

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

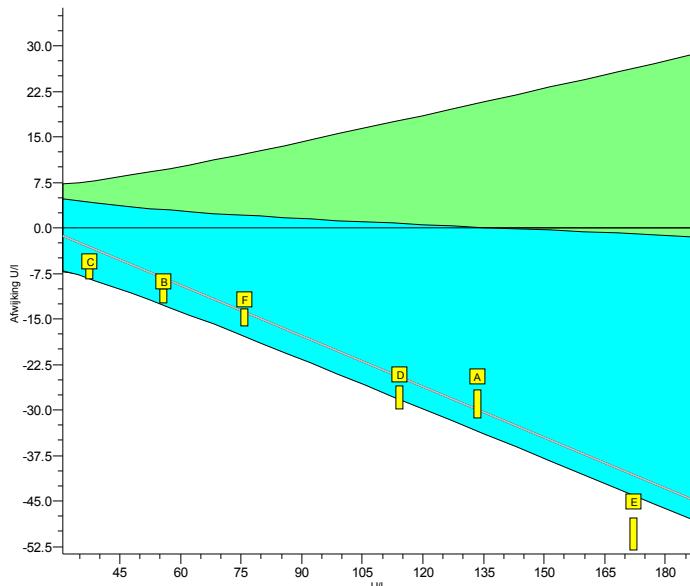
Scores	Your score	MAP	reported
Quantitative	27	38	

INPUTs 2013.1

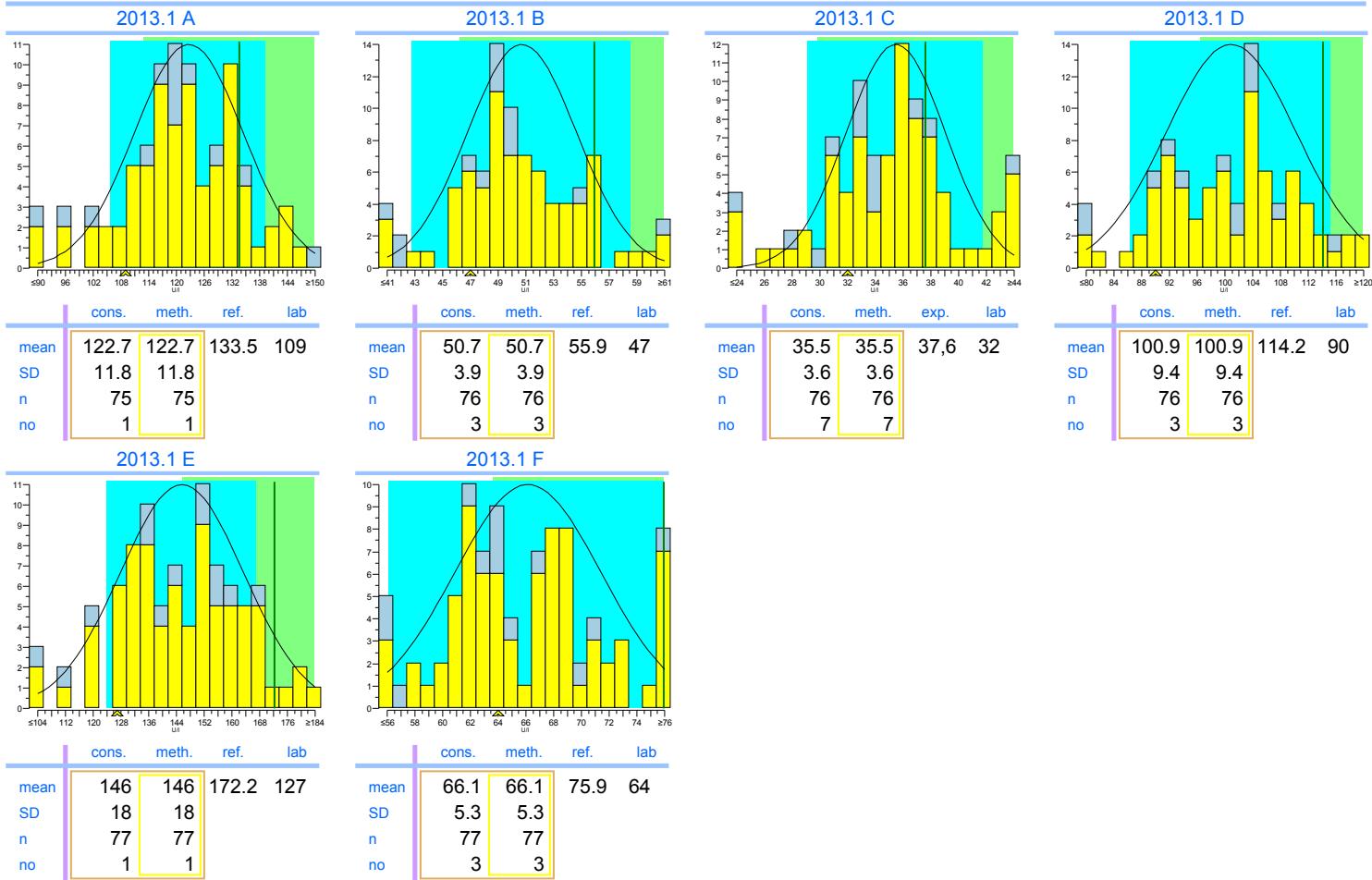
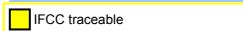
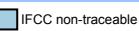
Analyte		Trueness				Precision		Performance			
		your mean	ref.	cons.	SDtl	your prec.	SDbl	this survey	PS	cumulative	PSc
ALAT	U/l	78.2	98.2	86.9	10.1	2.0	2.7		0		10
Alk. Phosphatase	U/l	174	188	173	15	2	2		2		12
Amylase	U/l	191	207	201	8	2	2		2		12
ASAT	U/l	68.2	78.6	71.5	6.8	2.8	1.6		0		10
Calcium	mmol/l	2.49	2.46	2.47	0.07	0.03	0.02		2		12
Chloride	mmol/l	104.2	101.4	102.3	1.8	0.8	0.9		1		11
CK	U/l	247	244	253	18	3	3		2		12
Creatinine	µmol/l	145.5	149.1	150.7	5.6	2.6	2.1		1		11
eGFR (F, 55, white)	ml/min/1,73m ²	33.4	32.0	32.2	2.6	0.3	0.6		2		12
Gamma-GT	U/l	90.8	87.7	86.3	8.6	2.0	1.2		2		12
Glucose	mmol/l	14.13	13.93	14.19	0.44	0.25	0.18		1		11
LD	U/l	498	532	523	33	15	9		2		12
Magnesium	mmol/l	1.29	1.22	1.23	0.04	0.04	0.02		1		11
Potassium	mmol/l	5.50	5.39	5.30	0.10	0.08	0.04		1		11
Sodium	mmol/l	145.3	144.0	142.7	1.7	1.5	0.9		1		11
Total Protein	g/l	68.0	66.8	66.0	1.8	0.7	0.7		2		12
Urate	µmol/l	389	376	374	15	6	4		2		12
Cholesterol	mmol/l	5.02	5.07	5.15	0.14	0.05	0.05		2		12
HDL-Cholesterol	mmol/l	1.22	1.14	1.17	0.06	0.04	0.03		1		11
		Total :				27		27			

INPUTS 2013.1**ALAT**

units: U/l



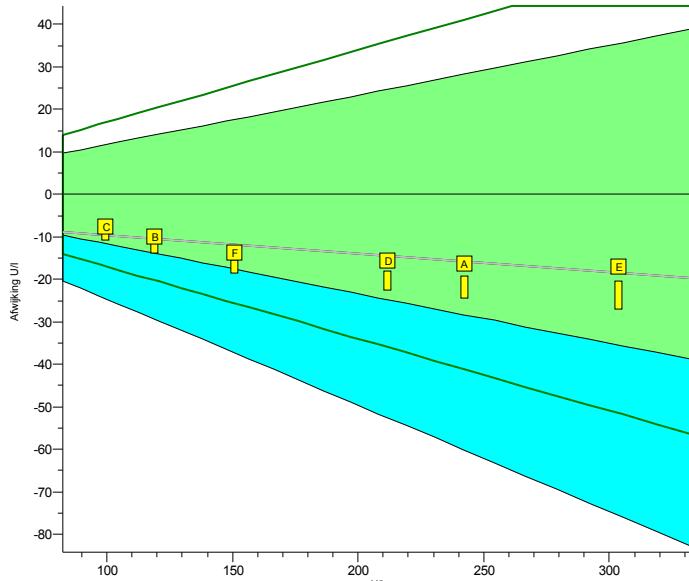
	2013.1	cumulative
Trueness	-20%	-20%
Precision	2.3%	2.3%
Number	6	6
Outliers	0	0
Sigma-TE	-0.1	-0.1
Sigma-SA	-1.3	-1.3
Score pictogram		
Regression line	$7.4 + 0.721 \cdot x$	$7.4 + 0.721 \cdot x$

 Consensus group IFCC traceerbaar
 Method IFCC traceable
**Legend**



INPUTs 2013.1

Alk. Phosphatase

units: U/l

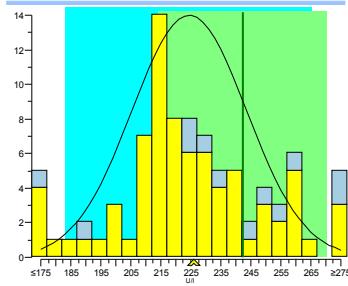


	2013.1	cumulative
Trueness	-7.2%	-7.2%
Precision	1.0%	1.0%
Number	6	6
Outliers	0	0
Sigma-TE	5.1	5.1
Sigma-SA	6.0 [2]	6.0 [2]
Score pictogram	[Yellow]	[Yellow]
Regression line	$-5 + 0.956 \cdot x$	$-5 + 0.956 \cdot x$

