



3714 dr. M. Bosomworth
 Leeds general Infirmary
 Leeds

Groot Brittannië

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	29	38	



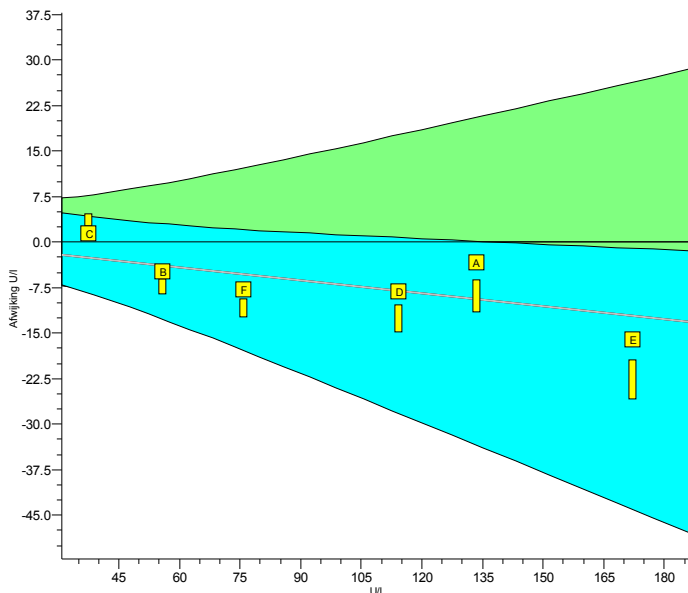
INPUTS 2013.1



Analyte		Trueness				Precision		Performance			
		your mean	ref.	cons.	SDtI	your prec.	SDbl	this survey	PS	cumulative PSc	
ALAT	U/l	91.7	98.2	86.9	10.1	1.9	2.7		2		2
Alk. Phosphatase	U/l	433	188	183	12	4	4		0		0
Amylase	U/l	191	207	201	8	1	2		2		2
ASAT	U/l	73.7	78.6	71.5	6.8	1.1	1.6		2		2
Calcium	mmol/l	2.42	2.46	2.47	0.07	0.02	0.02		2		2
Chloride	mmol/l	103.5	100.0	100.8	1.8	1.6	0.9		0		0
CK	U/l	247	244	253	18	2	3		2		2
Creatinine	µmol/l	145.3	149.1	150.7	5.6	0.8	2.1		2		2
eGFR (F, 55, white)	ml/min/1,73m ²	39.3	37.6	37.4	2.6	0.8	0.6		2		2
Gamma-GT	U/l	90.7	87.7	86.3	8.6	1.3	1.2		2		2
Glucose	mmol/l	14.47	13.93	14.19	0.44	0.18	0.18		1		1
LD	U/l	1069	532	895	321	18	16		0		0
Magnesium	mmol/l	1.22	1.22	1.23	0.04	0.01	0.02		2		2
Potassium	mmol/l	5.50	5.39	5.30	0.10	0.08	0.04		1		1
Sodium	mmol/l	144.5	144.0	142.7	1.7	1.1	0.9		2		2
Total Protein	g/l	65.8	66.8	66.0	1.8	0.7	0.7		2		2
Urate	µmol/l	374	376	374	15	2	4		2		2
Cholesterol	mmol/l	5.00	5.07	5.15	0.14	0.09	0.05		2		2
HDL-Cholesterol	mmol/l	1.25	1.14	1.17	0.06	0.06	0.03		1		1
Total :								29	29		

INPUTS 2013.1

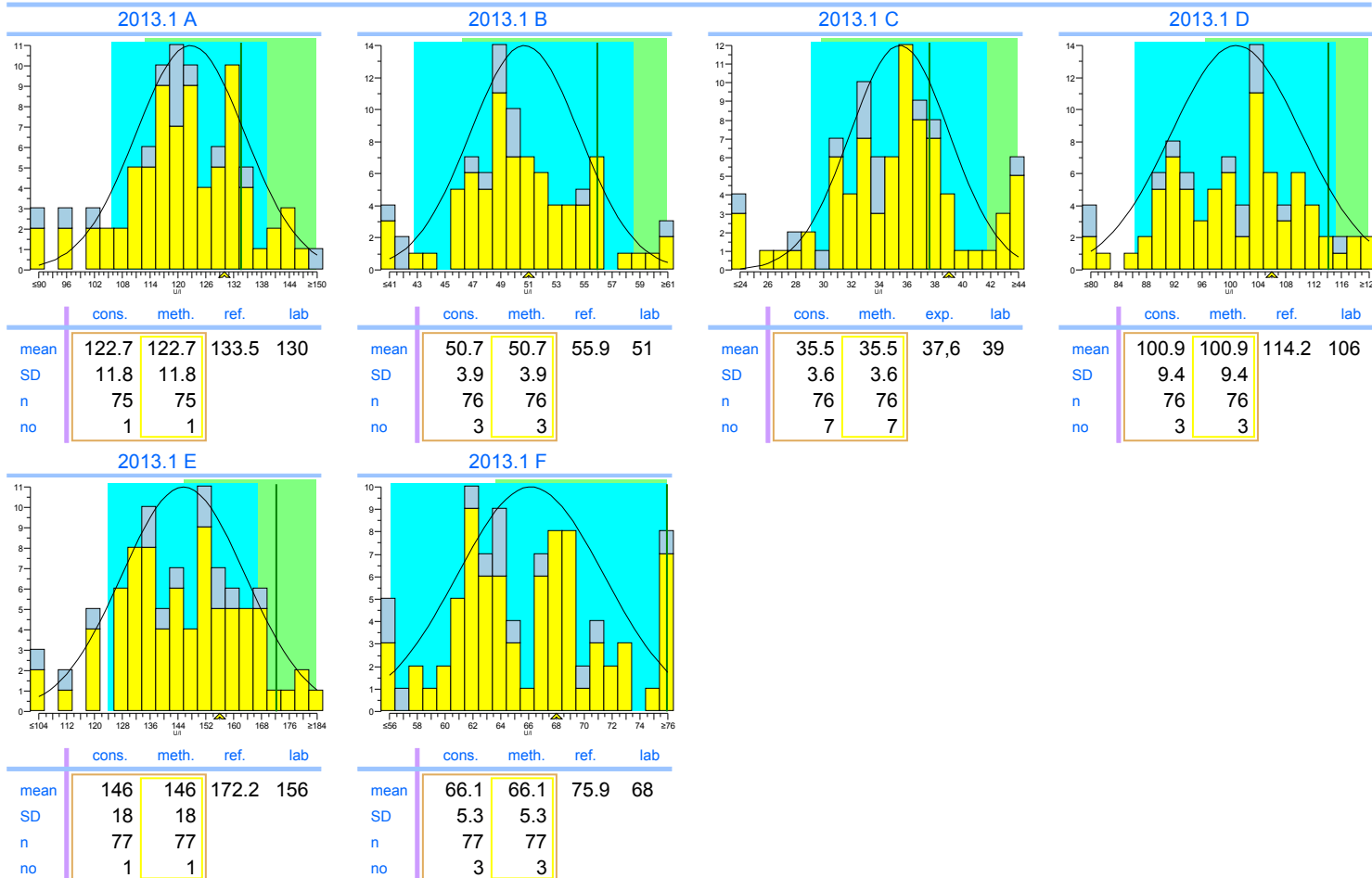
ALAT

units: U/I



	2013.1	cumulative
Trueness	-6.7%	-6.7%
Precision	2.2%	2.2%
Number	6	6
Outliers	0	0
Sigma-TE	5.9 2	5.9 2
Sigma-SA	4.3	4.3
Score pictogram		
Regression line	$0.0 + 0.929 \cdot x$	$0.0 + 0.929 \cdot x$

Consensus group IFCC traceerbaar
Method IFCC traceable



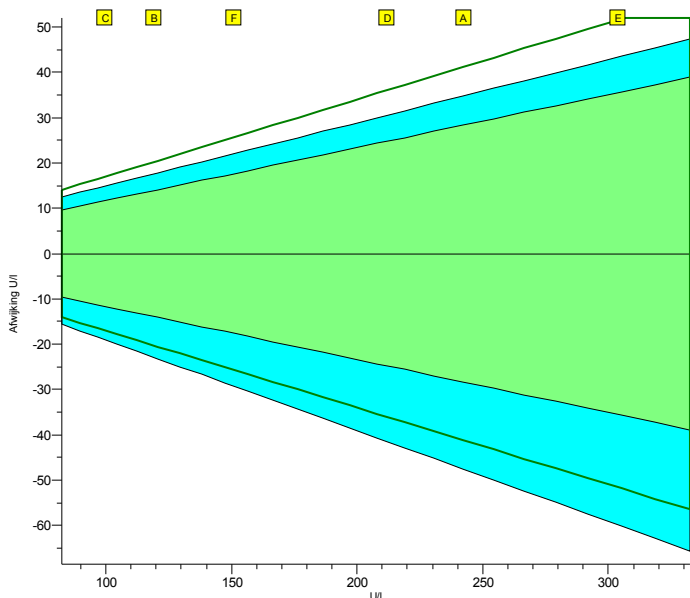
Legend

IFCC traceable IFCC non-traceable

INPUTS 2013.1

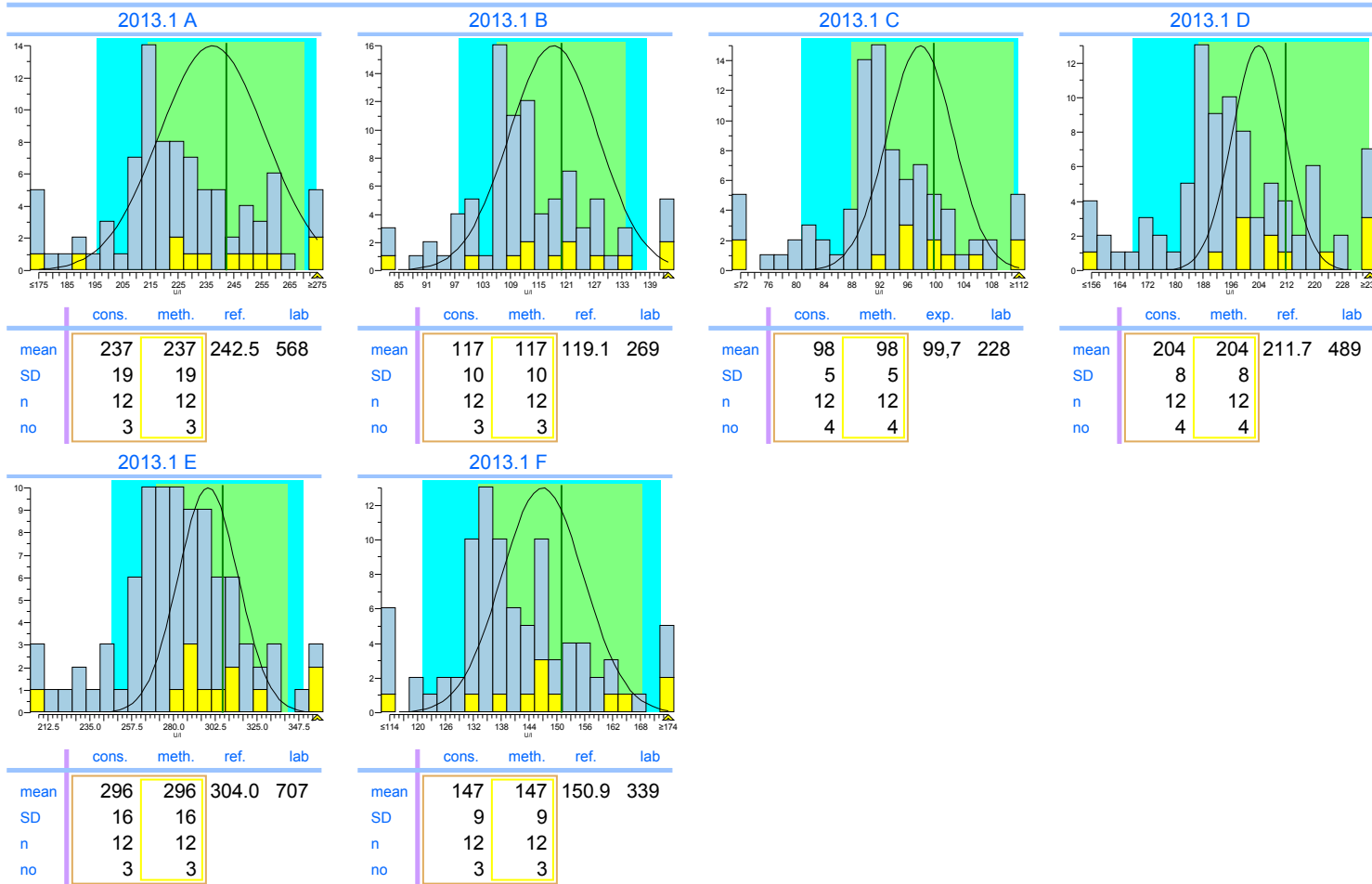
Alk. Phosphatase

units: U/l



	2013.1	cumulative
Trueness	+131%	+131%
Precision	2.4%	2.4%
Number	6	6
Outliers	0	0
Sigma-TE	-3.0	-3.0
Sigma-SA	-3.0 0	-3.0 0
Score pictogram	■ ■ ●	■ ■ ●
Regression line	$-13 + 2.373.x$	$-13 + 2.373.x$

