

## INPUTs 2013.1

Survey	INPUTs 2013.1
Start date	January, 1 2013
Supervisor	dr. C. Weykamp MCA Laboratory Streekziekenhuis Koningin Beatrix Winterswijk
Subscriptions	93
Result sets	91

Scores	Your score	MAP	reported
Quantitative	28	38	

## INPUTs 2013.1

Analyte		Trueness				Precision		Performance			
		your mean	ref.	cons.	SDtl	your prec.	SDbl	this survey	PS	cumulative	PSc
ALAT	U/l	92.2	98.2	86.9	10.1	4.0	2.7		2		2
Alk. Phosphatase	U/l	175	188	173	15	6	2		2		2
Amylase	U/l	160	207	175	17	8	3		0		0
ASAT	U/l	75.2	78.6	71.5	6.8	1.5	1.6		2		2
Calcium	mmol/l	2.50	2.46	2.47	0.07	0.05	0.02		2		2
Chloride	mmol/l	103.0	100.0	100.8	1.8	1.5	0.9		1		1
CK	U/l	245	244	249	14	16	3		2		2
Creatinine	µmol/l	149.3	149.1	150.9	7.1	1.9	4.2		1		1
eGFR (F, 55, white)	ml/min/1,73m <sup>2</sup>	37.3	37.6	37.6	2.7	0.5	1.5		2		2
Gamma-GT	U/l	91.5	87.7	86.3	8.6	2.7	1.2		2		2
Glucose	mmol/l	14.68	13.93	14.19	0.44	0.48	0.18		1		1
LD	U/l	1092	532	895	321	8	16		0		0
Magnesium	mmol/l	1.28	1.22	1.23	0.04	0.07	0.02		2		2
Potassium	mmol/l	5.33	5.39	5.30	0.10	0.07	0.04		2		2
Sodium	mmol/l	143.0	144.0	142.7	1.7	1.0	0.9		2		2
Total Protein	g/l	66.5	66.8	66.0	1.8	0.8	0.7		2		2
Urate	µmol/l	376	376	374	15	13	4		2		2
Cholesterol	mmol/l	5.40	5.01	5.09	0.14	0.04	0.05		0		0
HDL-Cholesterol	mmol/l	1.17	1.14	1.13	0.03	0.05	0.02		1		1

Total :

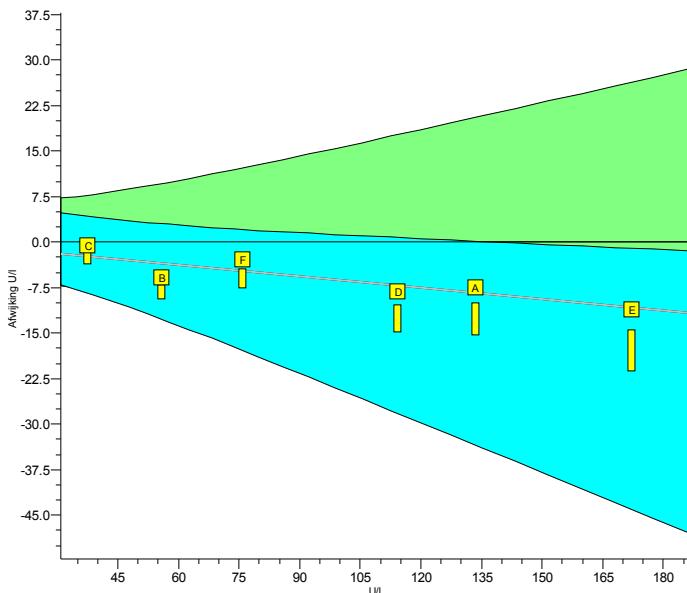
28

28

## INPUTs 2013.1

ALAT

units: U/l



	2013.1	cumulative
Trueness	-6.2%	-6.2%
Precision	4.7%	4.7%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 [2]	6.0 [2]
Sigma-SA	4.9	4.9
Score pictogram		
Regression line	$0.0 + 0.937 \cdot x$	$0.0 + 0.937 \cdot x$

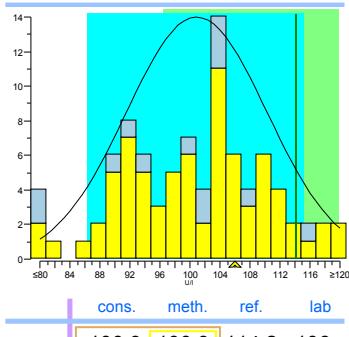
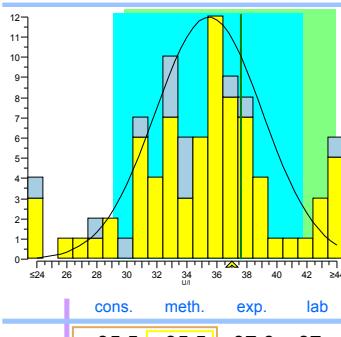
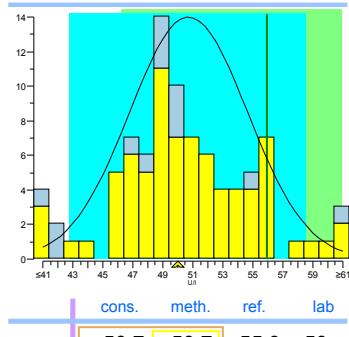
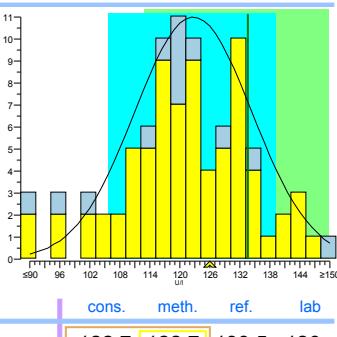
Consensus group IFCC traceerbaar  
Method IFCC traceable

2013.1 A

2013.1 B

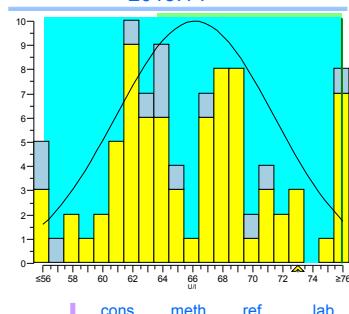
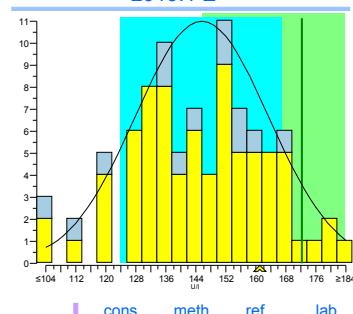
2013.1 C

2013.1 D

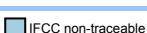


2013.1 E

2013.1 F



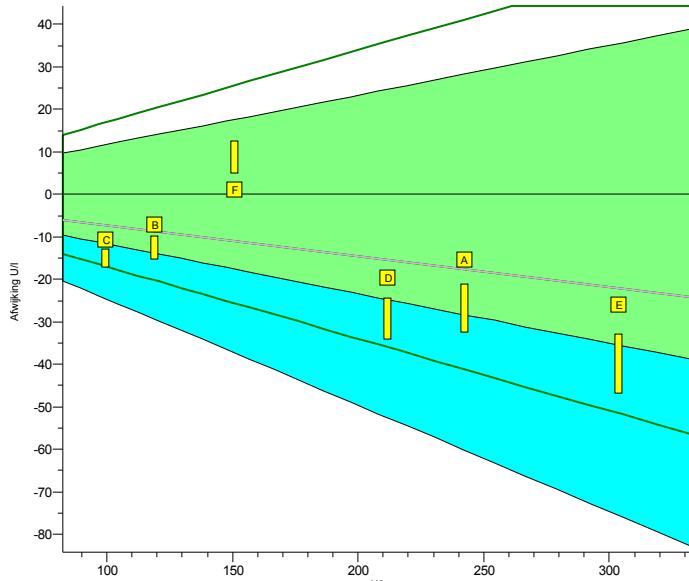
## Legend



## INPUTs 2013.1

Alk. Phosphatase

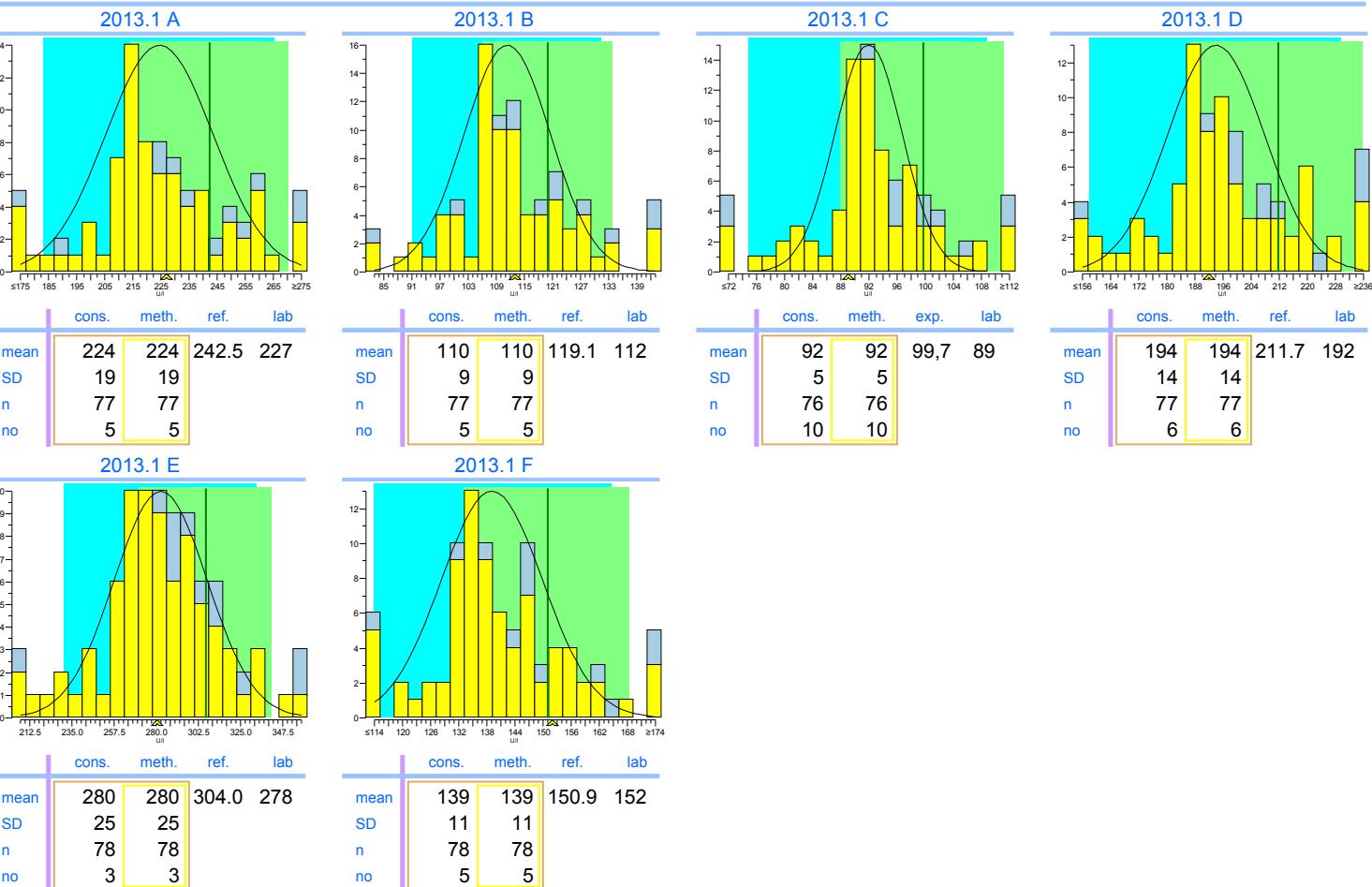
units: U/l



	2013.1	cumulative
Trueness	-6.9%	-6.9%
Precision	3.5%	3.5%
Number	6	6
Outliers	0	0
Sigma-TE	3.1	3.1
Sigma-SA	5.7	5.7
Score pictogram		
Regression line	$0 + 0.928 \cdot x$	$0 + 0.928 \cdot x$