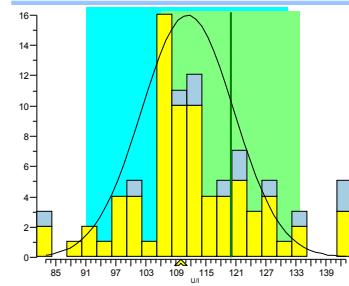
Consensus group
Method

IFCC traceerbaar
IFCC traceable

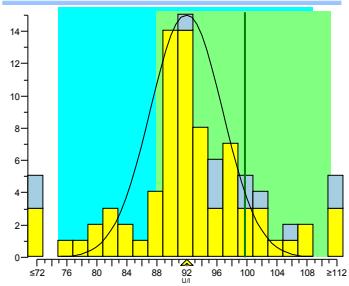
2013.1 A



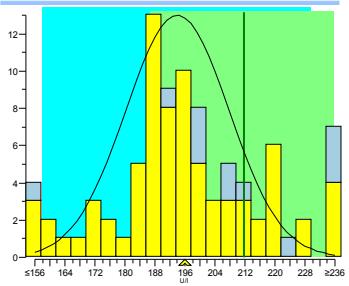
2013.1 B



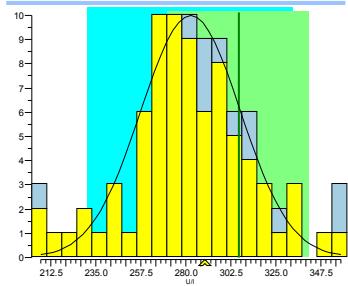
2013.1 C



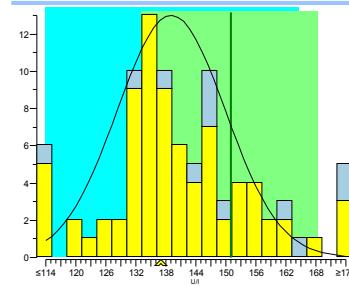
2013.1 D



2013.1 E



2013.1 F



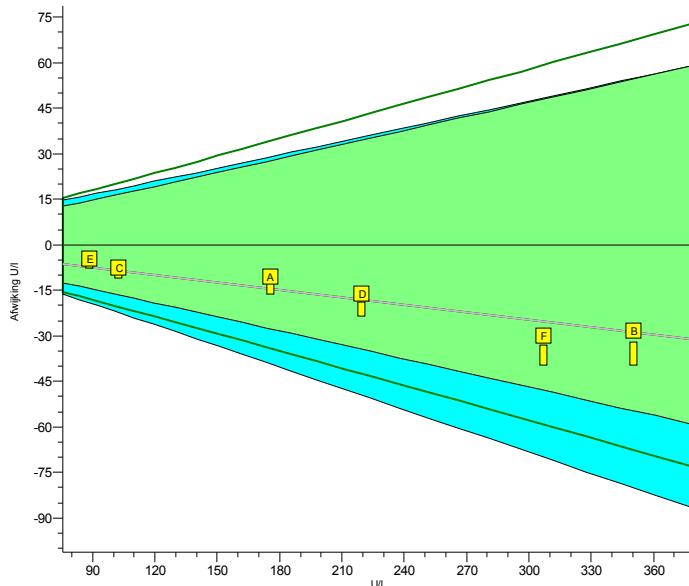
Legend

[Yellow] IFCC traceable

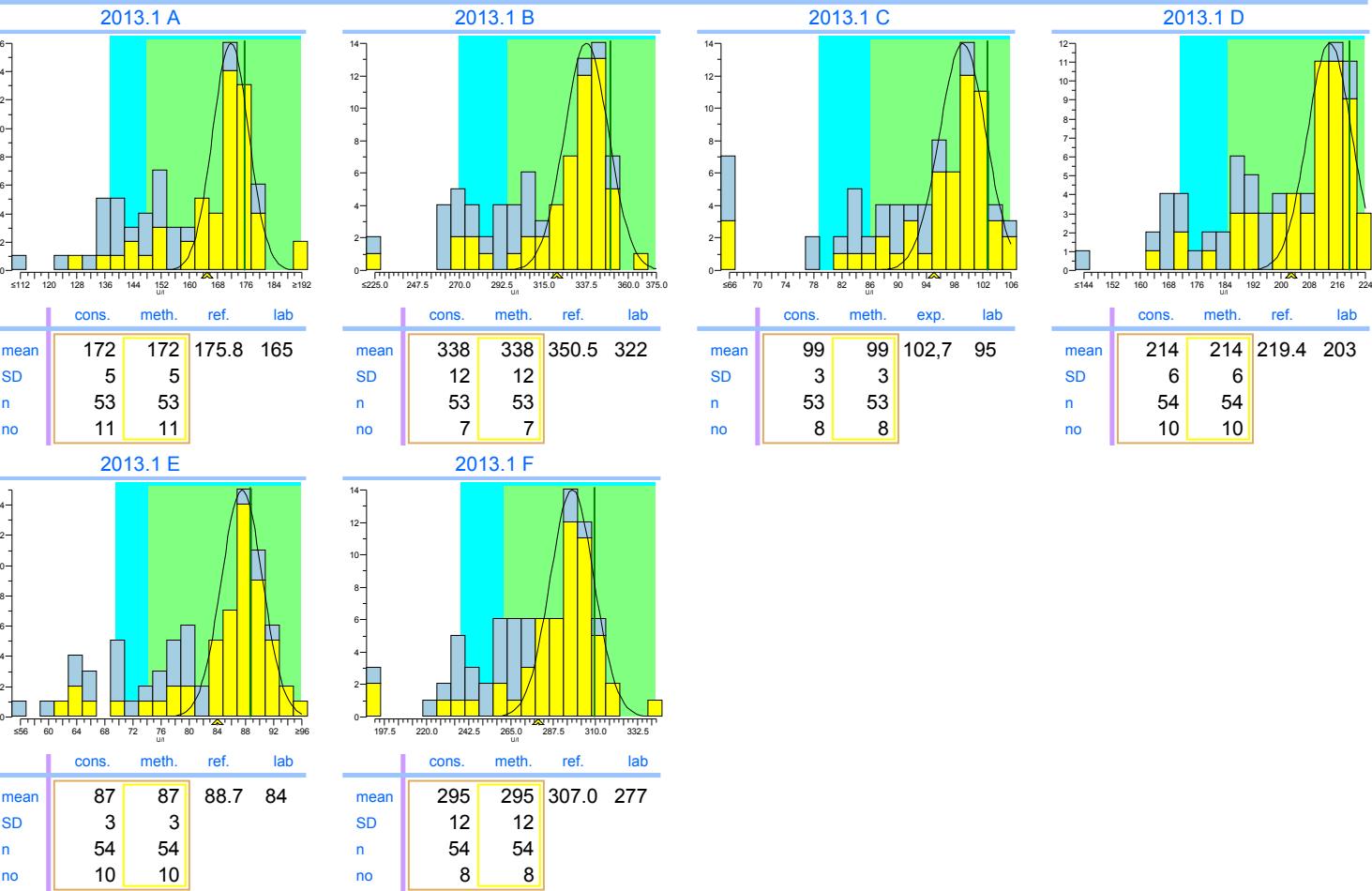
[Grey] IFCC non-traceable

INPUTs 2013.1**Amylase**

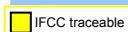
units: U/l

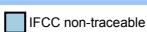


	2013.1	cumulative
Trueness	-7.9%	-7.9%
Precision	0.92%	0.92%
Number	6	6
Outliers	0	0
Sigma-TE	6.0	6.0
Sigma-SA	6.0 [2]	6.0 [2]
Score pictogram	[Yellow]	[Yellow]
Regression line	0 + 0.918.x	0 + 0.918.x

 Consensus group: IFCC traceerbaar
 Method: IFCC traceable


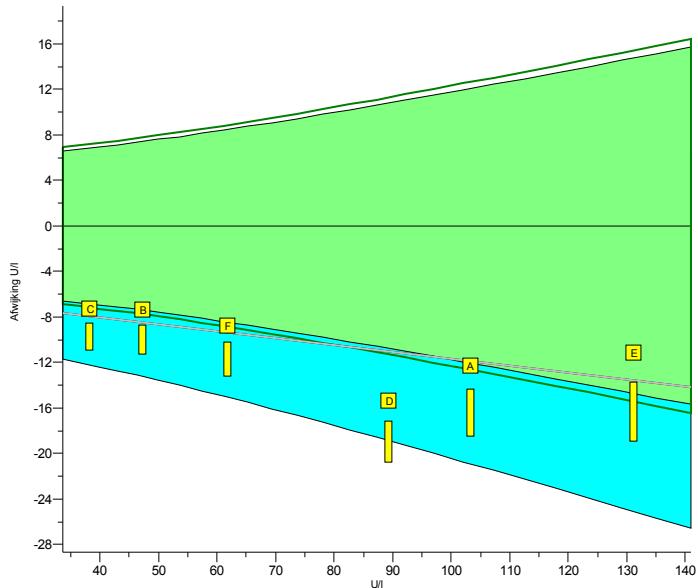
Legend

 IFCC traceable

 IFCC non-traceable

INPUTs 2013.1**ASAT**

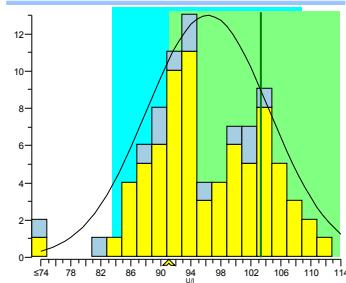
units: U/l



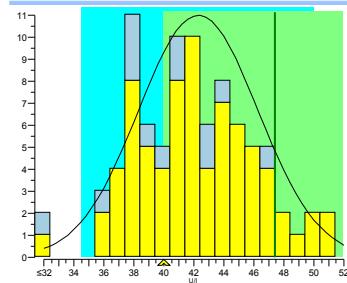
	2013.1	cumulative
Trueness	-13%	-13%
Precision	3.9%	3.9%
Number	6	6
Outliers	0	0
Sigma-TE	0.5	0.5
Sigma-SA	0.8 0	0.8 0
Score pictogram		
Regression line	$-5.6 + 0.939 \cdot x$	$-5.6 + 0.939 \cdot x$

 Consensus group: IFCC traceerbaar
 Method: IFCC traceable

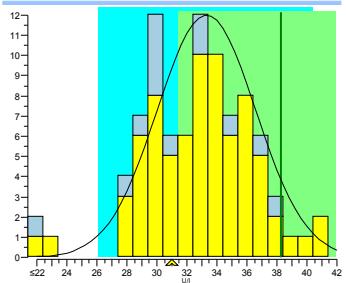
2013.1 A



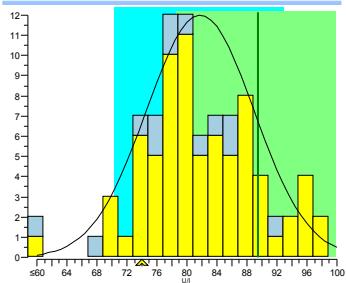
2013.1 B



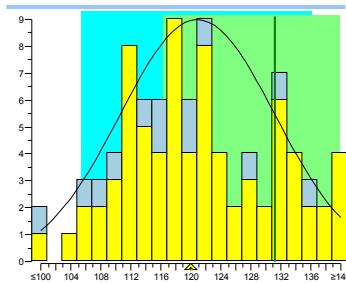
2013.1 C



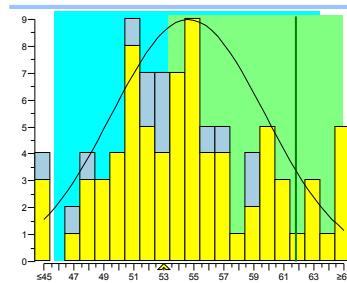
2013.1 D



2013.1 E



2013.1 F



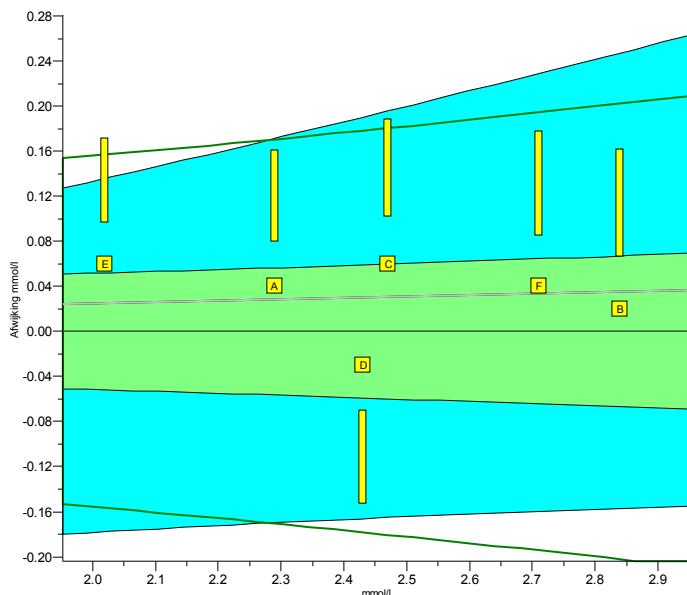
Legend

■ IFCC traceable■ IFCC non-traceable

INPUTs 2013.1

Calcium

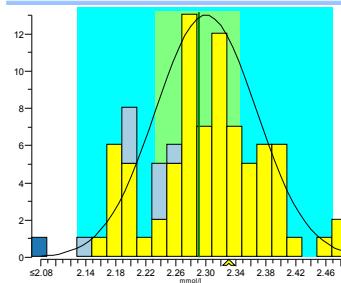
units: mmol/l



	2013.1	cumulative
Trueness	+1.3%	+1.3%
Precision	1.4%	1.4%
Number	6	6
Outliers	0	0
Sigma-TE	1.3	1.3
Sigma-SA	4.8 [2]	4.8 [2]
Score pictogram		
Regression line	$0.00 + 1.012 \cdot x$	$0.00 + 1.012 \cdot x$

