

# INPUTS 2013.1

Survey	INPUTS 2013.1		
Start date	January, 1 2013		
Supervisor	dr. C. Weykamp MCA Laboratory Streekziekenhuis Koningin Beatrix Winterswijk		
Subscriptions	93		
Result sets	91		
Scores	Your score	MAP	reported
Quantitative	28	38	



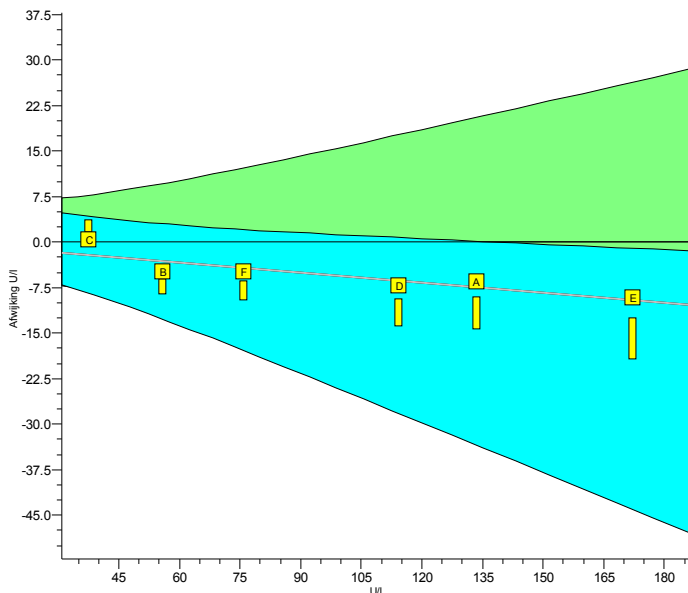
# INPUTS 2013.1

Analyte		Trueness				Precision		Performance			
		your mean	ref.	cons.	SDI	your prec.	SDbl	this survey	PS	cumulative PSc	
ALAT	U/l	92.8	98.2	86.9	10.1	4.0	2.7		2		2
Alk. Phosphatase	U/l	483	188	173	15	9	2		0		0
Amylase	U/l	154	187	182	8	1	2		1		1
ASAT	U/l	70.2	78.6	71.5	6.8	0.9	1.6		1		1
Calcium	mmol/l	2.49	2.46	2.47	0.07	0.04	0.02		2		2
Chloride	mmol/l	101.3	100.0	100.8	1.8	1.5	0.9		1		1
CK	U/l	246	244	253	18	2	3		2		2
Creatinine	µmol/l	150.0	149.1	150.9	7.1	4.0	4.2		1		1
eGFR (F, 55, white)	ml/min/1,73m <sup>2</sup>	37.7	37.6	37.6	2.7	0.6	1.5		2		2
Gamma-GT	U/l	86.7	87.7	86.3	8.6	1.0	1.2		2		2
Glucose	mmol/l	14.10	13.93	14.19	0.44	0.37	0.18		1		1
LD	U/l	985	532	523	33	17	9		0		0
Magnesium	mmol/l	1.26	1.22	1.23	0.04	0.02	0.02		2		2
Potassium	mmol/l	5.33	5.39	5.30	0.10	0.05	0.04		2		2
Sodium	mmol/l	142.5	144.0	142.7	1.7	1.5	0.9		1		1
Total Protein	g/l	65.5	66.8	66.0	1.8	0.8	0.7		2		2
Urate	µmol/l	380	376	374	15	3	4		2		2
Cholesterol	mmol/l	5.30	5.07	5.15	0.14	0.04	0.05		2		2
HDL-Cholesterol	mmol/l	1.13	1.14	1.13	0.03	0.01	0.02		2		2
<b>Total :</b>									<b>28</b>		<b>28</b>

# INPUTS 2013.1

ALAT

units: U/I



	2013.1	cumulative
Trueness	-5.5%	-5.5%
Precision	4.6%	4.6%
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	5.1	5.1
Score pictogram		
Regression line	$0.0 + 0.945.x$	$0.0 + 0.945.x$

Consensus group IFCC traceerbaar  
Method IFCC traceable

	2013.1 A				2013.1 B				2013.1 C				2013.1 D			
	cons.	meth.	ref.	lab	cons.	meth.	ref.	lab	cons.	meth.	exp.	lab	cons.	meth.	ref.	lab
mean	122.7	122.7	133.5	127	50.7	50.7	55.9	51	35.5	35.5	37,6	38	100.9	100.9	114.2	107
SD	11.8	11.8			3.9	3.9			3.6	3.6			9.4	9.4		
n	75	75			76	76			76	76			76	76		
no	1	1			3	3			7	7			3	3		

	2013.1 E				2013.1 F			
	cons.	meth.	ref.	lab	cons.	meth.	ref.	lab
mean	146	146	172.2	163	66.1	66.1	75.9	71
SD	18	18			5.3	5.3		
n	77	77			77	77		
no	1	1			3	3		