Consensus group  
Method

IFCC traceerbaar  
IFCC traceable



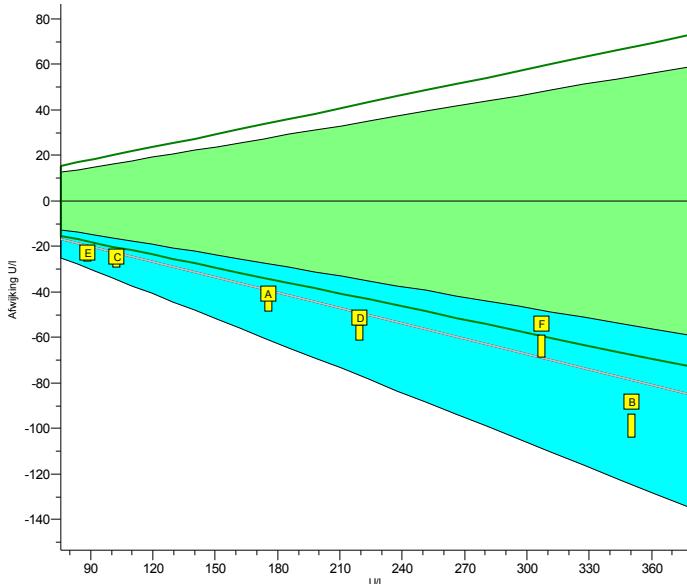
## Legend

IFCC traceable      IFCC non-traceable

## INPUTs 2013.1

Amylase

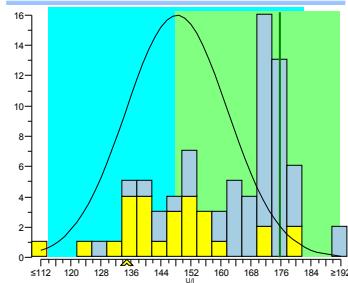
units: U/l



	2013.1	cumulative
Trueness	-23%	-23%
Precision	4.8%	4.8%
Number	6	6
Outliers	0	0
Sigma-TE	-2.6	-2.6
Sigma-SA	-1.4	-1.4
Score pictogram	0	0
Regression line	$0 + 0.776 \cdot x$	$0 + 0.776 \cdot x$

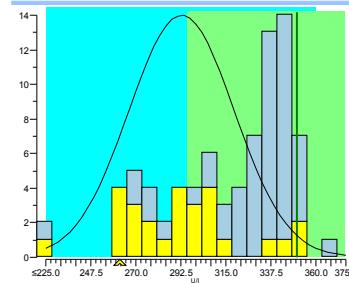
Consensus group: niet IFCC traceerbaar  
 Method: IFCC non-traceable

2013.1 A



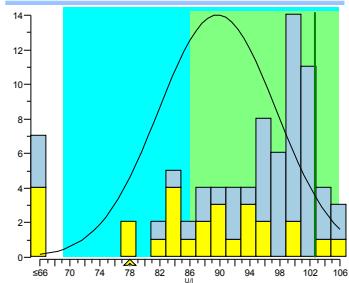
	cons.	meth.	ref.	lab
mean	148	148	175.8	135
SD	14	14		
n	27	27		
no	1	1		

2013.1 B



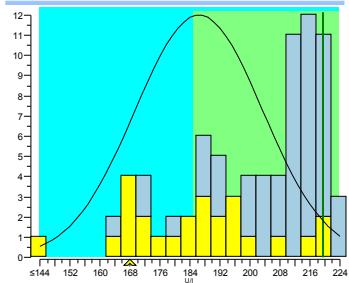
	cons.	meth.	ref.	lab
mean	293	293	350.5	262
SD	26	26		
n	27	27		
no	1	1		

2013.1 C



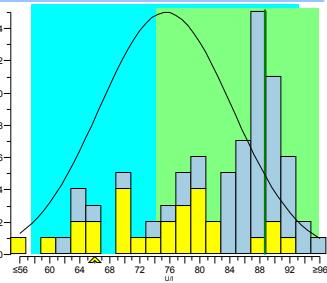
	cons.	meth.	exp.	lab
mean	90	90	102,7	78
SD	8	8		
n	27	27		
no	4	4		

2013.1 D



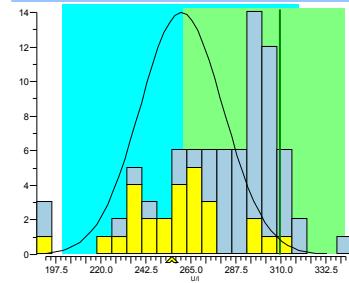
	cons.	meth.	ref.	lab
mean	186	186	219,4	168
SD	17	17		
n	25	25		
no	1	1		

2013.1 E



	cons.	meth.	ref.	lab
mean	75	75	88,7	66
SD	9	9		
n	27	27		
no	1	1		

2013.1 F



	cons.	meth.	ref.	lab
mean	258	258	307,0	253
SD	21	21		
n	27	27		
no	1	1		

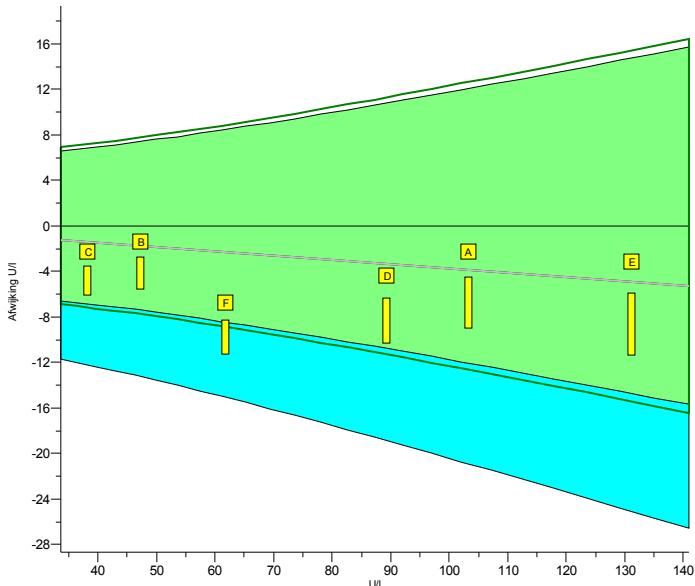
## Legend

 IFCC non-traceable

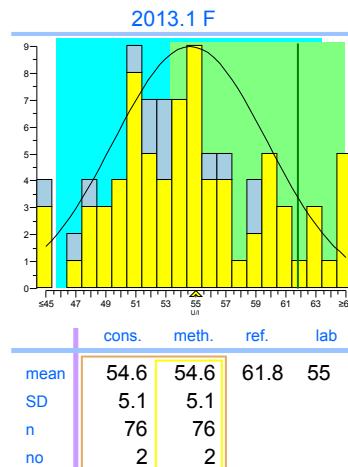
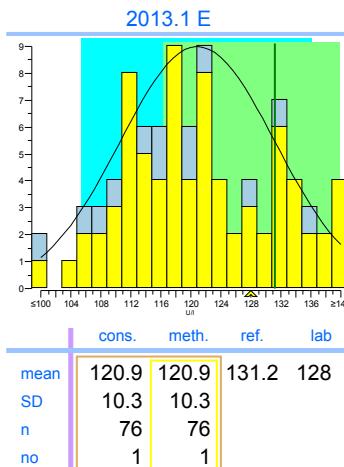
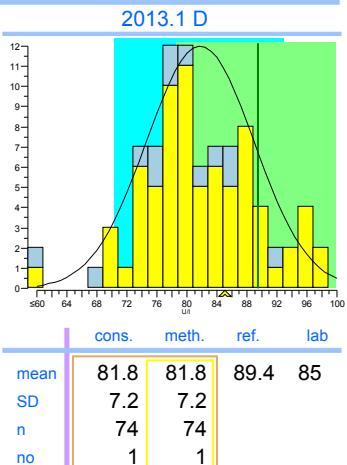
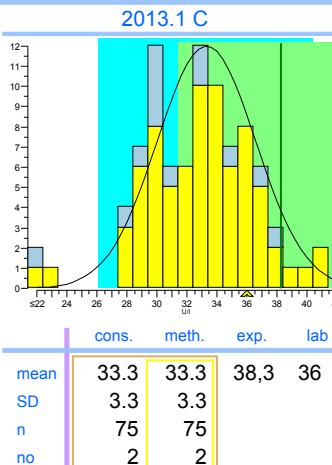
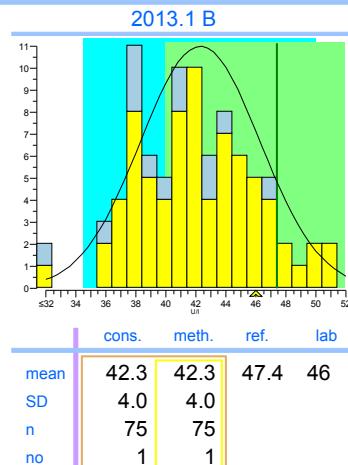
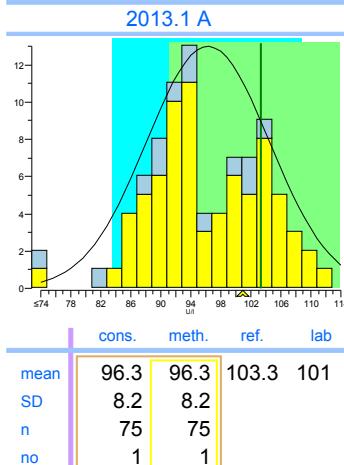
 IFCC traceable

**INPUTs 2013.1****ASAT**

units: U/l



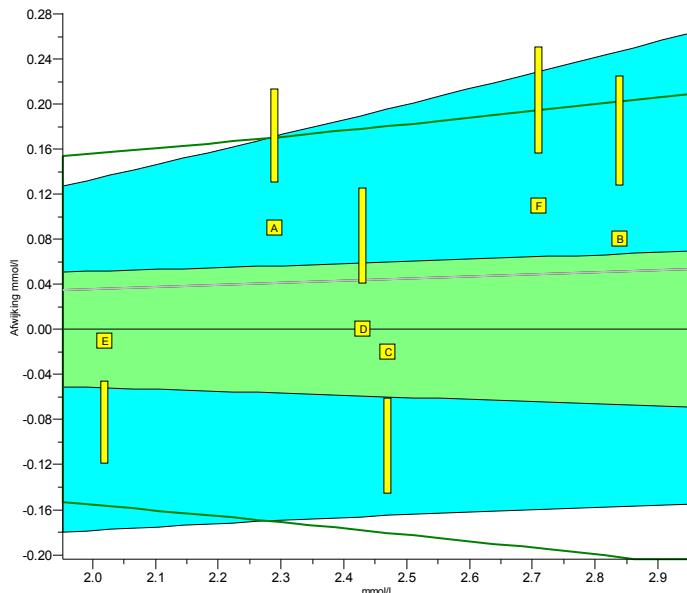
	2013.1	cumulative
Trueness	-4.3%	-4.3%
Precision	2.1%	2.1%
Number	6	6
Outliers	0	0
Sigma-TE	4.9	4.9
Sigma-SA	5.2	5.2
Score pictogram	[2]	[2]
Regression line	$0.0 + 0.963 \cdot x$	$0.0 + 0.963 \cdot x$

