



3703 J. Horner  
Glasgow Royal Infirmary  
-

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	28	36	



3703 J. Horner  
Glasgow Royal Infirmary

Groot Brittannië

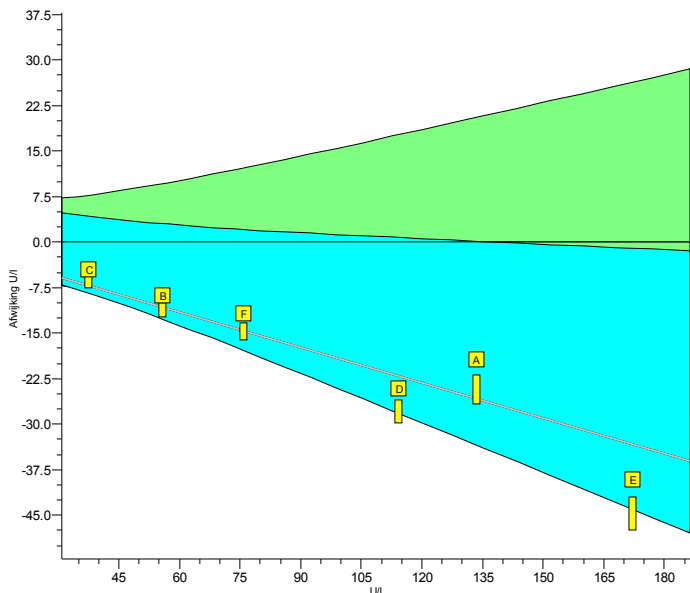
# INPUTS 2013.1

Analyte		Trueness				Precision		Performance			
		your mean	ref.	cons.	SDtI	your prec.	SDbl	this survey	PS	cumulative PSc	
ALAT	U/l	80.2	98.2	86.9	10.1	2.0	2.7		0		0
Alk. Phosphatase	U/l	179	188	173	15	4	2		2		2
Amylase	U/l	174	207	175	17	2	3		1		1
ASAT	U/l	65.5	78.6	71.5	6.8	1.6	1.6		0		0
Calcium	mmol/l	2.57	2.46	2.47	0.07	0.05	0.02		1		1
Chloride	mmol/l	101.5	100.0	100.8	1.8	0.6	0.9		2		2
CK	U/l	248	244	253	18	6	3		2		2
Creatinine	µmol/l	154.0	149.1	150.9	7.1	4.5	4.2		1		1
Gamma-GT	U/l	78.2	87.7	86.3	8.6	2.5	1.2		2		2
Glucose	mmol/l	14.52	13.93	14.19	0.44	0.43	0.18		1		1
LD	U/l	521	532	523	33	8	9		2		2
Magnesium	mmol/l	1.23	1.22	1.23	0.04	0.03	0.02		2		2
Potassium	mmol/l	5.28	5.39	5.30	0.10	0.03	0.04		2		2
Sodium	mmol/l	142.7	144.0	142.7	1.7	0.3	0.9		2		2
Total Protein	g/l	66.8	66.8	66.0	1.8	0.4	0.7		2		2
Urate	µmol/l	400	376	374	15	6	4		2		2
Cholesterol	mmol/l	5.15	5.07	5.15	0.14	0.09	0.05		2		2
HDL-Cholesterol	mmol/l	1.18	1.14	1.17	0.06	0.03	0.03		2		2
<b>Total :</b>									<b>28</b>		<b>28</b>

# INPUTS 2013.1

ALAT

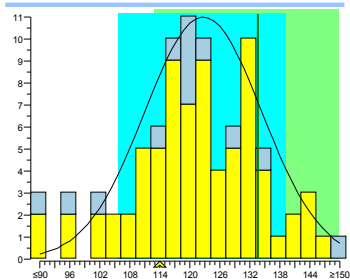
units: U/l



	2013.1	cumulative
Trueness	-18%	-18%
Precision	2.3%	2.3%
Number	6	6
Outliers	0	0
Sigma-TE	0.4 <span style="border: 1px solid red; padding: 0 2px;">0</span>	0.4 <span style="border: 1px solid red; padding: 0 2px;">0</span>
Sigma-SA	-0.7	-0.7
Score pictogram		
Regression line	$0.0 + 0.807 \cdot x$	$0.0 + 0.807 \cdot x$

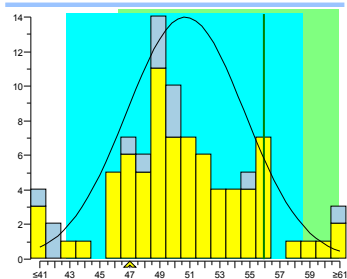
Consensus group IFCC traceerbaar  
Method IFCC traceable

2013.1 A



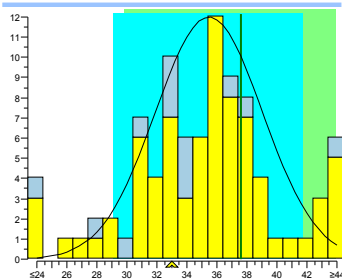
	cons.	meth.	ref.	lab
mean	122.7	122.7	133.5	114
SD	11.8	11.8		
n	75	75		
no	1	1		

2013.1 B



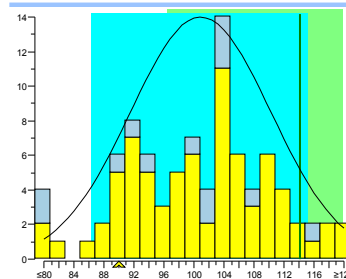
	cons.	meth.	ref.	lab
mean	50.7	50.7	55.9	47
SD	3.9	3.9		
n	76	76		
no	3	3		

2013.1 C



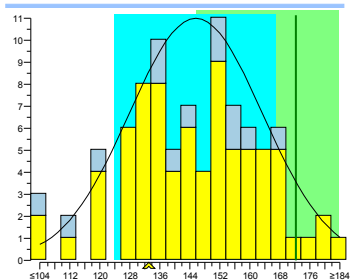
	cons.	meth.	exp.	lab
mean	35.5	35.5	37,6	33
SD	3.6	3.6		
n	76	76		
no	7	7		

2013.1 D



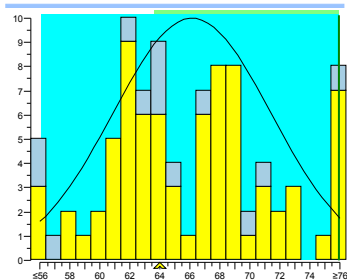
	cons.	meth.	ref.	lab
mean	100.9	100.9	114.2	90
SD	9.4	9.4		
n	76	76		
no	3	3		

2013.1 E



	cons.	meth.	ref.	lab
mean	146	146	172.2	133
SD	18	18		
n	77	77		
no	1	1		

2013.1 F



	cons.	meth.	ref.	lab
mean	66.1	66.1	75.9	64
SD	5.3	5.3		
n	77	77		
no	3	3		

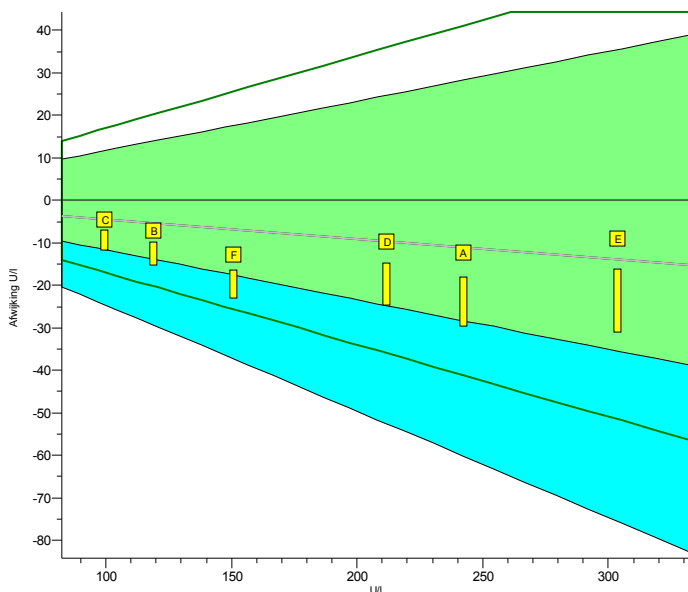
Legend



IFCC traceable     IFCC non-traceable

# INPUTS 2013.1

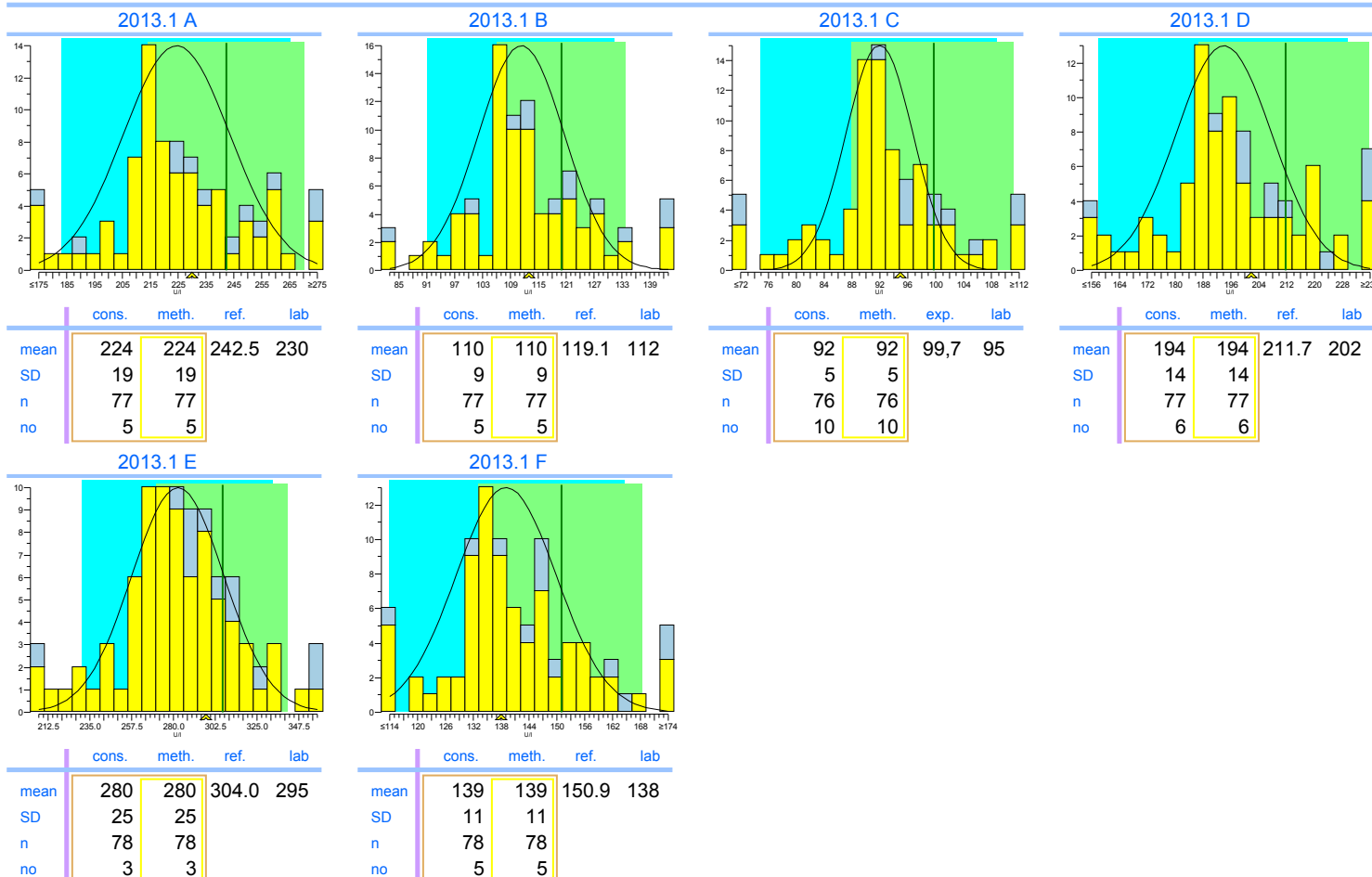
Alk. Phosphatase

units: U/l



	2013.1	cumulative
Trueness	-5.0%	-5.0%
Precision	2.2%	2.2%
Number	6	6
Outliers	0	0
Sigma-TE	3.9	3.9
Sigma-SA	6.0 <span style="background-color: green; color: white; padding: 2px;">2</span>	6.0 <span style="background-color: green; color: white; padding: 2px;">2</span>
Score pictogram		
Regression line	$0 + 0.954.x$	$0 + 0.954.x$