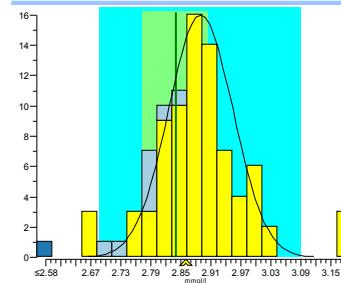
Consensus group: Colorimetrisch
 Method: Colorimetric, automatic, discrete

2013.1 A



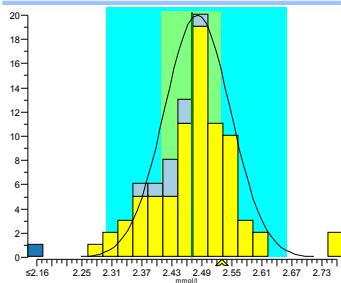
	cons.	meth.	ref.	lab
mean	2.30	2.30	2.29	2.33
SD	0.07	0.07		
n	80	80		
no	2	2		

2013.1 B



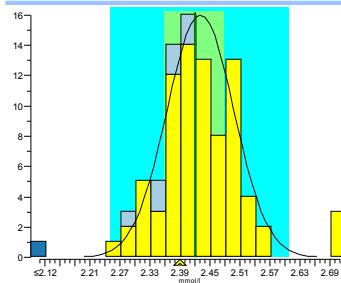
	cons.	meth.	ref.	lab
mean	2.89	2.89	2.84	2.86
SD	0.06	0.06		
n	80	80		
no	6	6		

2013.1 C



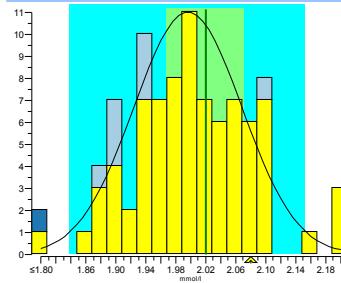
	cons.	meth.	exp.	lab
mean	2.48	2.48	2.47	2.53
SD	0.07	0.07		
n	79	79		
no	2	2		

2013.1 D



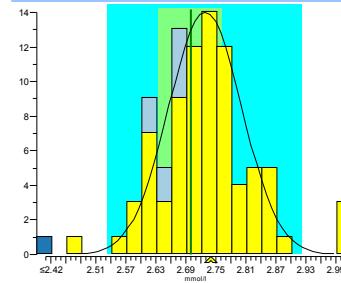
	cons.	meth.	ref.	lab
mean	2.44	2.44	2.43	2.40
SD	0.07	0.07		
n	80	80		
no	3	3		

2013.1 E



	cons.	meth.	ref.	lab
mean	2.00	2.00	2.02	2.08
SD	0.07	0.07		
n	81	81		
no	3	3		

2013.1 F



	cons.	meth.	ref.	lab
mean	2.74	2.74	2.71	2.75
SD	0.07	0.07		
n	80	80		
no	4	4		

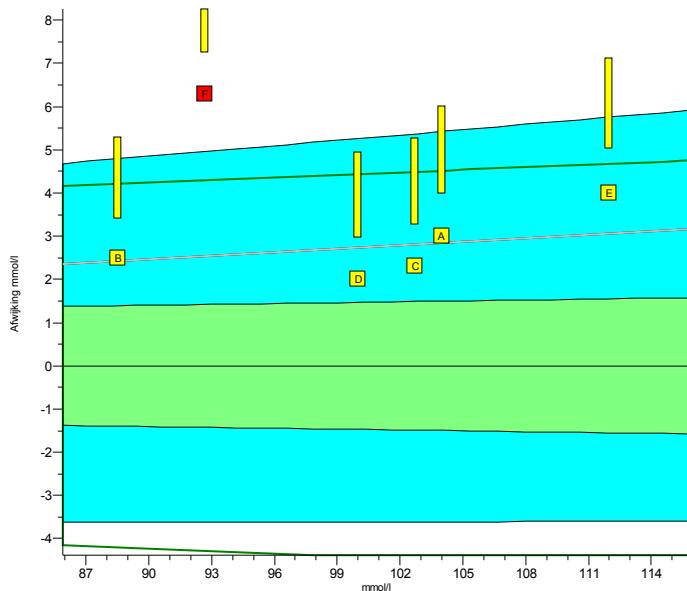
Legend

 Colorimetric, automatic, discrete ISE indirect (with predilution) Other methods

INPUTs 2013.1

Chloride

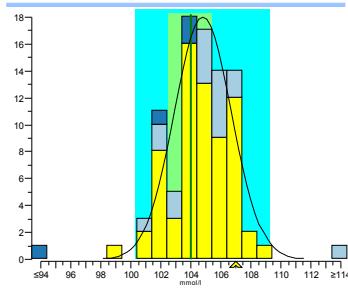
units: mmol/l



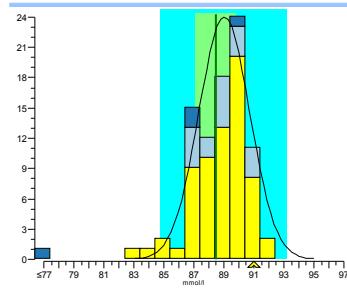
	2013.1	cumulative
Trueness	+2.7%	+2.7%
Precision	0.78%	0.78%
Number	6	6
Outliers	1	1
Sigma-TE	-1.2	-1.2
Sigma-SA	2.2	2.2
Score pictogram		
Regression line	$0.0 + 1.027 \cdot x$	$0.0 + 1.027 \cdot x$

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

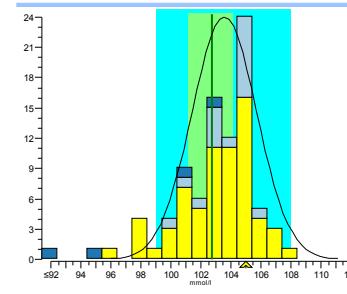
2013.1 A



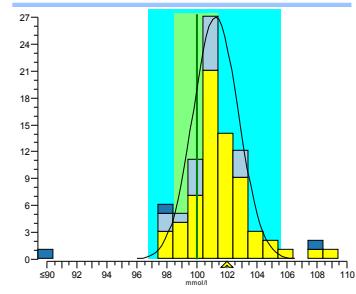
2013.1 B



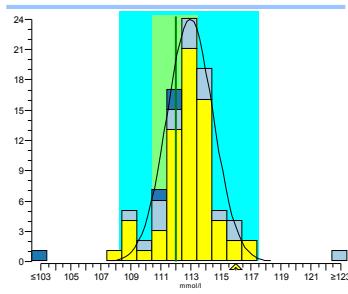
2013.1 C



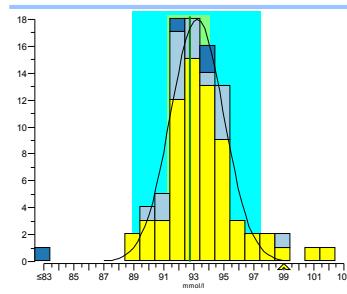
2013.1 D



2013.1 E



2013.1 F



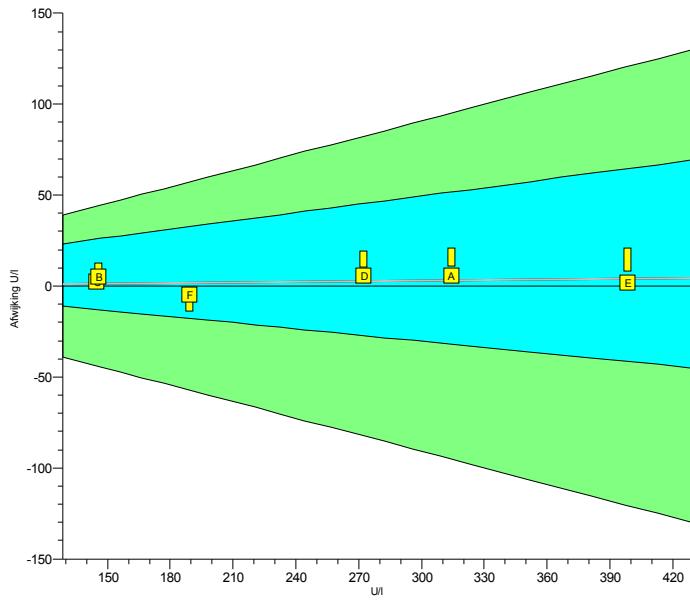
Legend

ISE indirect (with predilution)	ISE direct (no predilution)	Other methods
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INPUTS 2013.1

CK

units: U/l

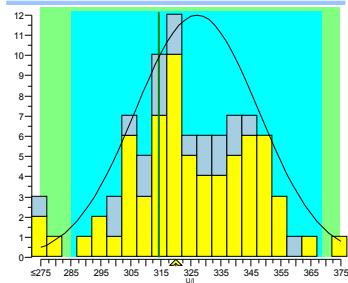


	2013.1	cumulative
Trueness	+1.0%	+1.0%
Precision	1.3%	1.3%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 [2]	6.0 [2]
Sigma-SA	6.0	6.0
Score pictogram	[Green box]	[Green box]
Regression line	$0 + 1.010 \cdot x$	$0 + 1.010 \cdot x$

Consensus group
Method

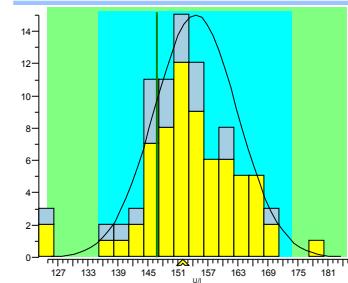
IFCC traceerbaar
IFCC traceable

2013.1 A



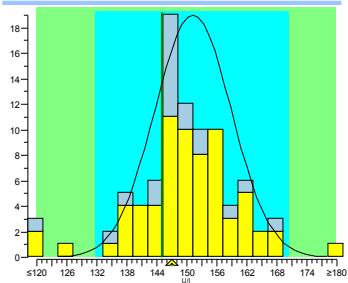
	cons.	meth.	ref.	lab
mean	327	327	314.3	320
SD	21	21		
n	68	68		
no	2	2		

2013.1 B



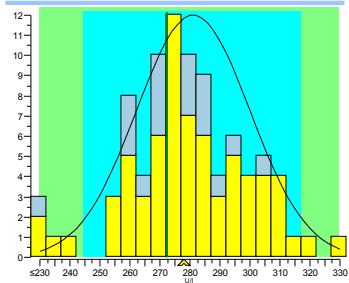
	cons.	meth.	ref.	lab
mean	154	154	145.8	151
SD	8	8		
n	67	67		
no	3	3		

2013.1 C



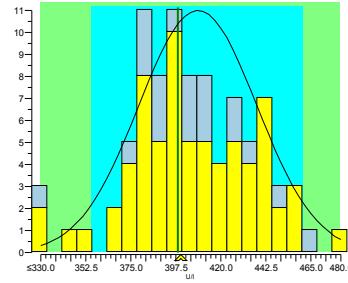
	cons.	meth.	exp.	lab
mean	151	151	145	147
SD	8	8		
n	68	68		
no	4	4		

2013.1 D



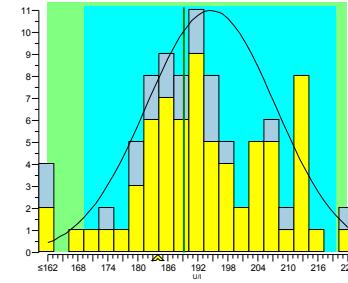
	cons.	meth.	ref.	lab
mean	281	281	272.3	278
SD	19	19		
n	69	69		
no	2	2		

