGT, Total Cholesterol, Creatinine kinase,	Creatinine kinase, Crea, GGT, Ferritin,
	Ferritin, Lipase, Triglyceride, Uric acid	Lipase, Triglyceride, Uric acid
Desirable Specification TEa	Amylase, AST, Glucose, LDH,	ALP, Calcium, HDL-Cholesterol, Glucose,
	Magnesium, Potassium, Phosphor, HDL-Cholesterol	LDH, Magnesium, Potassium, Phosphor
Minimum Specification TEa	-	Total Cholesterol

the CLIA protocols are presented in Table 3 according to the BV TEa acceptance criteria (minimum, desirable, and optimum) that they met. Table 4 shows the Level 1 and 2 control material tests that met the BV optimum acceptance criteria. Table 5 shows the tests that did not meet any BV acceptance criteria.

Although the analytical CV and/or bias for our laboratory was higher than the allowable CV and/or bias based on BV for both control levels, we principally targeted TAE<TEa based on the BV criteria, and all the evaluations were made accordingly.

In the present study, CVw, CVb, and the allowable CV values derived from these values were obtained from the BV data. For the Level 1 control material, 11 tests (alanine aminotransferase (ALT), direct bilirubin (DB), total bilirubin (TB), high-density lipoprotein cholesterol (HDL-Chol), total cholesterol (T-Chol), creatine kinase (CK), gamma-glutamyl transferase (GGT), ferritin, lipase (Lip), triglyceride (Tg), and uric acid (UA) met the BV optimum, and 8 tests (amylase (Amy), aspartate aminotransferase (AST), glucose (Glu), lactate dehydrogenase (LDH),

Table 4. The imprecision, bias, and TEa values of tests in Level 1 and Level 2 control materials that met the BV optimum acceptance criteria

BV Optimum	BV Optimum Imprecision (%)			Bias	s (%)		TAE	TE	
Q	uality Co	ntrol Level	Optimum	Quality Co	ntrol Level	Optimum	Quality Co	ntrol Level	Optimum
Test Name	CV ₁	CV ₂	cv	Bias ₁	Bias ₂	Bias	TAE,	TAE ₂	TE
ALT	3.0	1.6	4.9	6.33	1.5	5.7	11.4	4.1	13.7
Direct Bilirubin	3.0	2.5	9.2	0.2	3.8	7.1	5.1	8.0	22.3
Total Bilirubin	1.3	1.5	5.5	4.7	1.9	4.5	6.8	4.4	13.5
HDL-Cholesterol	2.1		1.8	0.03		2.8	3.4		5.8
Total Cholesterol	1.2		1.5	0.03		2.1	2.0		4.5
Creatinine kinase	1.4	1.0	5.7	5.4	0.5	5.8	7.7	2.1	15
GGT	1.6	1.6	3.4	2.4	1.1	5.5	5.0	3.7	11
Ferritin	1.4	2.7	6.7	5.8	6.1	4.4	8.0	10.6	15.3
Lipase	4.2	3.7	8.0	0.4	9.2	5.7	7.2	15.4	19
Triglyceride	3.3	3.5	5.0	5.0	0.8	4.8	10.5	6.6	13
Uric acid	1.6	2.0	2.2	1.4	1.2	2.4	4.1	4.5	6.0
Amylase		1.5	2.2		0.3	3.7		2.8	7.3
AST		1.6	3.1		0.9	3.3		3.7	8.3
Creatinine		2.1	2.8		3.5	3.2		7.0	7.7
BV Desirable	Imprec	ision (%)		В	ias (%)		TAE Quality Control Level		TE
 (Quality Co	ontrol Level	Desirable	Quality Co	ontrol Level	Desirable			Desirable
Test Name	CV ₁	CV ₂	cv	Bias,	Bias ₂	Bias	TAE,	TAE ₂	TE
Amylase	1.65		4.4	8.4		7.4	11		14.6
AST	2.6		6.1	8.3		6.5	12.6		16.7
Glucose	2	1.9	2.8	3.5	2.8	2.3	6.9	5.8	6.9
LDH	3.8	2.7	4.3	0.3	2.4	4.3	6.5	6.9	11.4
Magnesium	1.9	1.9	1.8	1.55	0.7	1.8	4.6	3.8	4.8
Potassium	1	1.3	2.3	2.5	1.6	1.8	4.2	3.7	5.6
Phosphor	2.7	2.5	4.1	2.6	1.4	3.4	7	5.5	10
LDL-Cholesterol	4.9		3.9	3.5		5.5	11.5		11.9
ALP	-	4.7	3.2	-	4.6	6.7	-	12.4	12
Calcium	-	1.3	1.1	-	0.2	0.8	-	2.4	2.6
HDL-Cholesterol	-	3.4	3.7	-	1.9	5.6	-	7.6	11.6
BV Minimum	Imprecis	mprecision (%)		Bias (%)			TAE		TE
Q	uality Co	ntrol Level	Minimum	Quality Control Level		Minimum	Quality Control Level		Minimum
Test Name	CV,	cv,	cv	Bias,	Bias	Bias	TAE, TAE,		TE