Consensus group	IFCC traceerbaar
Method	IFCC traceable



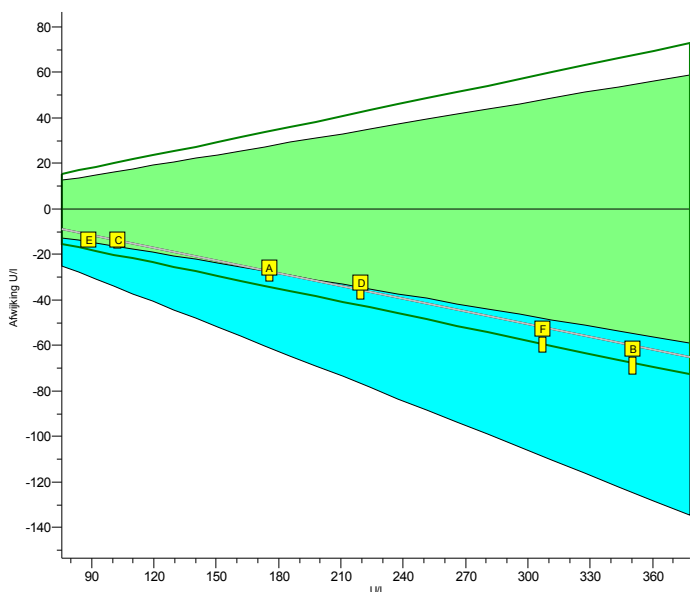
Legend

IFCC traceable     IFCC non-traceable

# INPUTS 2013.1

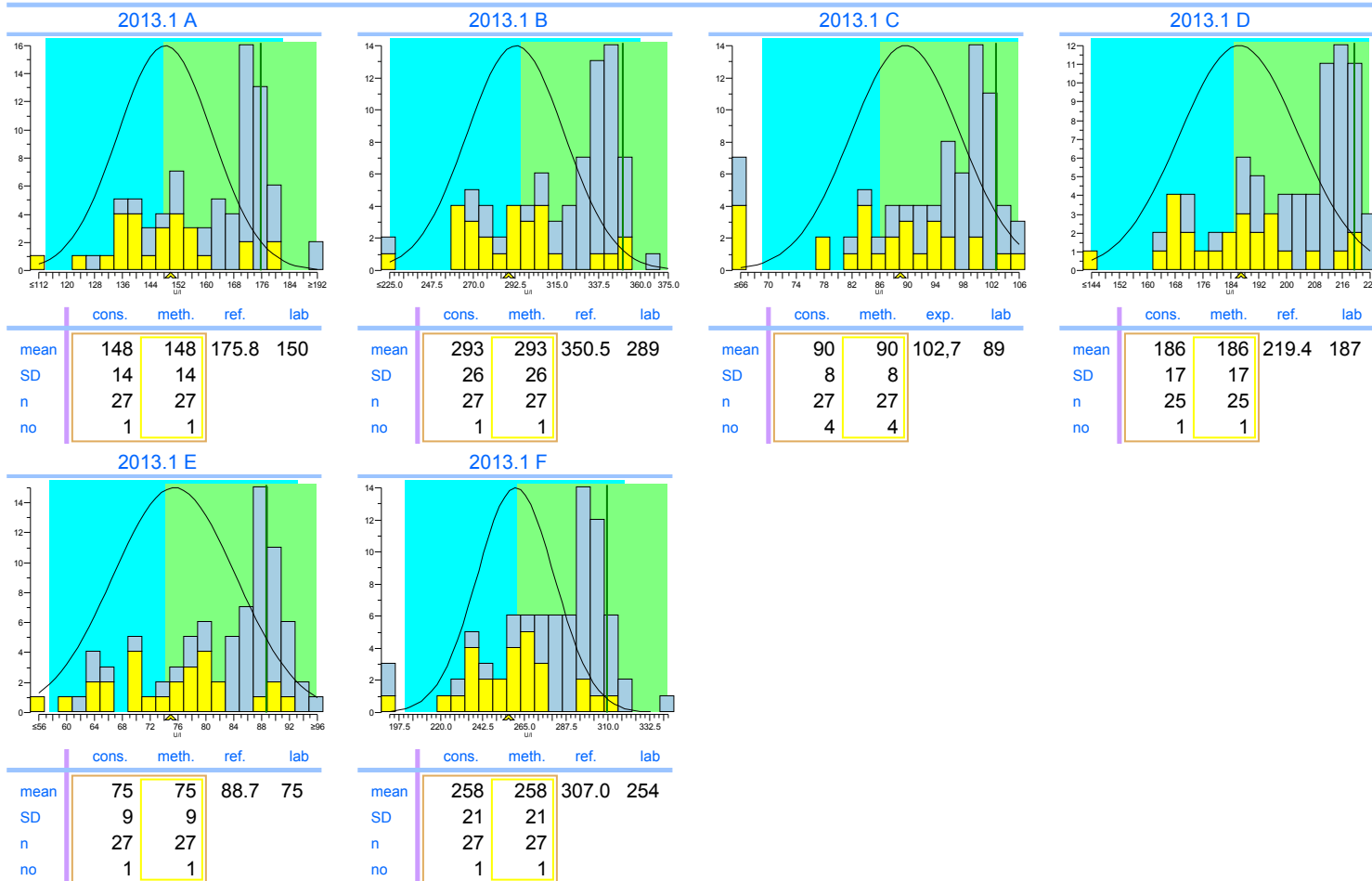
## Amylase

units: U/l



	2013.1	cumulative
Trueness	-16%	-16%
Precision	1.1%	1.1%
Number	6	6
Outliers	0	0
Sigma-TE	1.1	1.1
Sigma-SA	4.4	4.4
Score pictogram		
Regression line	$5 + 0.813.x$	$5 + 0.813.x$

Consensus group niet IFCC traceerbaar  
Method IFCC non-traceable



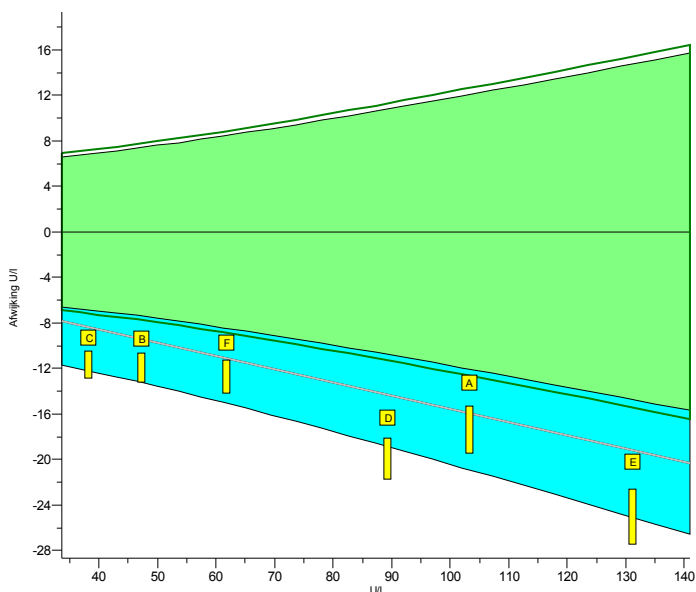
Legend

IFCC non-traceable IFCC traceable

# INPUTS 2013.1

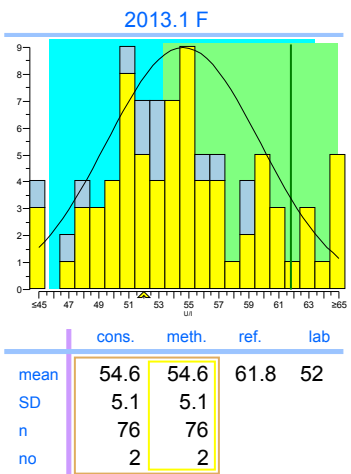
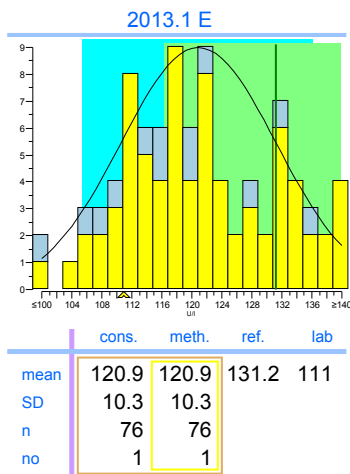
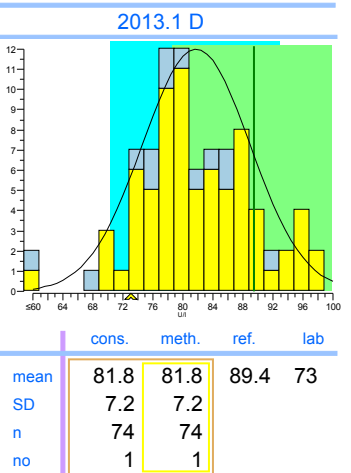
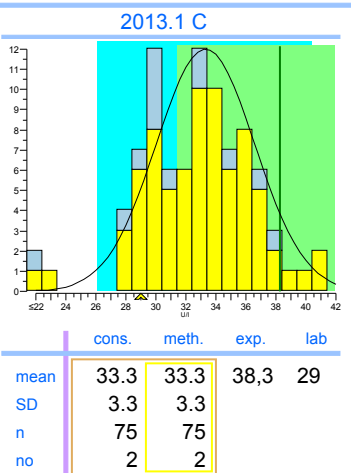
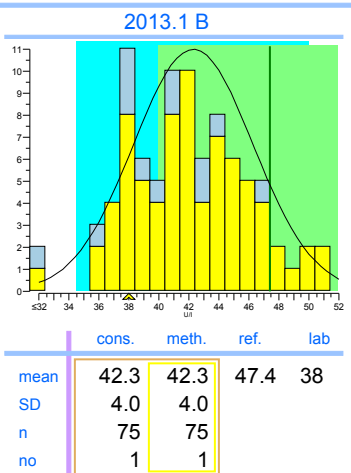
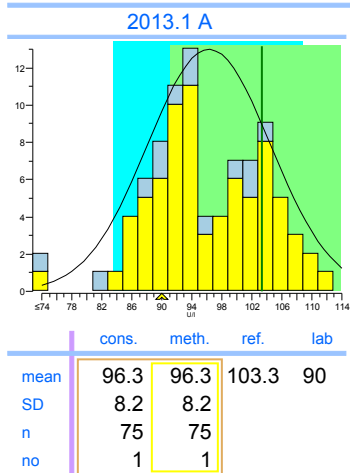
ASAT

units: U/I



	2013.1	cumulative
Trueness	-17%	-17%
Precision	2.3%	2.3%
Number	6	6
Outliers	0	0
Sigma-TE	-1.1	-1.1
Sigma-SA	-0.8 <span style="border: 1px solid red; padding: 0 2px;">0</span>	-0.8 <span style="border: 1px solid red; padding: 0 2px;">0</span>
Score pictogram		
Regression line	$-4.0 + 0.884.x$	$-4.0 + 0.884.x$