Consensus group niet IFCC traceerbaar
Method IFCC non-traceable



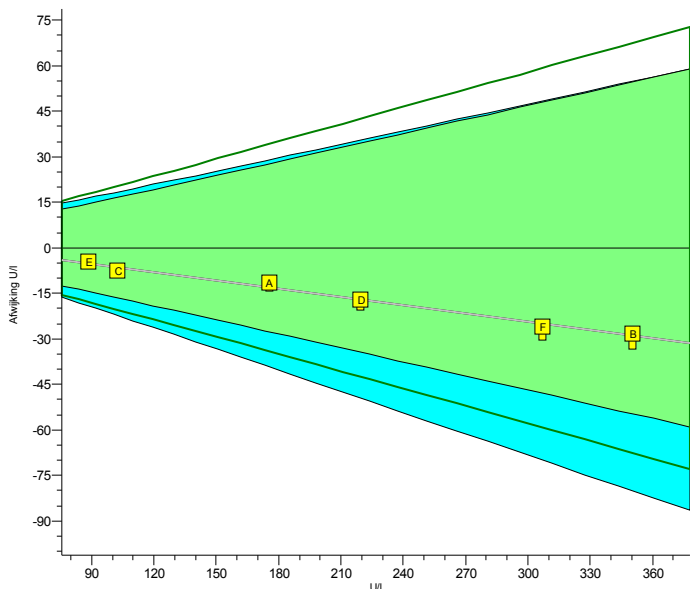
Legend

IFCC non-traceable IFCC traceable

INPUTS 2013.1

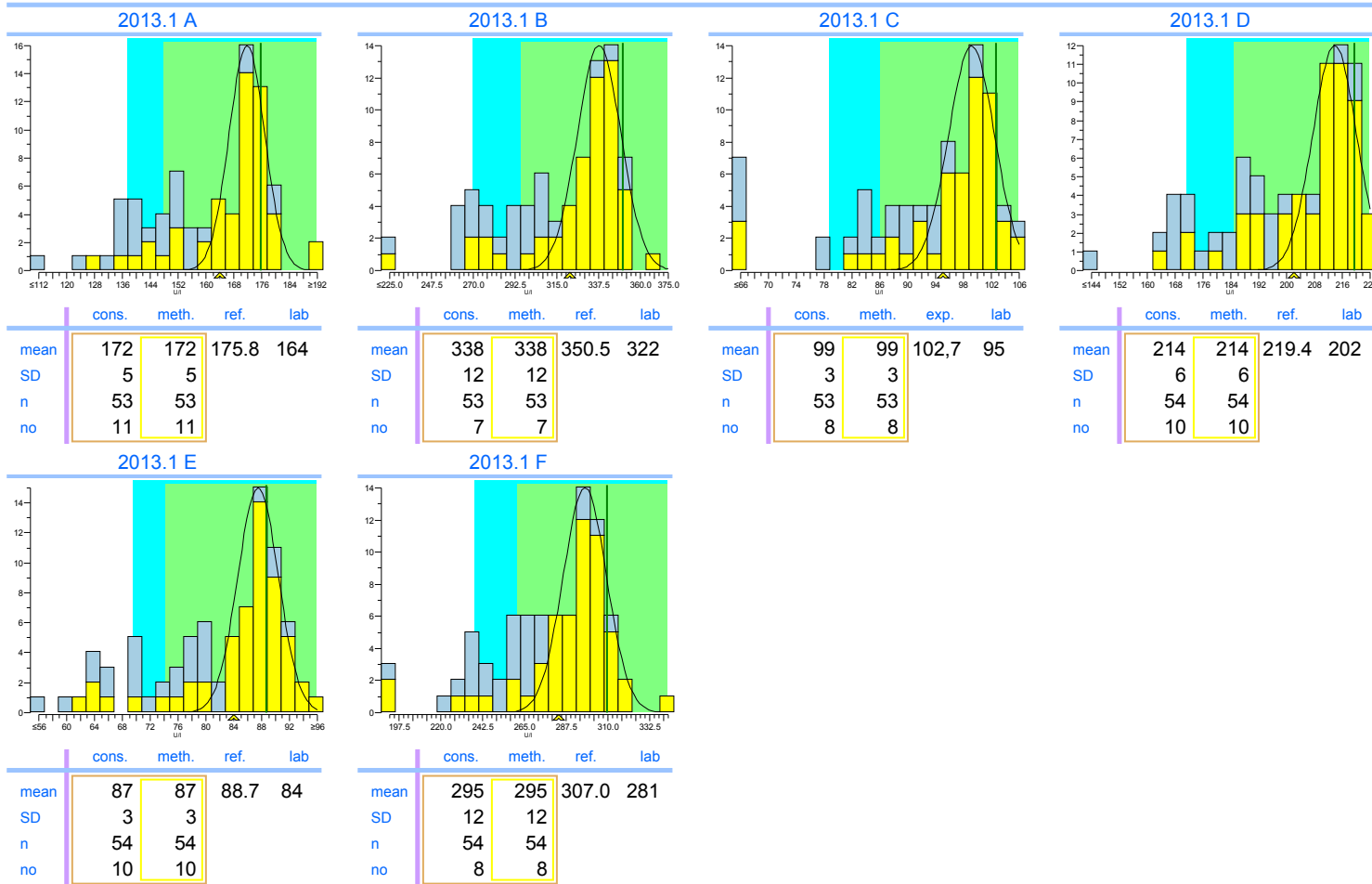
Amylase

units: U/l



	2013.1	cumulative
Trueness	-7.7%	-7.7%
Precision	0.43%	0.43%
Number	6	6
Outliers	0	0
Sigma-TE	6.0	6.0
Sigma-SA	6.0 2	6.0 2
Score pictogram		
Regression line	$3 + 0.909.x$	$3 + 0.909.x$

Consensus group IFCC traceerbaar
Method IFCC traceable



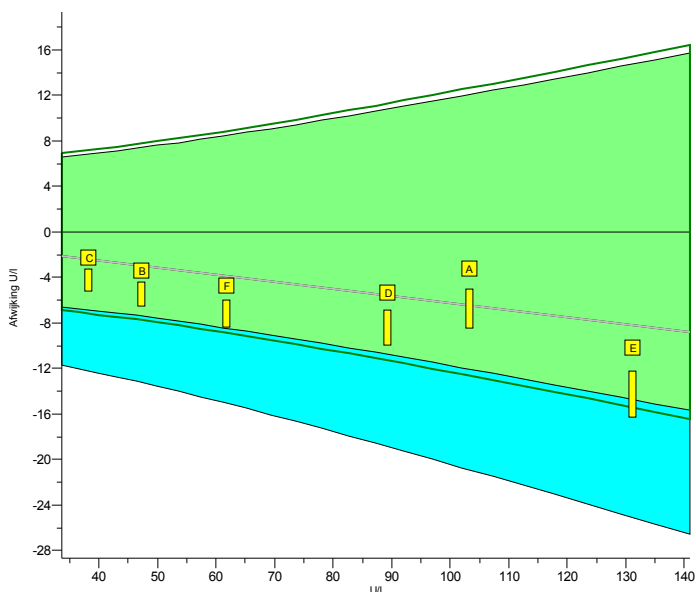
Legend



 IFCC traceable IFCC non-traceable

INPUTS 2013.1

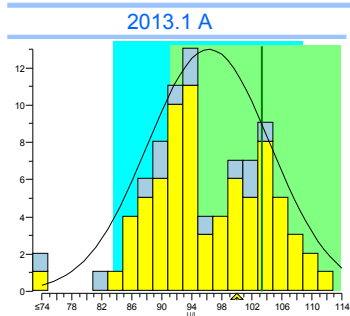
ASAT

units: U/l

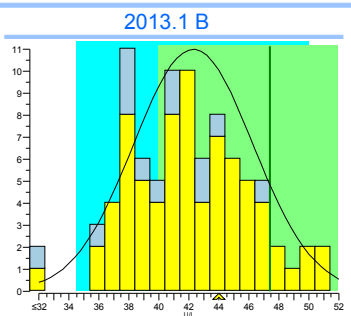


	2013.1	cumulative
Trueness	-6.2%	-6.2%
Precision	1.6%	1.6%
Number	6	6
Outliers	0	0
Sigma-TE	5.2	5.2
Sigma-SA	5.6 2	5.6 2
Score pictogram		
Regression line	<u>0.0 + 0.938.x</u>	<u>0.0 + 0.938.x</u>

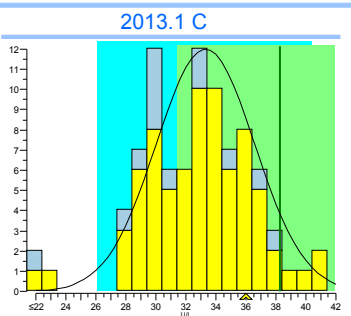
Consensus group IFCC traceerbaar
Method IFCC traceable



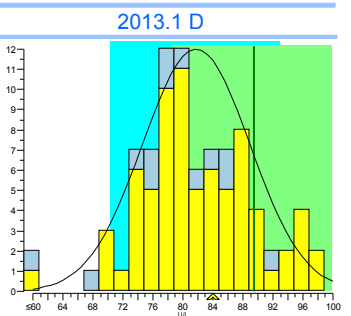
	cons.	meth.	ref.	lab
mean	96.3	96.3	103.3	100
SD	8.2	8.2		
n	75	75		
no	1	1		



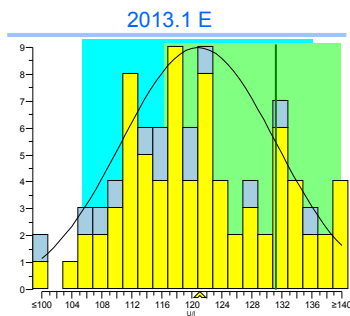
	cons.	meth.	ref.	lab
mean	42.3	42.3	47.4	44
SD	4.0	4.0		
n	75	75		
no	1	1		



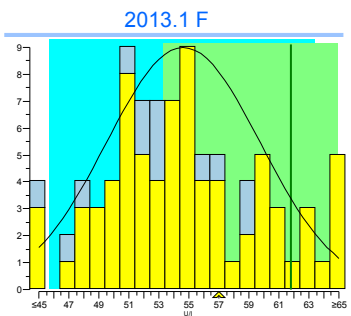
	cons.	meth.	exp.	lab
mean	33.3	33.3	38,3	36
SD	3.3	3.3		
n	75	75		
no	2	2		



	cons.	meth.	ref.	lab
mean	81.8	81.8	89.4	84
SD	7.2	7.2		
n	74	74		
no	1	1		



	cons.	meth.	ref.	lab
mean	120.9	120.9	131.2	121
SD	10.3	10.3		
n	76	76		
no	1	1		



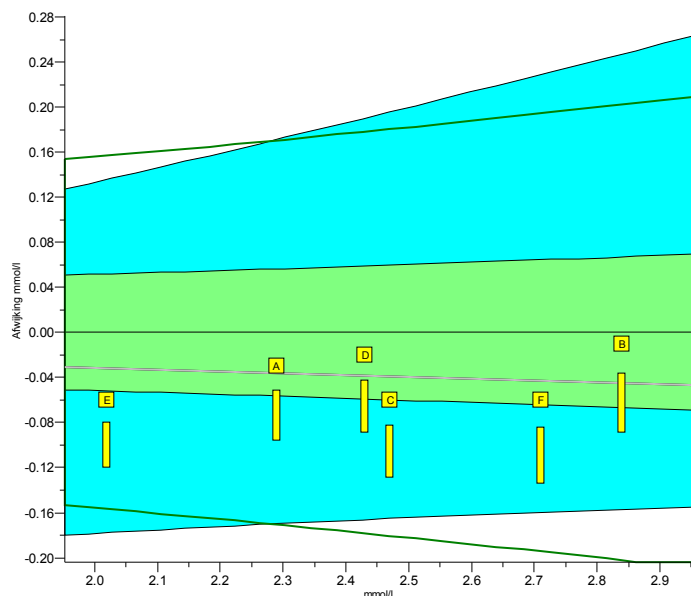
	cons.	meth.	ref.	lab
mean	54.6	54.6	61.8	57
SD	5.1	5.1		
n	76	76		
no	2	2		