2013.1 E



	cons.	meth.	ref.	lab
mean	409	409	398.4	400
SD	30	30		
n	69	69		
no	2	2		

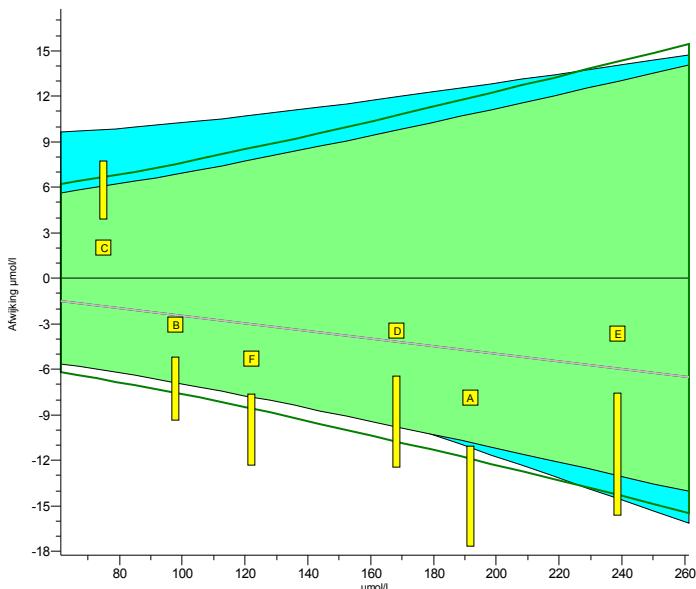
2013.1 F



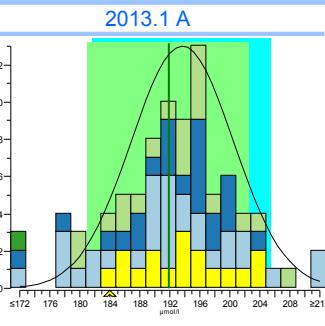
	cons.	meth.	ref.	lab
mean	195	195	189.2	184
SD	13	13		
n	69	69		
no	2	2		

Legend

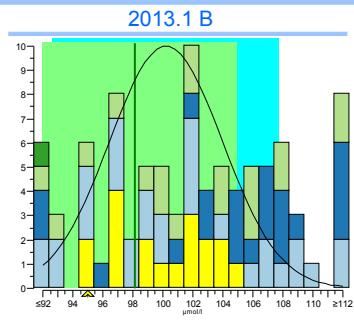


INPUTs 2013.1**Creatinine**units: $\mu\text{mol/l}$ 

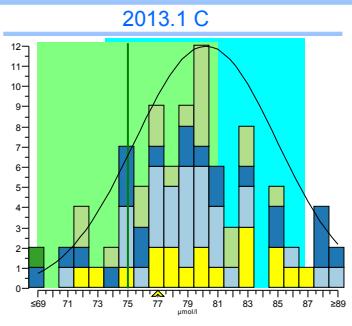
	2013.1	cumulative
Trueness	-2.4%	-2.4%
Precision	1.8%	1.8%
Number	6	6
Outliers	0	0
Sigma-TE	2.9	2.9
Sigma-SA	3.3	3.3
Score pictogram		
Regression line	$0.0 + 0.975 \cdot x$	$0.0 + 0.975 \cdot x$

 Consensus group: Enzymatische kreatinine
 Method: Enzymatic, automatic


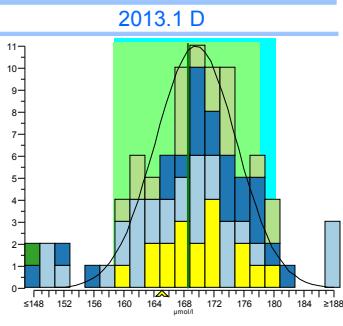
	cons.	meth.	ref.	lab
mean	193.7	193.7	191.9	184
SD	6.8	6.8		
n	17	17		
no	0	0		



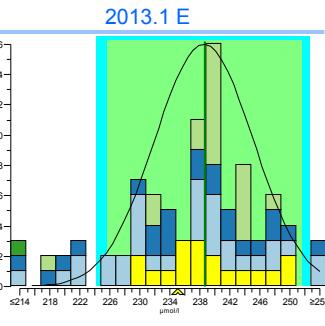
	cons.	meth.	ref.	lab
mean	100.2	100.2	98.11	95
SD	3.8	3.8		
n	18	18		
no	0	0		



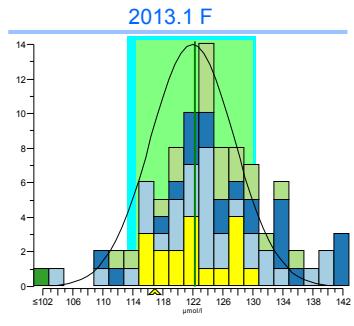
	cons.	meth.	exp.	lab
mean	80.2	80.2	75	77
SD	4.7	4.7		
n	18	18		
no	0	0		



	cons.	meth.	ref.	lab
mean	169.6	169.6	168.5	165
SD	5.4	5.4		
n	18	18		
no	0	0		



	cons.	meth.	ref.	lab
mean	238.5	238.5	238.7	235
SD	6.6	6.6		
n	18	18		
no	0	0		



	cons.	meth.	ref.	lab
mean	121.9	121.9	122.3	117
SD	5.6	5.6		
n	18	18		
no	0	0		

Legend
 Enzymatic, automatic

 Alk. Picrate, kinetic with compensation

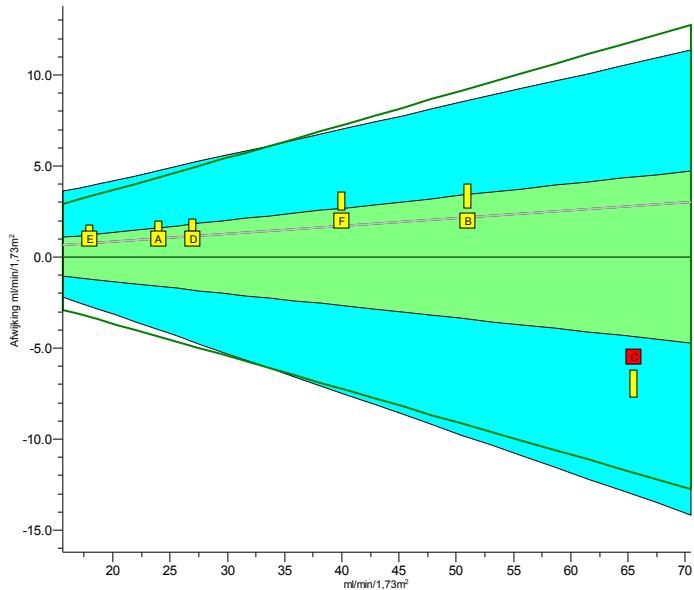
 Alk. Picrate, kinetic

 Alk. Picrate, endpoint

 Other methods

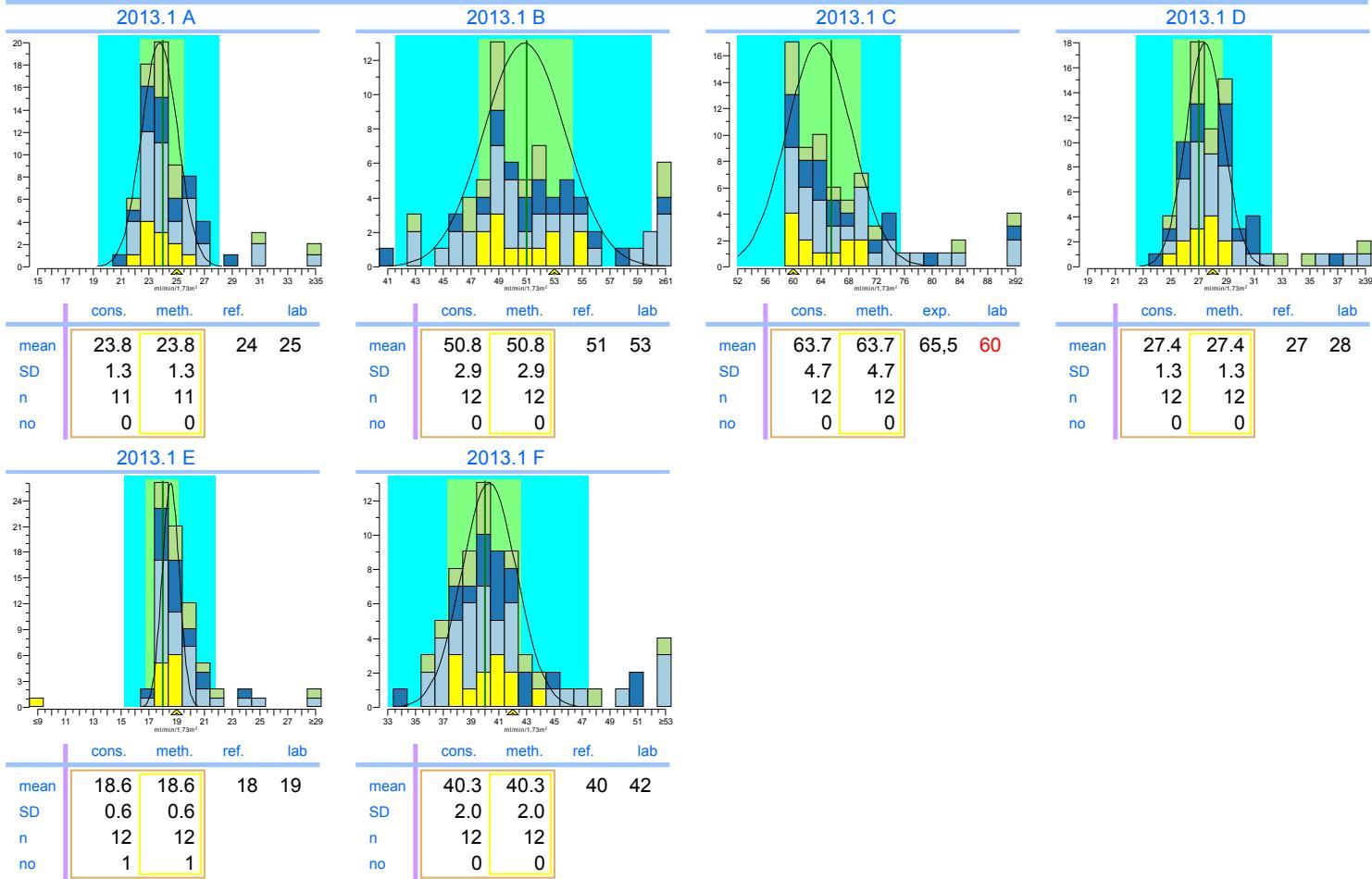
INPUTs 2013.1

eGFR (F, 55, white)

units: ml/min/1,73m²

	2013.1	cumulative
Trueness	+4.4%	+4.4%
Precision	1.1%	1.1%
Number	6	6
Outliers	1	1
Sigma-TE	2.4	2.4
Sigma-SA	6.0 [2]	6.0 [2]
Score pictogram	[Green, Yellow, Red]	[Green, Yellow, Red]
Regression line	0.0 + 1.043.x	0.0 + 1.043.x

Consensus group: Enzymatische kreatinine
Method: Enzymatic, automatic



Legend

[Yellow] Enzymatic, automatic

[Light Blue] Other methods

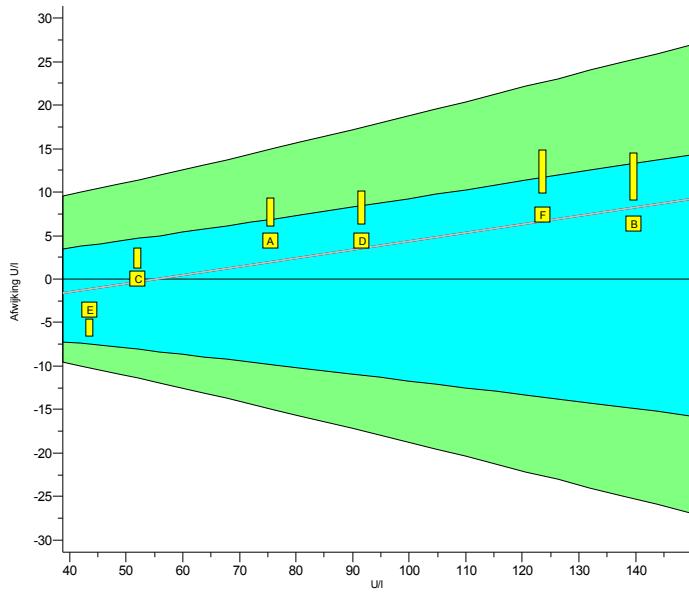
[Dark Blue] Alk. Picrate, kinetic with compensation