 Consensus group: IFCC traceerbaar  
 Method: IFCC traceable
**Legend** IFCC traceable IFCC non-traceable

## INPUTs 2013.1

Calcium

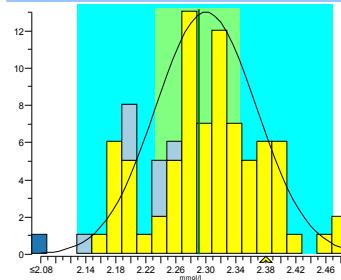
units: mmol/l



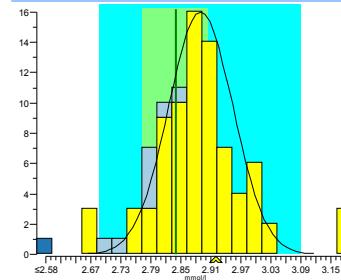
	2013.1	cumulative
Trueness	+1.7%	+1.7%
Precision	1.8%	1.8%
Number	6	6
Outliers	0	0
Sigma-TE	1.1	1.1
Sigma-SA	4.6 [2]	4.6 [2]
Score pictogram	[Green, Yellow, Red]	
Regression line	$0.00 + 1.018 \cdot x$	$0.00 + 1.018 \cdot x$

Consensus group: Colorimetrisch  
Method: Colorimetric, automatic, discrete

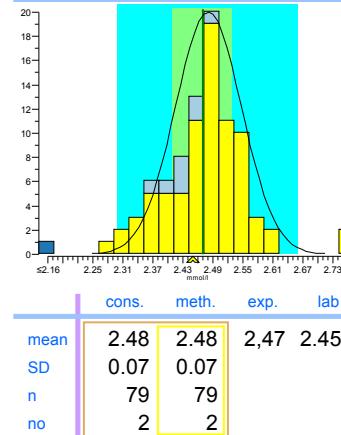
2013.1 A



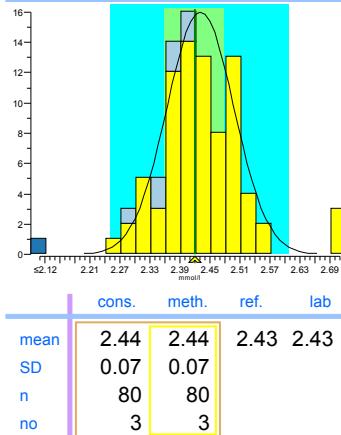
2013.1 B



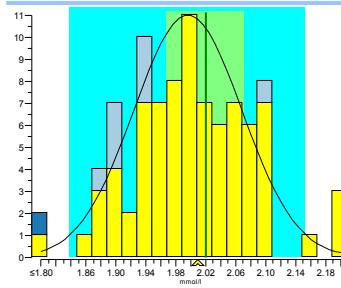
2013.1 C



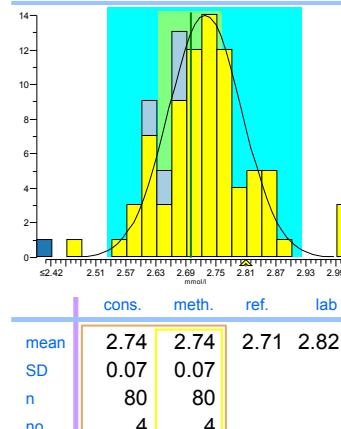
2013.1 D



2013.1 E



2013.1 F



## Legend

[Yellow] Colorimetric, automatic, discrete

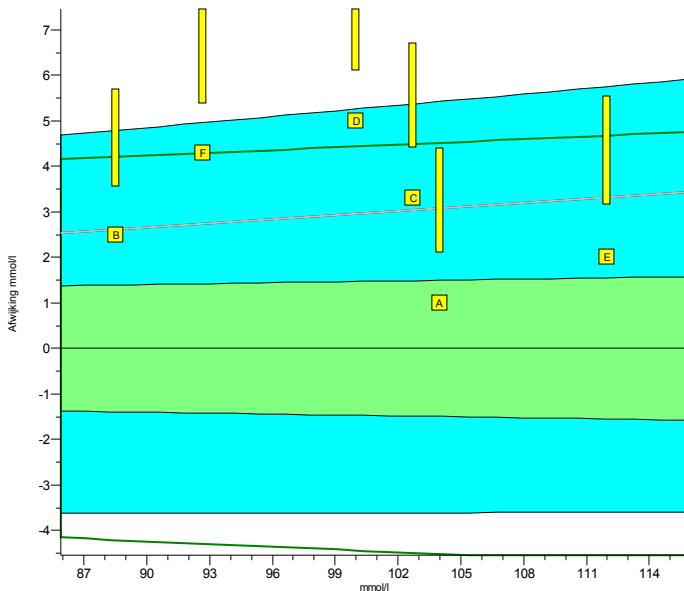
[Light Blue] ISE indirect (with predilution)

[Dark Blue] Other methods

## INPUTs 2013.1

Chloride

units: mmol/l



	2013.1	cumulative
Trueness	+3.0%	+3.0%
Precision	1.5%	1.5%
Number	6	6
Outliers	0	0
Sigma-TE	-0.9	-0.9
Sigma-SA	2.4	2.4
Score pictogram	1	1
Regression line	$0.0 + 1.030 \cdot x$	$0.0 + 1.030 \cdot x$

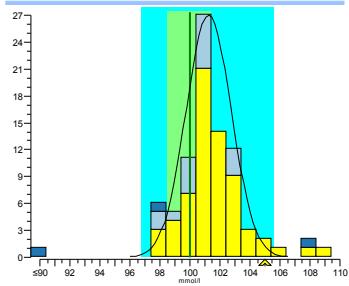
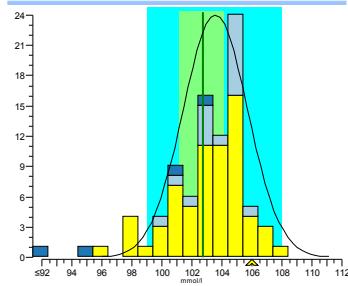
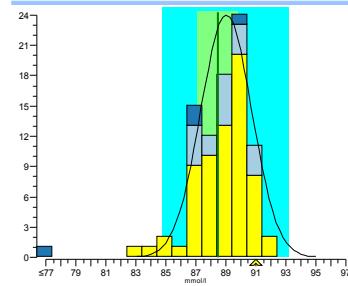
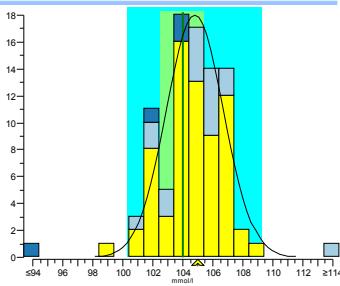
Consensus group ISE/Colorimetrie  
Method ISE indirect (with predilution)

2013.1 A

2013.1 B

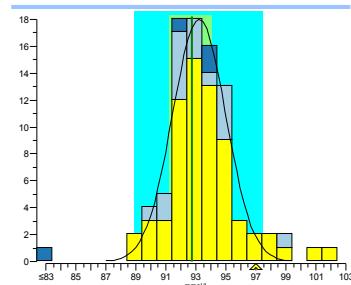
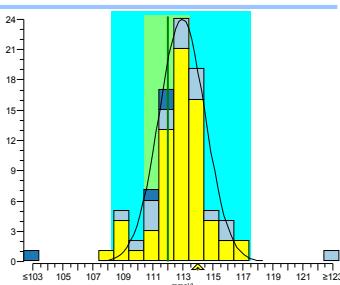
2013.1 C

2013.1 D



2013.1 E

2013.1 F



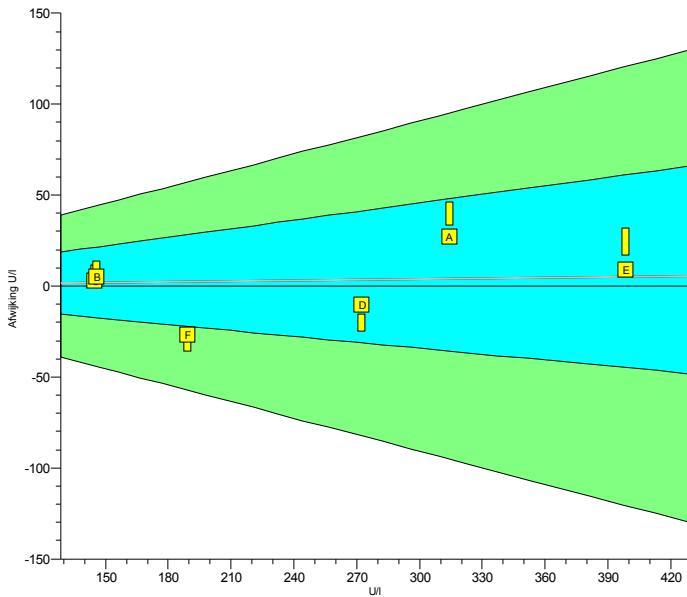
## Legend

- [Yellow square] ISE indirect (with predilution)
- [Light blue square] ISE direct (no predilution)
- [Dark blue square] Other methods

## INPUTs 2013.1

CK

units: U/l



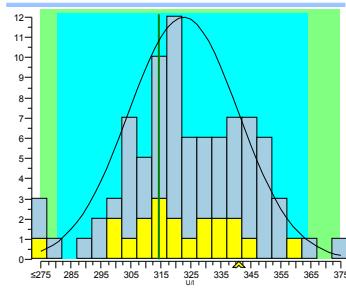
	2013.1	cumulative
Trueness	+0.41%	+0.41%
Precision	6.5%	6.5%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 [2]	6.0 [2]
Sigma-SA	5.6	5.6
Score pictogram	[Green box with yellow dot]	[Green box with yellow dot]
Regression line	$0 + 1.012.x$	$0 + 1.012.x$

Consensus group  
Method

niet IFCC traceerbaar

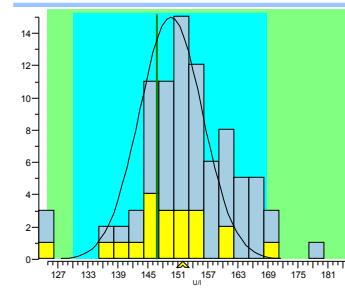
IFCC non-traceable

2013.1 A



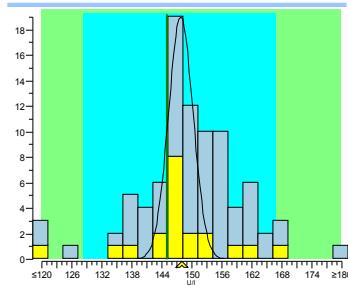
	cons.	meth.	ref.	lab
mean	323	323	314.3	341
SD	18	18		
n	20	20		
no	1	1		

2013.1 B



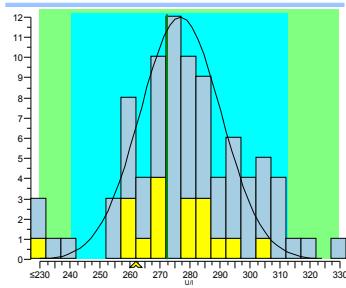
	cons.	meth.	ref.	lab
mean	149	149	145.8	151
SD	6	6		
n	20	20		
no	2	2		