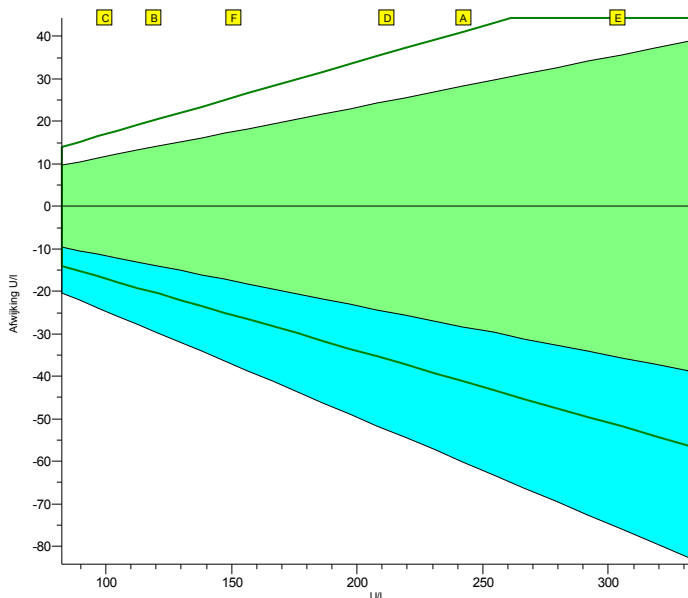
Legend

IFCC traceable     IFCC non-traceable

# INPUTS 2013.1

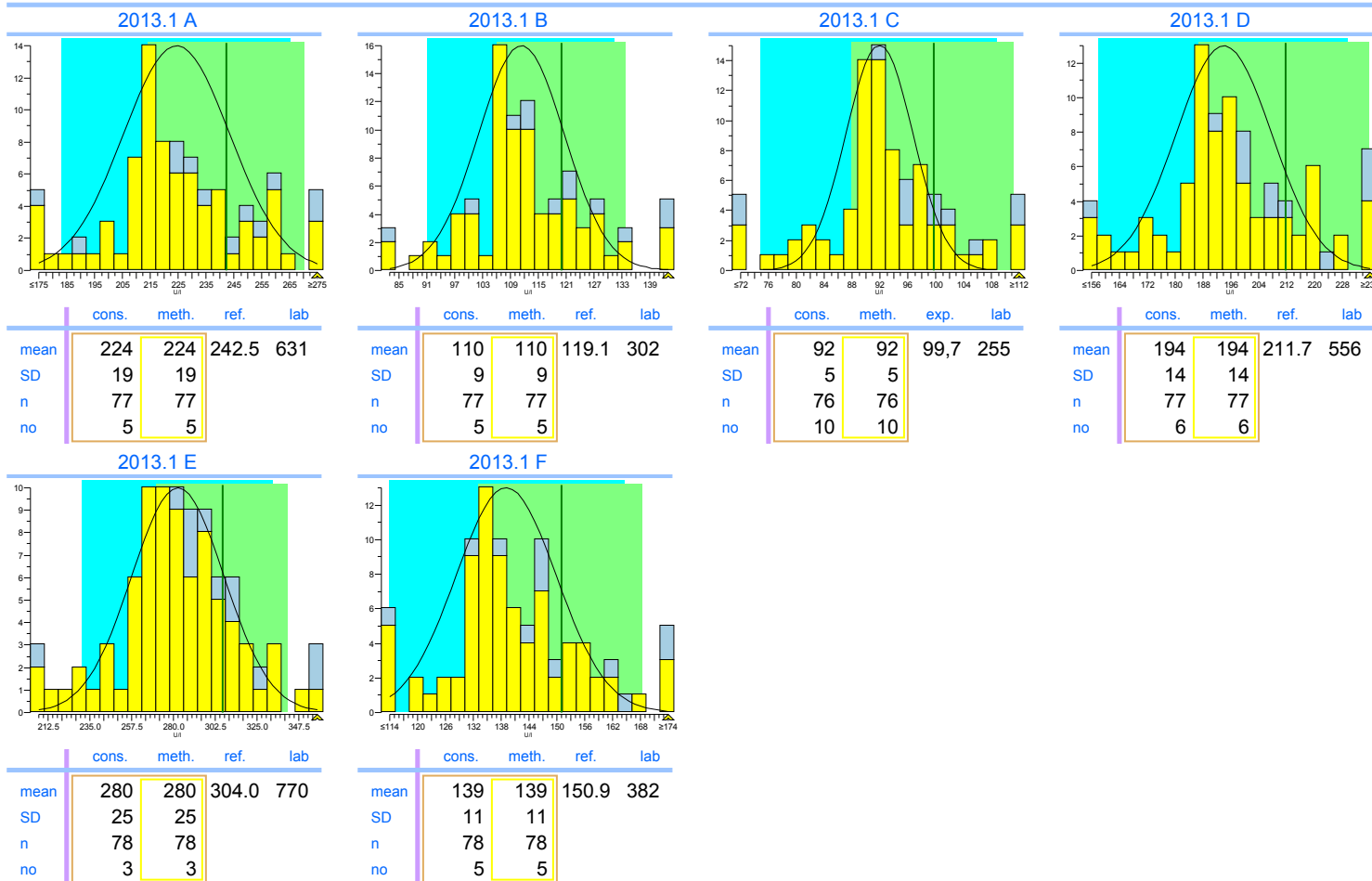
## Alk. Phosphatase

units: U/l



	2013.1	cumulative
Trueness	+157%	+157%
Precision	5.3%	5.3%
Number	6	6
Outliers	0	0
Sigma-TE	-3.0	-3.0
Sigma-SA	-3.0 <span style="border: 1px solid red; padding: 2px;">0</span>	-3.0 <span style="border: 1px solid red; padding: 2px;">0</span>
Score pictogram		
Regression line	$0 + 2.568.x$	$0 + 2.568.x$