7.7

6.2

magnesium (Mg), potassium (K), phosphor (P), and low-density lipoprotein cholesterol (LDL-Chol) met the BV desirable TEa criteria. Seven tests (albumin (Alb), alkaline phosphatase (ALP), chloride (Cl), calcium (Ca), creatinine (Crea), total protein (TP), and sodium (Na)) did not meet any BV acceptance criteria.

1.5

4.5

Total Cholesterol

Four tests (Alb, Cl, TP, and Na) did not meet any BV acceptance criteria.

10

13.5

In our study, the Alb, Cl, TP, and Na parameters showed unacceptable performance (TAE>TEa) for both control levels. Additionally, the test performances of ALP, Ca, and Crea were outside the acceptable limits (TAE>TEa) for Level 1 control material.

For the Level 2 control material, 12 tests (ALT, Amy, AST, DB, TB, CK, Crea, GGT, Fe, Lip, Tg, and UA) met the BV optimum, 8 tests (ALP, Ca, HDL-Chol, Glu, LDH, Mg, K, and P) met the BV desirable, and 1 test (T-Chol) met the BV minimum TEa criteria.

When compared with other specification values, the optimum specification TEa, CV, and bias values have the narrowest range and are the most difficult of all BV specifications to achieve.

BV Minimum Imprecision (%)			Bias (%)			TA				
	Quality Control Level		Minimum	Quality Control Level		Minimum	Quality Control Level		Minimum	
Test Name	CV ₁	CV ₂	CV	Bias ₁	Bias ₂	Bias	TAE ₁	TAE2	TAE	
Albumin	2.0	1.7	2.4	5.8	3.7	2.1	9.0	6.6	6.1	
Chlorid	0.9	1.1	0.9	2.1	0.5	0.7	3.6	2.4	2.2	
Total protein	1.9	2.2	2.1	3.0	4.0	2.0	6.2	7.6	5.4	
Sodium	0.9	1.0	0.5	2.5	1.6	0.3	4.0	3.3	1.1	
ALP	4.9	-	4.8	10.5	-	10	18.6	-	18.1	
Calcium	1.2	-	1.4	1.7	-	1.3	3.7	-	3.6	
Creatinine	2.7	-	4.5	12.9	-	5.9	17.3	-	13.4	

Table 5. Tests that did not meet any BV acceptance criteria according to quality control levels

Table 6. Recommended quality control procedure for Level 1 and 2 quality control tests meeting the BV optimum specifications