[Light Green] Alk. Picrate, kinetic

INPUTs 2013.1

Gamma-GT

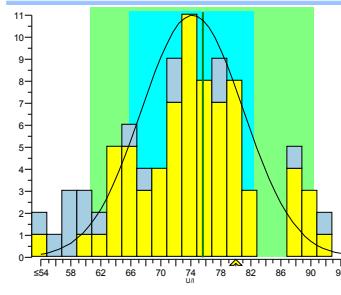
units: U/l



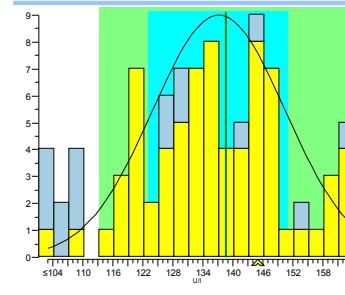
	2013.1	cumulative
Trueness	+3.6%	+3.6%
Precision	2.4%	2.4%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 [2]	6.0 [2]
Sigma-SA	4.5	4.5
Score pictogram	[Color-coded box]	[Color-coded box]
Regression line	$-5.4 + 1.097 \cdot x$	$-5.4 + 1.097 \cdot x$

 Consensus group: IFCC traceerbaar
 Method: IFCC traceable

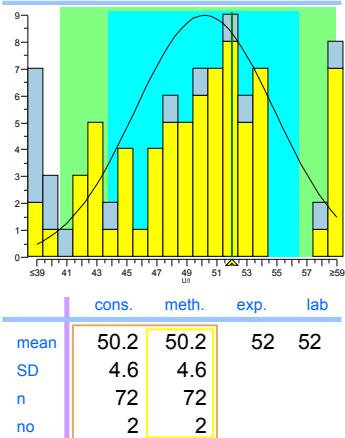
2013.1 A



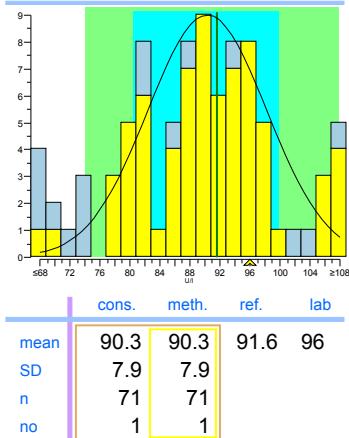
2013.1 B



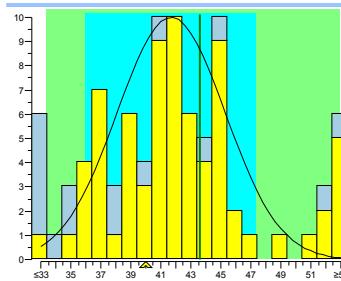
2013.1 C



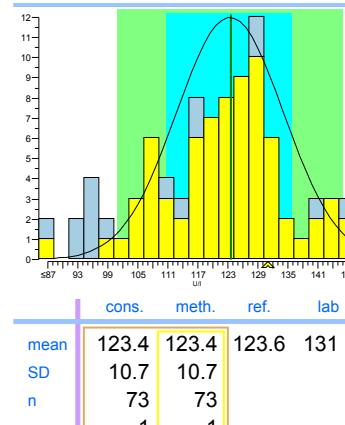
2013.1 D



2013.1 E



2013.1 F



Legend

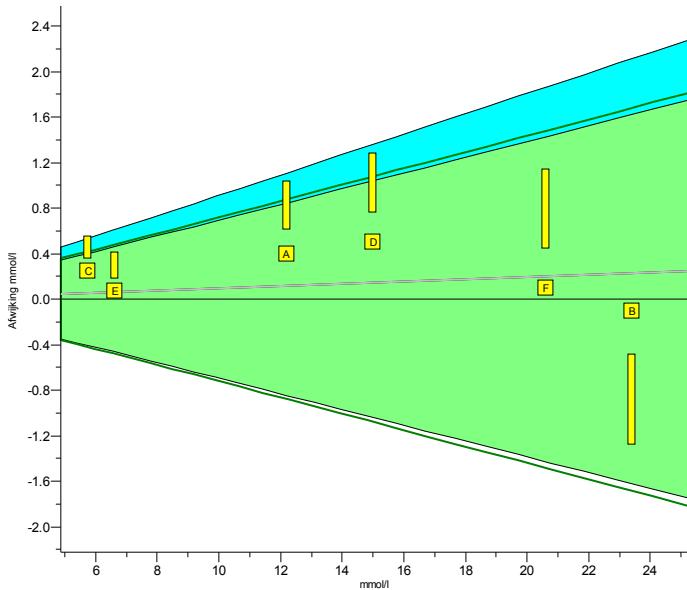
 IFCC traceable

 IFCC non-traceable

INPUTS 2013.1

Glucose

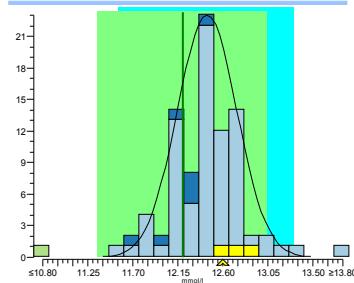
units: mmol/l



	2013.1	cumulative
Trueness	+1.5%	+1.5%
Precision	1.8%	1.8%
Number	6	6
Outliers	0	0
Sigma-TE	4.3	4.3
Sigma-SA	4.5 [1]	4.5 [1]
Score pictogram	[Green box with yellow dot]	[Green box with yellow dot]
Regression line	$0.00 + 1.010 \cdot x$	$0.00 + 1.010 \cdot x$

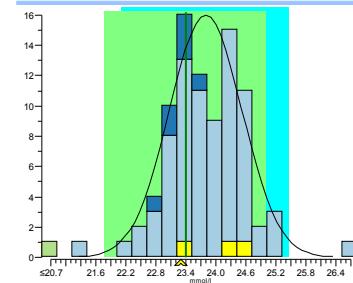
Consensus group: Natte chemie
Method: Glucose-oxydase/POD,automatic

2013.1 A



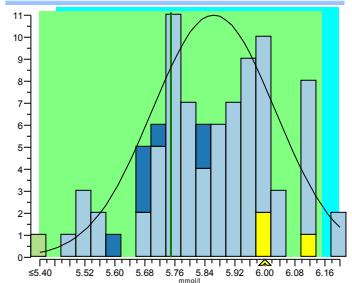
	cons.	meth.	ref.	lab
mean	12.44	12.76	12.2	12.6
SD	0.30			
n	88	3		
no	3	0		

2013.1 B



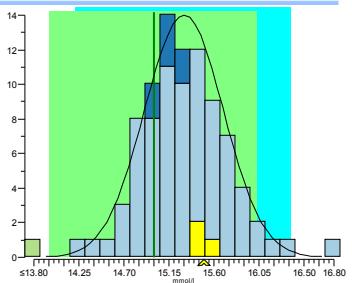
	cons.	meth.	ref.	lab
mean	23.8	24.1	23.4	23.3
SD	0.7			
n	88	3		
no	3	0		

2013.1 C



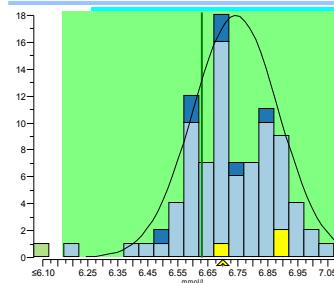
	cons.	meth.	exp.	lab
mean	5.86	6.03	5.75	6.0
SD	0.16			
n	88	3		
no	1	0		

2013.1 D



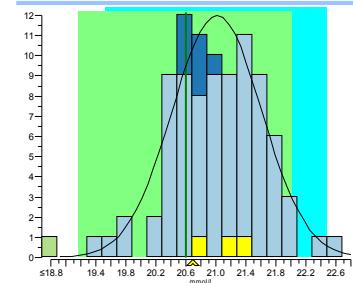
	cons.	meth.	ref.	lab
mean	15.30	15.55	15.0	15.5
SD	0.39			
n	88	3		
no	2	0		

2013.1 E



	cons.	meth.	ref.	lab
mean	6.74	6.83	6.63	6.7
SD	0.14			
n	89	3		
no	2	0		

2013.1 F



	cons.	meth.	ref.	lab
mean	21.01	21.04	20.6	20.7
SD	0.60			
n	89	3		
no	1	0		

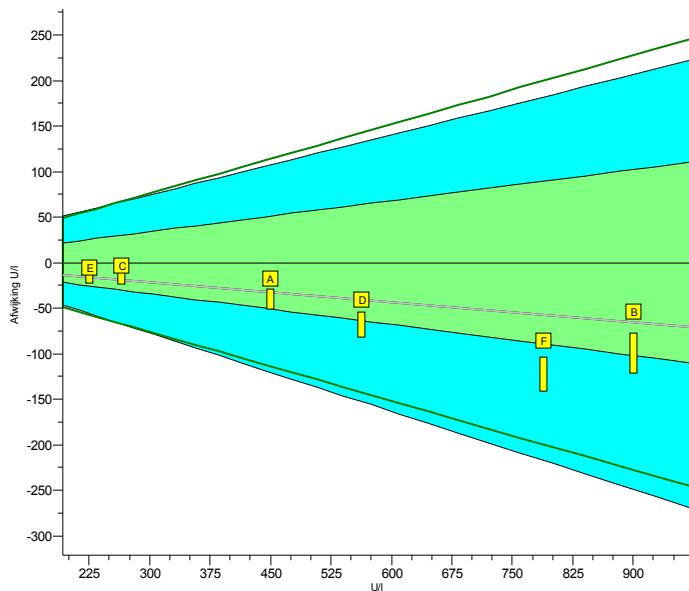
Legend

- [Yellow square] Glucose-oxydase/POD,automatic
- [Light blue square] Hexokinase, automatic
- [Dark blue square] Glucose-oxydase, amperometric, H₂O₂
- [Green square] Other methods

INPUTs 2013.1

LD

units: U/l



	2013.1	cumulative
Trueness	-6.6%	-6.6%
Precision	2.8%	2.8%
Number	6	6
Outliers	0	0
Sigma-TE	3.5	3.5
Sigma-SA	6.0 [2]	6.0 [2]
Score pictogram	[Yellow/Green]	[Yellow/Green]
Regression line	$0 + 0.927 \cdot x$	$0 + 0.927 \cdot x$