Consensus group IFCC traceerbaar  
Method IFCC traceable



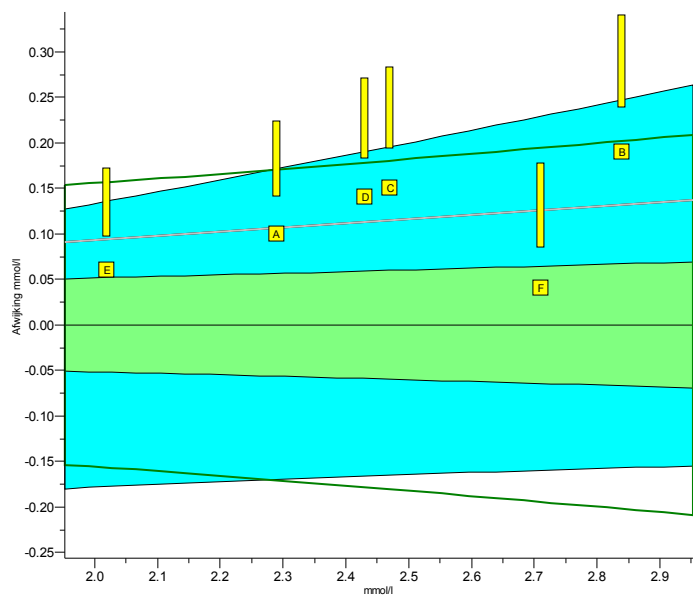
Legend



IFCC traceable     IFCC non-traceable

# INPUTS 2013.1

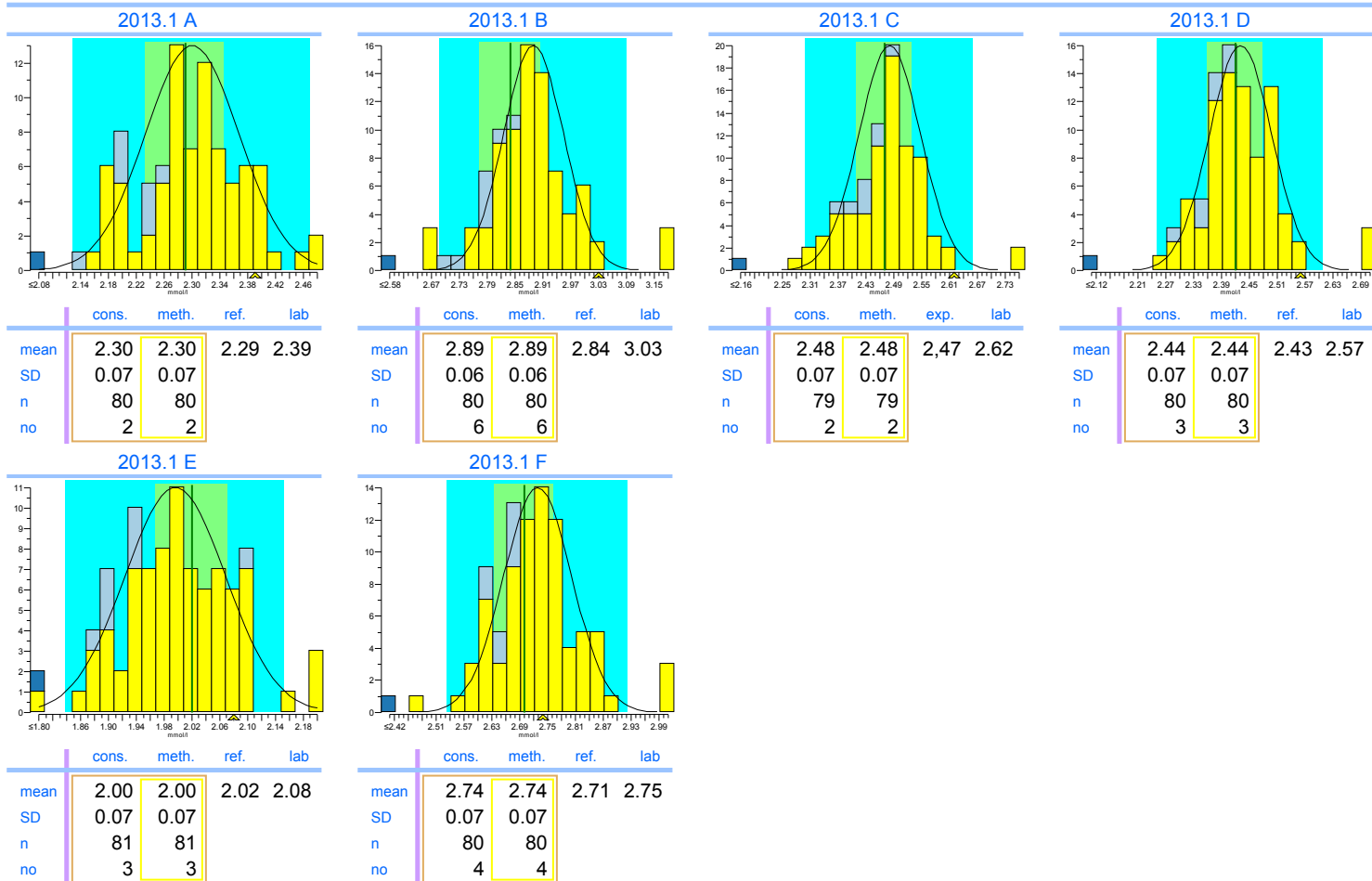
## Calcium

units: mmol/l



	2013.1	cumulative
Trueness	+4.6%	+4.6%
Precision	2.0%	2.0%
Number	6	6
Outliers	0	0
Sigma-TE	-0.7	-0.7
Sigma-SA	2.8 <span style="background-color: green; color: white; padding: 0 2px;">1</span>	2.8 <span style="background-color: green; color: white; padding: 0 2px;">1</span>
Score pictogram		
Regression line	<u>0.00 + 1.047.x</u>	<u>0.00 + 1.047.x</u>

Consensus group	Colorimetrisch
Method	Colorimetric, automatic, discrete



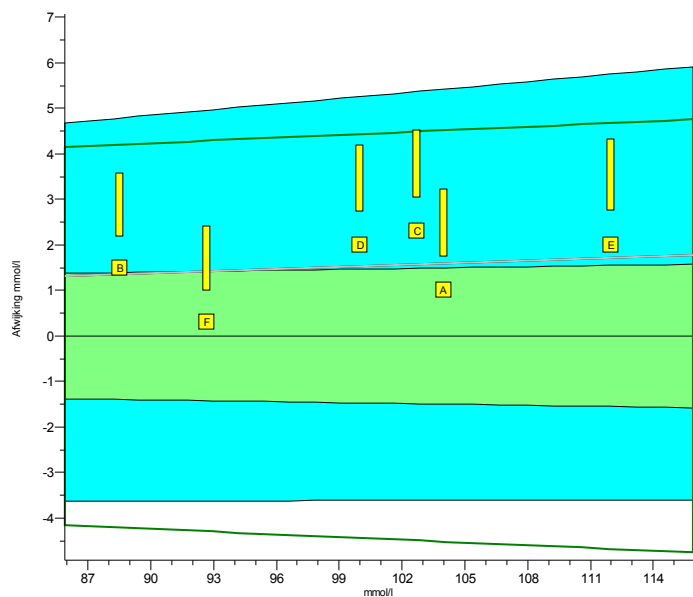
Legend

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

# INPUTS 2013.1

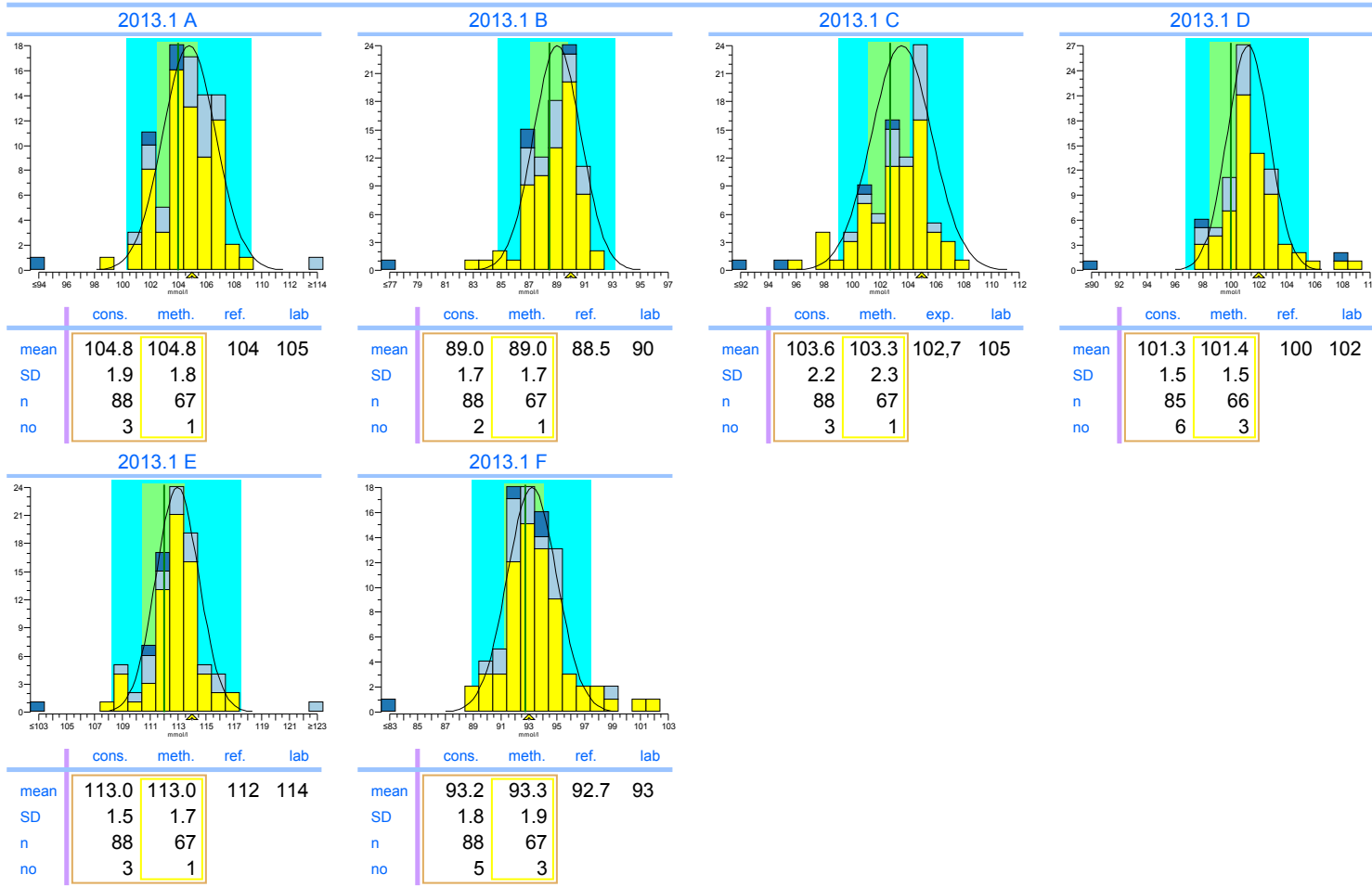
Chloride

units: mmol/l



	2013.1	cumulative
Trueness	+1.5%	+1.5%
Precision	0.59%	0.59%
Number	6	6
Outliers	0	0
Sigma-TE	0.7	0.7
Sigma-SA	5.8 <span style="background-color: green; color: white; padding: 2px;">2</span>	5.8 <span style="background-color: green; color: white; padding: 2px;">2</span>
Score pictogram		
Regression line	$0.0 + 1.015 \cdot x$	$0.0 + 1.015 \cdot x$

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



Legend

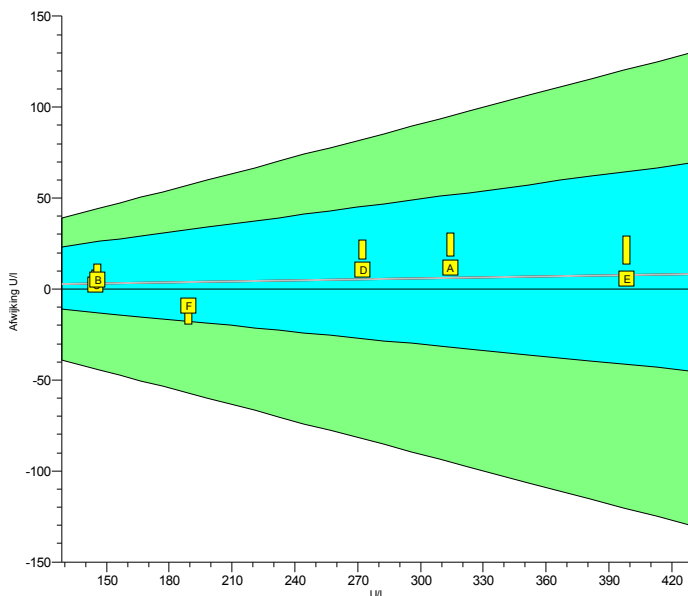
ISE indirect (with predilution)	ISE direct (no predilution)	Other methods
---------------------------------	-----------------------------	---------------