Test	Level	1		Level 2				
	Recommended rule	Ν	Ped	Pfr	Recommended rule	Ν	Ped	Pfr
ALT	1-2s	4	0.369	0.171	1-5s	2	0.984	0
Direct Bilirubin	1-5s	2	0.948	0	1-5s	2	0.933	0
Total Bilirubin	1-5s	2	0.998	0	1-5s	2	1	0
HDL-Cholesterol	1-3s/2-2s/R-4S/4-1s/8-X	4	0.500	0.039	1-3s/2-2s/R-4S/4-1s/8-X	4	NA	0.039
Creatinine kinase	1-4s	2	0.991	0	1-5s	2	1	0
GGT	1-3s	2	0.949	0.006	1-4s	2	0.947	0
Ferritin	1-4s	2	0.994	0	1-3s/2-2s/R-4S/3-1s/12-X	6	0.948	0.089
Lipase	1-3s/2-2s/R-4S/4-1s	2	0.955	0.012	1-2s	4	0.500	0.171
Triglyceride	1-2s	4	0.369	0.171	1-3s/2-2s/R-4S/3-1s/12-X	6	0.968	0.089
Uric acid	1-3s/2-2s/R-4S/4-1s/8-X	4	0.500	0.039	1-2s	4	0.369	0.171

Table 6 shows the recommended quality control rule(s), the required number of quality control measurements (N), the Ped values, and the Pfr values for the tests that meet these criteria to maintain test performance.

Discussion

In the present study, the results of the routine clinical biochemistry tests for Level 1 and Level 2 quality control materials were evaluated for performance according to the BV (minimum, desirable, and optimum) TEa acceptance criteria. In addition, the study aimed to determine the accuracy, precision, and reliability of the examined tests; to document sustainability of quality; and to improve quality.

Physicians request laboratory tests for approximately 80% of patients presenting to the hospital. Considering this, it is very important to prevent errors by improving the quality of each testing procedure. Problems, such as repeated test requisitions, unnecessary therapeutic interventions, and wasted money, time and labor, plus distrust in the laboratory can all be avoided by improving the quality of the total testing process (TTP) [9].

Laboratory managers have the responsibility of detecting the sources of error impeding the TTP and correcting them. Ana-

lytical errors account for 21% of all errors occurring in the TTP, but they may be easier to manage compared with pre-analytical or post-analytical errors [10, 11]. This is because quality control procedures and several quality planning tools have been devised for analytical quality management.

Three different quality specifications exist according to BV for the evaluation of analytical quality performance. Since BV minimum specification (min. imprecision/min. bias) has the widest range, it is regarded as the easiest target for laboratories to achieve. Furthermore, BV desirable specification (des. imprecision/des. bias) is a target for every laboratory in general. BV optimum specification (opt. imprecision/opt. bias) has the narrowest range and is the most difficult to achieve. When BV optimum specification targets are met, other specifications (minimum and desirable) are already achieved. Similarly, when BV desirable specification targets are met, BV minimum targets are already achieved. The performance test classifications used in the present study are based on this fact [12]. It is considered best for our laboratory to maintain the TEa target for these tests according to BV optimum specifications and according to BV desirable specifications when possible. For those tests that only meet BV minimum specifications or do not meet any BV-based criteria, we aim to develop and implement the necessary procedures to at least achieve the BV desirable specification criteria.

In the present study, we targeted the more stringent, BVbased analytical quality goals instead of the wider analytical quality targets (e.g., CLIA and Turkey). While it is easier to meet the general quality specifications with the existing methodology and technology for some tests, such as Tg using the threestage BV model [3], which is derived from imprecision, bias, and total error limits in BV, it may be harder for other tests, such as Ca, Na, and Cl.

Bal et al. [13] calculated imprecision values for 10 clinical biochemistry parameters using internal and external quality control data from three analyzers of the same brand and compared these values with the %TEa values for Fraser and CLIA. The Alb, Na, TP, and Crea tests did not meet the BV-based quality specifications in any of the three devices, and this was explained by low target TEa limits. Similarly, in our study, while Alb, Na, TP, and Cl tests were able to meet the CLIA TEa targets, the BV TEa targets could not be achieved.