Consensus group
Method

IFCC traceerbaar
IFCC traceable

2013.1 A

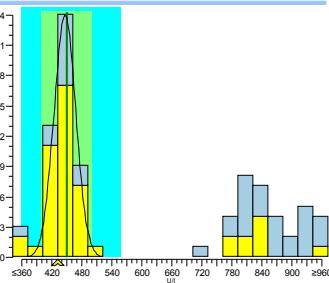
2013.1 B

2013.1 C

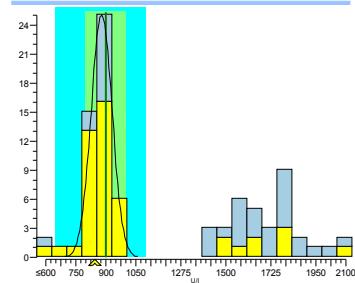
2013.1 D

2013.1 E

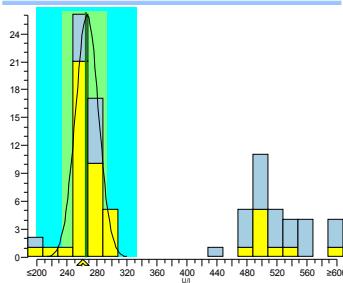
2013.1 F



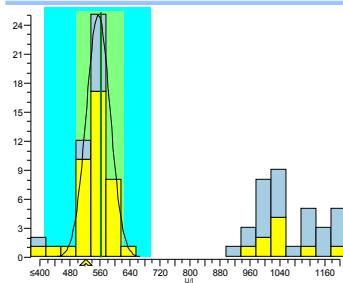
	cons.	meth.	ref.	lab
mean	446	446	450.7	433
SD	22	22		
n	48	48		
no	11	11		



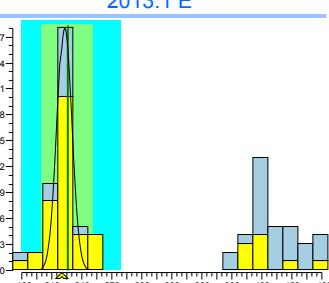
	cons.	meth.	ref.	lab
mean	879	879	900.8	846
SD	52	52		
n	47	47		
no	11	11		



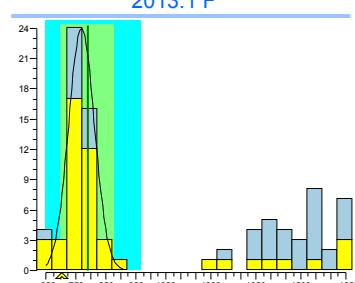
	cons.	meth.	exp.	lab
mean	267	267	265	261
SD	15	15		
n	48	48		
no	11	11		



	cons.	meth.	ref.	lab
mean	556	556	563.0	522
SD	32	32		
n	48	48		
no	11	11		



	cons.	meth.	ref.	lab
mean	223	223	226.1	220
SD	7	7		
n	48	48		
no	16	16		



	cons.	meth.	ref.	lab
mean	769	769	788.9	703
SD	43	43		
n	48	48		
no	11	11		

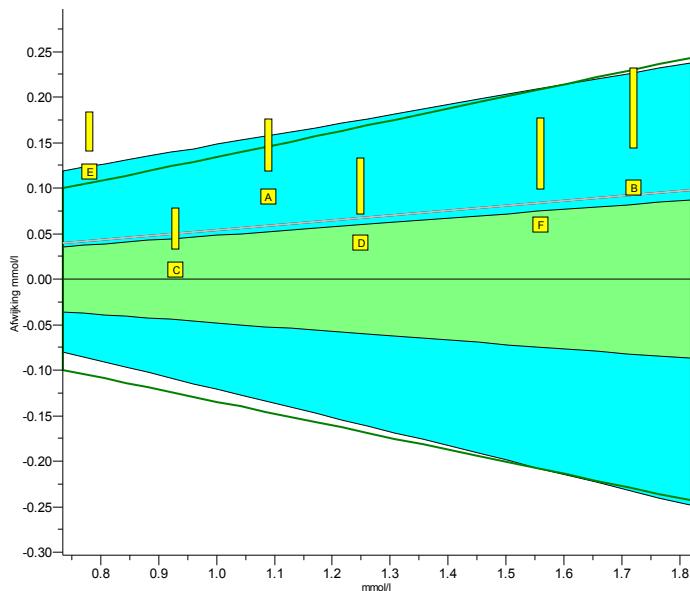
Legend

■ IFCC traceable■ IFCC non-traceable

INPUTs 2013.1

Magnesium

units: mmol/l

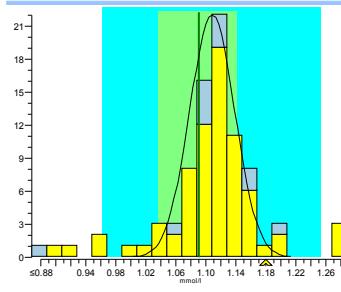


	2013.1	cumulative
Trueness	+5.7%	+5.7%
Precision	3.3%	3.3%
Number	6	6
Outliers	0	0
Sigma-TE	0.3	0.3
Sigma-SA	4.5 [1]	4.5 [1]
Score pictogram		
Regression line	$0.00 + 1.054 \cdot x$	$0.00 + 1.054 \cdot x$

Consensus group
Method

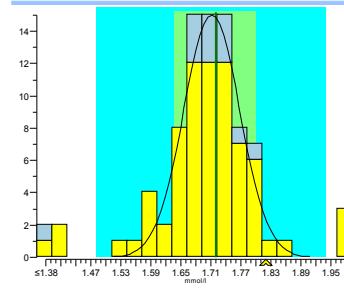
Overall
Colorimetric

2013.1 A



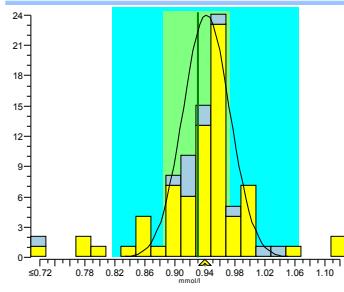
	cons.	meth.	ref.	lab
mean	1.11	1.11	1.09	1.18
SD	0.03	0.03		
n	85	73		
no	11	10		

2013.1 B



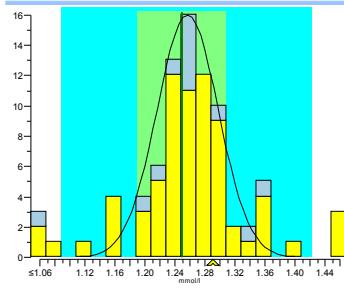
	cons.	meth.	ref.	lab
mean	1.71	1.71	1.72	1.82
SD	0.06	0.06		
n	85	73		
no	8	7		

2013.1 C



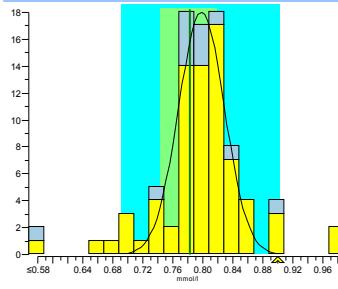
	cons.	meth.	exp.	lab
mean	0.94	0.94	0.93	0.94
SD	0.03	0.03		
n	85	73		
no	9	8		

2013.1 D



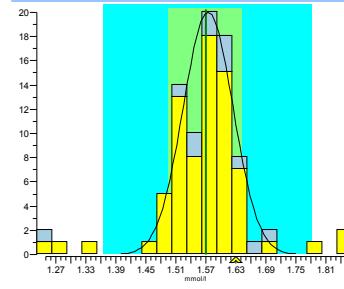
	cons.	meth.	ref.	lab
mean	1.26	1.26	1.25	1.29
SD	0.04	0.05		
n	83	71		
no	9	8		

2013.1 E



	cons.	meth.	ref.	lab
mean	0.798	0.799	0.782	0.9
SD	0.031	0.031		
n	86	74		
no	13	11		

2013.1 F



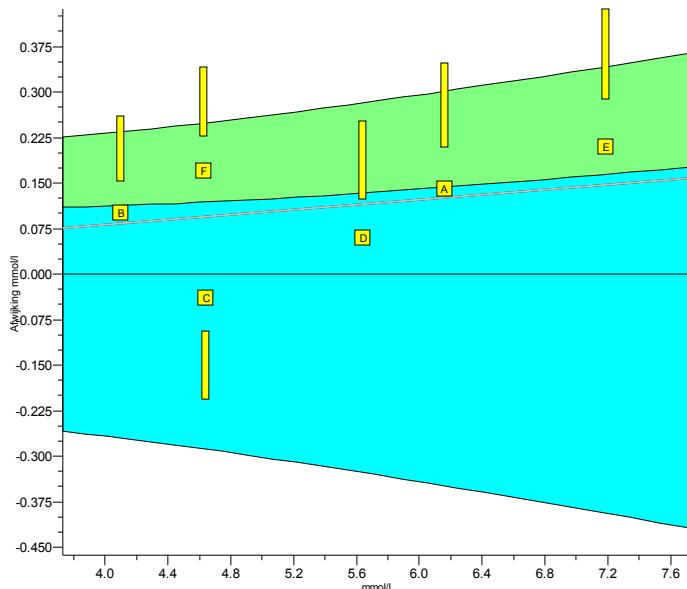
	cons.	meth.	ref.	lab
mean	1.56	1.56	1.56	1.62
SD	0.05	0.05		
n	86	74		
no	7	6		

Legend

Colorimetric	Other methods
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INPUTs 2013.1**Potassium**

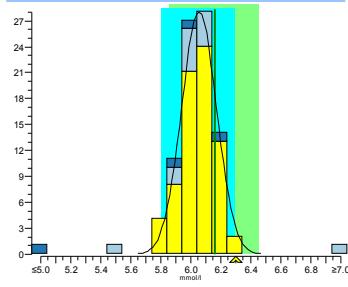
units: mmol/l



	2013.1	cumulative
Trueness	+2.0%	+2.0%
Precision	1.5%	1.5%
Number	6	6
Outliers	0	0
Sigma-TE	3.9	3.9
Sigma-SA	2.9	2.9
Score pictogram		
Regression line	$0.00 + 1.021 \cdot x$	$0.00 + 1.021 \cdot x$

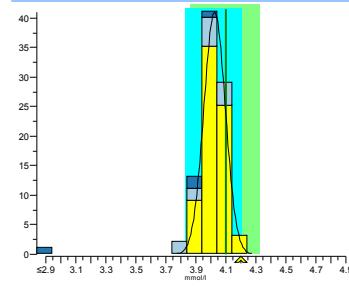
Consensus group ISE verdund/Vlamfotometrie
 Method ISE indirect (with predilution)

2013.1 A



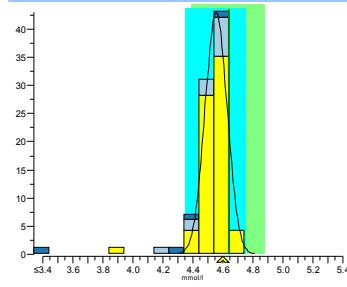
	cons.	meth.	ref.	lab
mean	6.05	6.05	6.16	6.3
SD	0.12	0.12		
n	72	72		
no	0	0		

2013.1 B



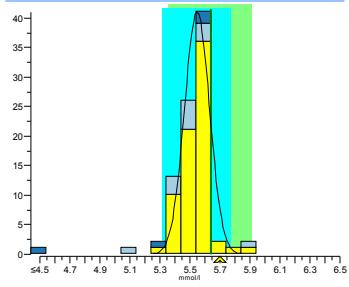
	cons.	meth.	ref.	lab
mean	4.02	4.02	4.10	4.2
SD	0.07	0.07		
n	72	72		
no	0	0		

2013.1 C



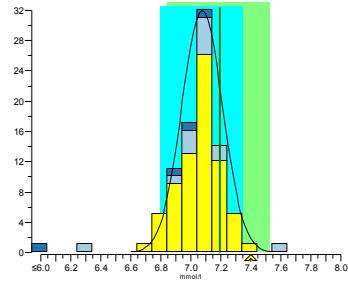
	cons.	meth.	ref.	lab
mean	4.55	4.55	4.64	4.6
SD	0.07	0.07		
n	72	72		
no	1	1		

2013.1 D



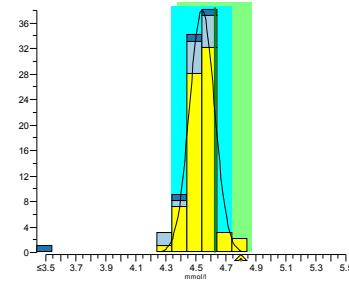
	cons.	meth.	ref.	lab
mean	5.55	5.55	5.64	5.7
SD	0.08	0.08		
n	72	72		
no	2	2		