Legend
 IFCC traceable
 IFCC non-traceable

INPUTS 2013.1

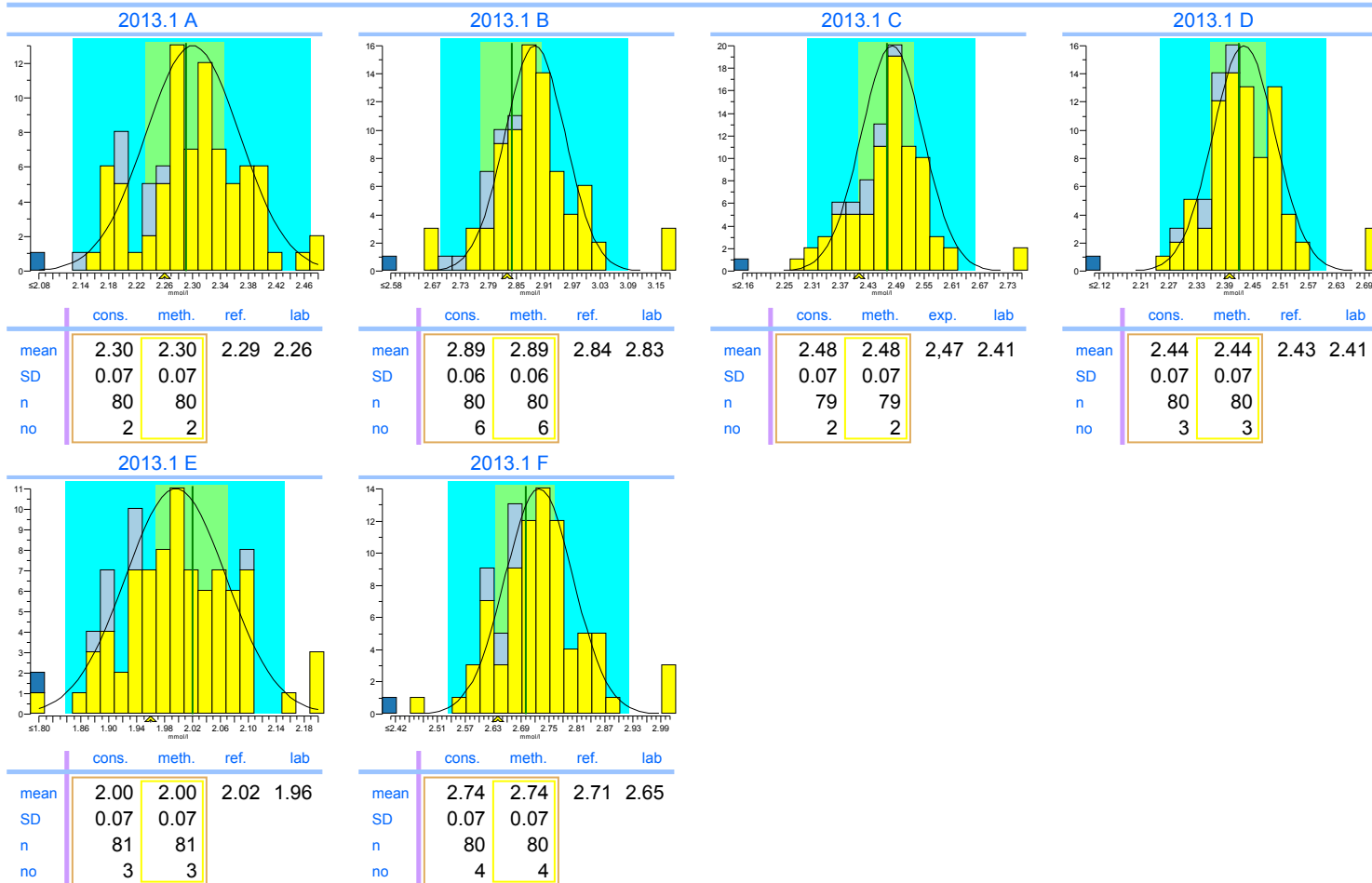
Calcium

units: mmol/l



	2013.1	cumulative
Trueness	-1.6%	-1.6%
Precision	0.76%	0.76%
Number	6	6
Outliers	0	0
Sigma-TE	1.8	1.8
Sigma-SA	6.0 2	6.0 2
Score pictogram		
Regression line	<u>0.00 + 0.984.x</u>	<u>0.00 + 0.984.x</u>

Consensus group	Colorimetrisch
Method	Colorimetric, automatic, discrete



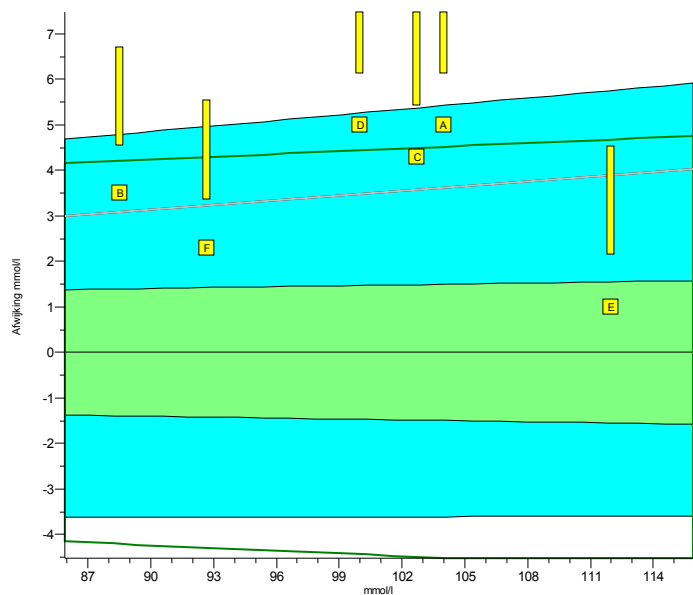
Legend

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

INPUTS 2013.1

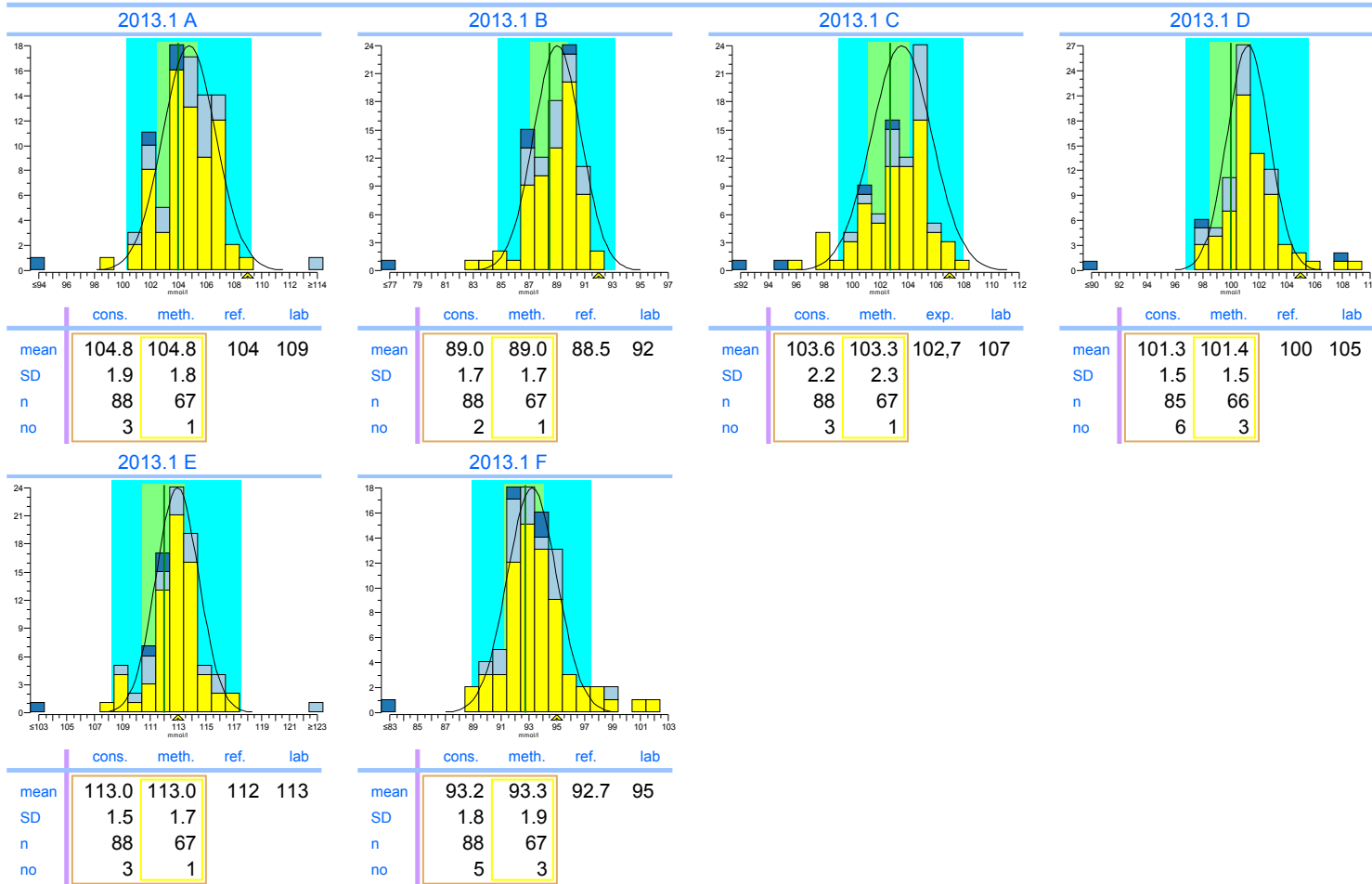
Chloride

units: mmol/l



	2013.1	cumulative
Trueness	+3.5%	+3.5%
Precision	1.6%	1.6%
Number	6	6
Outliers	0	0
Sigma-TE	-1.5	-1.5
Sigma-SA	1.8 0	1.8 0
Score pictogram		
Regression line	$0.0 + 1.035 \cdot x$	$0.0 + 1.035 \cdot x$

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



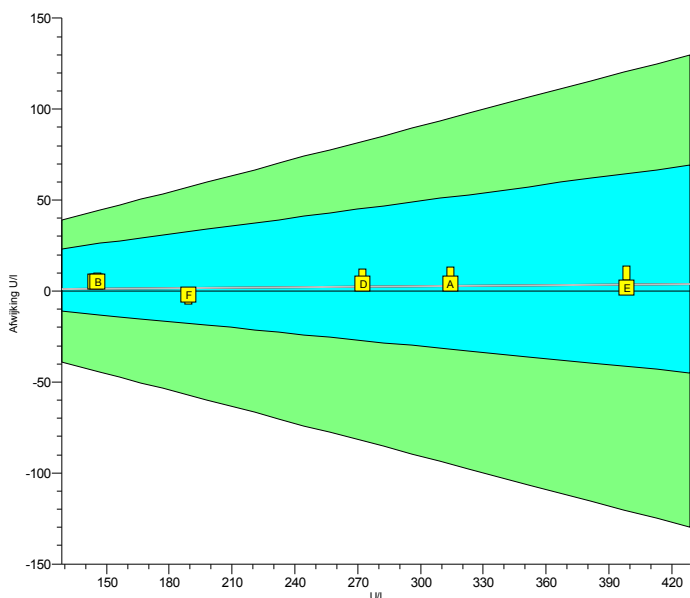
Legend





INPUTS 2013.1

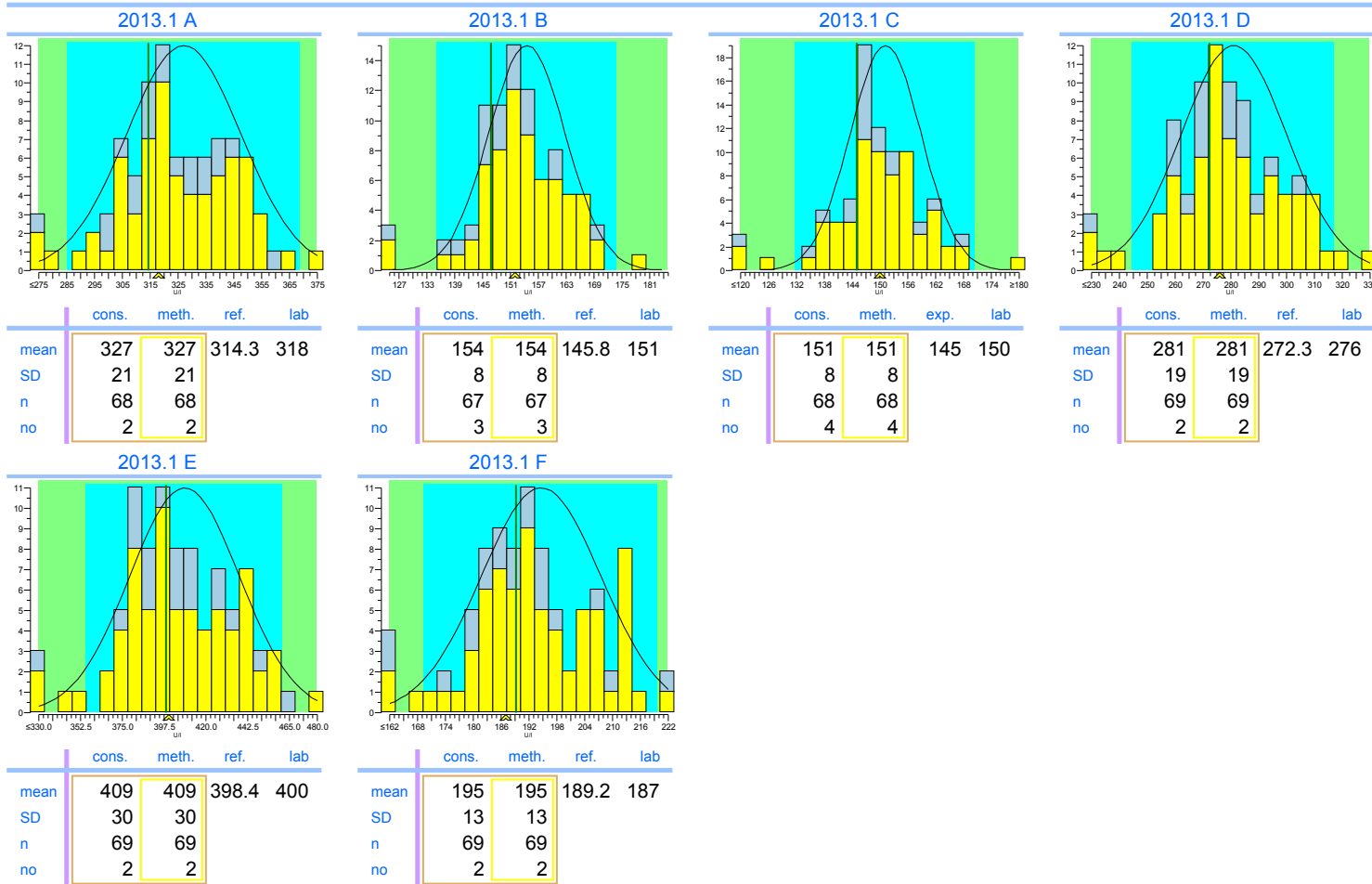
CK

units: U/l



	2013.1	cumulative
Trueness	+1.2%	+1.2%
Precision	0.81%	0.81%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 2	6.0 2
Sigma-SA	6.0	6.0
Score pictogram		
Regression line	$0 + 1.009 \cdot x$	$0 + 1.009 \cdot x$