Consensus group IFCC traceerbaar  
Method IFCC traceable



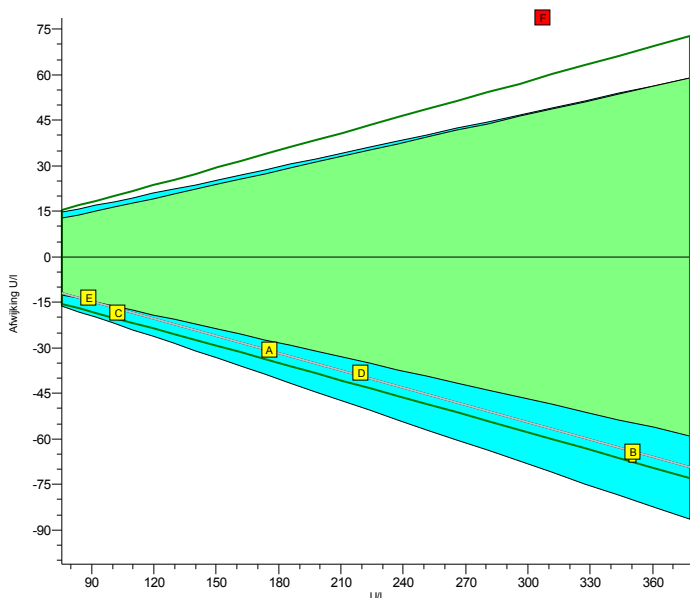
Legend

IFCC traceable     IFCC non-traceable

# INPUTS 2013.1

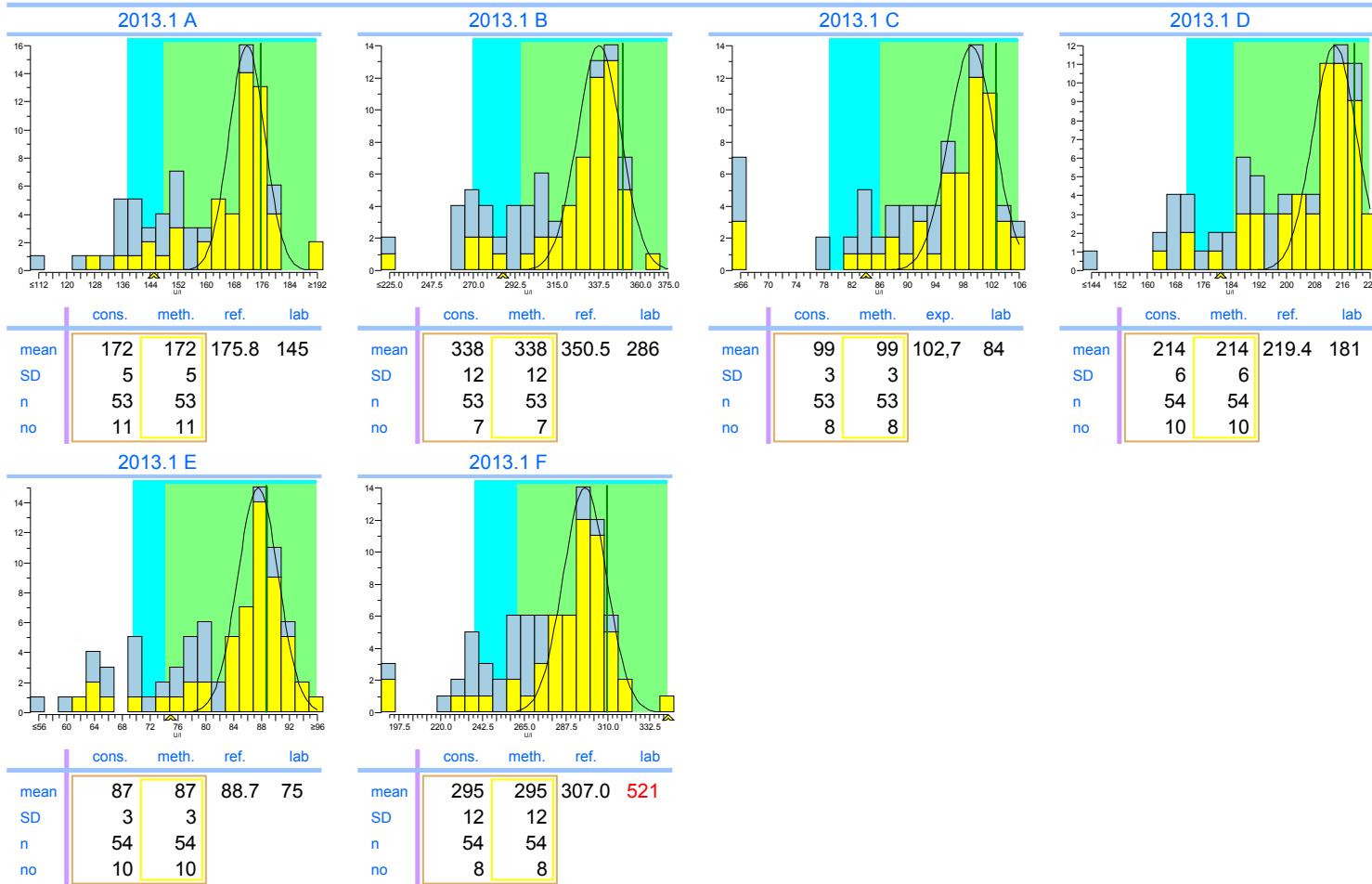
## Amylase

units: U/l

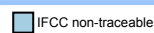


	2013.1	cumulative
Trueness	-18%	-18%
Precision	0.28%	0.28%
Number	6	6
Outliers	1	1
Sigma-TE	-2.0	-2.0
Sigma-SA	3.0	3.0
Score pictogram		
Regression line	$2 + 0.811 \cdot x$	$2 + 0.811 \cdot x$

Consensus group IFCC traceerbaar  
Method IFCC traceable



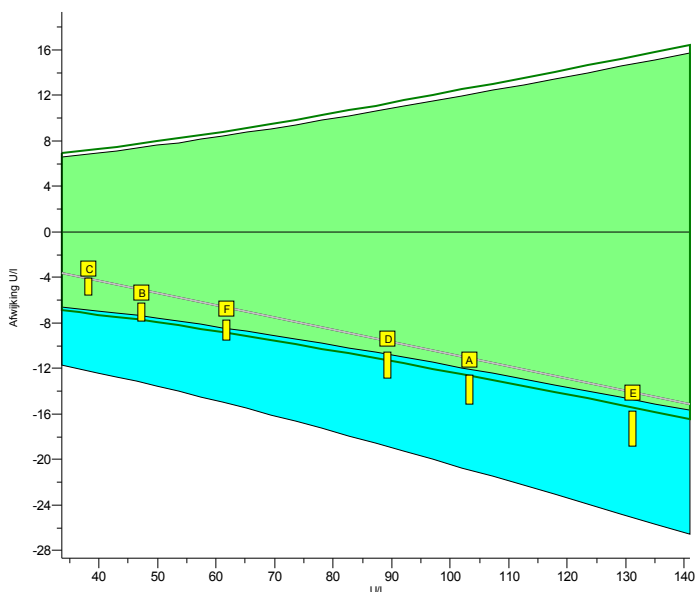
Legend



 IFCC traceable  IFCC non-traceable

# INPUTS 2013.1

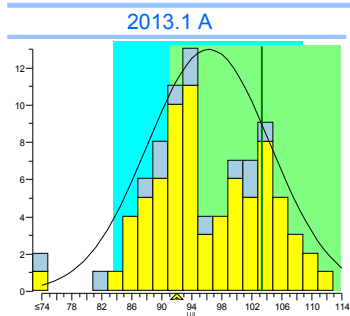
ASAT

units: U/I

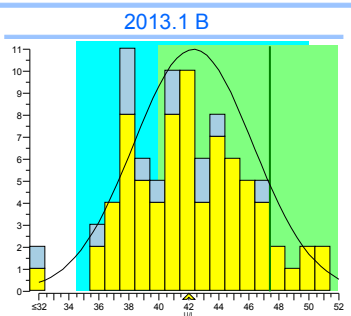


	2013.1	cumulative
Trueness	-11%	-11%
Precision	1.2%	1.2%
Number	6	6
Outliers	0	0
Sigma-TE	2.9	2.9
Sigma-SA	3.4 <span style="background-color: green; color: white;">1</span>	3.4 <span style="background-color: green; color: white;">1</span>
Score pictogram		
Regression line	$0.0 + 0.892.x$	$0.0 + 0.892.x$