2013.1 E



	cons.	meth.	ref.	lab
mean	7.08	7.08	7.19	7.4
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	4.8
SD	0.08	0.08		
n	73	73		
no	2	2		

Legend

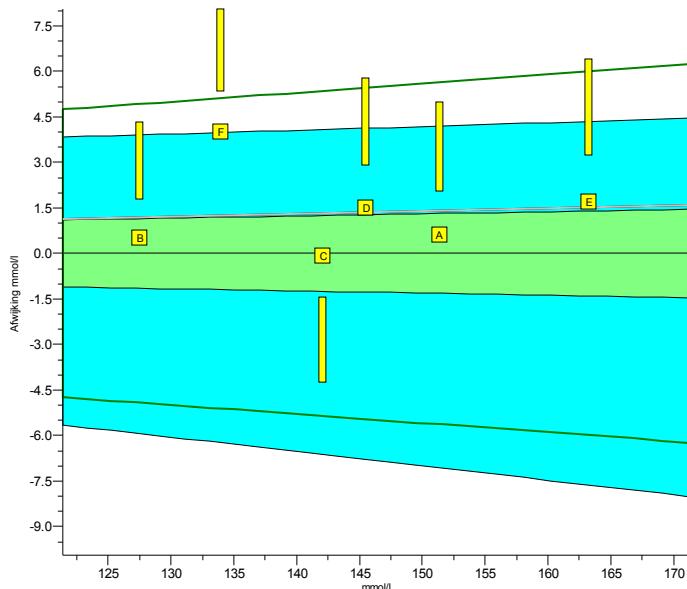
ISE indirect (with predilution)

ISE direct (no predilution)

Other methods

INPUTs 2013.1**Sodium**

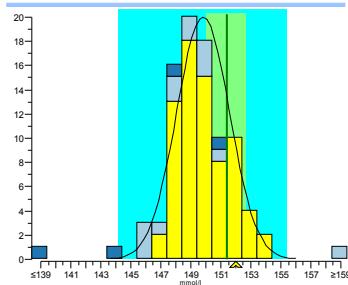
units: mmol/l



	2013.1	cumulative
Trueness	+0.95%	+0.95%
Precision	1.0%	1.0%
Number	6	6
Outliers	0	0
Sigma-TE	0.7	0.7
Sigma-SA	4.3 [1]	4.3 [1]
Score pictogram	[Yellow]	[Yellow]
Regression line	$0.0 + 1.009.x$	$0.0 + 1.009.x$

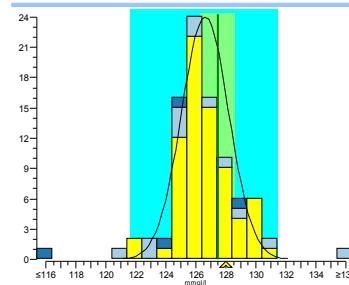
 Consensus group ISE verdund/Vlamfotometrie
 Method ISE indirect (with predilution)

2013.1 A



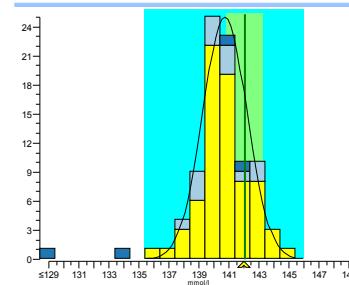
mean	149.8	149.8	151.4	152
SD	1.8	1.8		
n	72	72		
no	0	0		

2013.1 B



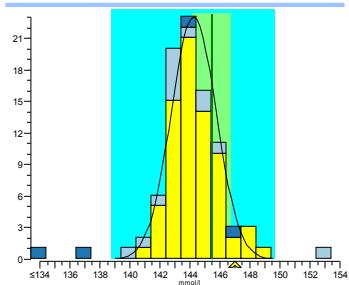
mean	126.6	126.6	127.5	128
SD	1.6	1.6		
n	72	72		
no	0	0		

2013.1 C



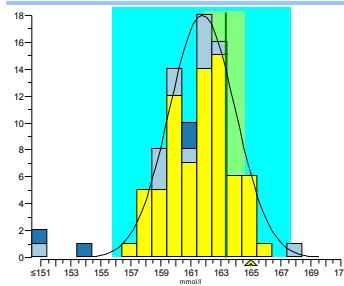
mean	140.7	140.7	142.1	142
SD	1.5	1.5		
n	72	72		
no	1	1		

2013.1 D



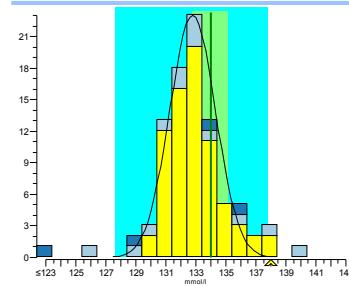
mean	144.3	144.3	145.5	147
SD	1.5	1.5		
n	72	72		
no	1	1		

2013.1 E



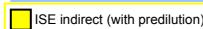
mean	161.8	161.8	163.3	165
SD	2.2	2.2		
n	72	72		
no	0	0		

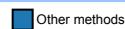
2013.1 F



mean	132.8	132.8	134.0	138
SD	1.5	1.5		
n	73	73		
no	2	2		

Legend

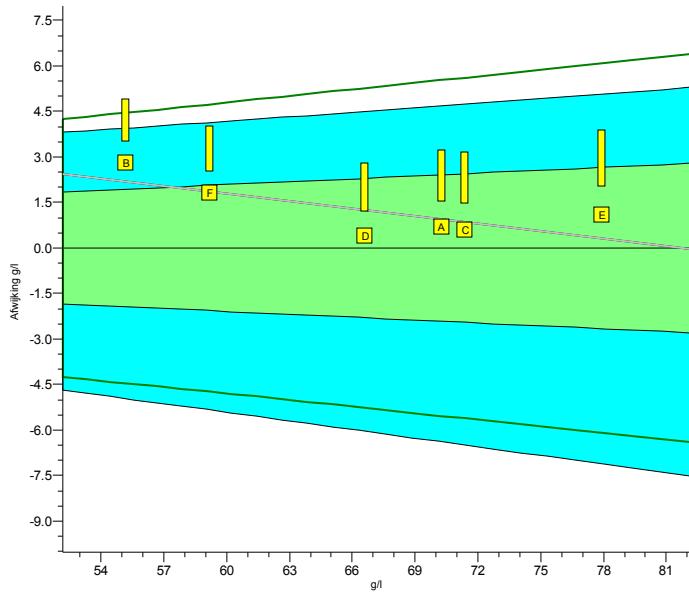




INPUTS 2013.1

Total Protein

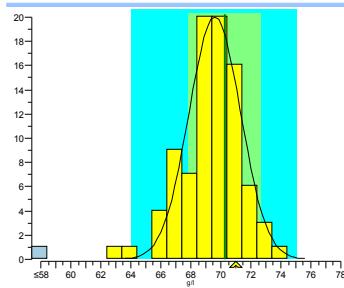
units: g/l



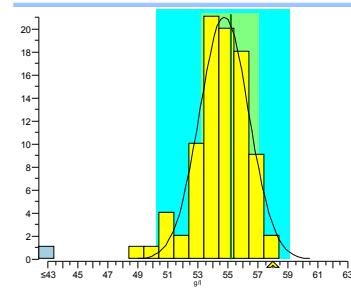
	2013.1	cumulative
Trueness	+1.8%	+1.8%
Precision	1.0%	1.0%
Number	6	6
Outliers	0	0
Sigma-TE	2.3	2.3
Sigma-SA	6.0 [2]	6.0 [2]
Score pictogram	[Green]	[Green]
Regression line	$6.7 + 0.918 \cdot x$	$6.7 + 0.918 \cdot x$

Consensus group: Biureet
Method: Biurete, automatic

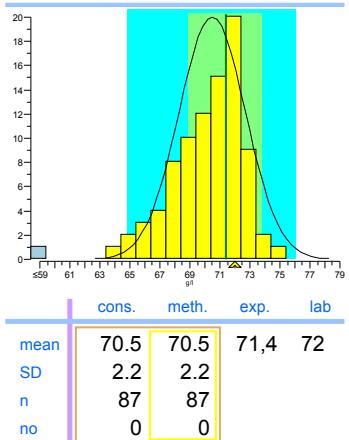
2013.1 A



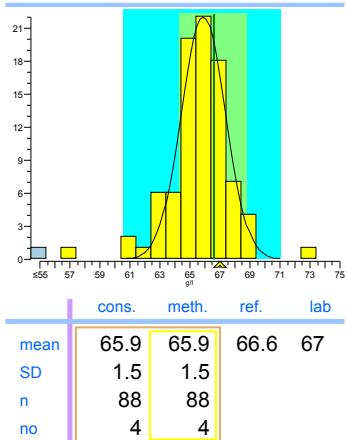
2013.1 B



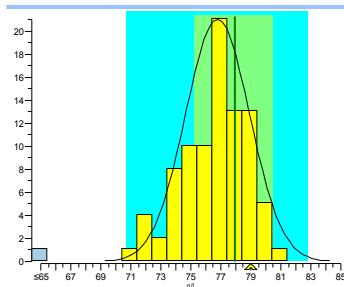
2013.1 C



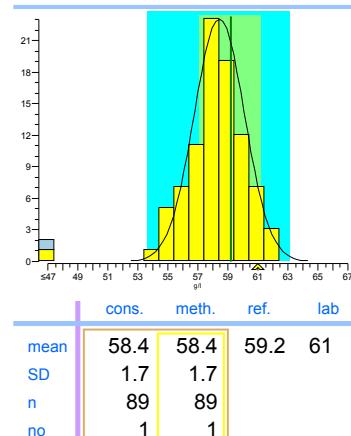
2013.1 D



2013.1 E



2013.1 F



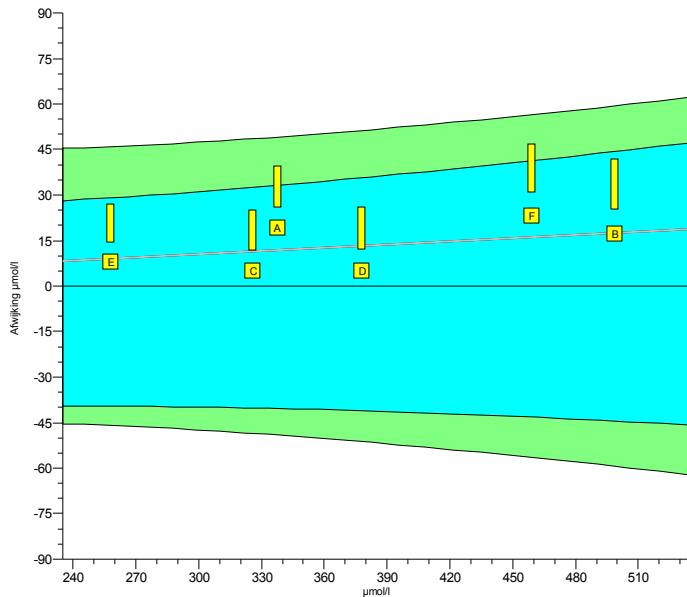
Legend

■ Biurete, automatic

■ Other methods

INPUTs 2013.1

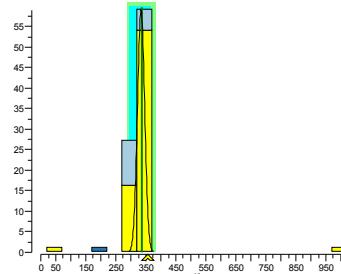
Urate

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

	2013.1	cumulative
Trueness	+3.4%	+3.4%
Precision	1.6%	1.6%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 [2]	6.0 [2]
Sigma-SA	5.4	5.4
Score pictogram	[Green box]	[Green box]
Regression line	$0 + 1.035.x$	$0 + 1.035.x$