Consensus group IFCC traceerbaar
Method IFCC traceable



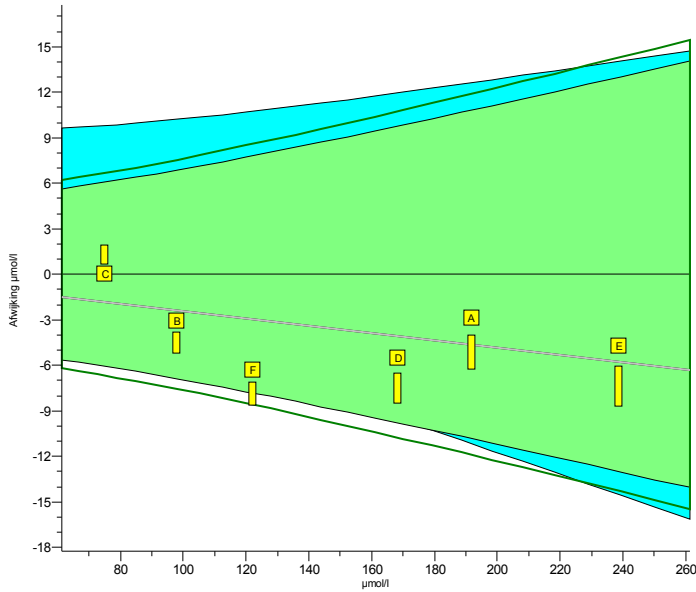
Legend

IFCC traceable IFCC non-traceable

INPUTS 2013.1

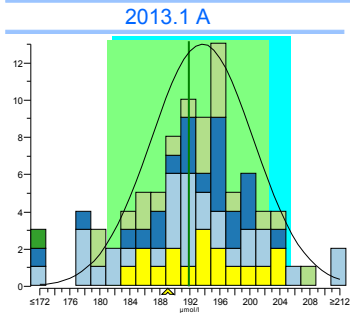
Creatinine

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

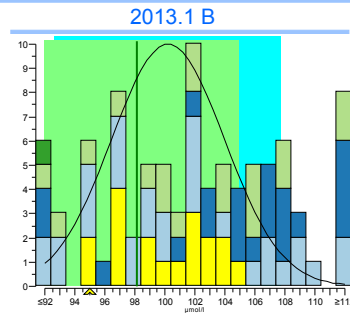


	2013.1	cumulative
Trueness	-2.5%	-2.5%
Precision	0.50%	0.50%
Number	6	6
Outliers	0	0
Sigma-TE	6.0	6.0
Sigma-SA	6.0 2	6.0 2
Score pictogram		
Regression line	$0.0 + 0.976.x$	$0.0 + 0.976.x$

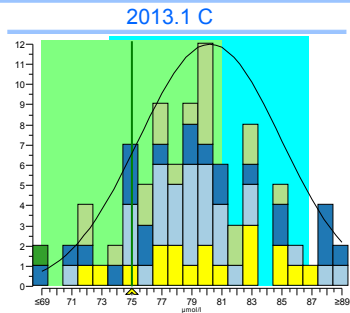
Consensus group: Enzymatische kreatinine
Method: Enzymatic, automatic



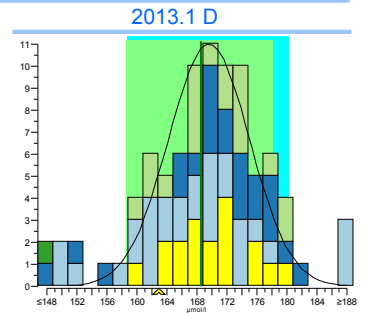
	cons.	meth.	ref.	lab
mean	193.7	193.7	191.9	189
SD	6.8	6.8		
n	17	17		
no	0	0		



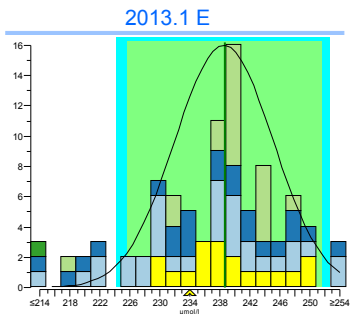
	cons.	meth.	ref.	lab
mean	100.2	100.2	98.11	95
SD	3.8	3.8		
n	18	18		
no	0	0		



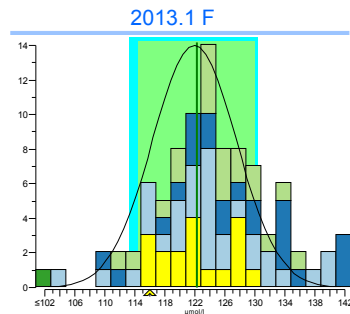
	cons.	meth.	exp.	lab
mean	80.2	80.2	75	75
SD	4.7	4.7		
n	18	18		
no	0	0		



	cons.	meth.	ref.	lab
mean	169.6	169.6	168.5	163
SD	5.4	5.4		
n	18	18		
no	0	0		



	cons.	meth.	ref.	lab
mean	238.5	238.5	238.7	234
SD	6.6	6.6		
n	18	18		
no	0	0		



	cons.	meth.	ref.	lab
mean	121.9	121.9	122.3	116
SD	5.6	5.6		
n	18	18		
no	0	0		

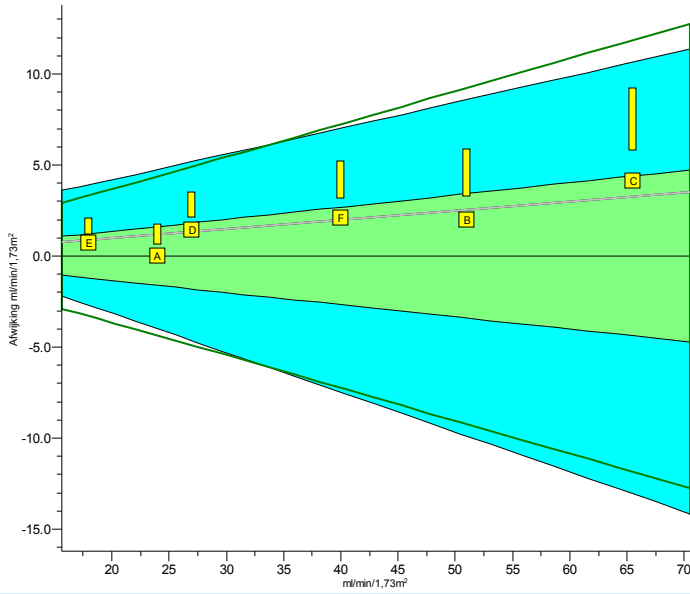
Legend



- Enzymatic, automatic
- Alk. Picrate, kinetic with compensation
- Alk. Picrate, kinetic
- Alk. Picrate, endpoint
- Other methods

INPUTs 2013.1

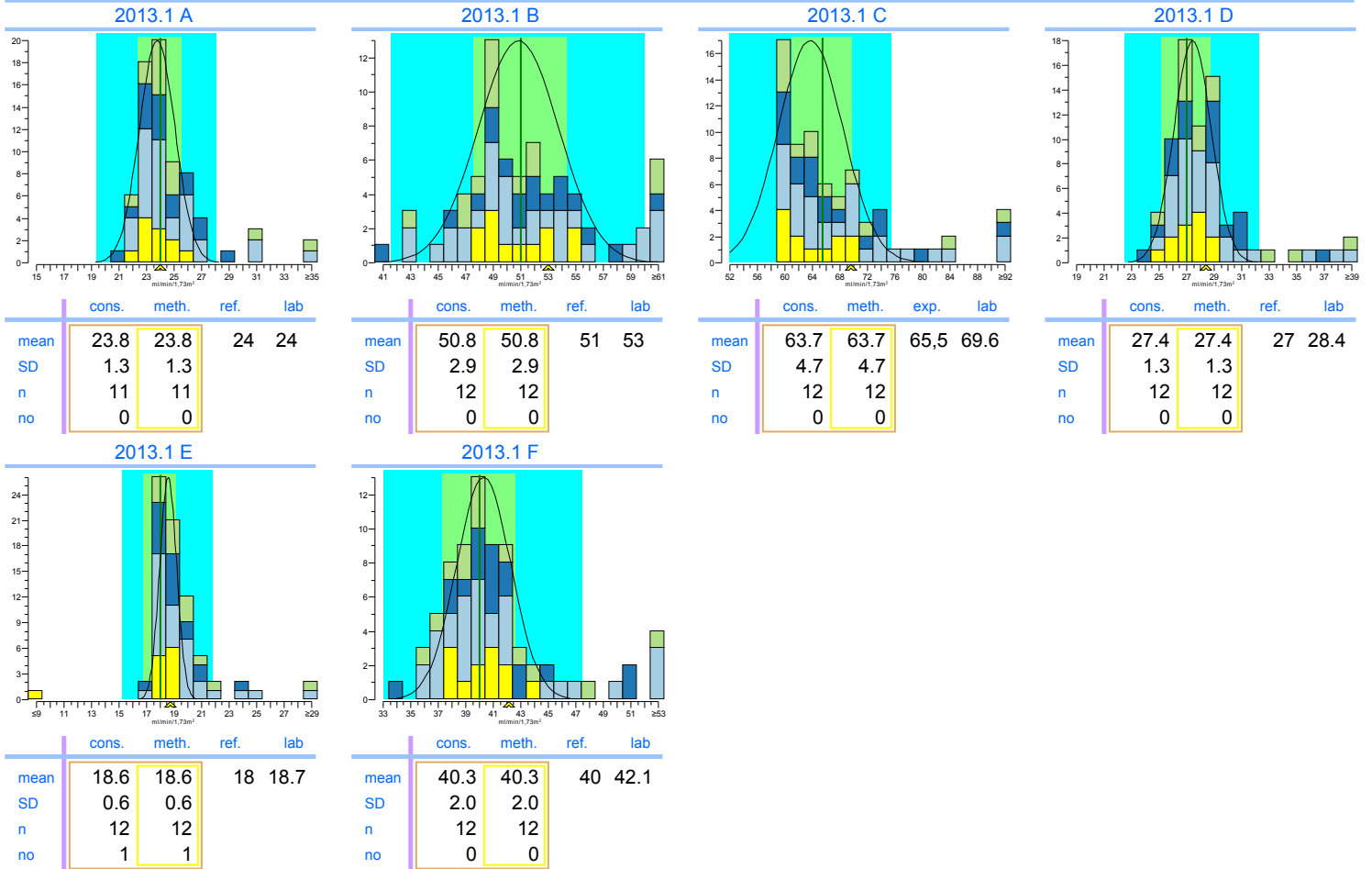
eGFR (F, 55, white)

units: ml/min/1,73m²



	2013.1	cumulative
Trueness	+4.6%	+4.6%
Precision	2.1%	2.1%
Number	6	6
Outliers	0	0
Sigma-TE	2.1	2.1
Sigma-SA	6.0 2	6.0 2
Score pictogram		
Regression line	<u>0.0 + 1.050.x</u>	<u>0.0 + 1.050.x</u>