Studies have reported that it is quite difficult to meet the BV TEa criteria for electrolytes that are analyzed via the indirect ion selective electrode method, such as Na and Cl [14, 15]. Indeed, we were not able to meet the BV TEa criteria for these electrolytes in our study. Regardless, it may be possible to achieve the desired targets for these parameters by meticulously checking various details, such as inspecting the cleaning system, following a stringent maintenance protocol, and checking the water purifying systems and consumables.

In our study, different TAE values were obtained for the two levels of control samples depending on the analyte concentrations. BV TEa quality specifications could not be achieved in seven tests for Level 1 control materials and four tests for Level 2 control materials. For instance, Level 1 control results for the ALP and Ca test did not meet the minimum BV-based TEa criteria. One study, which applied Six Sigma to evaluate the performance of analytical phase and calculated bias values from external quality assurance results over a 6-month period, found the sigma values of the internal quality sample for the ALT test to be 2.79 for the normal level and 4.24 for the pathological level [16].

Although calculation of the CV and TAE values for the same analyte at different concentrations is cumbersome and costly, it allows for a more reliable assessment of that analyte's performance. In our study, we calculated the monthly bias values using the daily mean peer-group values of the inter-laboratory internal quality control comparison program. Such an assessment of bias allows any deviation to be noticed earlier than the reported bias in external quality control results. Therefore, we believe that it is very useful and reliable for developing prospective corrective actions.

In their study, Coskun et al. [17] presented their internal and external quality assessment using control samples of the same matrix, and they reported that T-Chol, Glu, Ca, Cl, K, Mg, Na, and GGT for Level 1 control material and T-Chol, Glu, UA, Crea, Ca, Cl, Na, and CK for Level 2 control material do not meet any BV TEa criteria. Their results are consistent with the results of the present study.

In another study, Fernandez-Calle et al. [15] performed a validation assessment of the two new clinical chemistry analyzers brought to their laboratory. For this purpose, they calculated the CV, %B, and total error values of 30 analytes, and they chose the BV criteria as their target. Similar to our study, they used the Bio-Rad control materials and Unity Real Time software, and they used the cumulative peer-group mean from the third-party controls as the reference value when evaluating bias. For each test, the BV criteria were determined (minimum, desirable, and optimum), and bias and imprecision were evaluated separately. The authors emphasized that the control materials that are to be used in such studies should have a matrix similar to the patient samples and should contain an analyte concentration at clinical decision level according to the CLSI guidelines. In their BV-based ISO 15189 laboratory accreditation study, the Na, Cl, and Mg tests did not meet the TEa criteria in either the Level 1 or Level 2 control materials. Although the Alb, ALP, Ca, Crea, Glu, K, and P tests were able to meet the TEa criteria in that study, these tests did not meet any of the bias or CV values.

Despite some weaknesses, the TAE methods are widely used for performance calculations in laboratory medicine. Regardless, other areas of metrology have shifted to the measurement uncertainty (MU) methods. The Guide to Expression of Uncertainty in Measurements (GUM) and the International Vocabulary of Metrology (VIM) do not use the concept of "true value." However, the TAE methods are based on the concept of "true value." The closer the true value of the measurement result is, the smaller the error that will occur. By contrast, MU is not based on "true value" [18, 19].

When evaluating analytical performance, the uncertainty of the whole test process must be included in the account. The analytical variation constitutes only a part of the total uncertainty. BV, pre- and post-analytical variation all need to be included. MU includes these variations, but there are few instances of information on the causes of these variations being applied to the MU theory. According to the GUM, the MU methods in laboratory medicine have not developed sufficiently to cope with the diagnostic uncertainty. The development of analytical performance characteristics for diagnostic uncertainty has resulted in qualitative improvement. Today, the error and uncertainty methods are used as complementary methods when evaluating measurement results in laboratory medicine [20, 21].

Conclusion

Determining CV, bias, and TEa limits for the targeted analytical performance quality specifications based on BV is very valuable. Since these values are determined depending on the "within-individual" and "between-individual" variations of the test, they define a target that is specific for the test. Moreover, defining allowable bias and CV limits separately for each specification of BV in addition to TEa evaluation can give a more objective idea about the source of error.

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