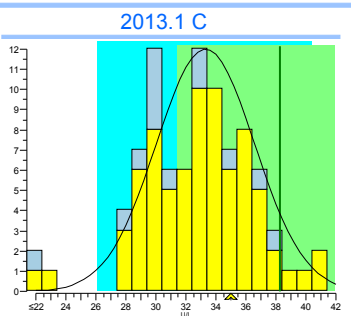
Consensus group IFCC traceerbaar  
Method IFCC traceable



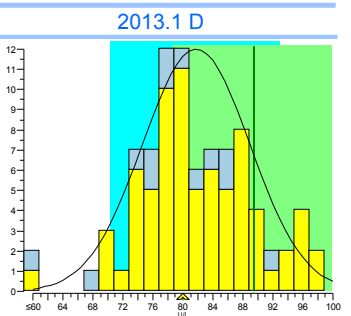
	cons.	meth.	ref.	lab
mean	96.3	96.3	103.3	92
SD	8.2	8.2		
n	75	75		
no	1	1		



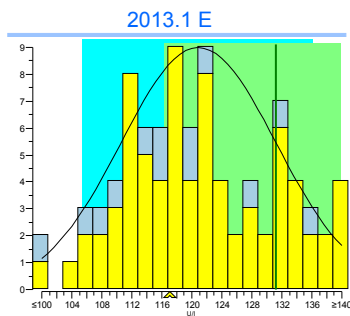
	cons.	meth.	ref.	lab
mean	42.3	42.3	47.4	42
SD	4.0	4.0		
n	75	75		
no	1	1		



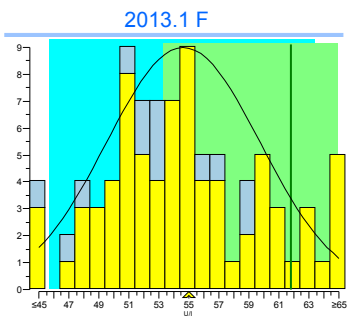
	cons.	meth.	exp.	lab
mean	33.3	33.3	38,3	35
SD	3.3	3.3		
n	75	75		
no	2	2		