2013.1 C



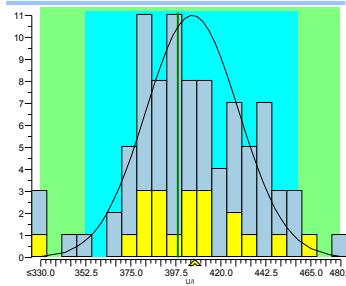
	cons.	meth.	exp.	lab
mean	148	148	145	148
SD	2	2		
n	20	20		
no	6	6		

2013.1 D



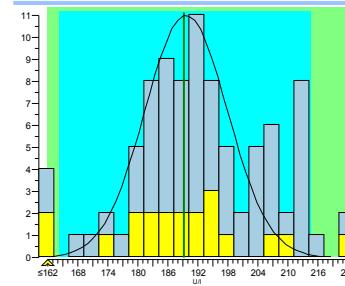
	cons.	meth.	ref.	lab
mean	277	277	272.3	262
SD	13	13		
n	18	18		
no	1	1		

2013.1 E



	cons.	meth.	ref.	lab
mean	406	406	398.4	407
SD	23	23		
n	20	20		
no	1	1		

2013.1 F



	cons.	meth.	ref.	lab
mean	190	190	189.2	162
SD	8	8		
n	20	20		
no	3	3		

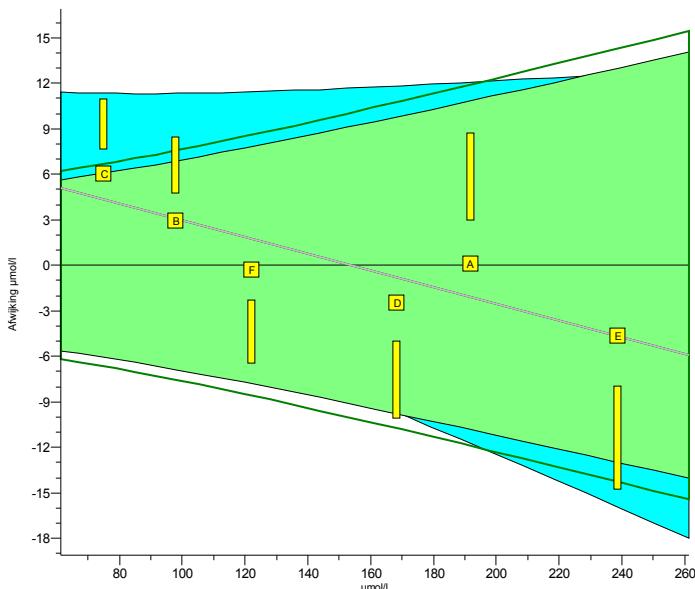
Legend

■ IFCC non-traceable

■ IFCC traceable

## INPUTs 2013.1

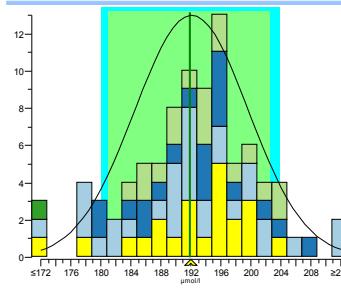
## Creatinine

units:  $\mu\text{mol/l}$ 

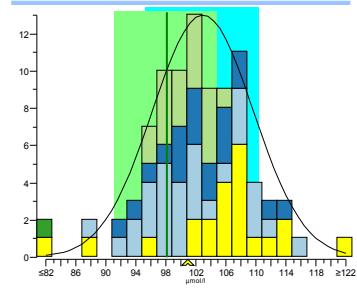
	2013.1	cumulative
Trueness	+0.17%	+0.17%
Precision	1.3%	1.3%
Number	6	6
Outliers	0	0
Sigma-TE	3.9	3.9
Sigma-SA	4.3 [1]	4.3 [1]
Score pictogram	[Green circle]	[Green circle]
Regression line	$8.5 + 0.945 \cdot x$	$8.5 + 0.945 \cdot x$

Consensus group: Jaffe  
Method: Alk. Picrate, kinetic

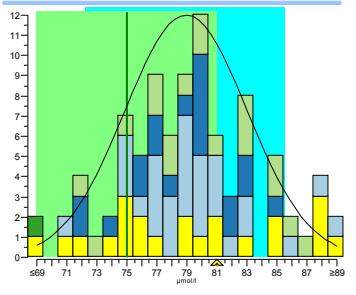
2013.1 A



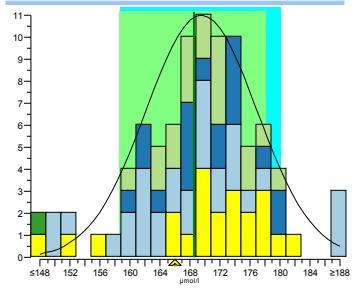
2013.1 B



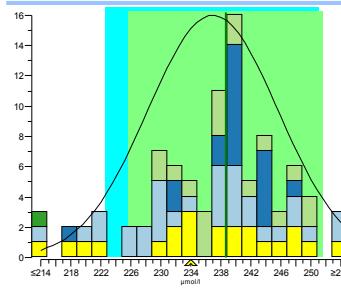
2013.1 C



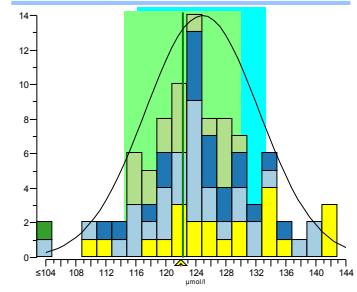
2013.1 D



2013.1 E



2013.1 F

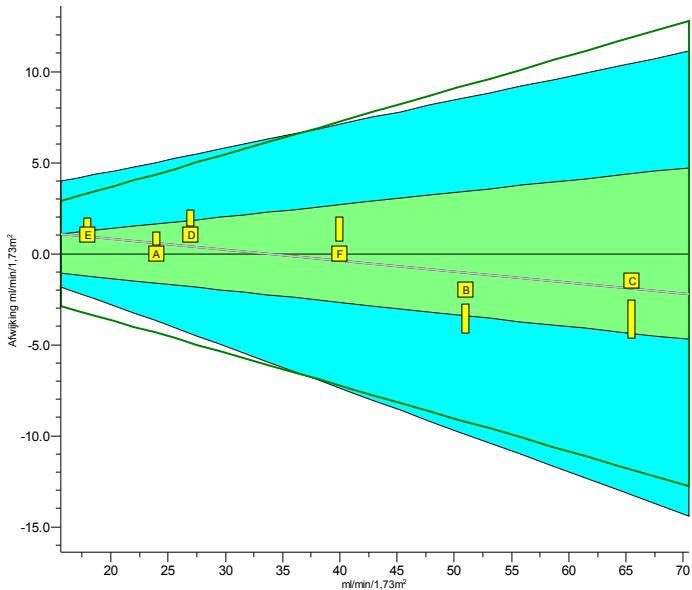


## Legend

- [Yellow square] Alk. Picrate, kinetic
- [Light blue square] Alk. Picrate, kinetic with compensation
- [Dark blue square] Alk. Picrate, endpoint
- [Light green square] Enzymatic, automatic
- [Dark green square] Other methods

## INPUTs 2013.1

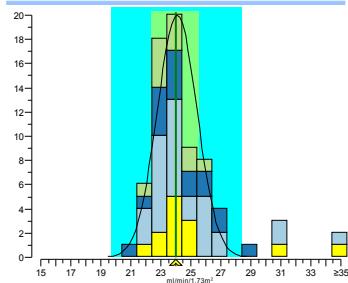
eGFR (F, 55, white)

units: ml/min/1,73m<sup>2</sup>

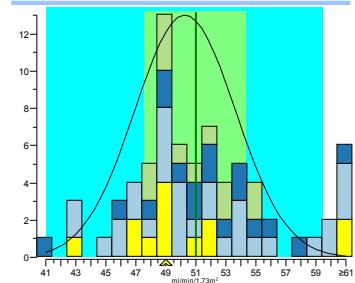
	2013.1	cumulative
Trueness	-0.67%	-0.67%
Precision	1.3%	1.3%
Number	6	6
Outliers	0	0
Sigma-TE	3.9	3.9
Sigma-SA	6.0 [2]	6.0 [2]
Score pictogram		
Regression line	$2.0 + 0.940 \cdot x$	$2.0 + 0.940 \cdot x$

Consensus group: Jaffe  
Method: Alk. Picrate, kinetic

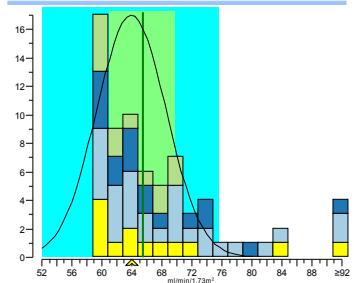
2013.1 A



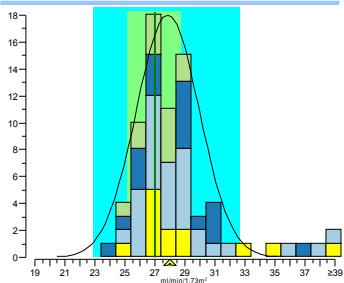
2013.1 B



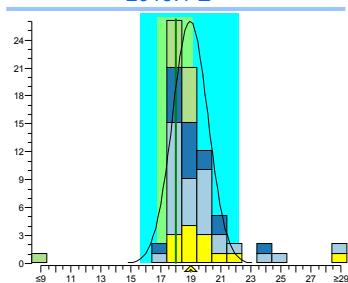
2013.1 C



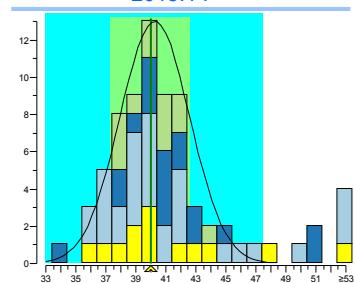
2013.1 D



2013.1 E



2013.1 F



## Legend

Alk. Picrate, kinetic

Other methods

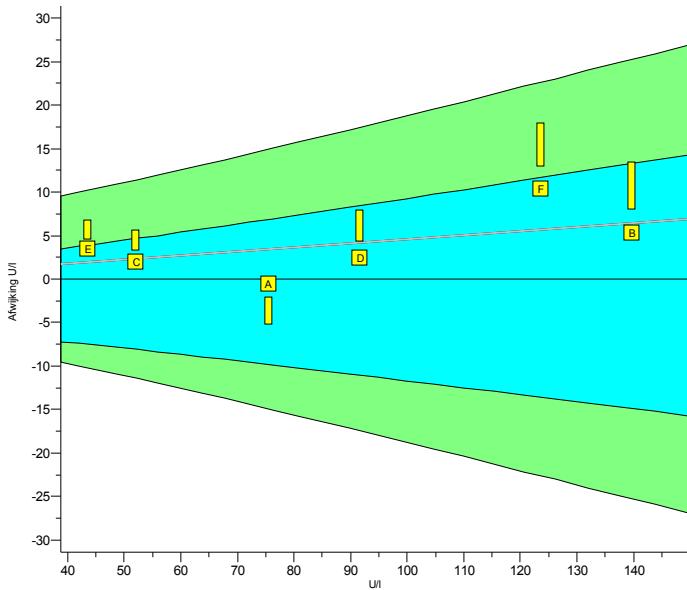
Alk. Picrate, kinetic with compensation

Enzymatic, automatic

## INPUTs 2013.1

Gamma-GT

units: U/l

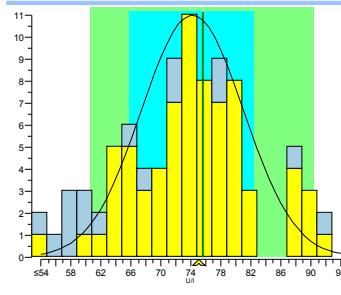


	2013.1	cumulative
Trueness	+4.4%	+4.4%
Precision	3.2%	3.2%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 [2]	6.0 [2]
Sigma-SA	4.7	4.7
Score pictogram	[Score pictogram]	[Score pictogram]
Regression line	$0.0 + 1.046 \cdot x$	$0.0 + 1.046 \cdot x$