# INPUTS 2013.1

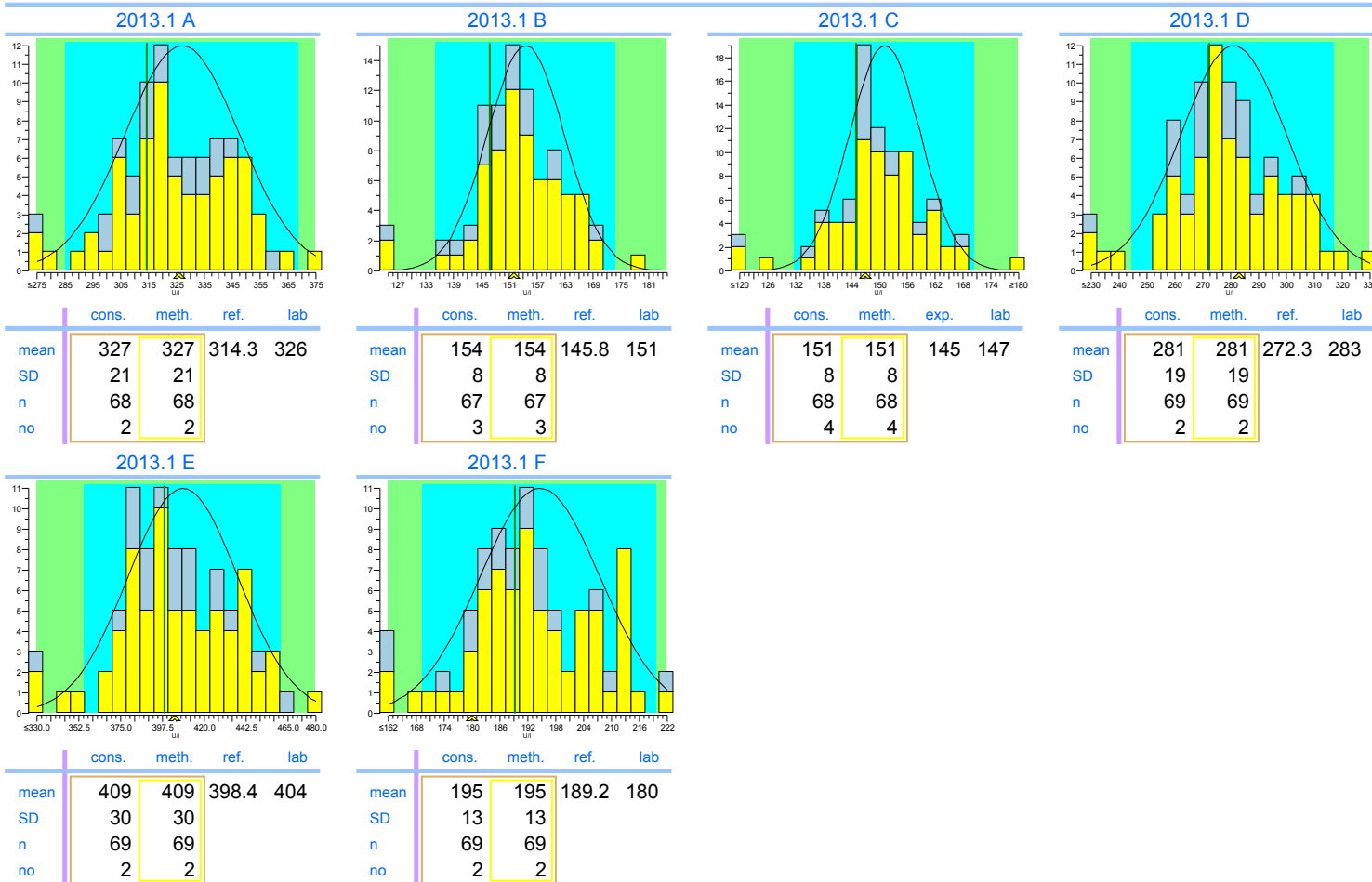
CK

units: U/l



	2013.1	cumulative
Trueness	+1.8%	+1.8%
Precision	2.5%	2.5%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 <span style="background-color: green; color: white;">2</span>	6.0 <span style="background-color: green; color: white;">2</span>
Sigma-SA	6.0	6.0
Score pictogram		
Regression line	$0 + 1.020 \cdot x$	$0 + 1.020 \cdot x$

Consensus group IFCC traceerbaar  
Method IFCC traceable

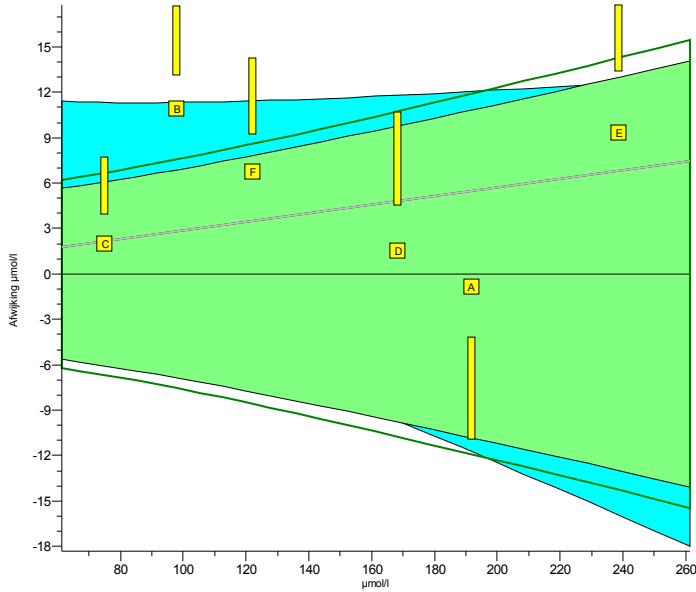






Legend  
 IFCC traceable  
 IFCC non-traceable

# INPUTS 2013.1

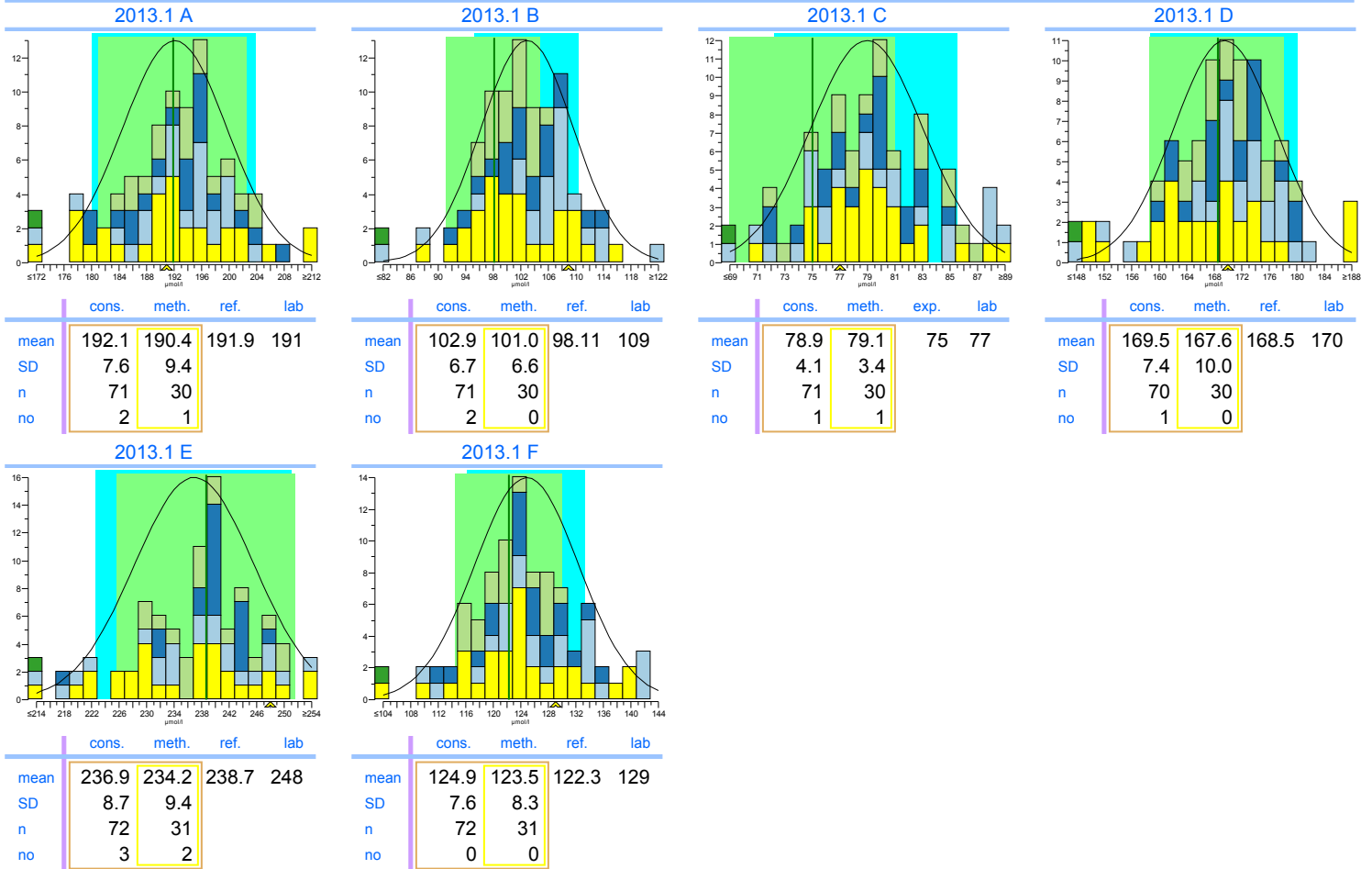
## Creatinine

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



	2013.1	cumulative
Trueness	+3.3%	+3.3%
Precision	3.0%	3.0%
Number	6	6
Outliers	0	0
Sigma-TE	2.3	2.3
Sigma-SA	2.7 	2.7 
Score pictogram		
Regression line	$0.0 + 1.028.x$	$0.0 + 1.028.x$

Consensus group Jaffe  
Method Alk. Picrate, kinetic with compensation



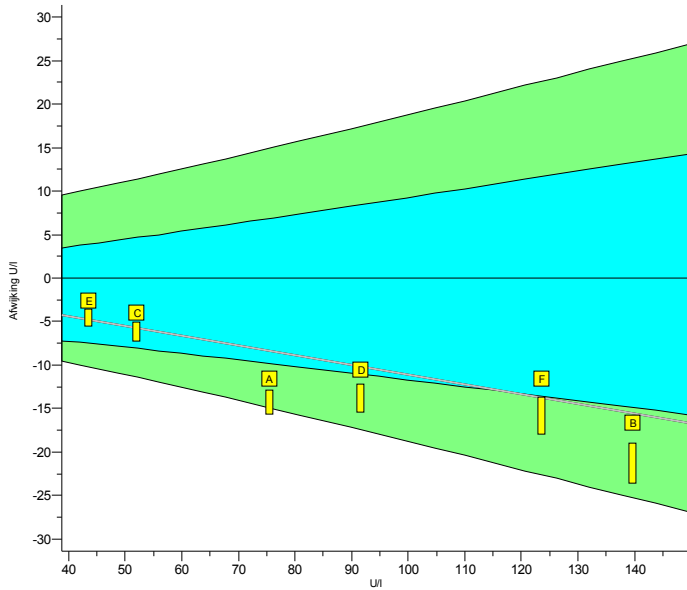
Legend

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

# INPUTS 2013.1

Gamma-GT

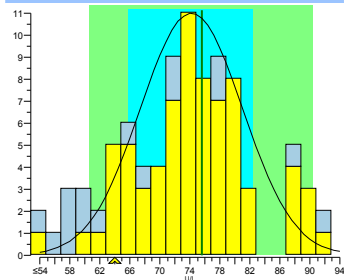
units: U/I



	2013.1	cumulative
Trueness	-11%	-11%
Precision	2.9%	2.9%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 <span style="background-color: green; color: white;">2</span>	6.0 <span style="background-color: green; color: white;">2</span>
Sigma-SA	1.2	1.2
Score pictogram		
Regression line	$0.0 + 0.889.x$	$0.0 + 0.889.x$

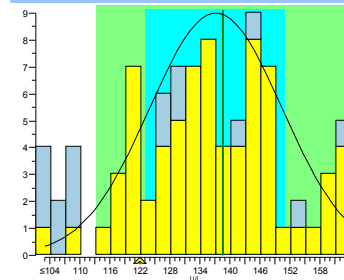
Consensus group IFCC traceerbaar  
Method IFCC traceable

2013.1 A



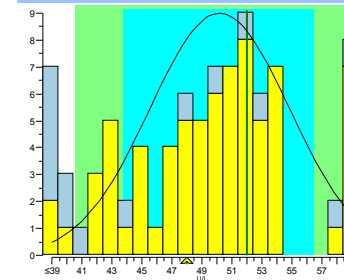
	cons.	meth.	ref.	lab
mean	74.2	74.2	75.6	64
SD	6.8	6.8		
n	72	72		
no	1	1		

2013.1 B



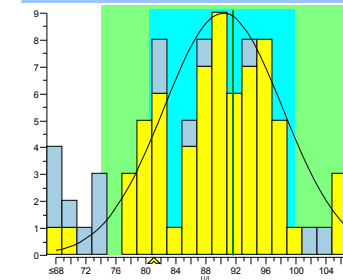
	cons.	meth.	ref.	lab
mean	138.2	138.2	139.7	123
SD	13.3	13.3		
n	72	72		
no	1	1		

2013.1 C



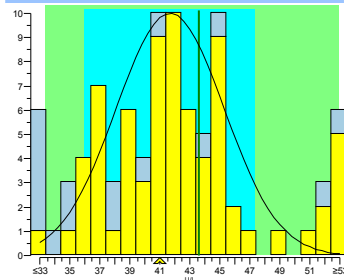
	cons.	meth.	exp.	lab
mean	50.2	50.2	52	48
SD	4.6	4.6		
n	72	72		
no	2	2		

2013.1 D



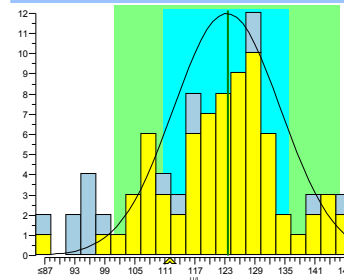
	cons.	meth.	ref.	lab
mean	90.3	90.3	91.6	81
SD	7.9	7.9		
n	71	71		
no	1	1		