	cons.	meth.	ref.	lab
mean	81.8	81.8	89.4	80
SD	7.2	7.2		
n	74	74		
no	1	1		



	cons.	meth.	ref.	lab
mean	120.9	120.9	131.2	117
SD	10.3	10.3		
n	76	76		
no	1	1		



	cons.	meth.	ref.	lab
mean	54.6	54.6	61.8	55
SD	5.1	5.1		
n	76	76		
no	2	2		

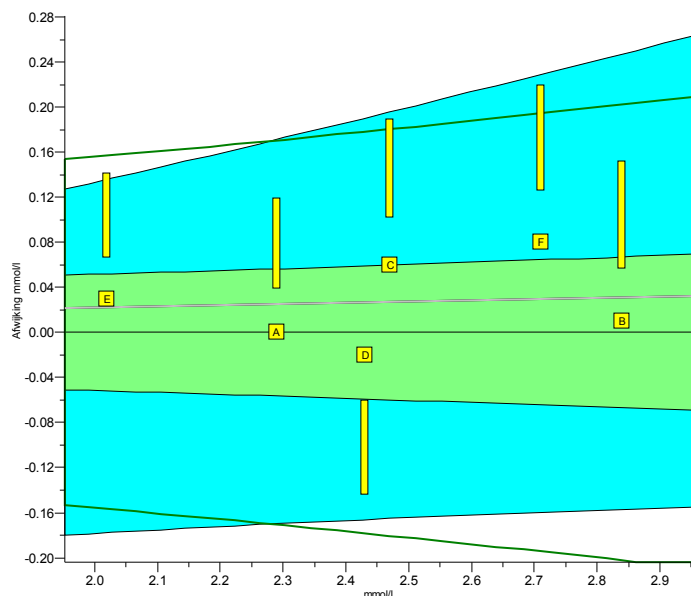
Legend

IFCC traceable     IFCC non-traceable

# INPUTS 2013.1

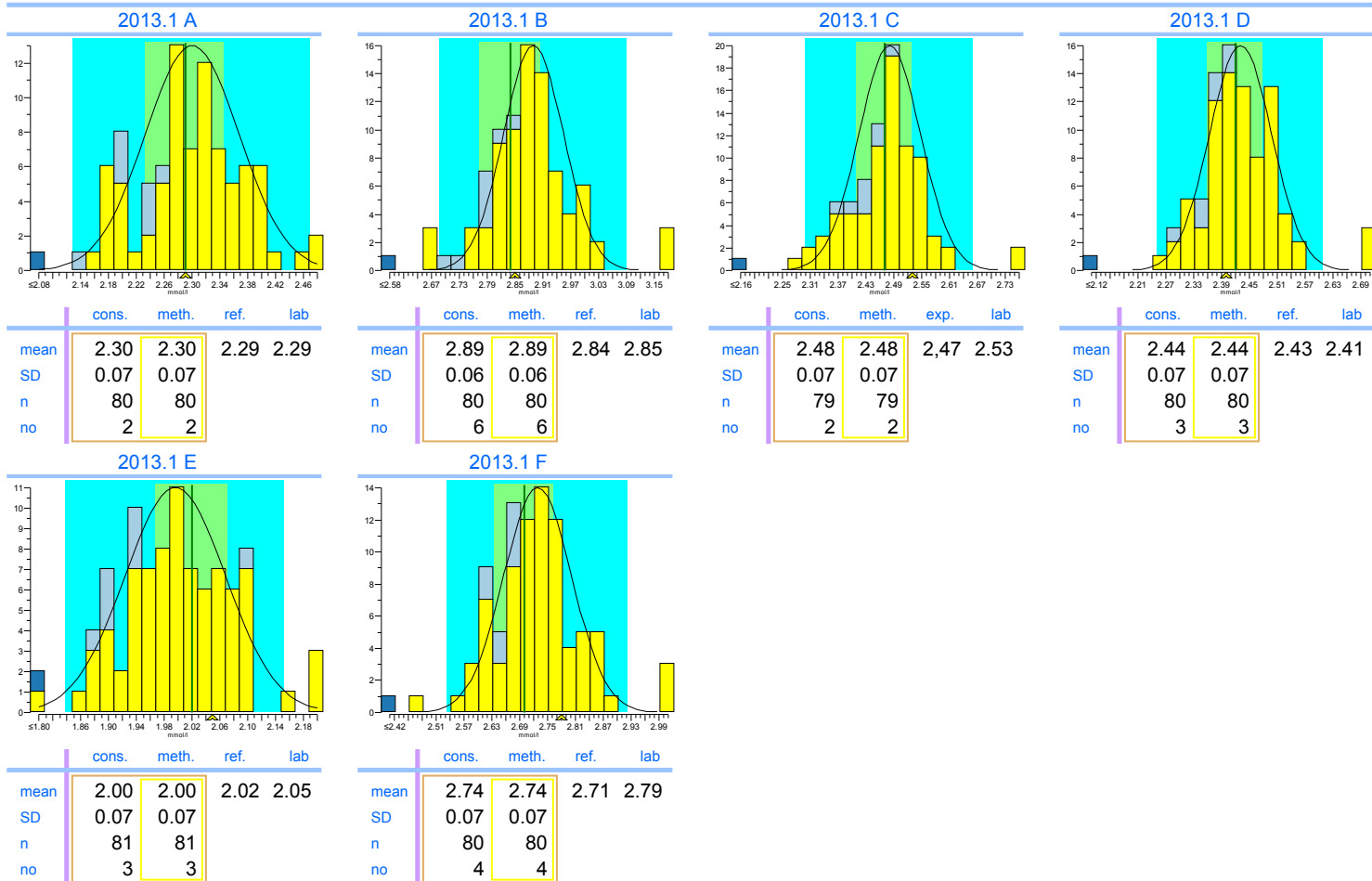
## Calcium

units: mmol/l



	2013.1	cumulative
Trueness	+1.1%	+1.1%
Precision	1.8%	1.8%
Number	6	6
Outliers	0	0
Sigma-TE	1.6	1.6
Sigma-SA	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>
Score pictogram		
Regression line	$0.00 + 1.011 \cdot x$	$0.00 + 1.011 \cdot x$