Consensus group Enzymatische kreatinine
Method Enzymatic, automatic



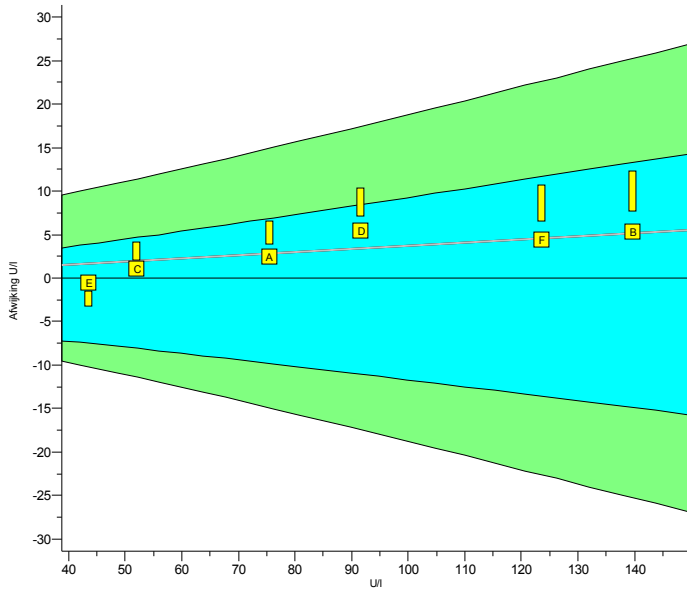
Legend

- Enzymatic, automatic
- Other methods
- Alk. Picrate, kinetic with compensation
- Alk. Picrate, kinetic

INPUTS 2013.1

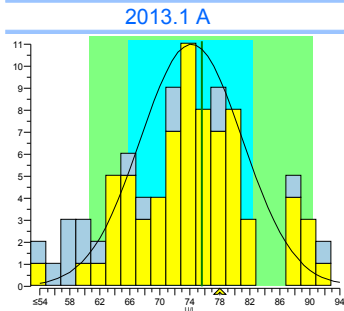
Gamma-GT

units: U/I

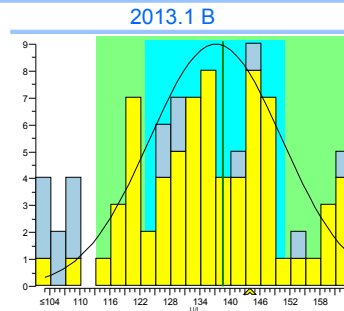


	2013.1	cumulative
Trueness	+3.4%	+3.4%
Precision	1.5%	1.5%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 2	6.0 2
Sigma-SA	6.0	6.0
Score pictogram		
Regression line	$0.0 + 1.037.x$	$0.0 + 1.037.x$

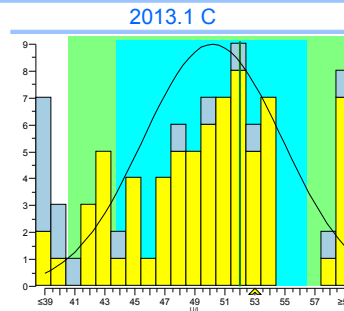
Consensus group IFCC traceerbaar
Method IFCC traceable



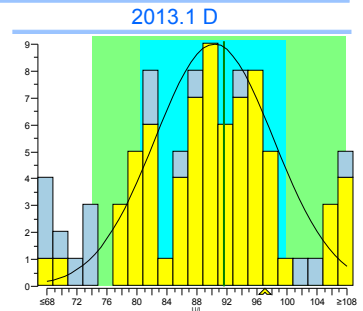
	cons.	meth.	ref.	lab
mean	74.2	74.2	75.6	78
SD	6.8	6.8		
n	72	72		
no	1	1		



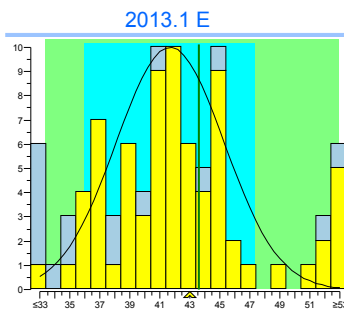
	cons.	meth.	ref.	lab
mean	138.2	138.2	139.7	145
SD	13.3	13.3		
n	72	72		
no	1	1		



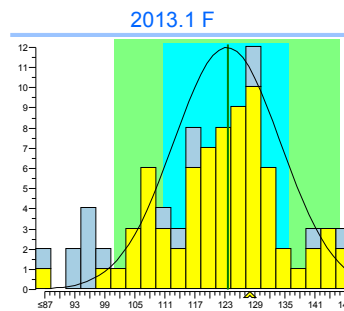
	cons.	meth.	exp.	lab
mean	50.2	50.2	52	53
SD	4.6	4.6		
n	72	72		
no	2	2		



	cons.	meth.	ref.	lab
mean	90.3	90.3	91.6	97
SD	7.9	7.9		
n	71	71		
no	1	1		



	cons.	meth.	ref.	lab
mean	41.7	41.7	43.6	43
SD	3.6	3.6		
n	73	73		
no	6	6		



	cons.	meth.	ref.	lab
mean	123.4	123.4	123.6	128
SD	10.7	10.7		
n	73	73		
no	1	1		

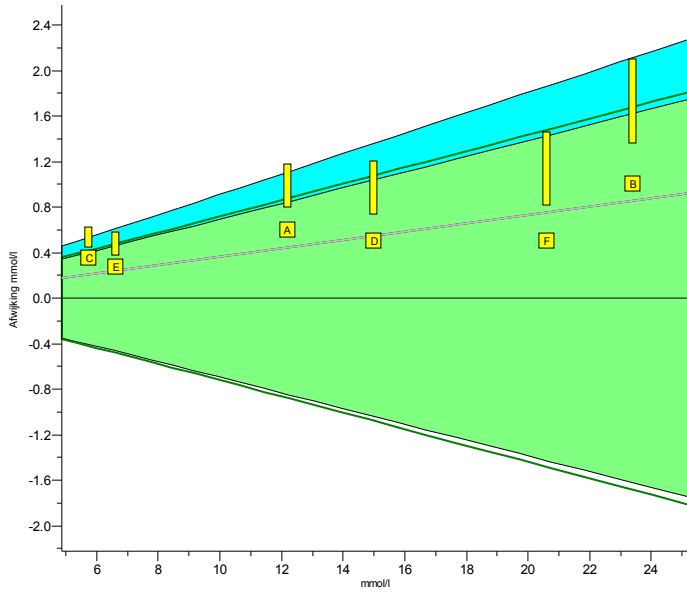
Legend



IFCC traceable IFCC non-traceable

INPUTS 2013.1

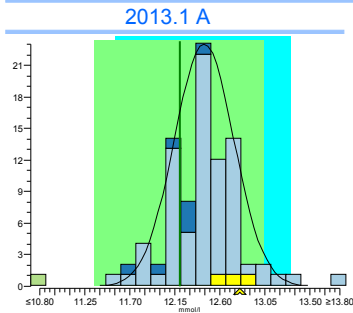
Glucose

units: mmol/l

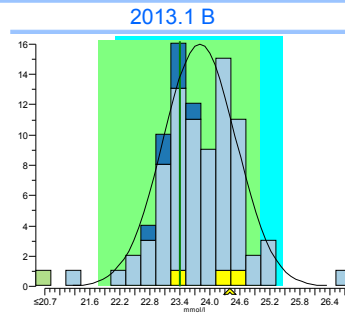


	2013.1	cumulative
Trueness	+3.9%	+3.9%
Precision	1.2%	1.2%
Number	6	6
Outliers	0	0
Sigma-TE	3.1	3.1
Sigma-SA	3.3	3.3
Score pictogram		
Regression line	$0.00 + 1.037 \cdot x$	$0.00 + 1.037 \cdot x$

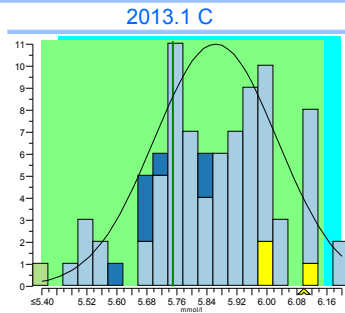
Consensus group: Natte chemie
Method: Glucose-oxidase/POD,automatic



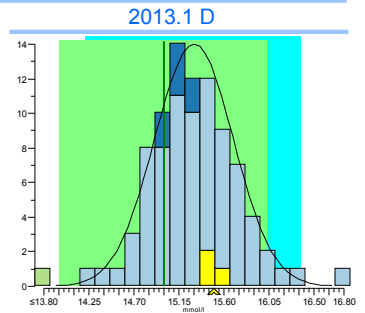
	cons.	meth.	ref.	lab
mean	12.44	12.76	12.2	12.8
SD	0.30			
n	88	3		
no	3	0		



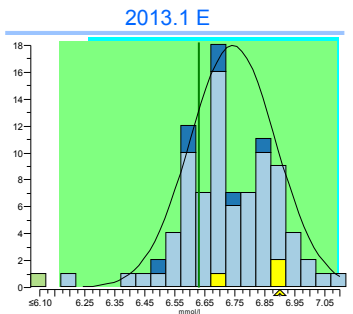
	cons.	meth.	ref.	lab
mean	23.8	24.1	23.4	24.4
SD	0.7			
n	88	3		
no	3	0		



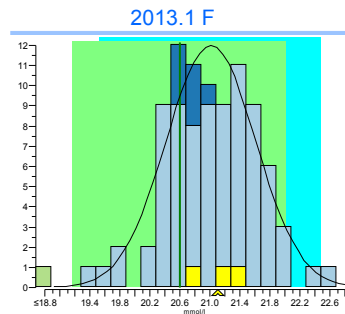
	cons.	meth.	exp.	lab
mean	5.86	6.03	5.75	6.1
SD	0.16			
n	88	3		
no	1	0		



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



	cons.	meth.	ref.	lab
mean	6.74	6.83	6.63	6.9
SD	0.14			
n	89	3		
no	2	0		



	cons.	meth.	ref.	lab
mean	21.01	21.04	20.6	21.1
SD	0.60			
n	89	3		
no	1	0		

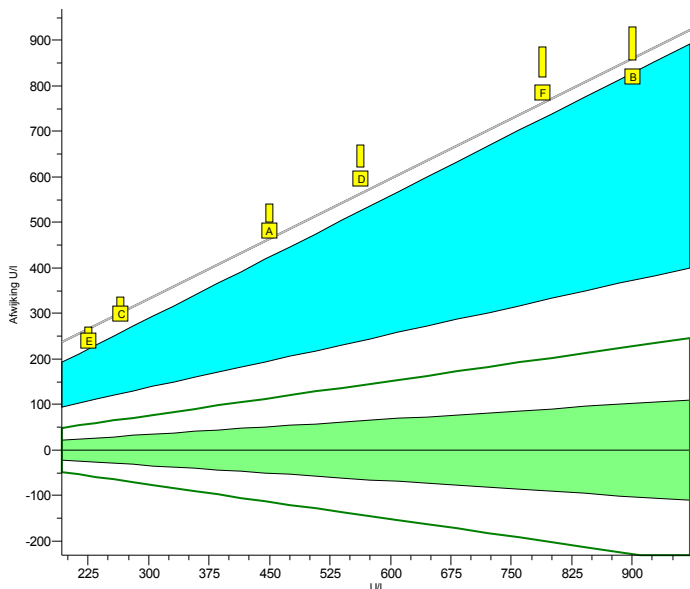
Legend

- Glucose-oxidase/POD,automatic
- Hexokinase, automatic
- Glucose-oxidase, amperometric, H2O2
- Other methods