2013.1 E



	cons.	meth.	ref.	lab
mean	41.7	41.7	43.6	41
SD	3.6	3.6		
n	73	73		
no	6	6		

2013.1 F



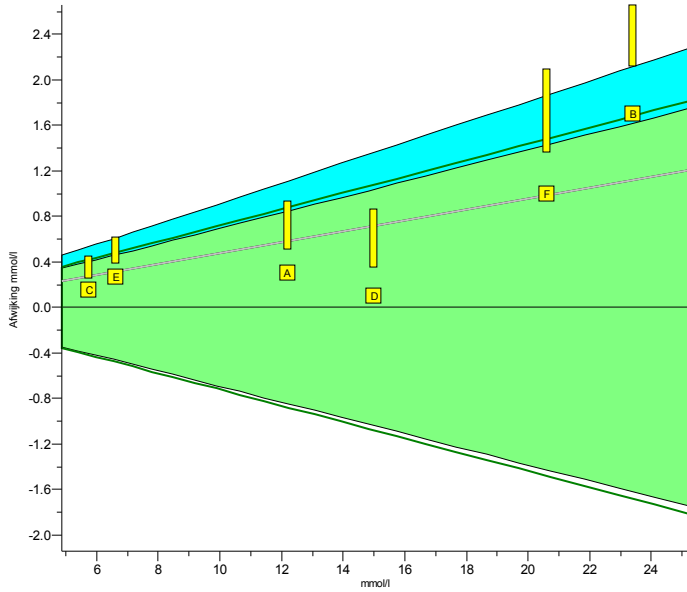
	cons.	meth.	ref.	lab
mean	123.4	123.4	123.6	112
SD	10.7	10.7		
n	73	73		
no	1	1		



Legend  
 IFCC traceable  
 IFCC non-traceable

# INPUTS 2013.1

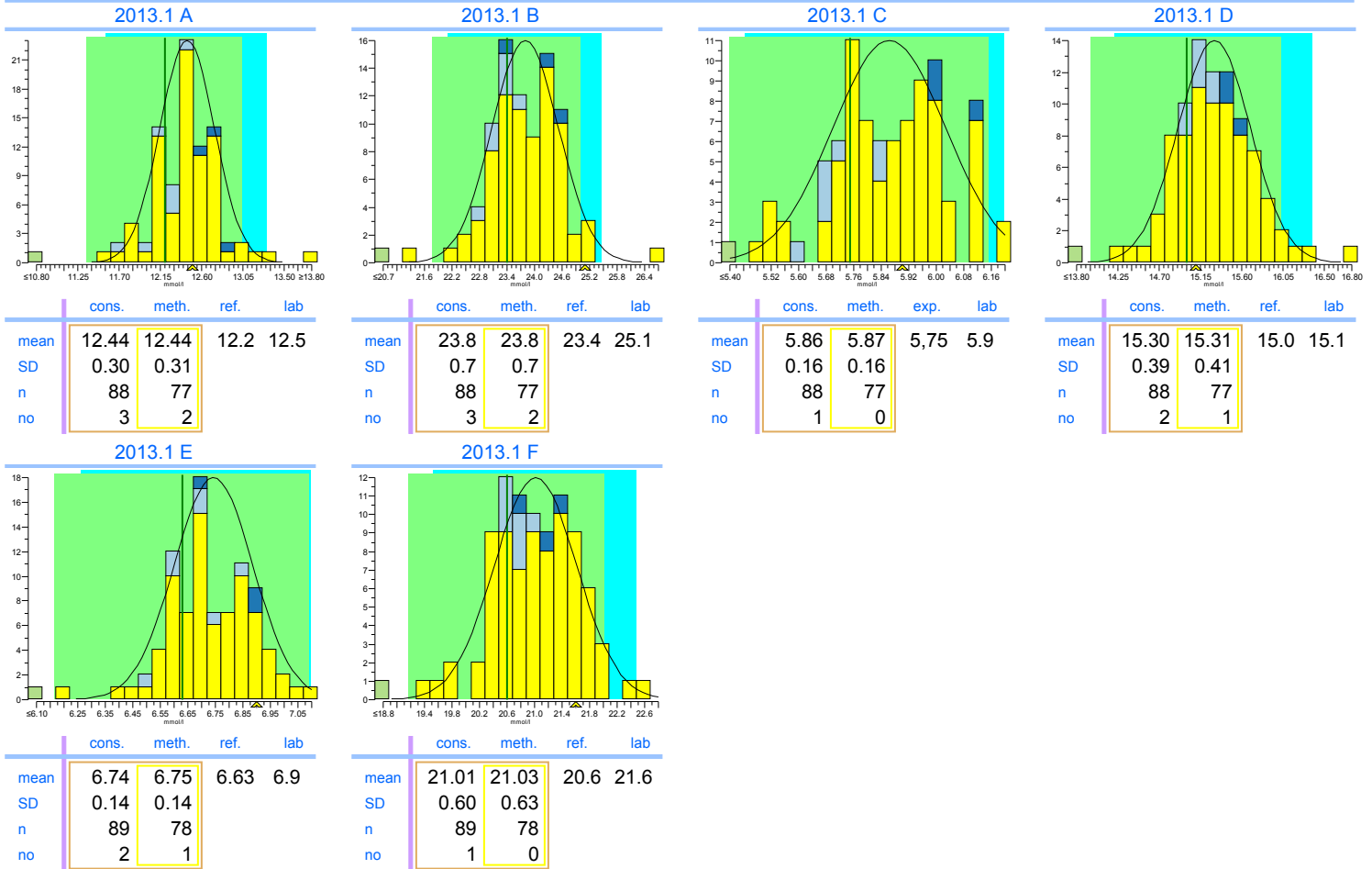
## Glucose

units: mmol/l



	2013.1	cumulative
Trueness	+4.2%	+4.2%
Precision	3.0%	3.0%
Number	6	6
Outliers	0	0
Sigma-TE	3.2	3.2
Sigma-SA	3.4	3.4
Score pictogram		
Regression line	$0.00 + 1.048.x$	$0.00 + 1.048.x$

Consensus group: Natte chemie  
Method: Hexokinase, automatic



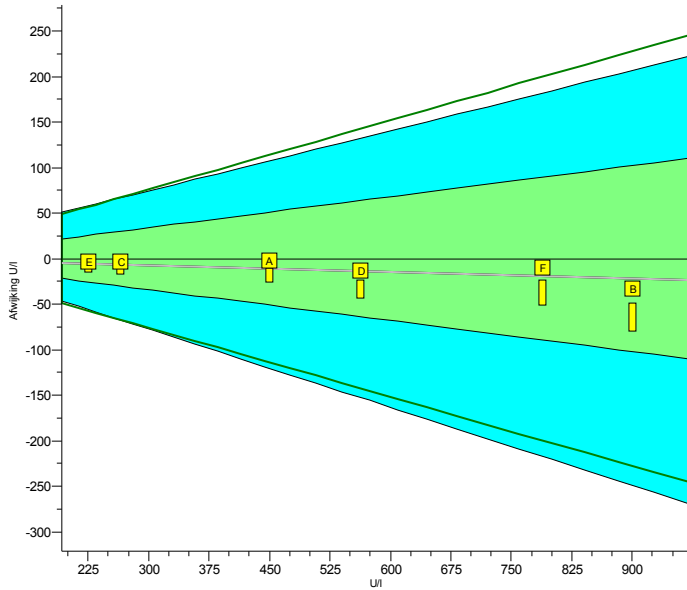
Legend

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

# INPUTS 2013.1

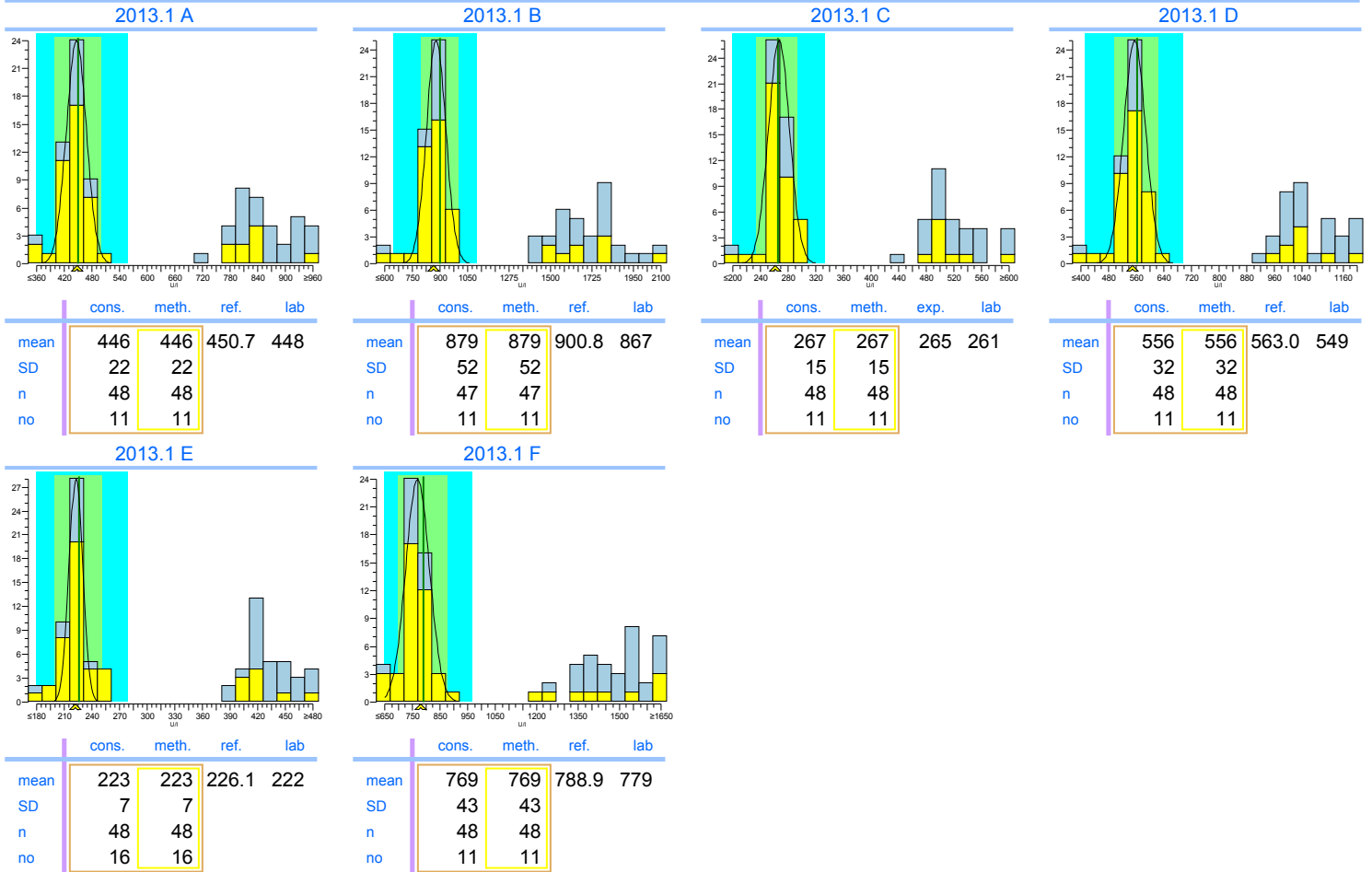
LD

units: U/I



	2013.1	cumulative
Trueness	-2.1%	-2.1%
Precision	1.5%	1.5%
Number	6	6
Outliers	0	0
Sigma-TE	6.0	6.0
Sigma-SA	6.0 <span style="background-color: green; color: white; padding: 2px;">2</span>	6.0 <span style="background-color: green; color: white; padding: 2px;">2</span>
Score pictogram		
Regression line	$0 + 0.976.x$	$0 + 0.976.x$

Consensus group IFCC tracerbaar  
Method IFCC traceable

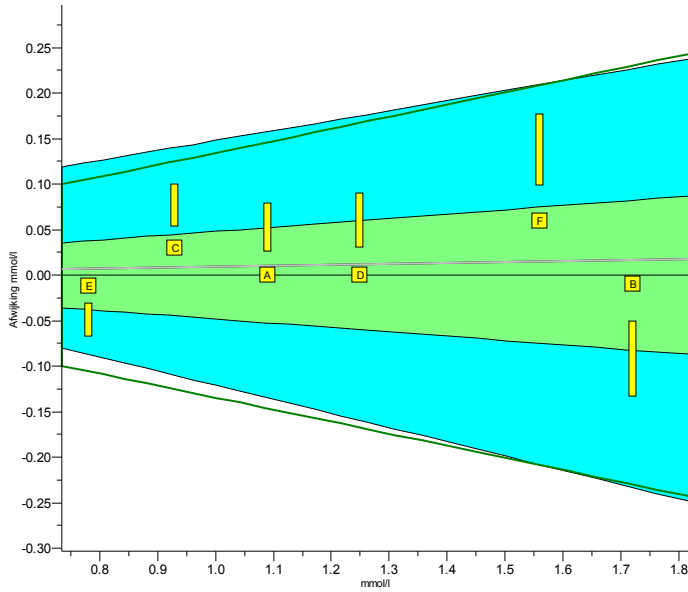


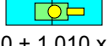
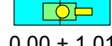
Legend  
 IFCC traceable  
 IFCC non-traceable