Consensus group	Colorimetrisch
Method	Colorimetric, automatic, discrete



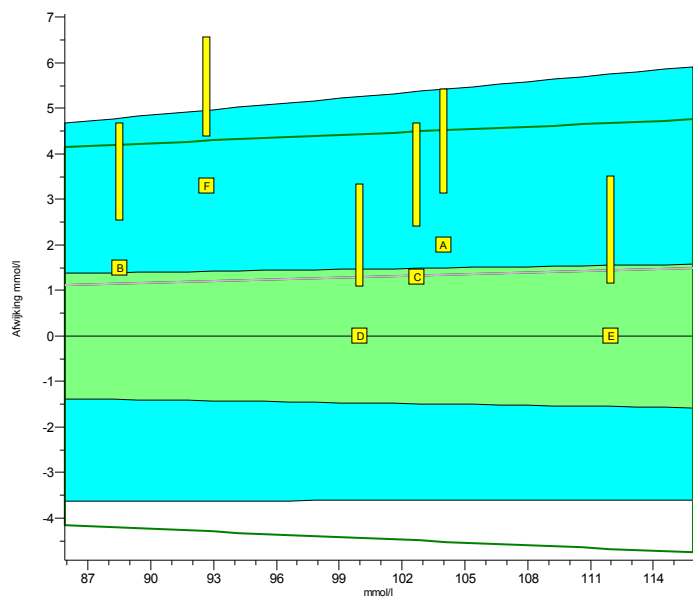
Legend

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

# INPUTS 2013.1

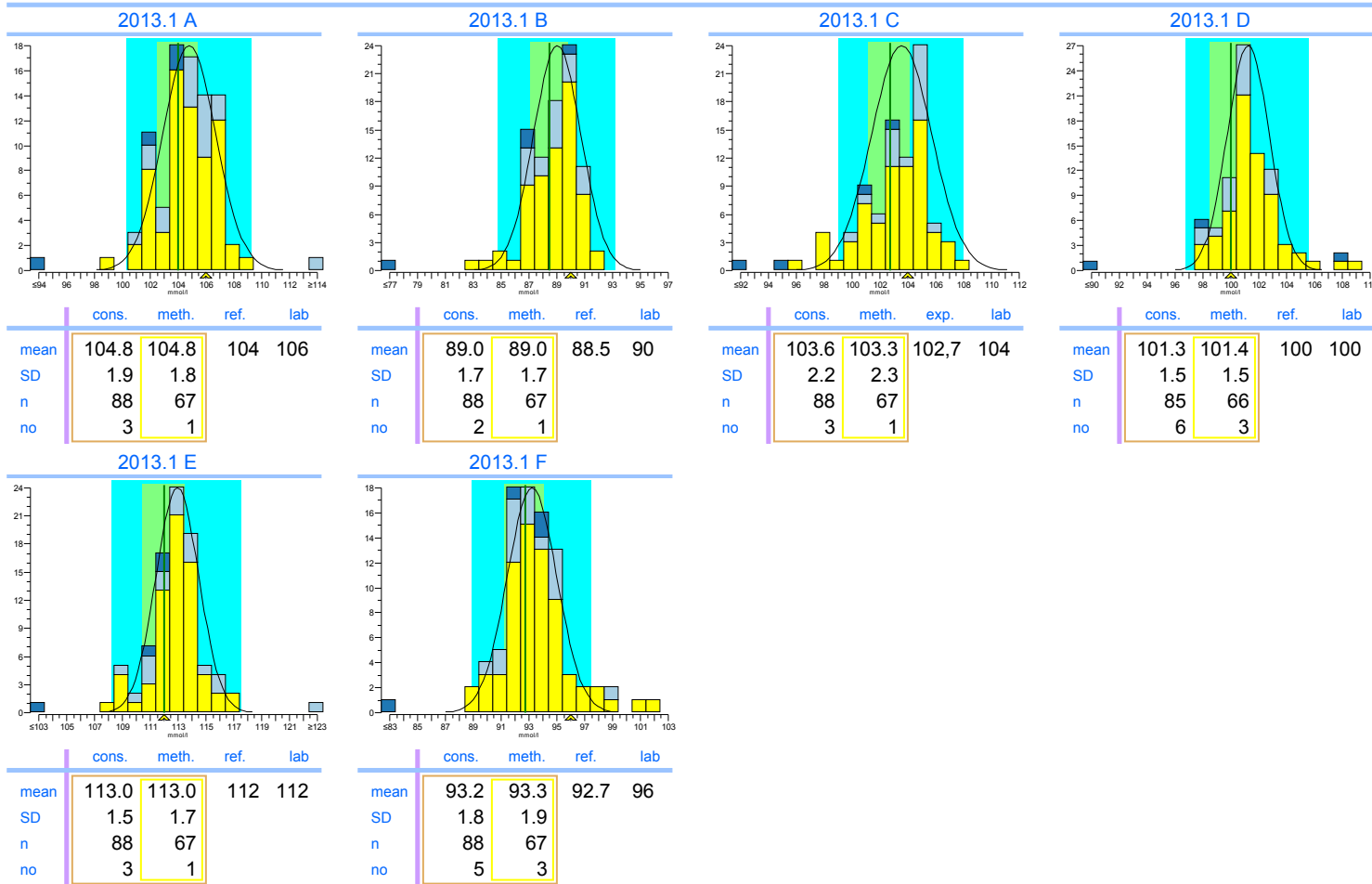
Chloride

units: mmol/l



	2013.1	cumulative
Trueness	+1.4%	+1.4%
Precision	1.5%	1.5%
Number	6	6
Outliers	0	0
Sigma-TE	0.9	0.9
Sigma-SA	4.2	4.2
Score pictogram		
Regression line	$0.0 + 1.013 \cdot x$	$0.0 + 1.013 \cdot x$

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



Legend

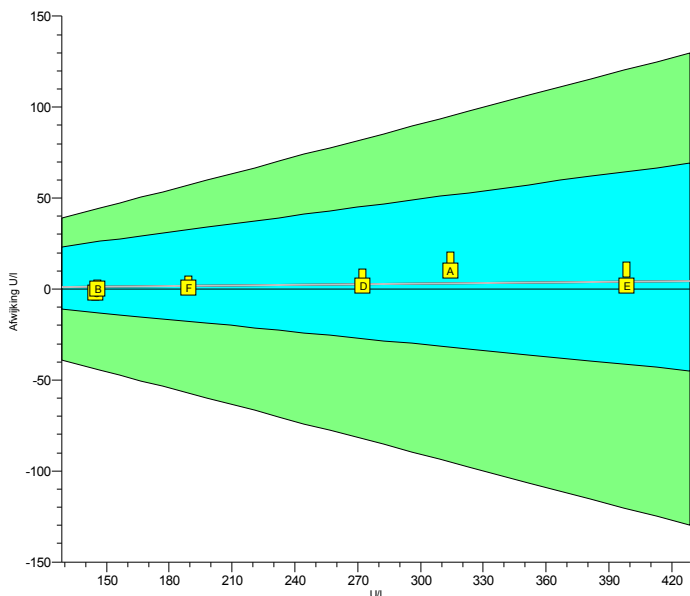






# INPUTS 2013.1

CK

units: U/l



	2013.1	cumulative
Trueness	+0.82%	+0.82%
Precision	0.88%	0.88%
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	6.0	6.0
Score pictogram		
Regression line	$0 + 1.010 \cdot x$	$0 + 1.010 \cdot x$

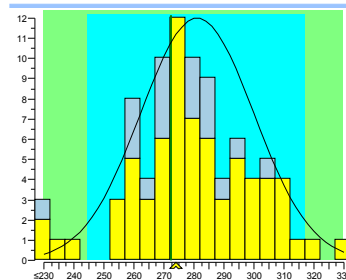
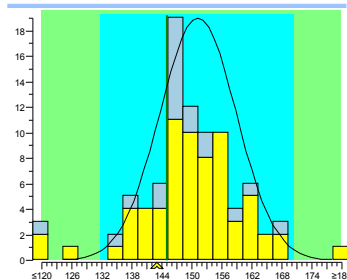
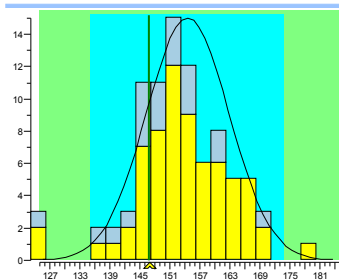
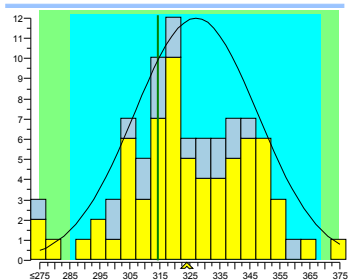
Consensus group IFCC traceerbaar  
Method IFCC traceable