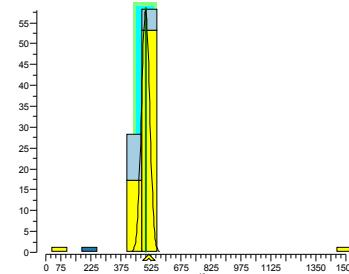
Consensus group: Colorimetrisch
Method: Uricase, colorim., automatic

2013.1 A



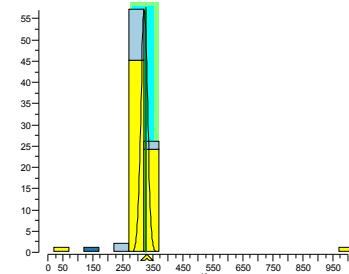
	cons.	meth.	ref.	lab
mean	334	334	0.338	357
SD	12	12		
n	88	72		
no	18	2		

2013.1 B



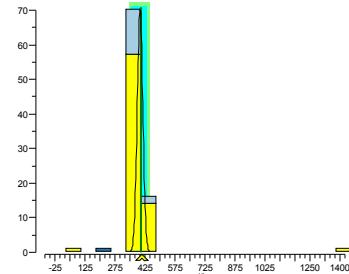
	cons.	meth.	ref.	lab
mean	499	499	0.499	516
SD	20	20		
n	88	72		
no	18	2		

2013.1 C



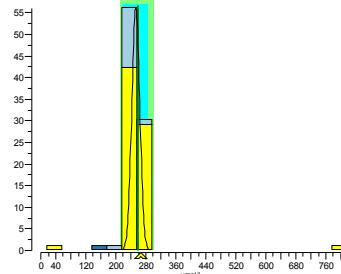
	cons.	meth.	exp.	lab
mean	321	321	0.326	331
SD	11	11		
n	87	71		
no	18	2		

2013.1 D



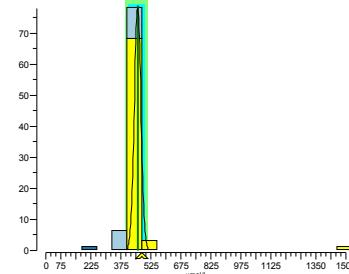
	cons.	meth.	ref.	lab
mean	376	376	0.378	383
SD	14	14		
n	88	73		
no	17	2		

2013.1 E



	cons.	meth.	ref.	lab
mean	253	253	0.258	266
SD	11	11		
n	89	73		
no	18	2		

2013.1 F



	cons.	meth.	ref.	lab
mean	459	459	0.459	482
SD	17	17		
n	88	72		
no	17	1		

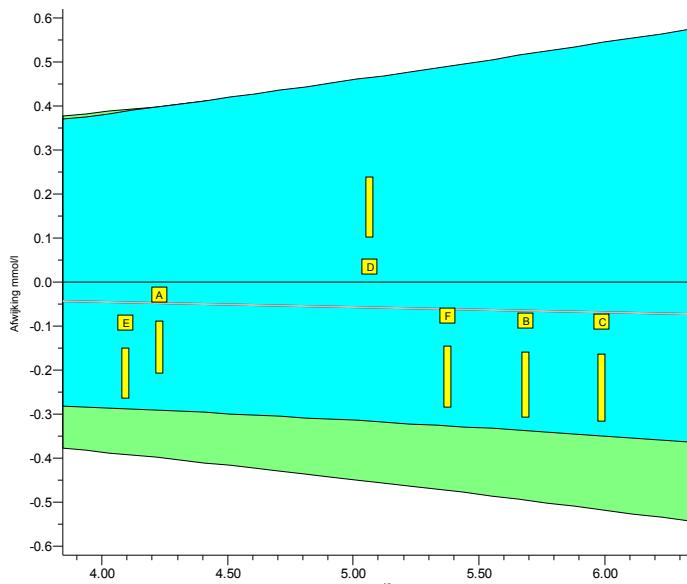
Legend

[Yellow box] Uricase, colorim., automatic [Light blue box] Uricase, differential UV, automatic [Dark blue box] Oudege methoden

INPUTs 2013.1

Cholesterol

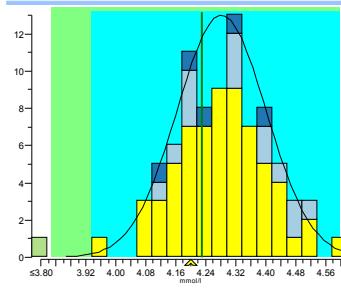
units: mmol/l



	2013.1	cumulative
Trueness	-1.1%	-1.1%
Precision	1.1%	1.1%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 [2]	6.0 [2]
Sigma-SA	6.0	6.0
Score pictogram	[Score pictogram]	[Score pictogram]
Regression line	$0.00 + 0.989 \cdot x$	$0.00 + 0.989 \cdot x$

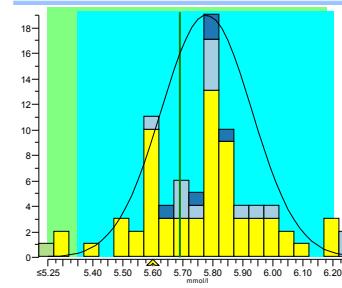
Consensus group: Enzymatisch
Method: Enzymatic, automatic, discrete

2013.1 A



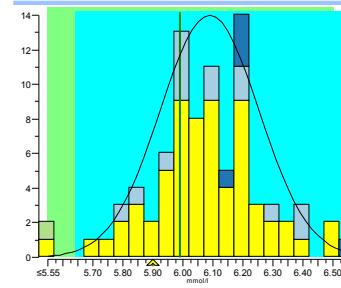
	cons.	meth.	ref.	lab
mean	4.28	4.28	4.230	4.2
SD	0.12	0.12		
n	83	64		
no	0	0		

2013.1 B



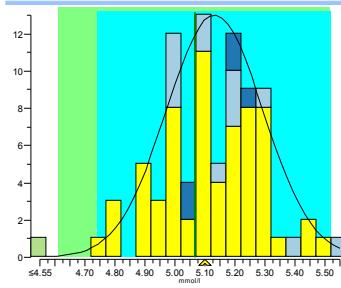
	cons.	meth.	ref.	lab
mean	5.78	5.78	5.689	5.6
SD	0.15	0.18		
n	83	64		
no	4	2		

2013.1 C



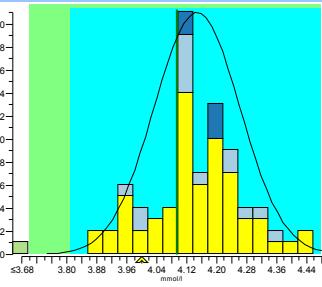
	cons.	meth.	exp.	lab
mean	6.09	6.08	5.99	5.9
SD	0.16	0.17		
n	83	64		
no	1	1		

2013.1 D



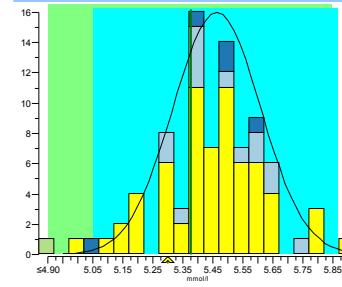
	cons.	meth.	ref.	lab
mean	5.13	5.12	5.066	5.1
SD	0.16	0.17		
n	82	64		
no	0	0		

2013.1 E



	cons.	meth.	ref.	lab
mean	4.15	4.14	4.094	4
SD	0.11	0.13		
n	84	65		
no	0	0		

2013.1 F



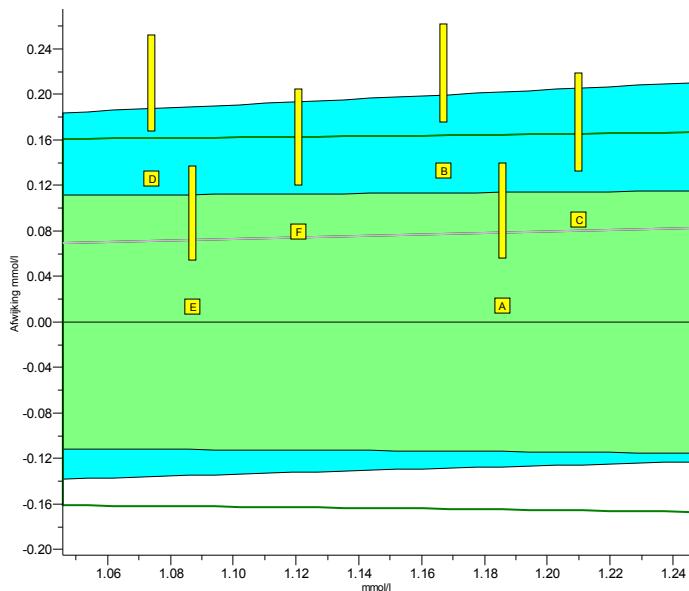
	cons.	meth.	ref.	lab
mean	5.46	5.45	5.377	5.3
SD	0.15	0.16		
n	84	65		
no	2	2		

Legend

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

INPUTs 2013.1**HDL-Cholesterol**

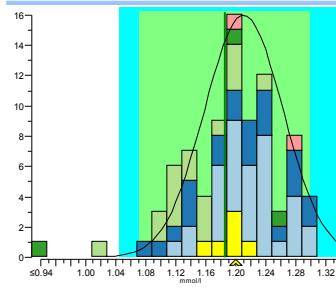
units: mmol/l



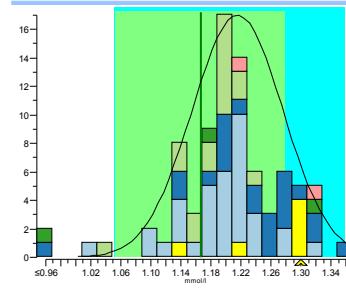
	2013.1	cumulative
Trueness	+6.6%	+6.6%
Precision	3.7%	3.7%
Number	6	6
Outliers	0	0
Sigma-TE	1.9	1.9
Sigma-SA	3.4	3.4
Score pictogram		
Regression line	$0.00 + 1.066 \cdot x$	$0.00 + 1.066 \cdot x$

Consensus group Direct
 Method Catalase method (Denka Seiken)

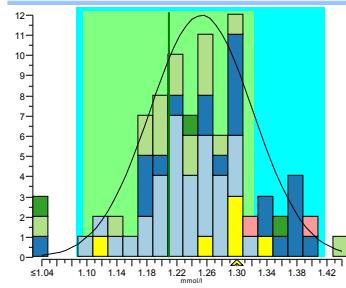
2013.1 A



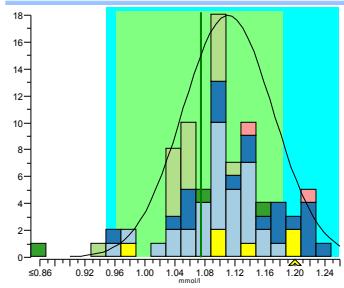
2013.1 B



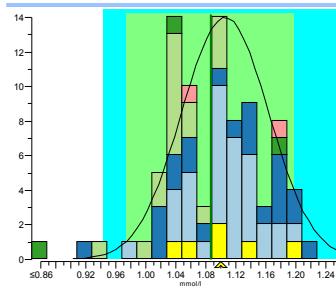
2013.1 C



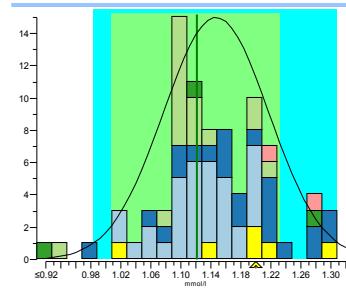
2013.1 D



2013.1 E



2013.1 F

**Legend** Catalase method (Denka Seiken) PEG modified enzyme, PEGME (Kyowa Medex) Accelerator Selective Detergent ("Ultra HDL") Immunoinhibition Overige methoden Precipitation Technique