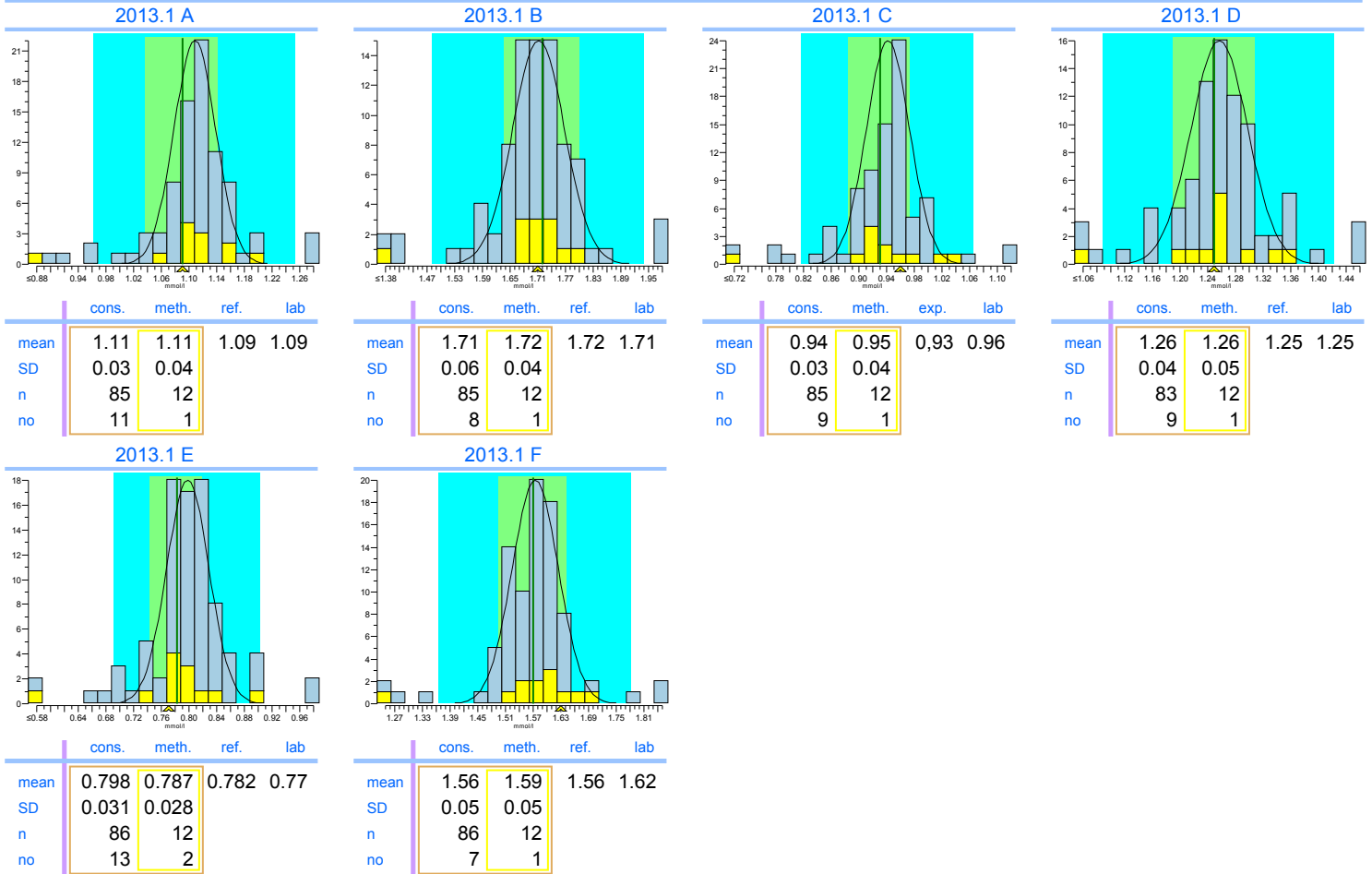
# INPUTS 2013.1

## Magnesium

units: mmol/l



	2013.1	cumulative
Trueness	+0.93%	+0.93%
Precision	2.4%	2.4%
Number	6	6
Outliers	0	0
Sigma-TE	2.5	2.5
Sigma-SA	6.0 <span style="background-color: green; color: white; padding: 2px;">2</span>	6.0 <span style="background-color: green; color: white; padding: 2px;">2</span>
Score pictogram		
Regression line	<u>0.00 + 1.010.x</u>	<u>0.00 + 1.010.x</u>
Consensus group	Overall	
Method	Other methods	



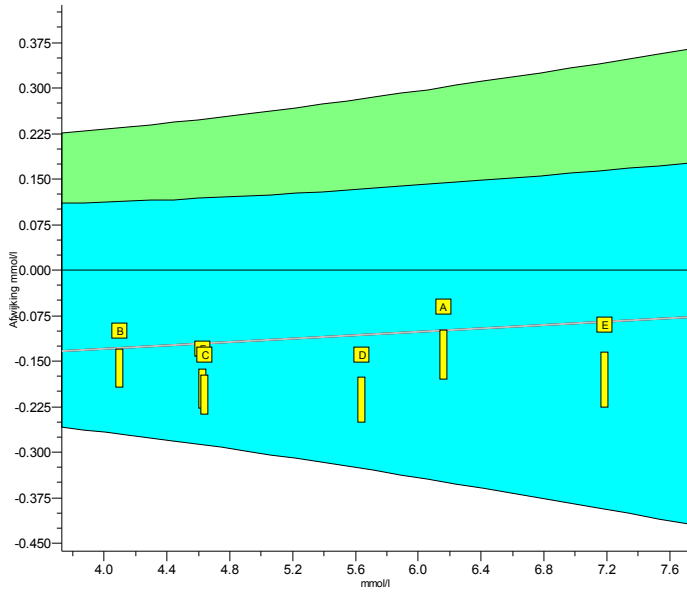
Legend



# INPUTS 2013.1

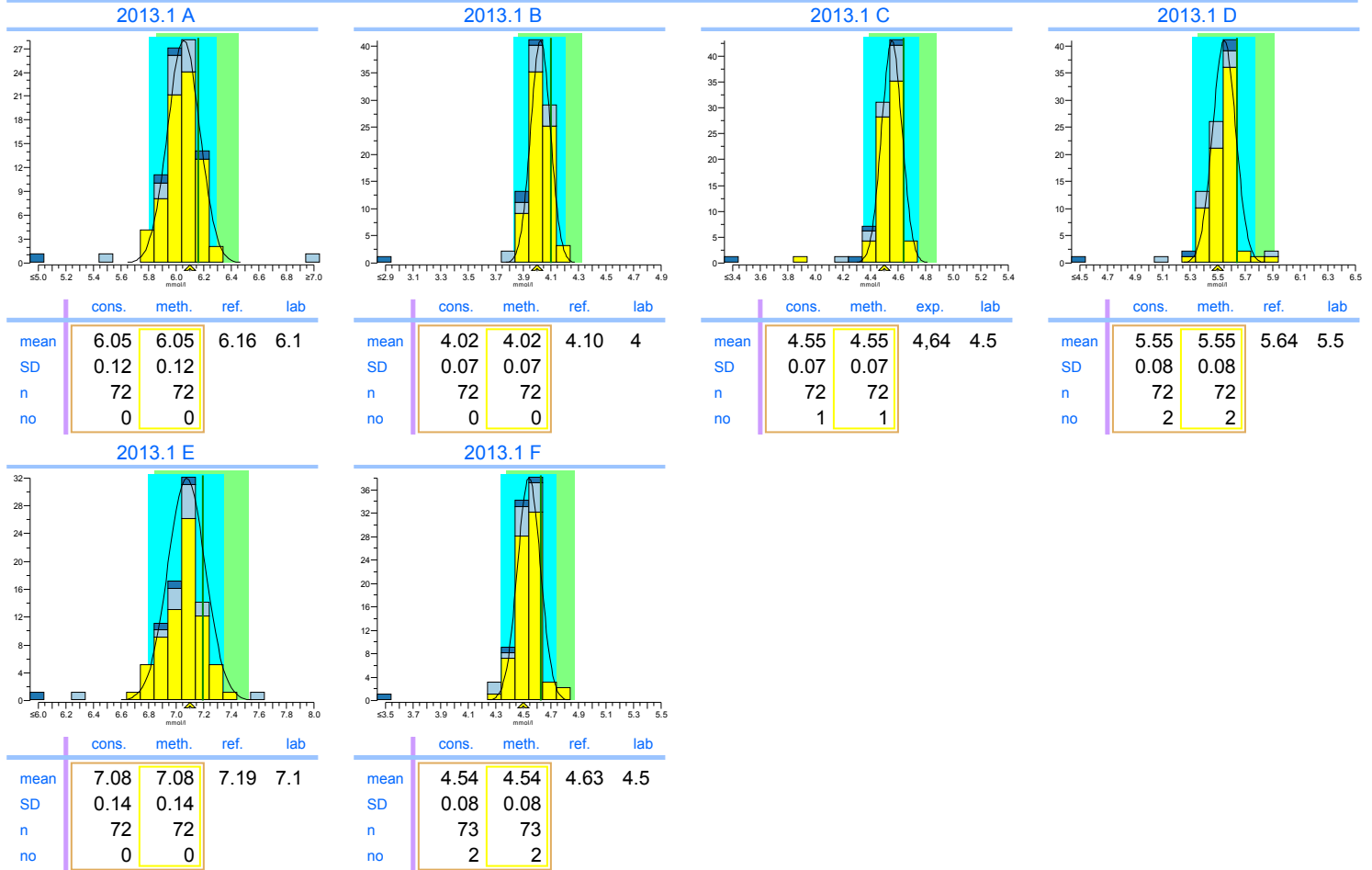
Potassium

units: mmol/l



	2013.1	cumulative
Trueness	-2.0%	-2.0%
Precision	0.56%	0.56%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 <span style="border: 1px solid green; padding: 2px;">2</span>	6.0 <span style="border: 1px solid green; padding: 2px;">2</span>
Sigma-SA	4.5	4.5
Score pictogram		
Regression line	$-0.18 + 1.014 \cdot x$	$-0.18 + 1.014 \cdot x$