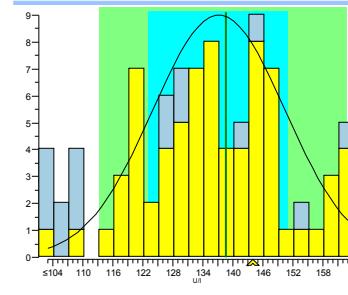
Consensus group  
Method

IFCC traceerbaar  
IFCC traceable

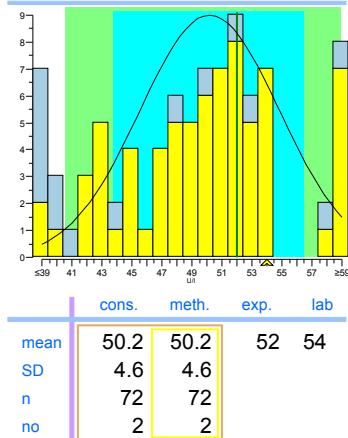
2013.1 A



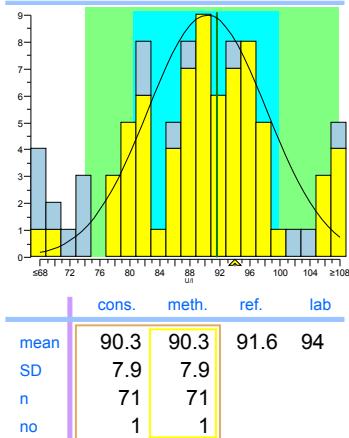
2013.1 B



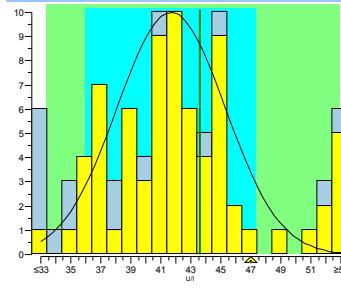
2013.1 C



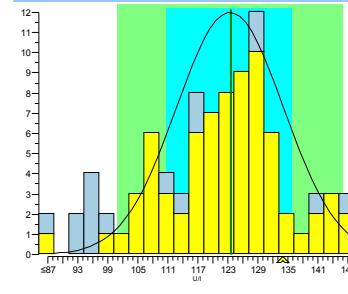
2013.1 D



2013.1 E



2013.1 F



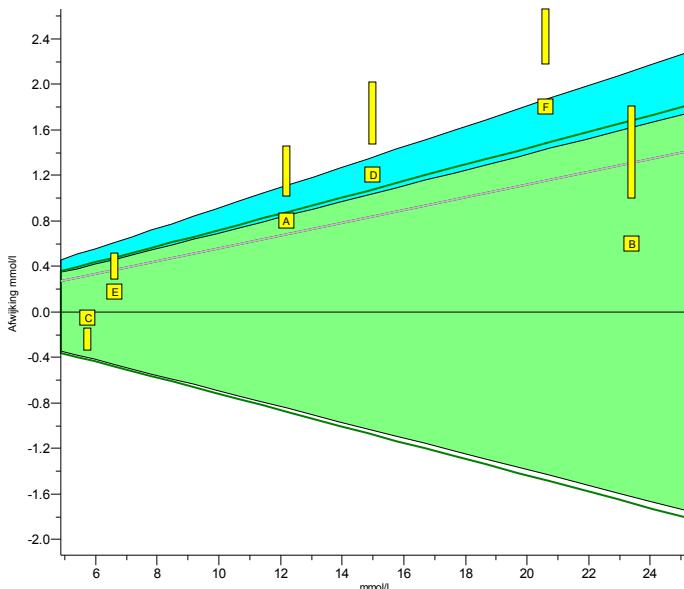
## Legend



## INPUTs 2013.1

Glucose

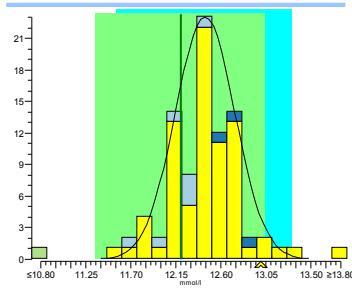
units: mmol/l



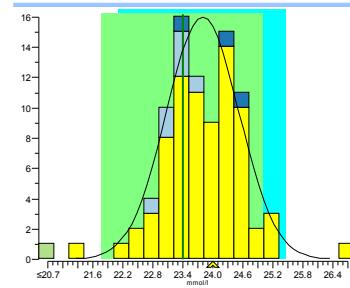
	2013.1	cumulative
Trueness	+5.4%	+5.4%
Precision	3.4%	3.4%
Number	6	6
Outliers	0	0
Sigma-TE	2.3	2.3
Sigma-SA	2.5	2.5
Score pictogram		
Regression line	$0.00 + 1.056 \cdot x$	$0.00 + 1.056 \cdot x$

Consensus group: Natte chemie  
Method: Hexokinase, automatic

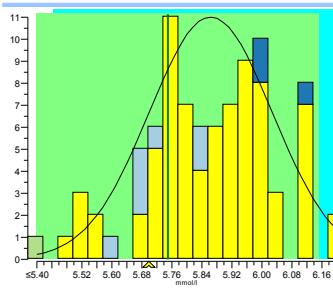
2013.1 A



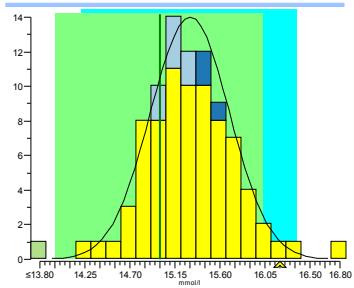
2013.1 B



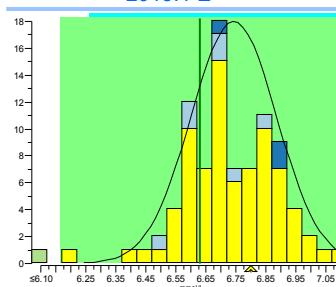
2013.1 C



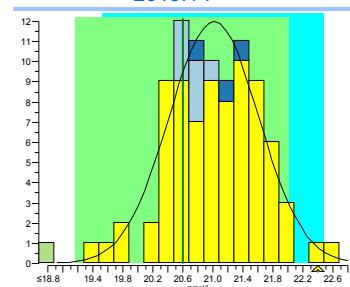
2013.1 D



2013.1 E



2013.1 F



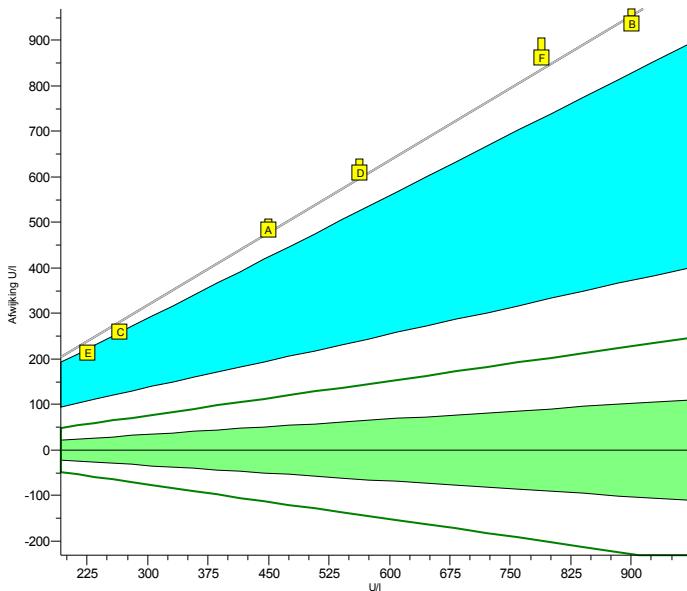
## Legend

- Hexokinase, automatic
- Glucose-oxydase, amperometric, H<sub>2</sub>O<sub>2</sub>
- Glucose-oxydase/POD,automatic
- Other methods

## INPUTS 2013.1

LD

units: U/l



	2013.1	cumulative
Trueness	+105%	+105%
Precision	0.88%	0.88%
Number	6	6
Outliers	0	0
Sigma-TE	-3.0	-3.0
Sigma-SA	-3.0	-3.0
Score pictogram	0	0
Regression line	$0 + 2.060 \cdot x$	$0 + 2.060 \cdot x$

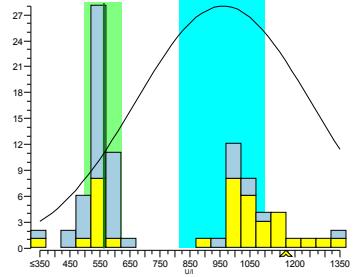
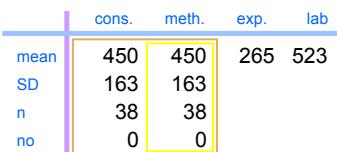
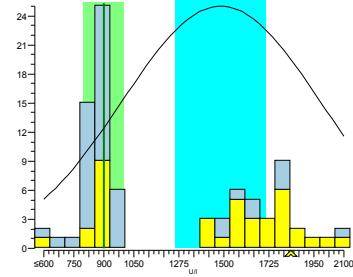
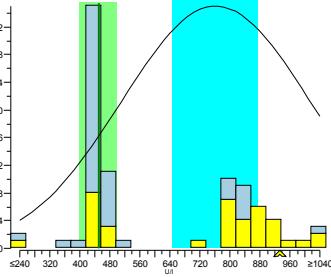
Consensus group: niet IFCC traceerbaar  
Method: IFCC non-traceable

2013.1 A

2013.1 B

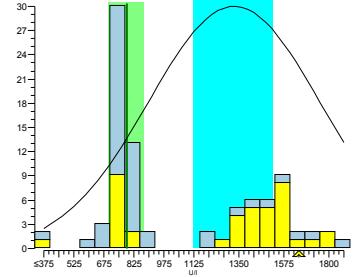
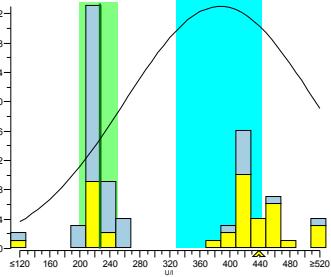
2013.1 C