INPUTs 2013.1

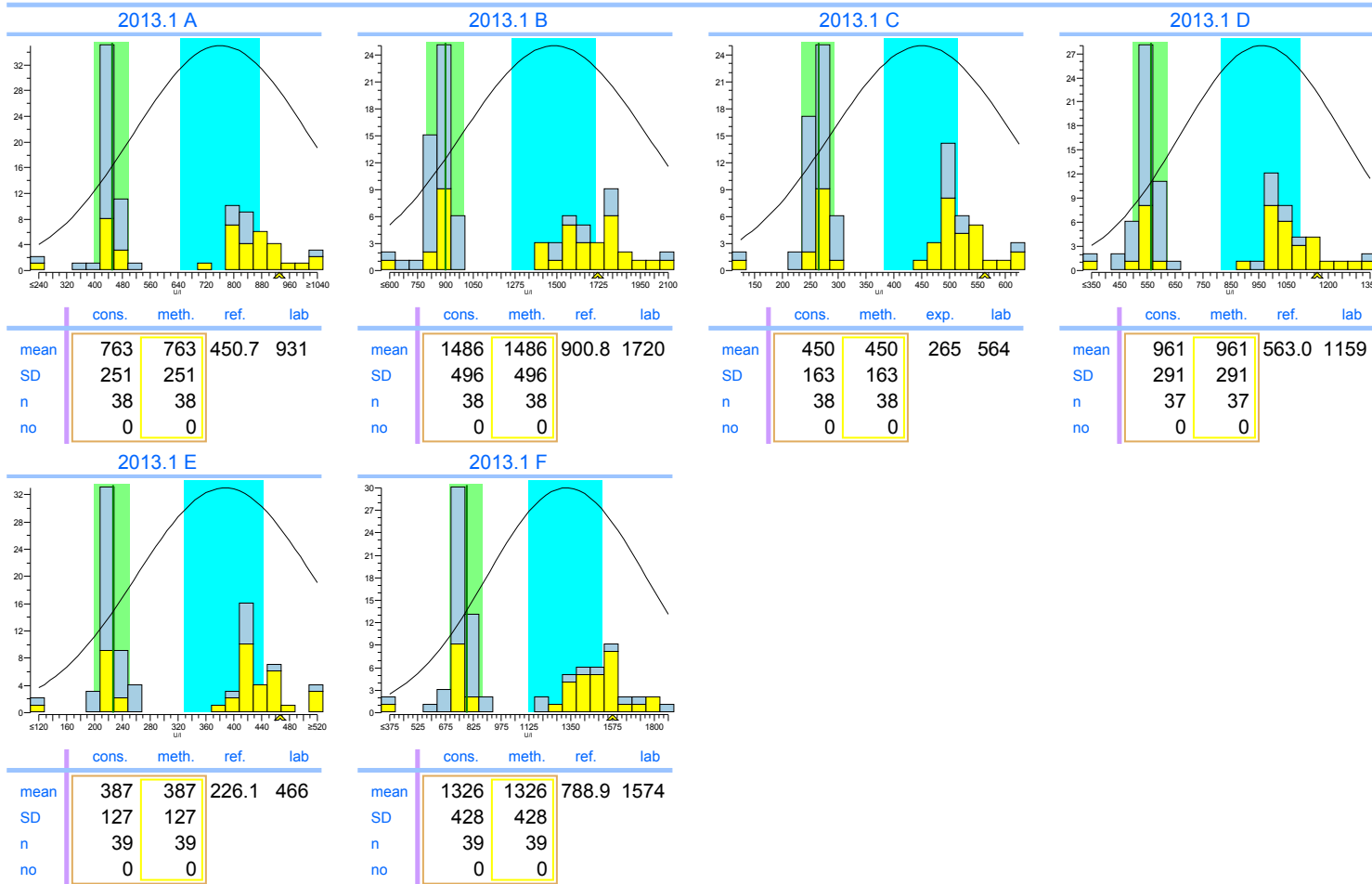
LD

units: U/I



	2013.1	cumulative
Trueness	+101%	+101%
Precision	2.1%	2.1%
Number	6	6
Outliers	0	0
Sigma-TE	-3.0	-3.0
Sigma-SA	-3.0 0	-3.0 0
Score pictogram		
Regression line	$69 + 1.878.x$	$69 + 1.878.x$

Consensus group niet IFCC traceerbaar
Method IFCC non-traceable



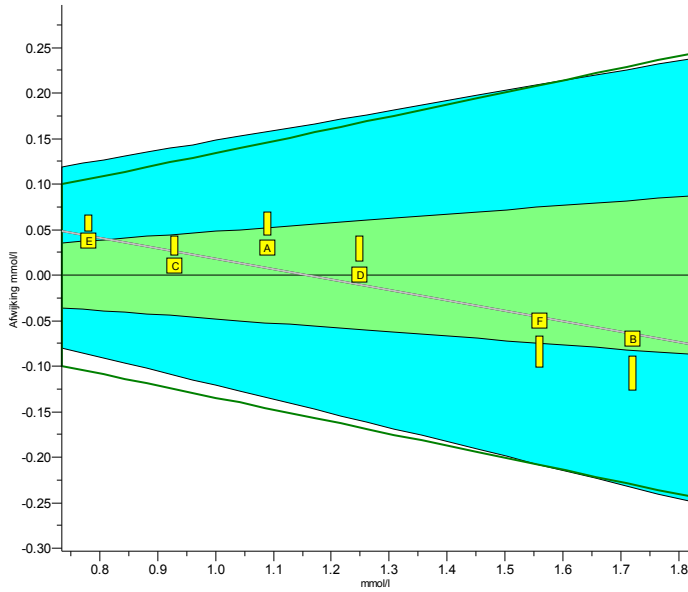
Legend

 IFCC non-traceable IFCC traceable

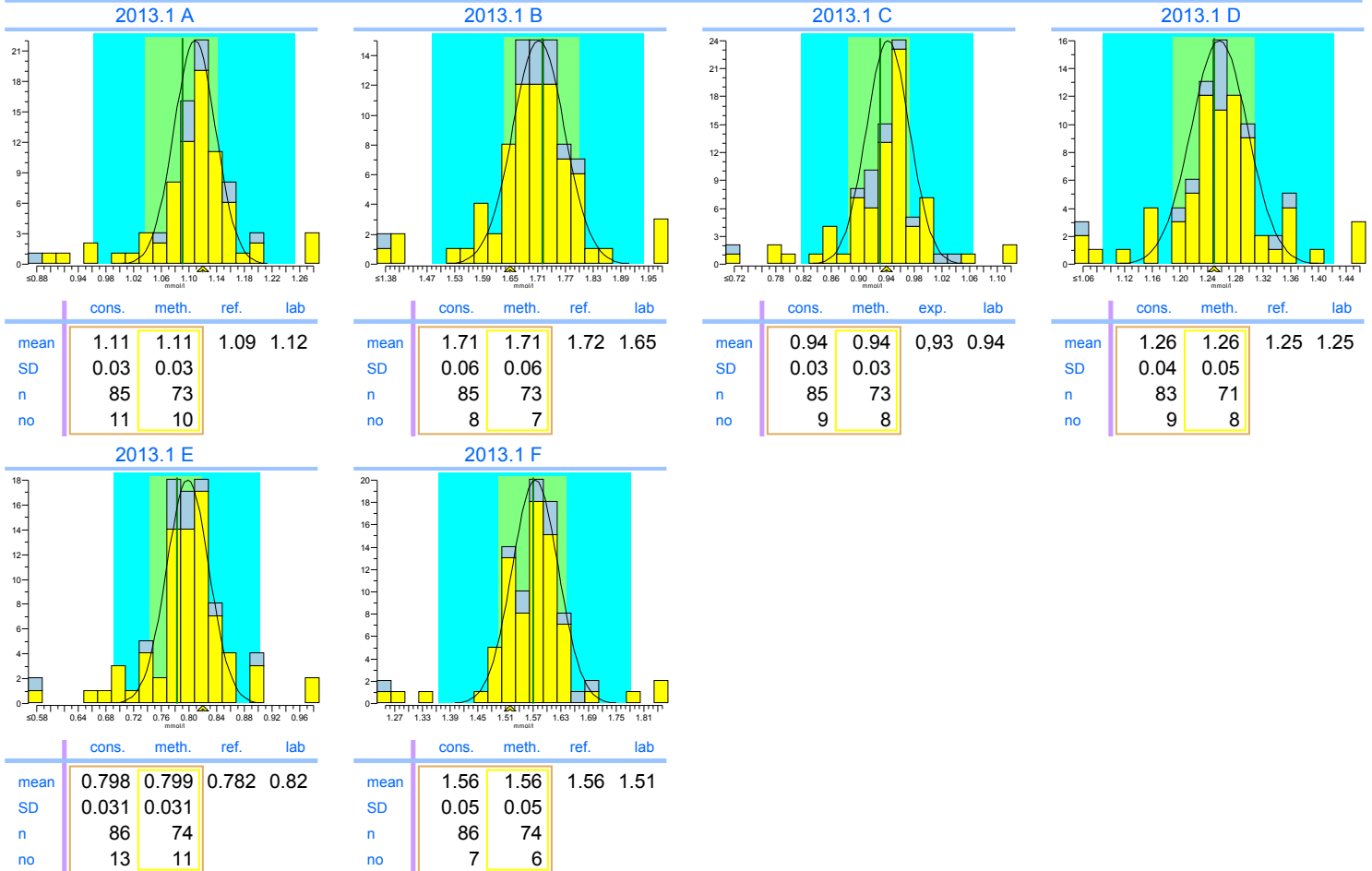
INPUTS 2013.1

Magnesium

units: mmol/l



	2013.1	cumulative
Trueness	-0.57%	-0.57%
Precision	0.92%	0.92%
Number	6	6
Outliers	0	0
Sigma-TE	3.1	3.1
Sigma-SA	6.0 2	6.0 2
Score pictogram		
Regression line	$0.13 + 0.886.x$	$0.13 + 0.886.x$
Consensus group	Overall	
Method	Colorimetric	



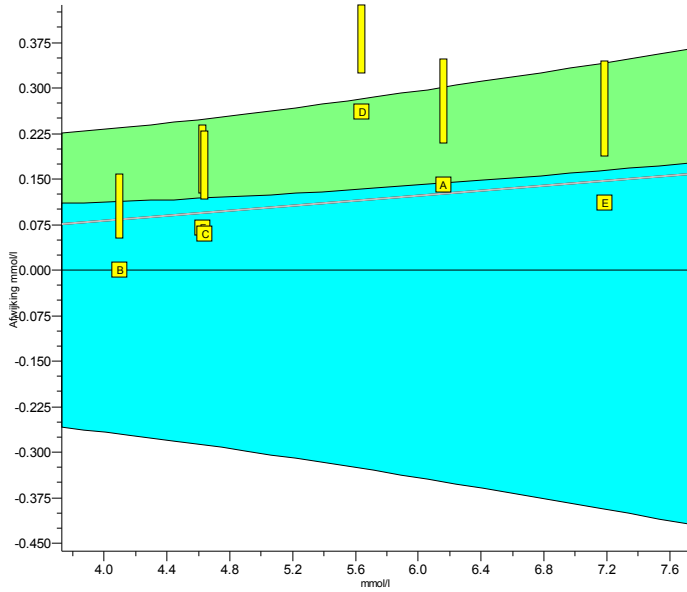
Legend



INPUTS 2013.1

Potassium

units: mmol/l



	2013.1	cumulative
Trueness	+2.0%	+2.0%
Precision	1.4%	1.4%
Number	6	6
Outliers	0	0
Sigma-TE	4.2 1	4.2 1
Sigma-SA	3.2	3.2
Score pictogram		
Regression line	<u>0.00 + 1.021.x</u>	<u>0.00 + 1.021.x</u>

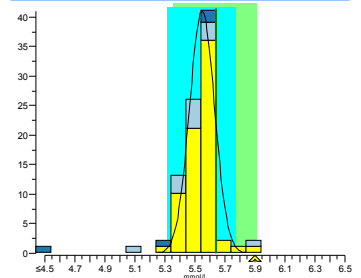
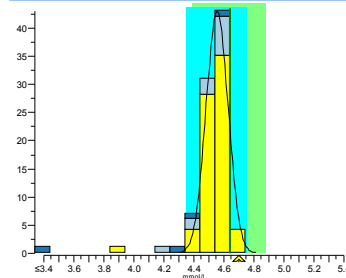
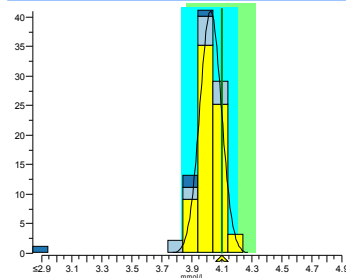
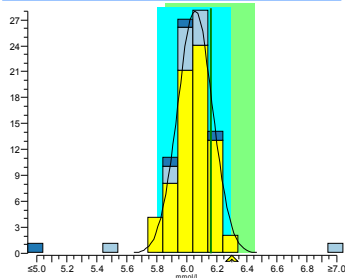
Consensus group ISE verdund/Vlamfotometrie
Method ISE indirect (with predilution)

2013.1 A

2013.1 B

2013.1 C

2013.1 D



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

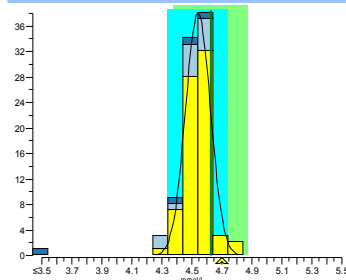
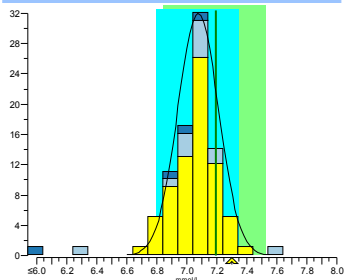
	cons.	meth.	ref.	lab
mean	4.02	4.02	4.10	4.1
SD	0.07	0.07		
n	72	72		
no	0	0		

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

	cons.	meth.	ref.	lab
mean	5.55	5.55	5.64	5.9
SD	0.08	0.08		
n	72	72		
no	2	2		

2013.1 E

2013.1 F



	cons.	meth.	ref.	lab
mean	7.08	7.08	7.19	7.3
SD	0.14	0.14		
n	72	72		
no	0	0		

	cons.	meth.	ref.	lab
mean	4.54	4.54	4.63	4.7
SD	0.08	0.08		
n	73	73		
no	2	2		

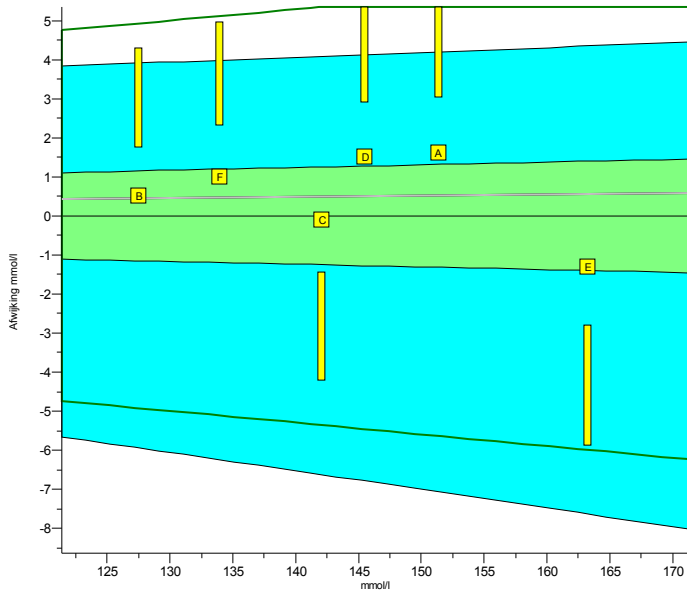
Legend

- ISE indirect (with predilution)
- ISE direct (no predilution)
- Other methods

INPUTS 2013.1

Sodium

units: mmol/l



	2013.1	cumulative
Trueness	+0.37%	+0.37%
Precision	0.79%	0.79%
Number	6	6
Outliers	0	0
Sigma-TE	1.0	1.0
Sigma-SA	4.7 2	4.7 2
Score pictogram		
Regression line	$0.0 + 1.003.x$	$0.0 + 1.003.x$