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



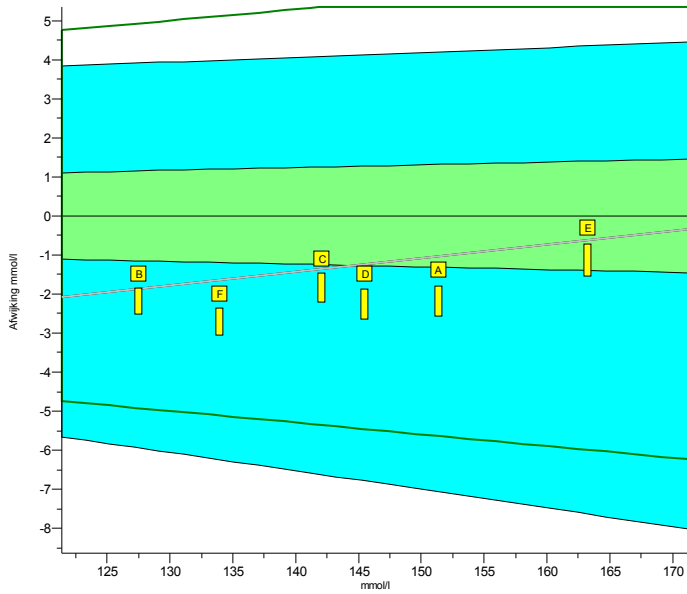
Legend

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

# INPUTS 2013.1

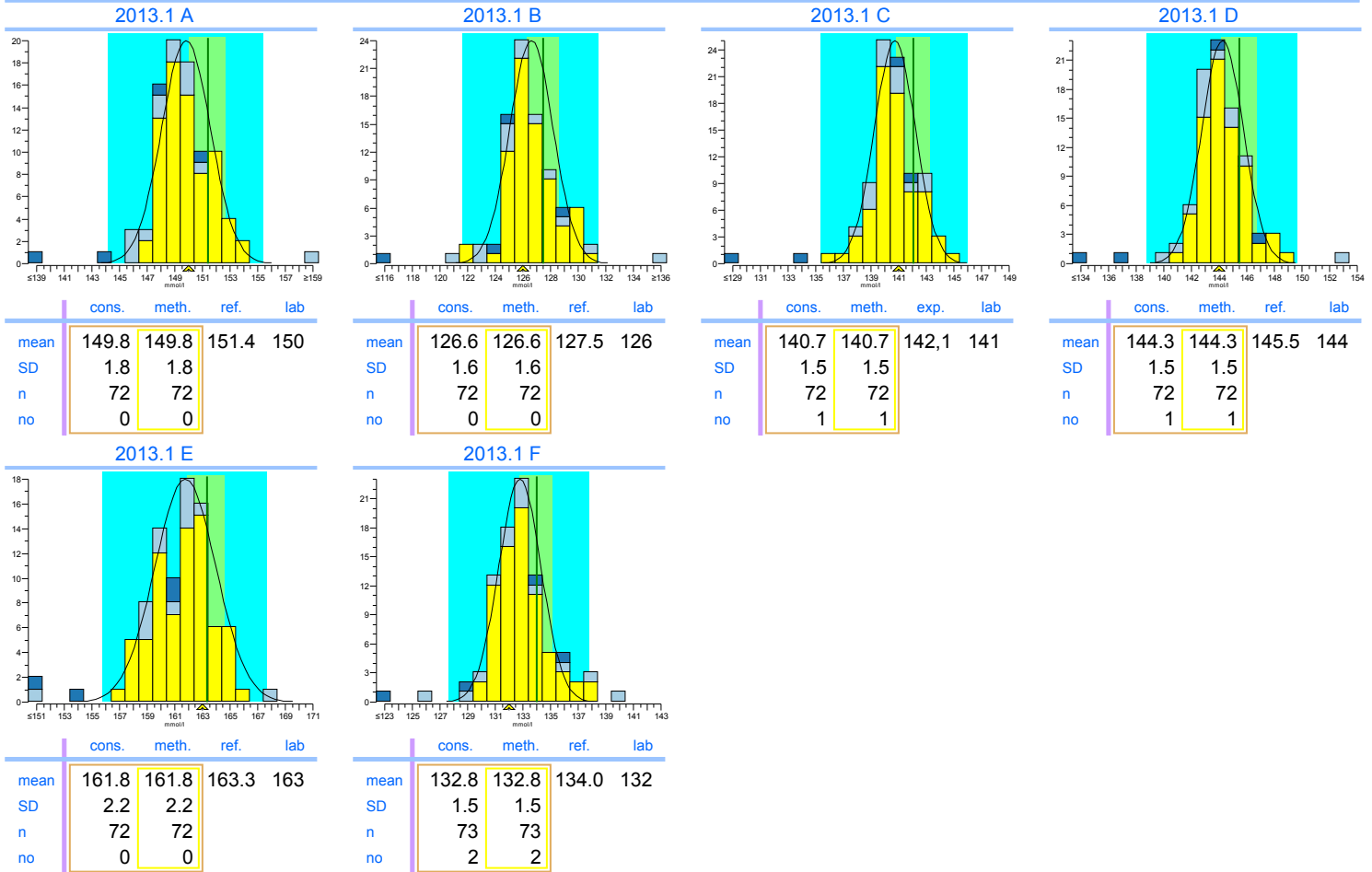
## Sodium

units: mmol/l



	2013.1	cumulative
Trueness	-0.90%	-0.90%
Precision	0.22%	0.22%
Number	6	6
Outliers	0	0
Sigma-TE	0.6	0.6
Sigma-SA	6.0 <span style="border: 1px solid black; padding: 0 2px;">2</span>	6.0 <span style="border: 1px solid black; padding: 0 2px;">2</span>
Score pictogram		
Regression line	$-6.3 + 1.035 \cdot x$	$-6.3 + 1.035 \cdot x$

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



Legend

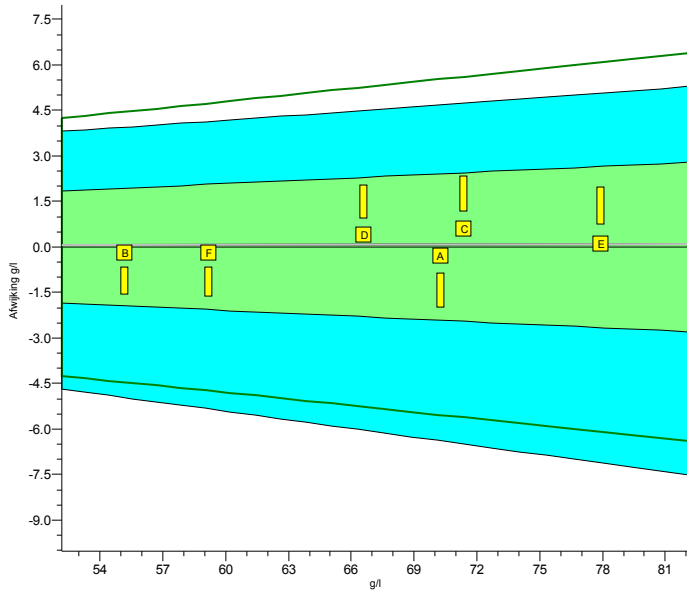
- ISE indirect (with predilution)
- ISE direct (no predilution)
- Other methods





# INPUTS 2013.1

Total Protein

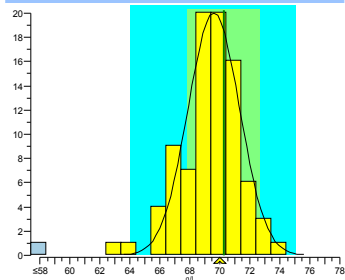
units: g/l



	2013.1	cumulative
Trueness	+0.10%	+0.10%
Precision	0.67%	0.67%
Number	6	6
Outliers	0	0
Sigma-TE	5.4	5.4
Sigma-SA	6.0 <span style="background-color: green; color: white; padding: 2px;">2</span>	6.0 <span style="background-color: green; color: white; padding: 2px;">2</span>
Score pictogram		
Regression line	$0.0 + 1.001 \cdot x$	$0.0 + 1.001 \cdot x$

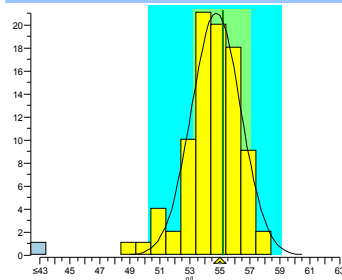
Consensus group Biureet  
Method Biurete, automatic

2013.1 A



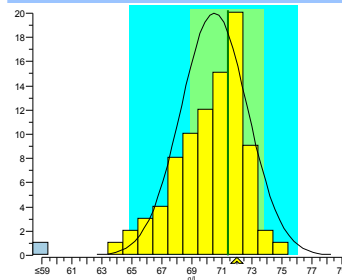
	cons.	meth.	ref.	lab
mean	69.6	69.6	70.3	70
SD	1.7	1.7		
n	88	88		
no	2	2		

2013.1 B



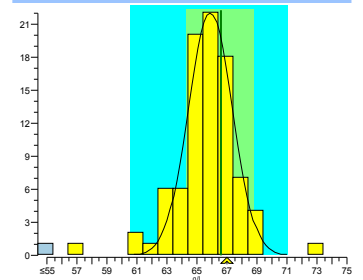
	cons.	meth.	ref.	lab
mean	54.8	54.8	55.2	55
SD	1.6	1.6		
n	88	88		
no	1	1		

2013.1 C



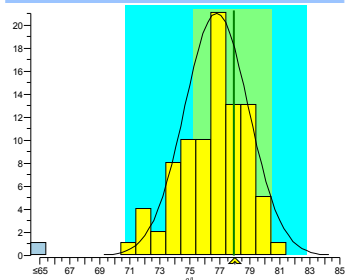
	cons.	meth.	exp.	lab
mean	70.5	70.5	71.4	72
SD	2.2	2.2		
n	87	87		
no	0	0		

2013.1 D



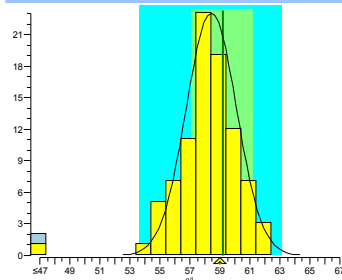
	cons.	meth.	ref.	lab
mean	65.9	65.9	66.6	67
SD	1.5	1.5		
n	88	88		
no	4	4		

2013.1 E



	cons.	meth.	ref.	lab
mean	76.8	76.8	77.9	78
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	59
SD	1.7	1.7		
n	89	89		
no	1	1		

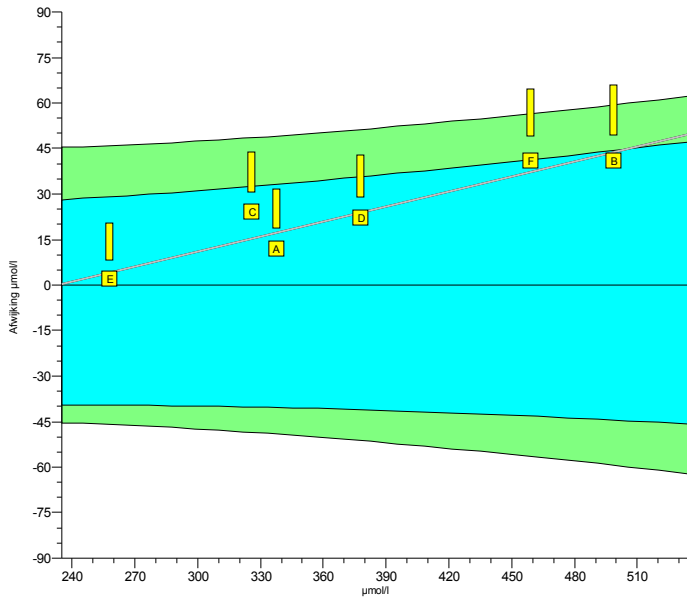
Legend

  Biurete, automatic      Other methods

# INPUTS 2013.1

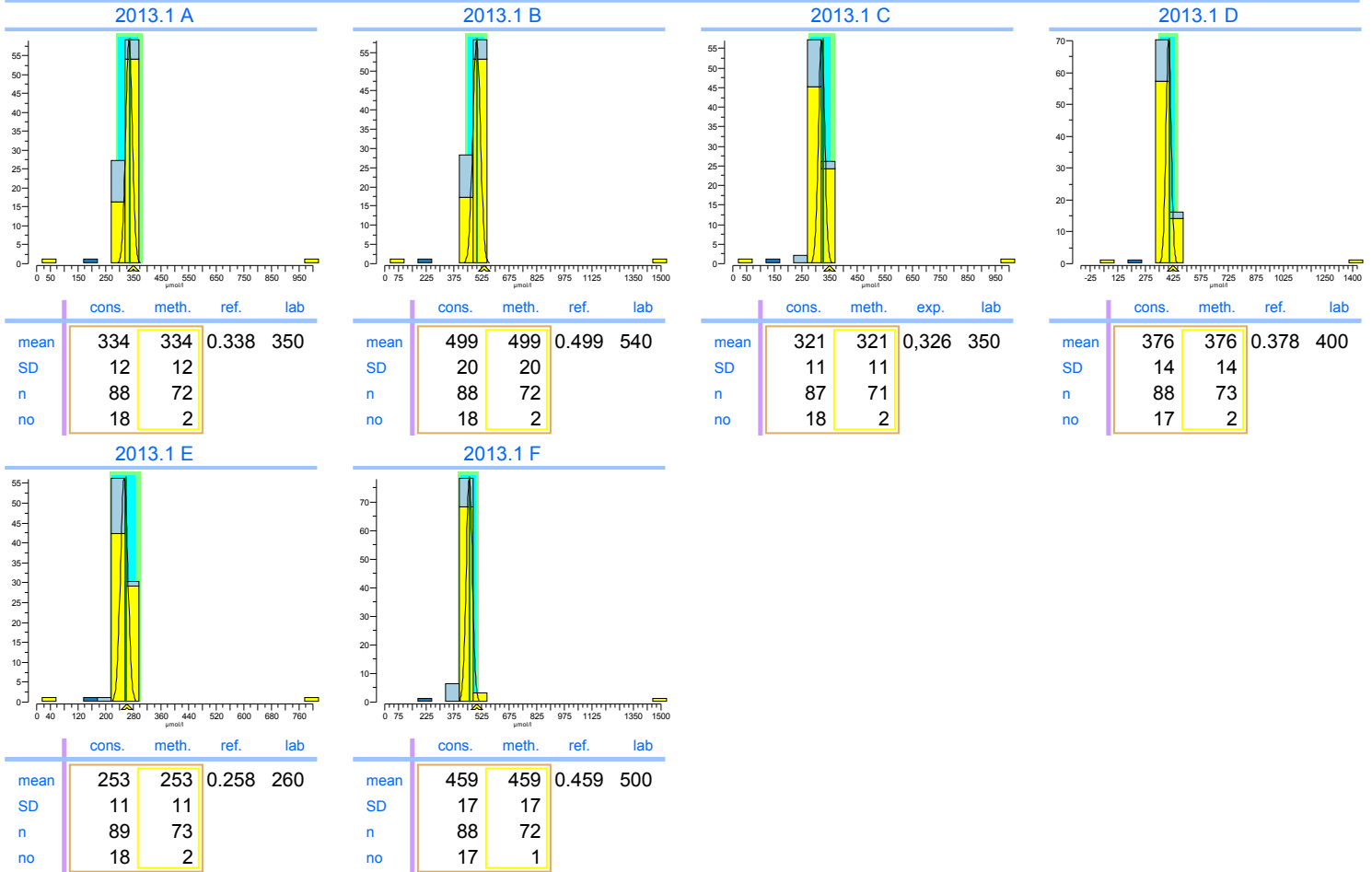
## Urate

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



	2013.1	cumulative
Trueness	+6.3%	+6.3%
Precision	1.5%	1.5%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 <span style="border: 1px solid black; padding: 0 2px;">2</span>	6.0 <span style="border: 1px solid black; padding: 0 2px;">2</span>
Sigma-SA	3.7	3.7
Score pictogram		
Regression line	$0 + 1.164.x$	$0 + 1.164.x$