2013.1 D



2013.1 E

2013.1 F



## Legend

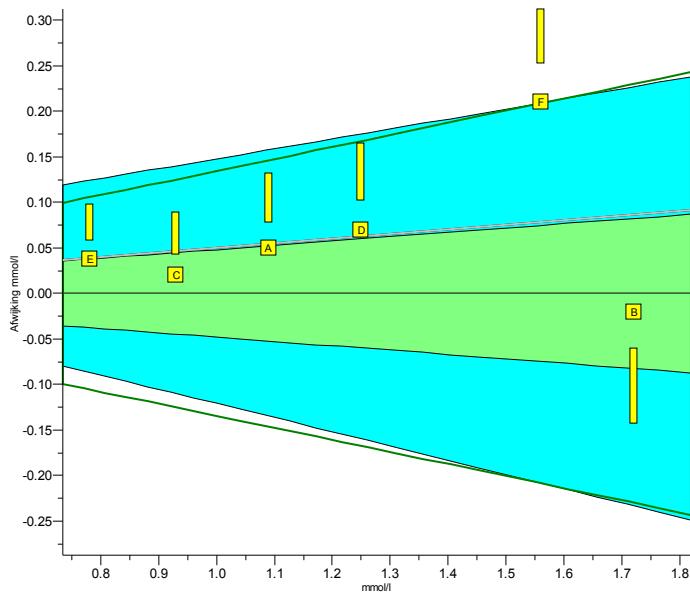
 IFCC non-traceable

 IFCC traceable

## INPUTs 2013.1

## Magnesium

units: mmol/l

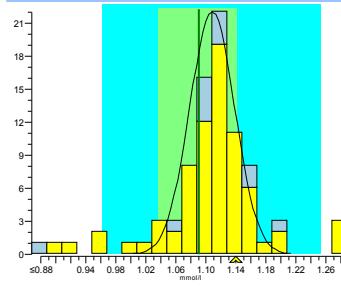


	2013.1	cumulative
Trueness	+5.0%	+5.0%
Precision	6.1%	6.1%
Number	6	6
Outliers	0	0
Sigma-TE	0.6	0.6
Sigma-SA	4.9 [2]	4.9 [2]
Score pictogram		
Regression line	$0.00 + 1.050 \cdot x$	$0.00 + 1.050 \cdot x$

Consensus group  
Method

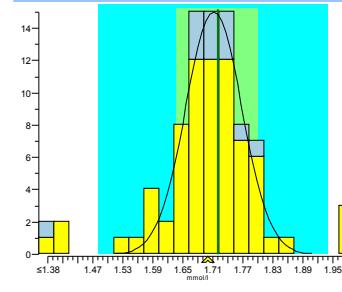
Overall  
Colorimetric

2013.1 A



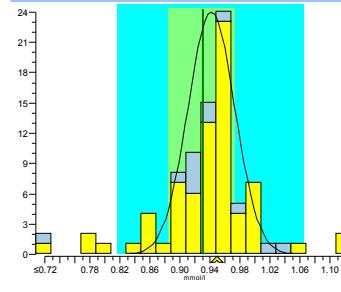
	cons.	meth.	ref.	lab
mean	1.11	1.11	1.09	1.14
SD	0.03	0.03		
n	85	73		
no	11	10		

2013.1 B



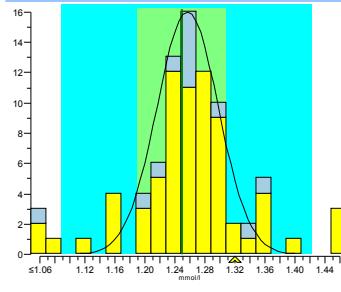
	cons.	meth.	ref.	lab
mean	1.71	1.71	1.72	1.7
SD	0.06	0.06		
n	85	73		
no	8	7		

2013.1 C



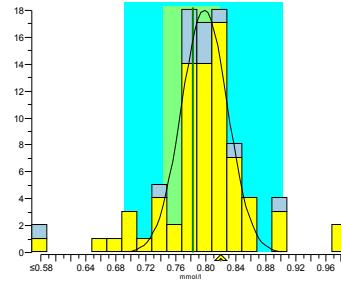
	cons.	meth.	exp.	lab
mean	0.94	0.94	0.93	0.95
SD	0.03	0.03		
n	85	73		
no	9	8		

2013.1 D



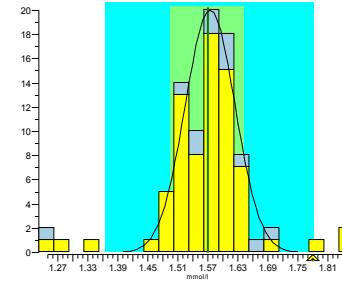
	cons.	meth.	ref.	lab
mean	1.26	1.26	1.25	1.32
SD	0.04	0.05		
n	83	71		
no	9	8		

2013.1 E



	cons.	meth.	ref.	lab
mean	0.798	0.799	0.782	0.82
SD	0.031	0.031		
n	86	74		
no	13	11		

2013.1 F



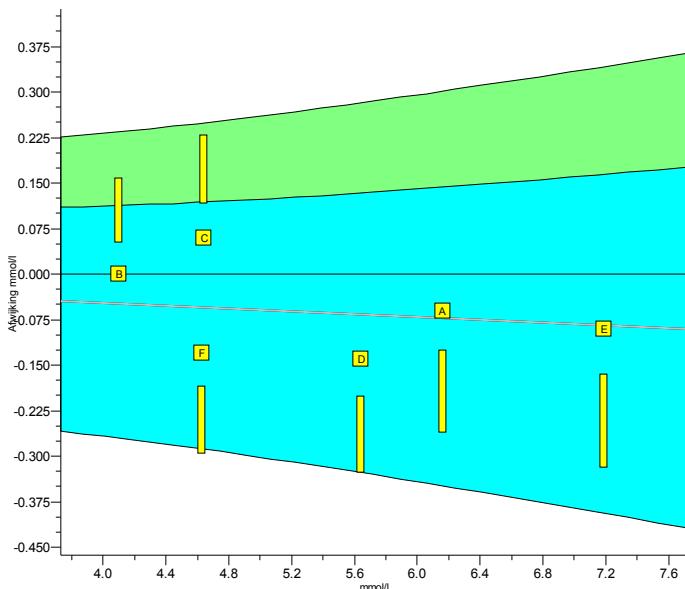
	cons.	meth.	ref.	lab
mean	1.56	1.56	1.56	1.77
SD	0.05	0.05		
n	86	74		
no	7	6		

## Legend

 Colorimetric	 Other methods
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**INPUTs 2013.1****Potassium**

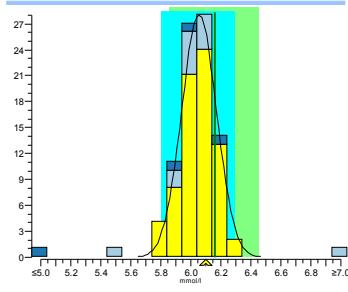
units: mmol/l



	2013.1	cumulative
Trueness	-1.1%	-1.1%
Precision	1.4%	1.4%
Number	6	6
Outliers	0	0
Sigma-TE	4.7 [2]	4.7 [2]
Sigma-SA	3.7	3.7
Score pictogram	[Color-coded boxes]	[Color-coded boxes]
Regression line	$0.00 + 0.988.x$	$0.00 + 0.988.x$

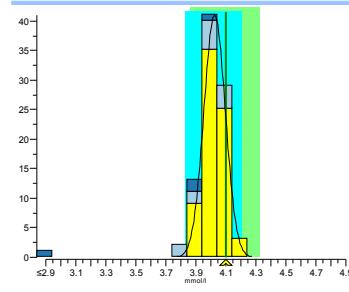
Consensus group: ISE verdund/Vlamfotometrie  
Method: ISE indirect (with predilution)

2013.1 A



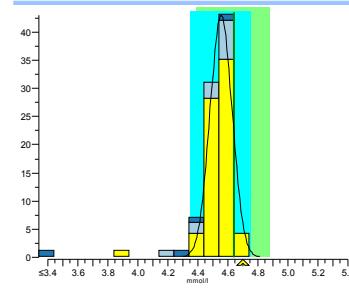
	cons.	meth.	ref.	lab
mean	6.05	6.05	6.16	6.1
SD	0.12	0.12		
n	72	72		
no	0	0		

2013.1 B



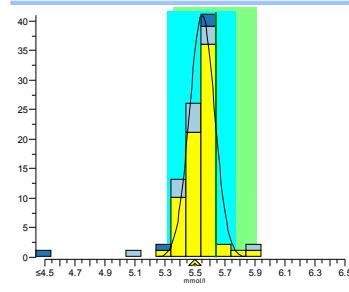
	cons.	meth.	ref.	lab
mean	4.02	4.02	4.10	4.1
SD	0.07	0.07		
n	72	72		
no	0	0		

2013.1 C



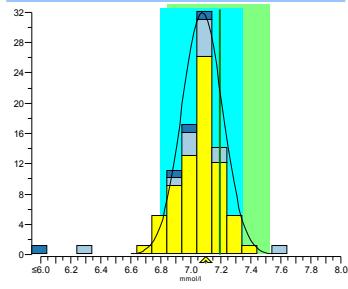
	cons.	meth.	exp.	lab
mean	4.55	4.55	4.64	4.7
SD	0.07	0.07		
n	72	72		
no	1	1		

2013.1 D



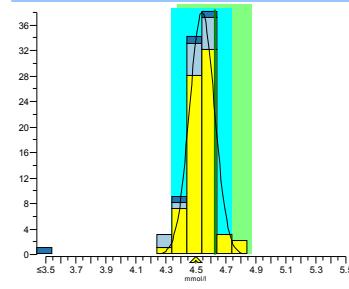
	cons.	meth.	ref.	lab
mean	5.55	5.55	5.64	5.5
SD	0.08	0.08		
n	72	72		
no	2	2		

2013.1 E



	cons.	meth.	ref.	lab
mean	7.08	7.08	7.19	7.1
SD	0.14	0.14		
n	72	72		
no	0	0		

2013.1 F



	cons.	meth.	ref.	lab
mean	4.54	4.54	4.63	4.5
SD	0.08	0.08		
n	73	73		
no	2	2		

## Legend

ISE indirect (with predilution)

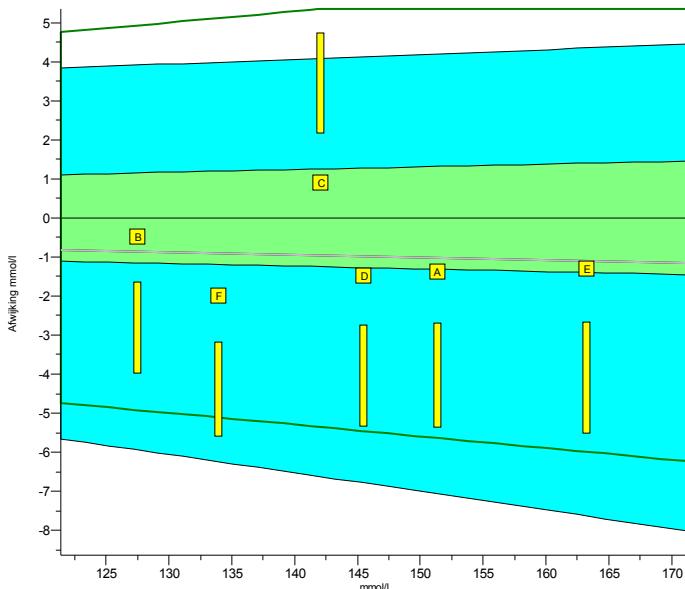
ISE direct (no predilution)