2013.1 A

2013.1 B

2013.1 C

2013.1 D



	cons.	meth.	ref.	lab
mean	327	327	314.3	324
SD	21	21		
n	68	68		
no	2	2		

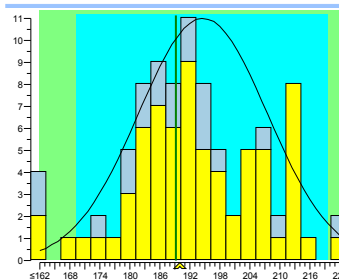
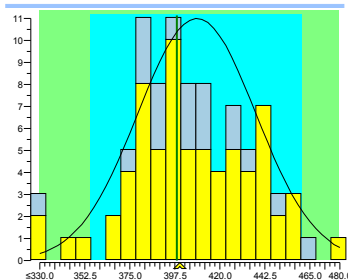
	cons.	meth.	ref.	lab
mean	154	154	145.8	146
SD	8	8		
n	67	67		
no	3	3		

	cons.	meth.	exp.	lab
mean	151	151	145	143
SD	8	8		
n	68	68		
no	4	4		

	cons.	meth.	ref.	lab
mean	281	281	272.3	274
SD	19	19		
n	69	69		
no	2	2		

2013.1 E

2013.1 F



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

	cons.	meth.	ref.	lab
mean	195	195	189.2	190
SD	13	13		
n	69	69		
no	2	2		

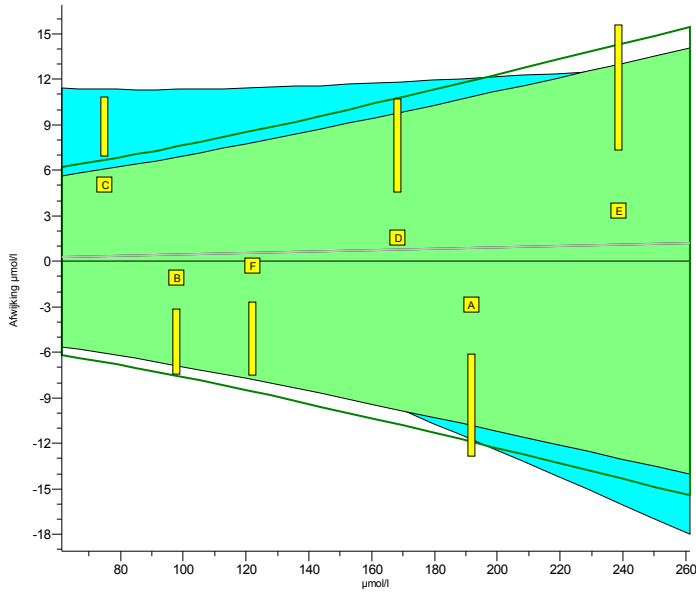
Legend

IFCC traceable     IFCC non-traceable

# INPUTS 2013.1

## Creatinine

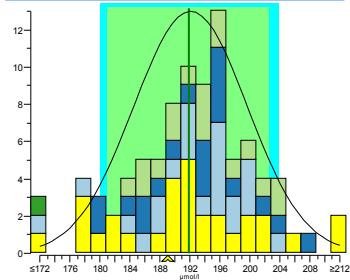
units:  $\mu\text{mol/l}$



	2013.1	cumulative
Trueness	+0.61%	+0.61%
Precision	2.6%	2.6%
Number	6	6
Outliers	0	0
Sigma-TE	3.6	3.6
Sigma-SA	4.0	4.0
Score pictogram		
Regression line	$0.0 + 1.005 \cdot x$	$0.0 + 1.005 \cdot x$

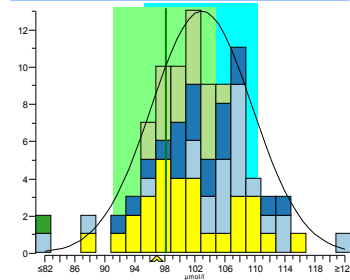
Consensus group: Jaffe  
Method: Alk. Picrate, kinetic with compensation

2013.1 A



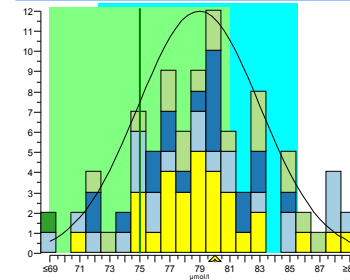
	cons.	meth.	ref.	lab
mean	192.1	190.4	191.9	189
SD	7.6	9.4		
n	71	30		
no	2	1		

2013.1 B



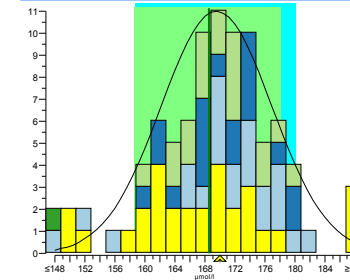
	cons.	meth.	ref.	lab
mean	102.9	101.0	98.11	97
SD	6.7	6.6		
n	71	30		
no	2	0		

2013.1 C



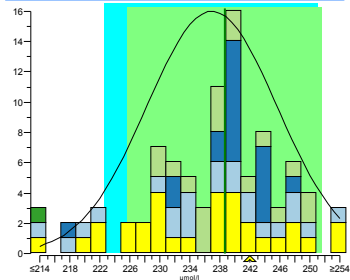
	cons.	meth.	exp.	lab
mean	78.9	79.1	75	80
SD	4.1	3.4		
n	71	30		
no	1	1		

2013.1 D



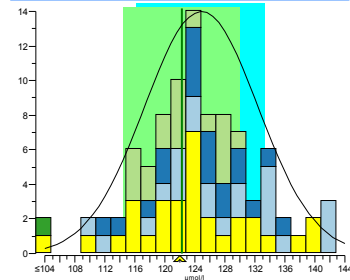
	cons.	meth.	ref.	lab
mean	169.5	167.6	168.5	170
SD	7.4	10.0		
n	70	30		
no	1	0		

2013.1 E



	cons.	meth.	ref.	lab
mean	236.9	234.2	238.7	242
SD	8.7	9.4		
n	72	31		
no	3	2		

2013.1 F



	cons.	meth.	ref.	lab
mean	124.9	123.5	122.3	122
SD	7.6	8.3		
n	72	31		
no	0	0		

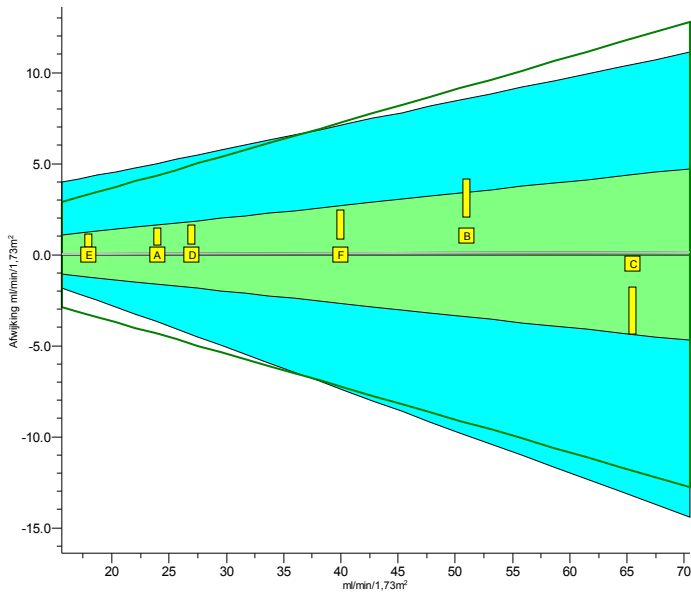
Legend

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

# INPUTS 2013.1

eGFR (F, 55, white)