Consensus group	Colorimetrisch
Method	Uricase, colorim., automatic



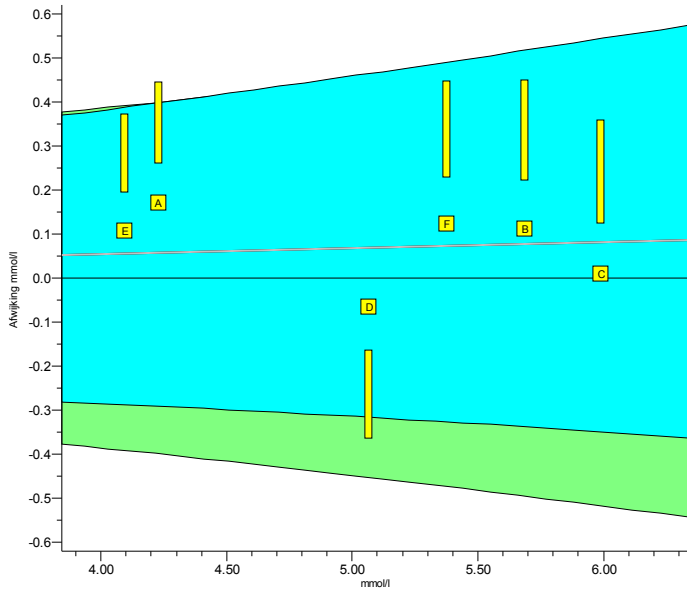
Legend

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

# INPUTS 2013.1

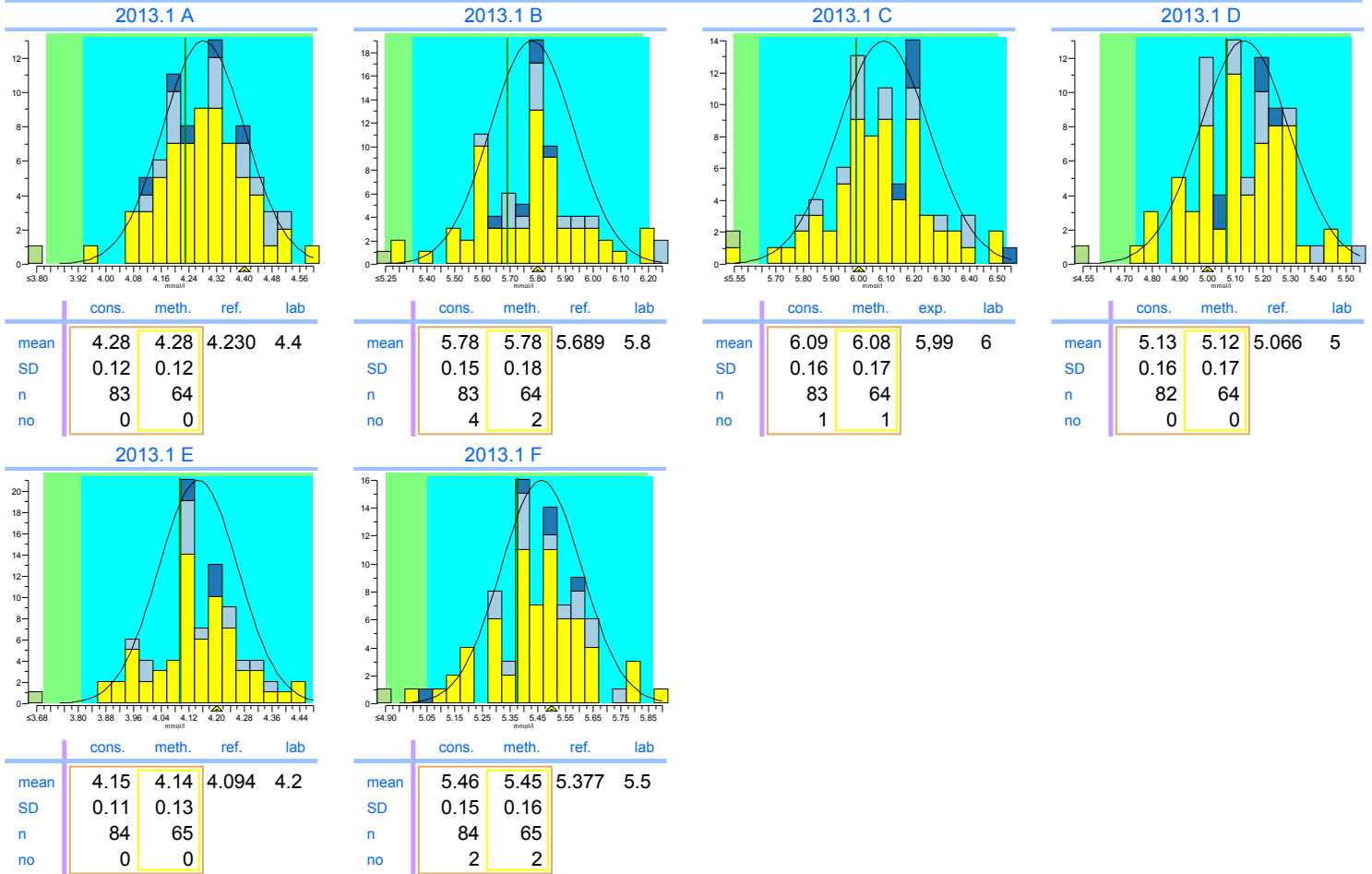
## Cholesterol

units: mmol/l



	2013.1	cumulative
Trueness	+1.5%	+1.5%
Precision	1.8%	1.8%
Number	6	6
Outliers	0	0
Sigma-TE	5.1 <span style="border: 1px solid black; padding: 0 2px;">2</span>	5.1 <span style="border: 1px solid black; padding: 0 2px;">2</span>
Sigma-SA	4.3	4.3
Score pictogram		
Regression line	<u>0.00 + 1.014.x</u>	<u>0.00 + 1.014.x</u>

Consensus group	Enzymatisch
Method	Enzymatic, automatic, discrete



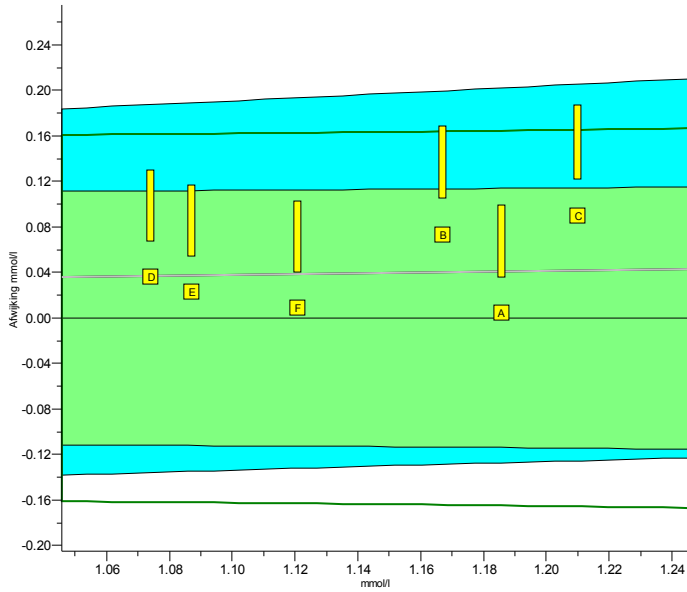
Legend

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

# INPUTS 2013.1

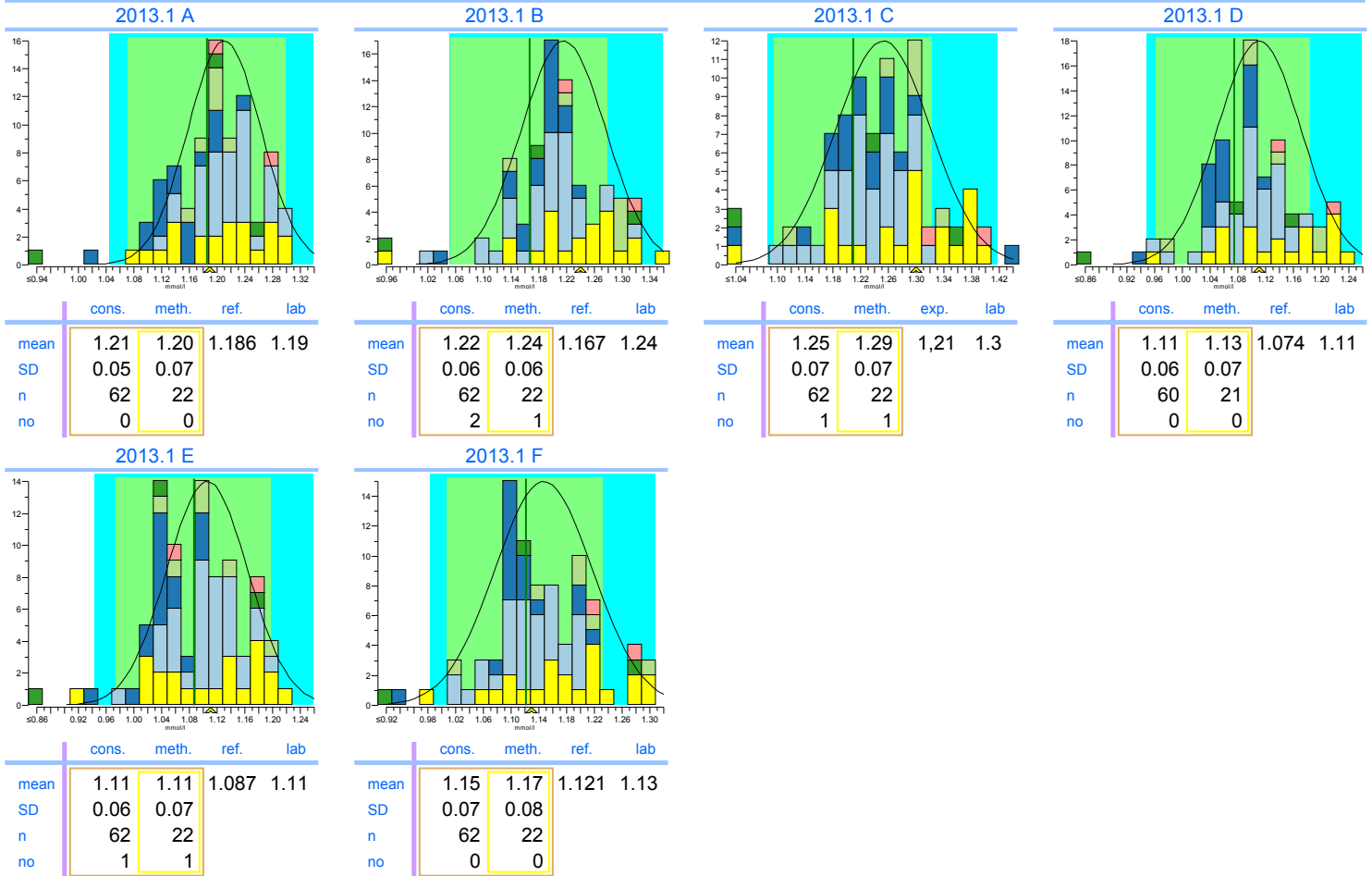
## HDL-Cholesterol

units: mmol/l



	2013.1	cumulative
Trueness	+3.4%	+3.4%
Precision	2.2%	2.2%
Number	6	6
Outliers	0	0
Sigma-TE	3.7	3.7
Sigma-SA	5.7 <span style="background-color: green; color: white; padding: 2px;">2</span>	5.7 <span style="background-color: green; color: white; padding: 2px;">2</span>
Score pictogram		
Regression line	<u>0.00 + 1.035.x</u>	<u>0.00 + 1.035.x</u>

Consensus group	Direct
Method	Accelerator Selective Detergent ("Ultra HDL")



Legend

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