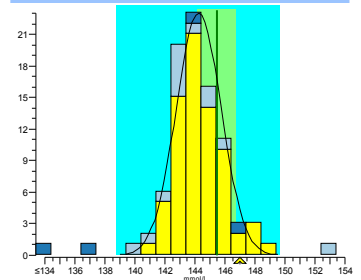
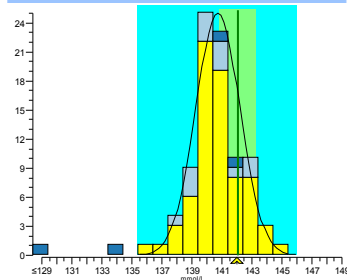
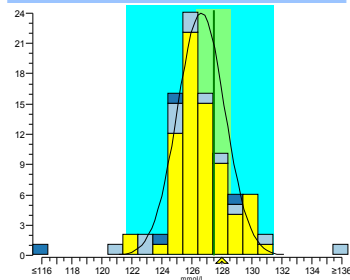
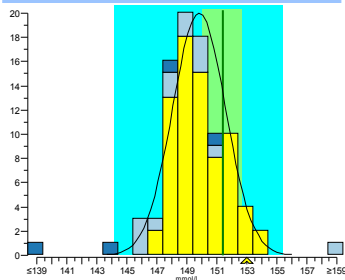
Consensus group ISE verdund/Vlamfotometrie
Method ISE indirect (with predilution)

2013.1 A

2013.1 B

2013.1 C

2013.1 D



	cons.	meth.	ref.	lab
mean	149.8	149.8	151.4	153
SD	1.8	1.8		
n	72	72		
no	0	0		

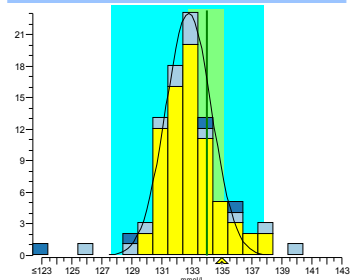
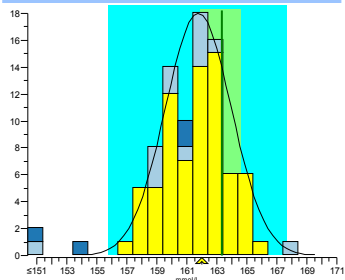
	cons.	meth.	ref.	lab
mean	126.6	126.6	127.5	128
SD	1.6	1.6		
n	72	72		
no	0	0		

	cons.	meth.	exp.	lab
mean	140.7	140.7	142.1	142
SD	1.5	1.5		
n	72	72		
no	1	1		

	cons.	meth.	ref.	lab
mean	144.3	144.3	145.5	147
SD	1.5	1.5		
n	72	72		
no	1	1		

2013.1 E

2013.1 F



	cons.	meth.	ref.	lab
mean	161.8	161.8	163.3	162
SD	2.2	2.2		
n	72	72		
no	0	0		

	cons.	meth.	ref.	lab
mean	132.8	132.8	134.0	135
SD	1.5	1.5		
n	73	73		
no	2	2		

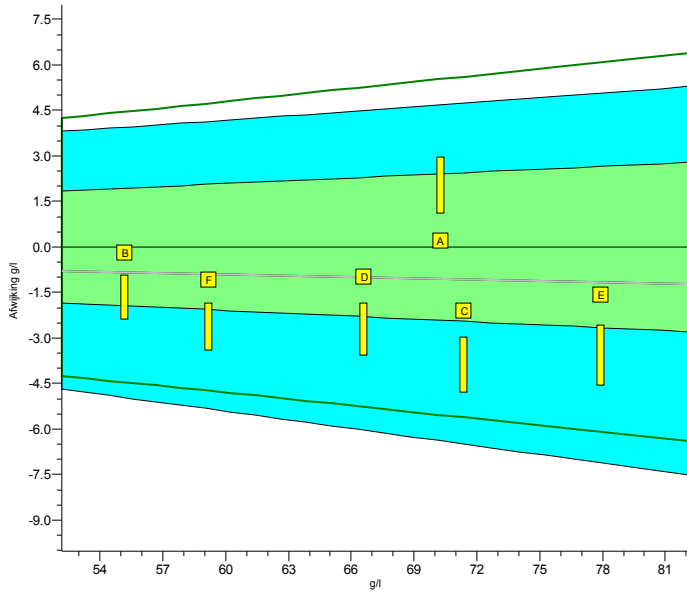
Legend

- ISE indirect (with predilution)
- ISE direct (no predilution)
- Other methods

INPUTS 2013.1

Total Protein

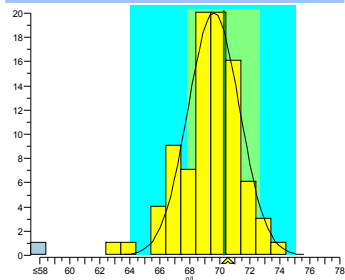
units: g/l



	2013.1	cumulative
Trueness	-1.4%	-1.4%
Precision	1.1%	1.1%
Number	6	6
Outliers	0	0
Sigma-TE	2.6	2.6
Sigma-SA	6.0 2	6.0 2
Score pictogram		
Regression line	$0.0 + 0.985.x$	$0.0 + 0.985.x$

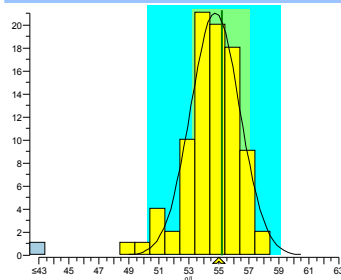
Consensus group Biureet
Method Biurete, automatic

2013.1 A



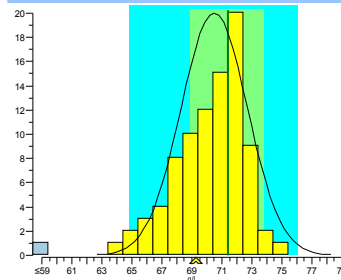
	cons.	meth.	ref.	lab
mean	69.6	69.6	70.3	70.5
SD	1.7	1.7		
n	88	88		
no	2	2		

2013.1 B



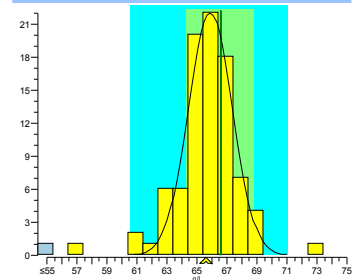
	cons.	meth.	ref.	lab
mean	54.8	54.8	55.2	55
SD	1.6	1.6		
n	88	88		
no	1	1		

2013.1 C



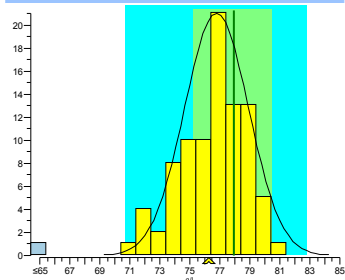
	cons.	meth.	exp.	lab
mean	70.5	70.5	71.4	69.3
SD	2.2	2.2		
n	87	87		
no	0	0		

2013.1 D



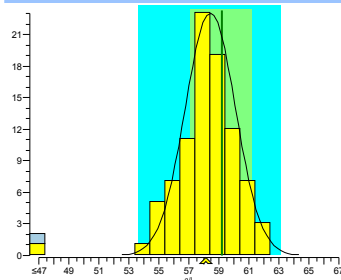
	cons.	meth.	ref.	lab
mean	65.9	65.9	66.6	65.6
SD	1.5	1.5		
n	88	88		
no	4	4		

2013.1 E



	cons.	meth.	ref.	lab
mean	76.8	76.8	77.9	76.3
SD	2.1	2.1		
n	88	88		
no	0	0		

2013.1 F



	cons.	meth.	ref.	lab
mean	58.4	58.4	59.2	58.1
SD	1.7	1.7		
n	89	89		
no	1	1		

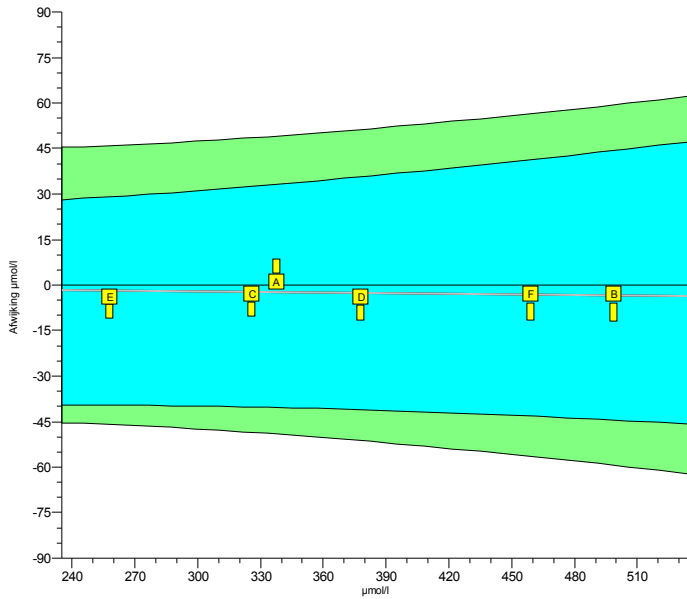
Legend

 Biurete, automatic Other methods

INPUTS 2013.1

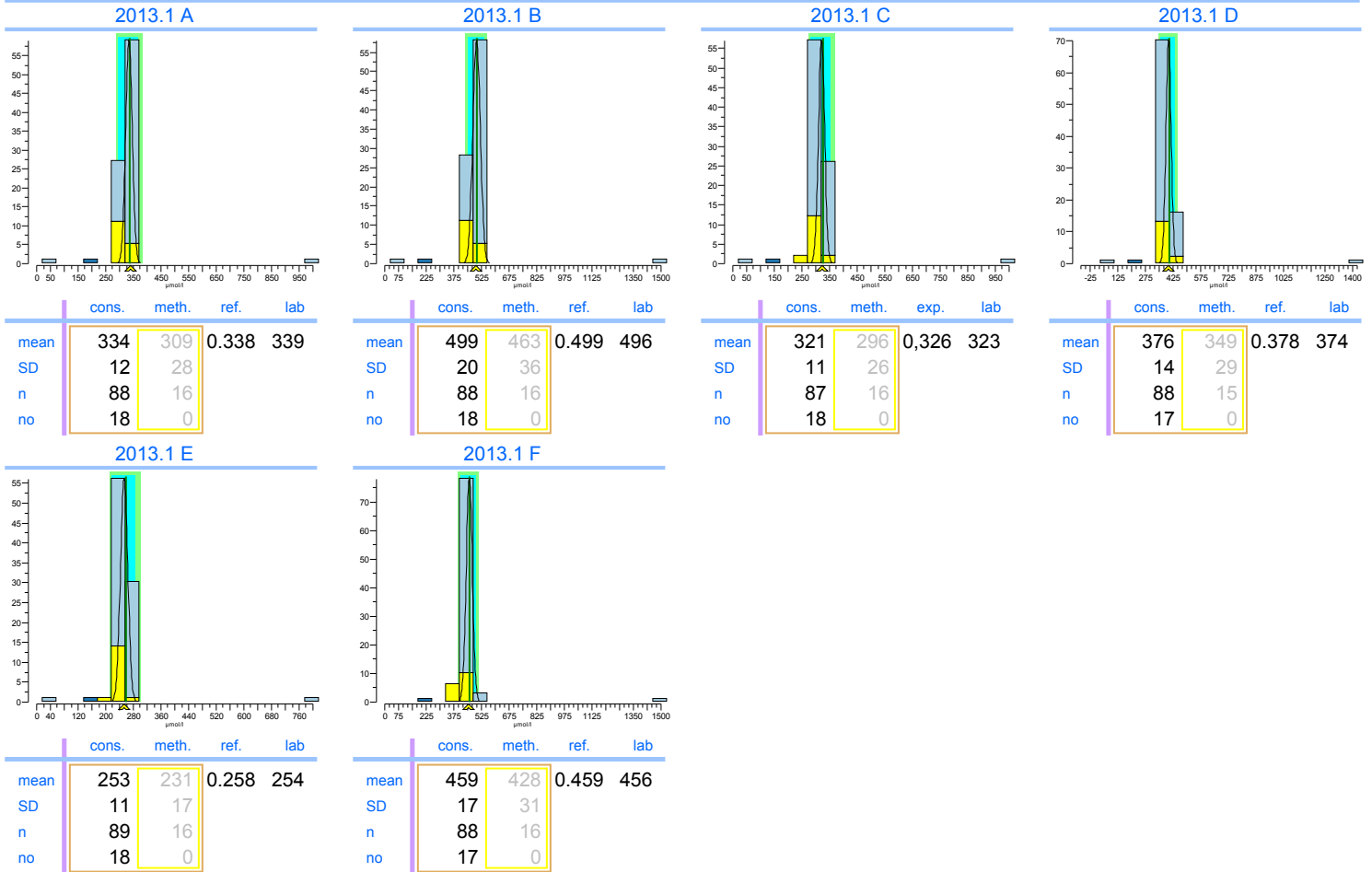
Urate

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



	2013.1	cumulative
Trueness	-0.71%	-0.71%
Precision	0.58%	0.58%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 2	6.0 2
Sigma-SA	6.0	6.0
Score pictogram		
Regression line	$0 + 0.993 \cdot x$	$0 + 0.993 \cdot x$