Other methods

## INPUTs 2013.1

Sodium

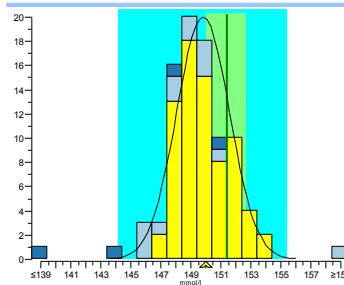
units: mmol/l



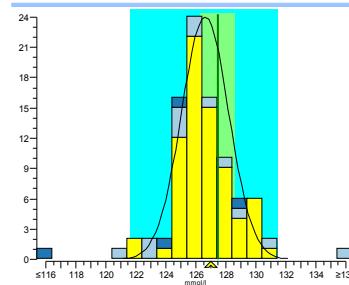
	2013.1	cumulative
Trueness	-0.67%	-0.67%
Precision	0.73%	0.73%
Number	6	6
Outliers	0	0
Sigma-TE	0.8	0.8
Sigma-SA	4.8 [2]	4.8 [2]
Score pictogram	[Yellow]	[Yellow]
Regression line	$0.0 + 0.993 \cdot x$	$0.0 + 0.993 \cdot x$

Consensus group ISE verdund/Vlamfotometrie  
Method ISE indirect (with predilution)

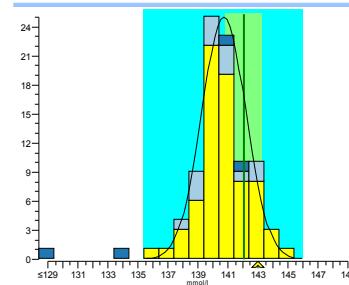
2013.1 A



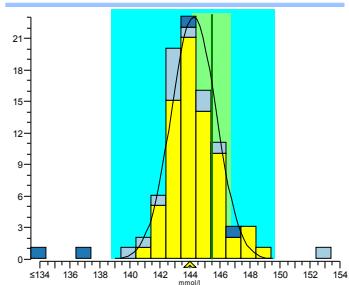
2013.1 B



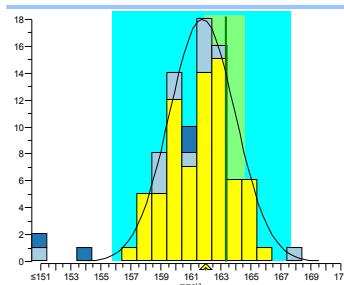
2013.1 C



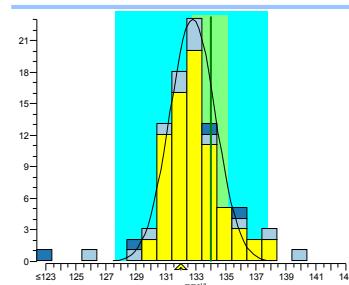
2013.1 D



2013.1 E



2013.1 F



## Legend

[Yellow] ISE indirect (with predilution)

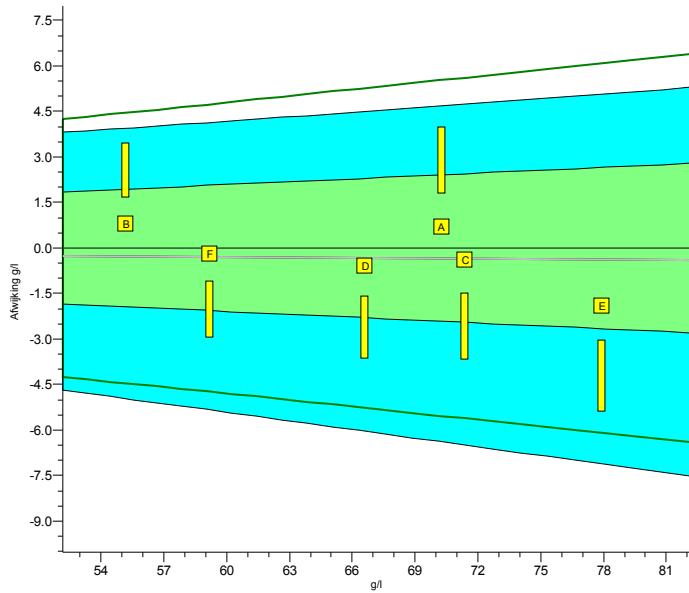
[Grey] ISE direct (no predilution)

[Blue] Other methods

## INPUTs 2013.1

## Total Protein

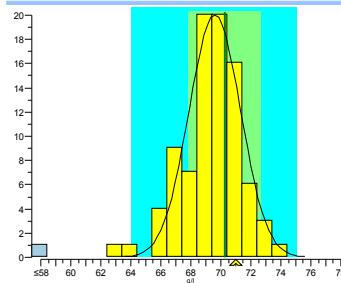
units: g/l



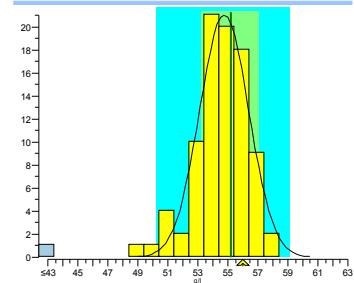
	2013.1	cumulative
Trueness	-0.40%	-0.40%
Precision	1.3%	1.3%
Number	6	6
Outliers	0	0
Sigma-TE	2.7	2.7
Sigma-SA	6.0 [2]	6.0 [2]
Score pictogram	[Yellow-Green]	[Yellow-Green]
Regression line	$0.0 + 0.995 \cdot x$	$0.0 + 0.995 \cdot x$

Consensus group: Biureet  
Method: Biurete, automatic

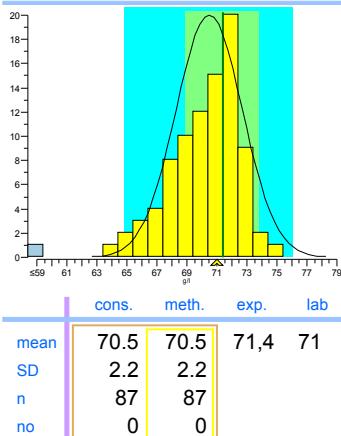
2013.1 A



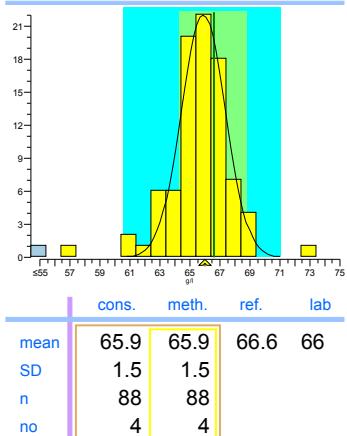
2013.1 B



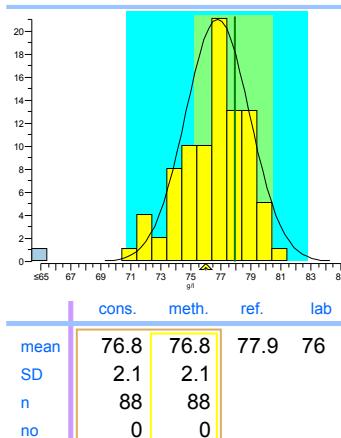
2013.1 C



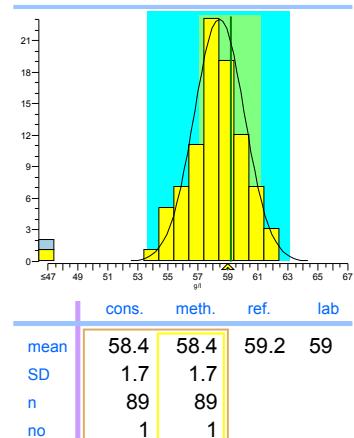
2013.1 D



2013.1 E



2013.1 F



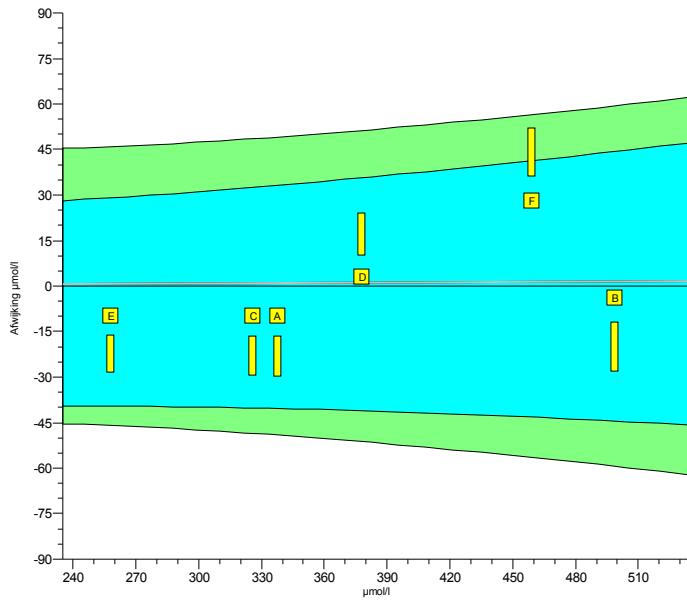
## Legend

 Biurete, automatic

 Other methods

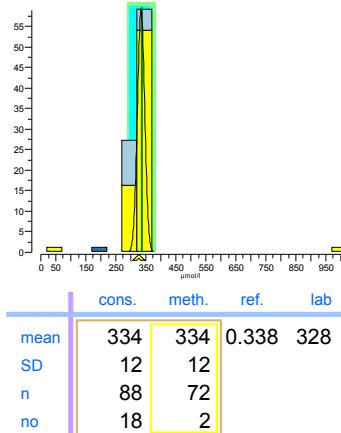
**INPUTs 2013.1**

Urate

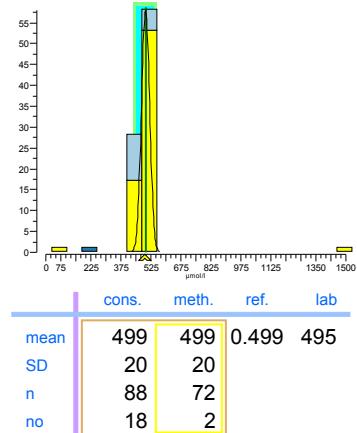
units:  $\mu\text{mol/l}$ 

	2013.1	cumulative
Trueness	-0.13%	-0.13%
Precision	3.5%	3.5%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 [2]	6.0 [2]
Sigma-SA	5.7	5.7
Score pictogram	[Yellow]	[Yellow]
Regression line	$0 + 1.003 \cdot x$	$0 + 1.003 \cdot x$
Consensus group	Colorimetrisch	
Method	Uricase, colorim., automatic	

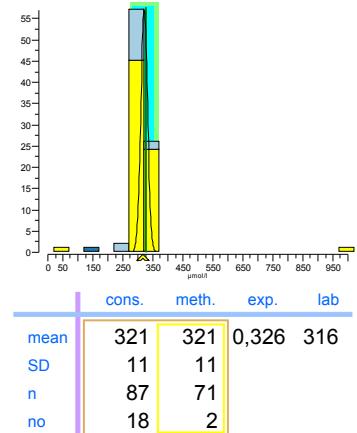
2013.1 A



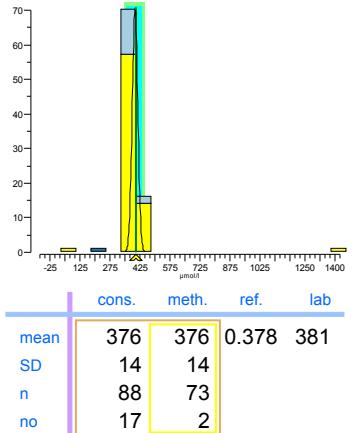
2013.1 B



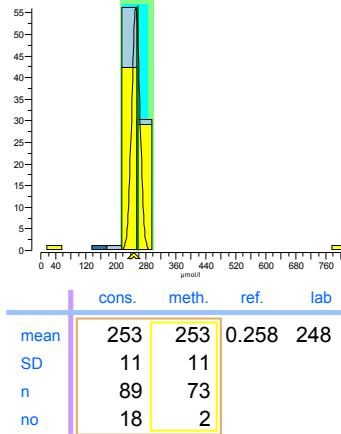
2013.1 C



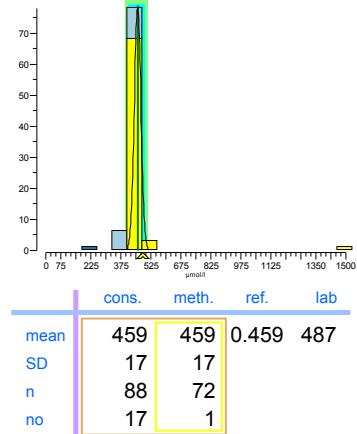
2013.1 D



2013.1 E



2013.1 F

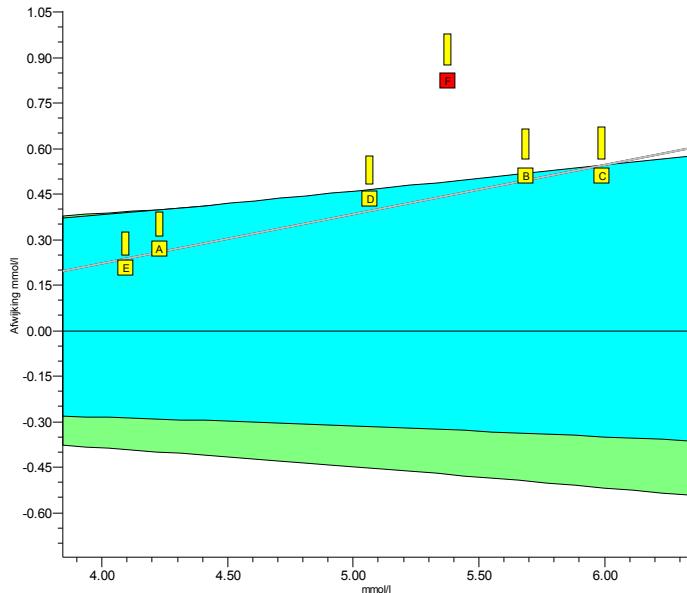


## Legend

- [Yellow] Uricase, colorim., automatic
- [Light Blue] Uricase, differential UV, automatic
- [Dark Blue] Oude methoden

**INPUTs 2013.1****Cholesterol**

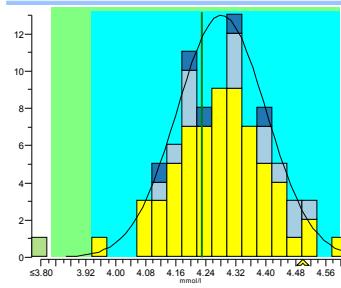
units: mmol/l



	2013.1	cumulative
Trueness	+7.7%	+7.7%
Precision	0.73%	0.73%
Number	6	6
Outliers	1	1
Sigma-TE	1.9 <span style="background-color: red; color: white;">0</span>	1.9 <span style="background-color: red; color: white;">0</span>
Sigma-SA	0.5	0.5
Score pictogram		
Regression line	$-0.42 + 1.162 \cdot x$	$-0.42 + 1.162 \cdot x$

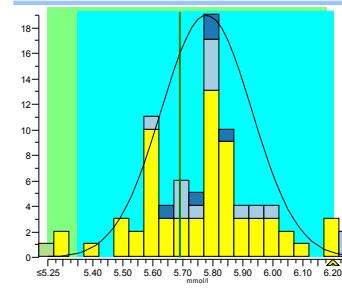
Consensus group: Enzymatisch  
 Method: Enzymatic, automatic, discrete

2013.1 A



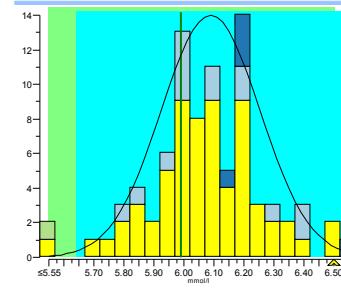
	cons.	meth.	ref.	lab
mean	4.28	4.28	4.230	4.5
SD	0.12	0.12		
n	83	64		
no	0	0		

2013.1 B



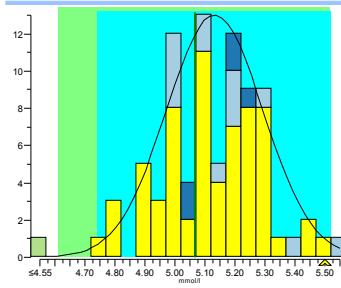
	cons.	meth.	ref.	lab
mean	5.78	5.78	5.689	6.2
SD	0.15	0.18		
n	83	64		
no	4	2		

2013.1 C



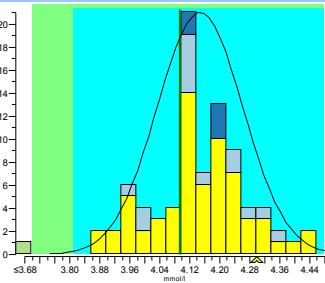
	cons.	meth.	exp.	lab
mean	6.09	6.08	5.99	6.5
SD	0.16	0.17		
n	83	64		
no	1	1		

2013.1 D



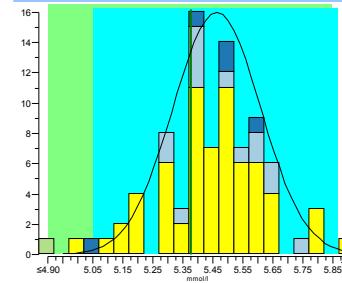
	cons.	meth.	ref.	lab
mean	5.13	5.12	5.066	5.5
SD	0.16	0.17		
n	82	64		
no	0	0		

2013.1 E



	cons.	meth.	ref.	lab
mean	4.15	4.14	4.094	4.3
SD	0.11	0.13		
n	84	65		
no	0	0		

2013.1 F



	cons.	meth.	ref.	lab
mean	5.46	5.45	5.377	6.2
SD	0.15	0.16		
n	84	65		
no	2	2		

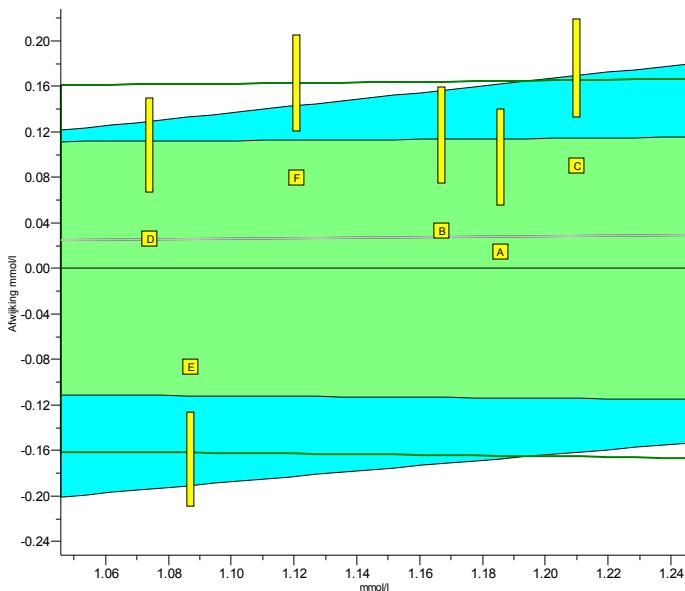
## Legend

- Enzymatic, automatic, discrete
- Enzymatic, automatic, kinetic
- Abell-Kendall reference values
- Overige methoden

## INPUTs 2013.1

HDL-Cholesterol

units: mmol/l



	2013.1	cumulative
Trueness	+2.3%	+2.3%
Precision	4.5%	4.5%
Number	6	6
Outliers	0	0
Sigma-TE	2.5	2.5
Sigma-SA	4.0 [1]	4.0 [1]
Score pictogram	[Green box with yellow dot]	[Green box with yellow dot]
Regression line	$0.00 + 1.024 \cdot x$	$0.00 + 1.024 \cdot x$

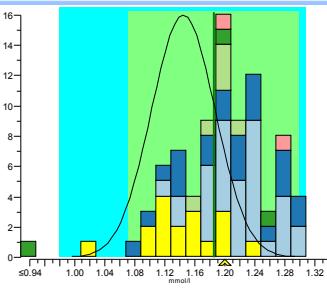
 Consensus group: Overall  
 Method: Immunoinhibition

2013.1 A

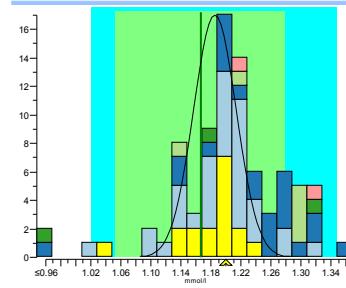
2013.1 B

2013.1 C

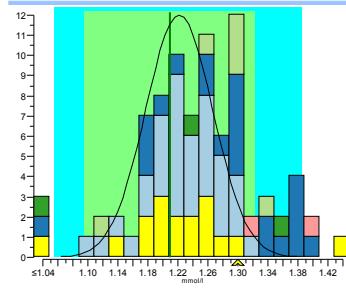
2013.1 D



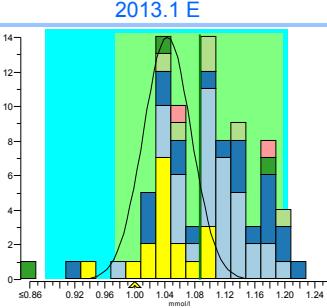
	cons.	meth.	ref.	lab
mean	1.14	1.14	1.186	1.2
SD	0.04	0.04		
n	17	17		
no	1	1		



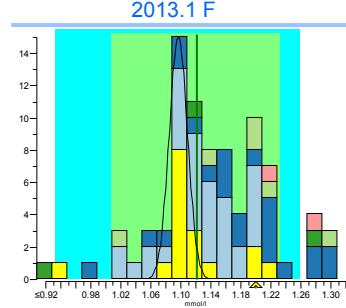
	cons.	meth.	ref.	lab
mean	1.19	1.19	1.167	1.2
SD	0.03	0.03		
n	17	17		
no	1	1		



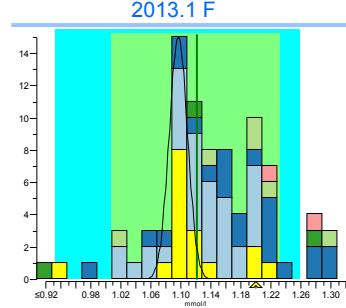
	cons.	meth.	exp.	lab
mean	1.22	1.22	1.21	1.3
SD	0.04	0.04		
n	17	17		
no	2	2		



	cons.	meth.	ref.	lab
mean	1.07	1.07	1.074	1.1
SD	0.03	0.03		
n	17	17		
no	1	1		



	cons.	meth.	ref.	lab
mean	1.04	1.04	1.087	1
SD	0.03	0.03		
n	17	17		
no	1	1		



	cons.	meth.	ref.	lab
mean	1.10	1.10	1.121	1.2
SD	0.01	0.01		
n	17	17		
no	5	5		

## Legend

- [Yellow square] Immunoinhibition
- [Light blue square] PEG modified enzyme, PEGME (Kyowa Medex)
- [Dark blue square] Accelerator Selective Detergent ("Ultra HDL")
- [Green square] Catalase method (Denka Seiken)
- [Dark green square] Overige methoden

[Pink square] Precipitation Technique