Consensus group: Colorimetrisch
Method: Uricase, differential UV, automatic



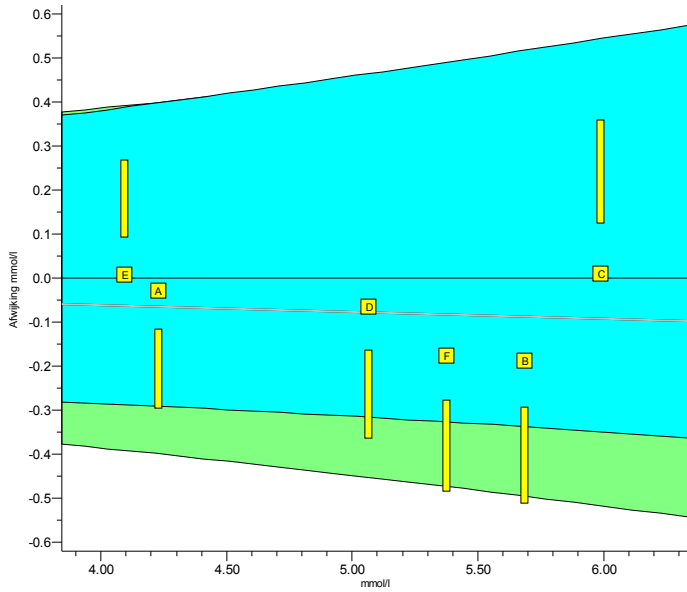
Legend

- Uricase, differential UV, automatic
- Uricase, colorim., automatic
- Overige methoden

INPUTS 2013.1

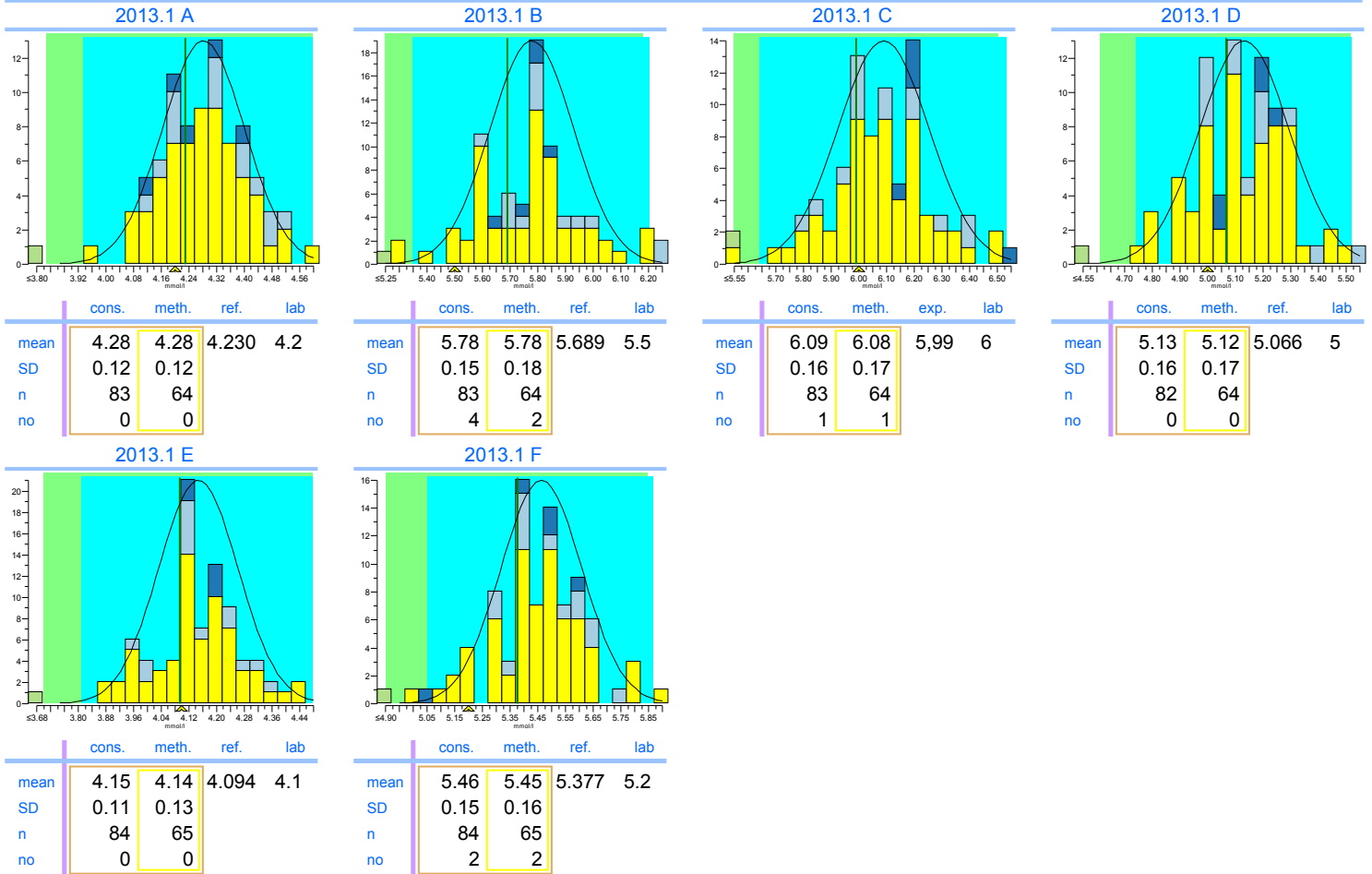
Cholesterol

units: mmol/l



	2013.1	cumulative
Trueness	-1.5%	-1.5%
Precision	1.7%	1.7%
Number	6	6
Outliers	0	0
Sigma-TE	5.4 2	5.4 2
Sigma-SA	4.6	4.6
Score pictogram		
Regression line	<u>0.00 + 0.985.x</u>	<u>0.00 + 0.985.x</u>

Consensus group Enzymatisch
Method Enzymatic, automatic, discrete



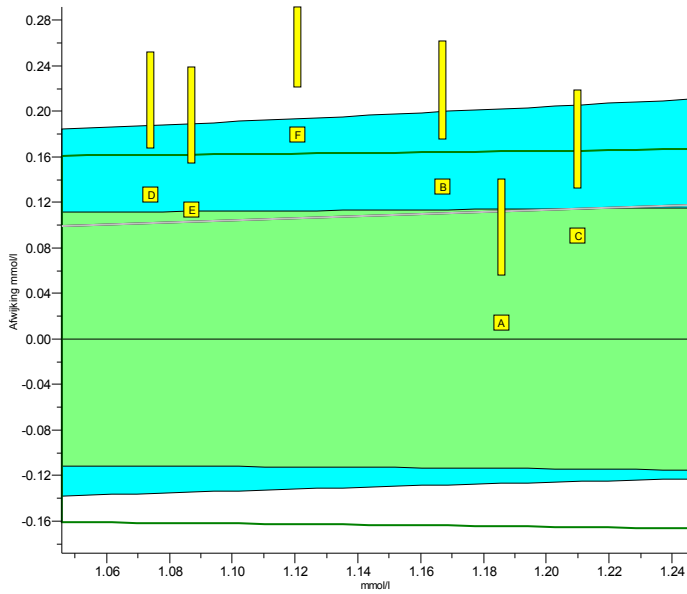
Legend

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

INPUTS 2013.1

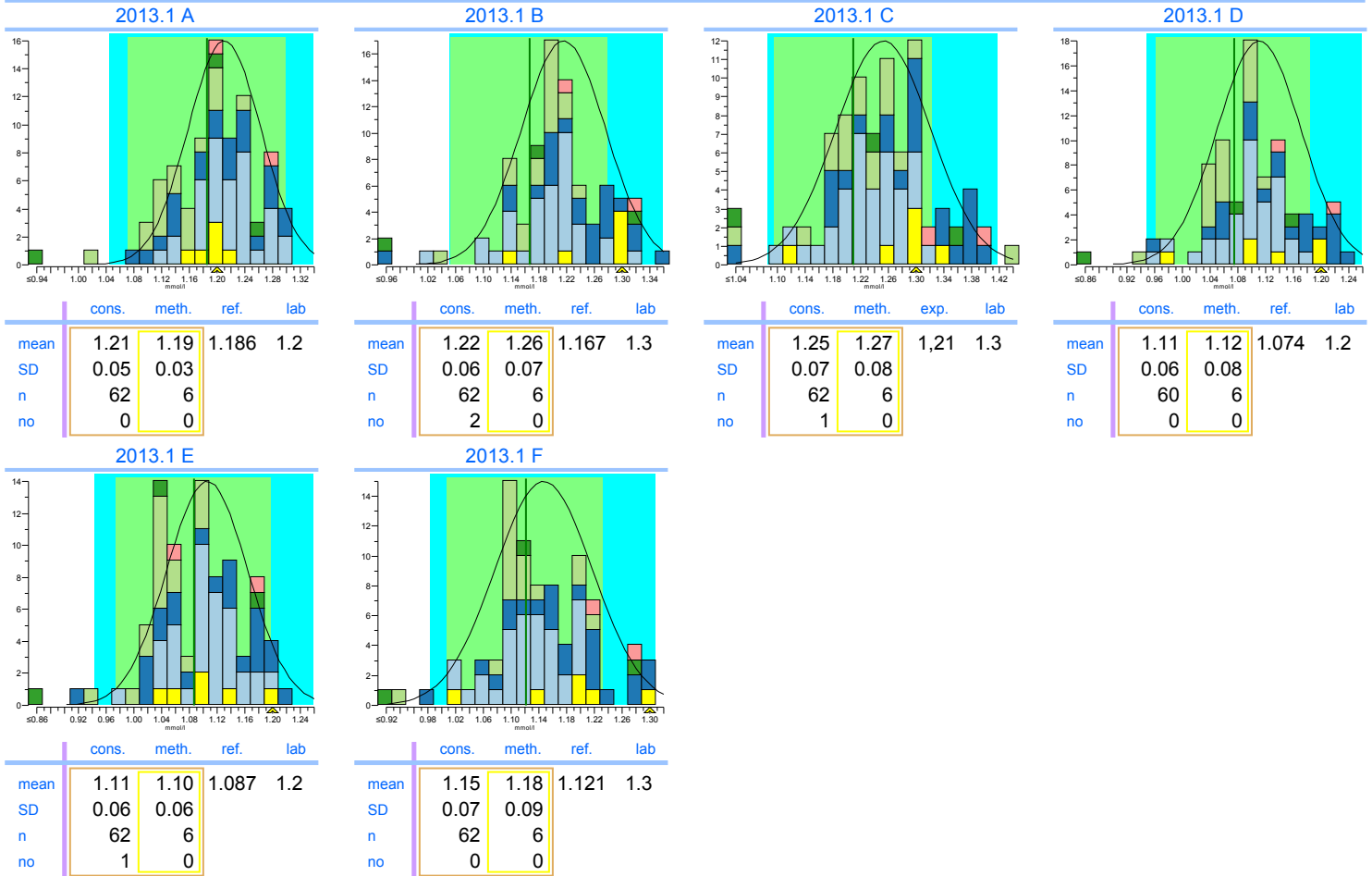
HDL-Cholesterol

units: mmol/l



	2013.1	cumulative
Trueness	+10%	+10%
Precision	4.8%	4.8%
Number	6	6
Outliers	0	0
Sigma-TE	0.9	0.9
Sigma-SA	2.4	2.4
Score pictogram		
Regression line	$0.00 + 1.095.x$	$0.00 + 1.095.x$

Consensus group	Direct
Method	Catalase method (Denka Seiken)



Legend

- Catalase method (Denka Seiken)
- PEG modified enzyme, PEGME (Kyowa Medex)
- Accelerator Selective Detergent ("Ultra HDL")
- Immuno-inhibition
- Overige methoden
- Precipitation Technique