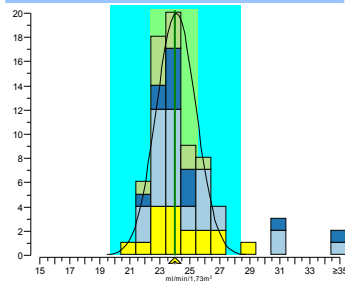
units: ml/min/1,73m<sup>2</sup>



	2013.1	cumulative
Trueness	+0.22%	+0.22%
Precision	1.6%	1.6%
Number	6	6
Outliers	0	0
Sigma-TE	4.6	4.6
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	$0.0 + 1.002 \cdot x$	$0.0 + 1.002 \cdot x$

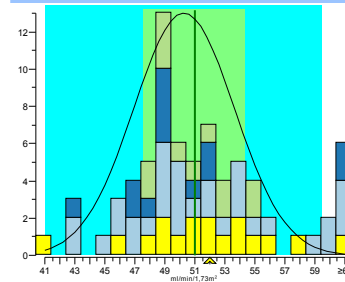
Consensus group Jaffe  
Method Alk. Picrate, kinetic with compensation

2013.1 A



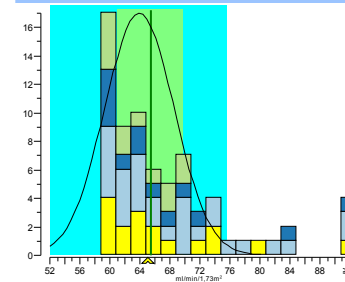
	cons.	meth.	ref.	lab
mean	24.1	24.2	24	24
SD	1.3	1.7		
n	30	17		
no	3	1		

2013.1 B



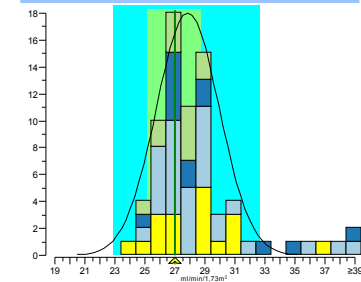
	cons.	meth.	ref.	lab
mean	50.3	51.2	51	52
SD	3.4	4.1		
n	30	17		
no	3	1		

2013.1 C



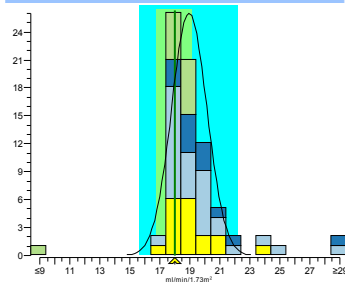
	cons.	meth.	exp.	lab
mean	63.9	64.7	65,5	65
SD	4.6	4.8		
n	30	17		
no	4	2		

2013.1 D



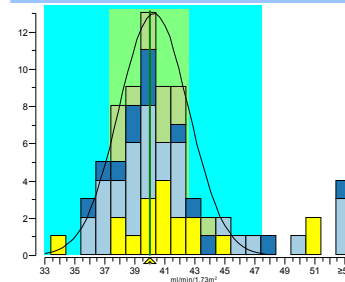
	cons.	meth.	ref.	lab
mean	27.9	28.0	27	27
SD	2.1	2.1		
n	31	18		
no	3	1		

2013.1 E



	cons.	meth.	ref.	lab
mean	19.0	18.9	18	18
SD	1.2	1.2		
n	31	18		
no	2	1		

2013.1 F



	cons.	meth.	ref.	lab
mean	40.3	40.5	40	40
SD	2.3	2.5		
n	31	18		
no	4	2		

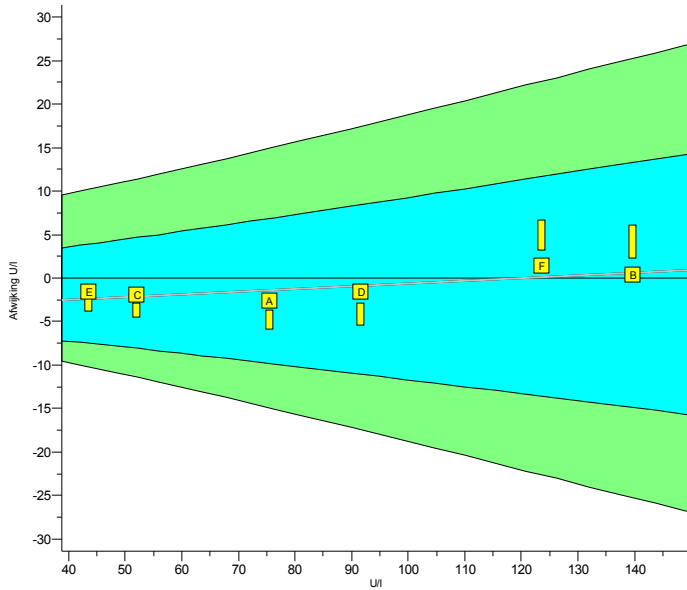
Legend



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

# INPUTS 2013.1

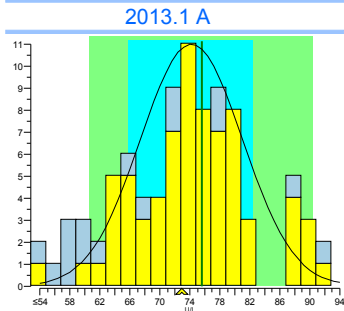
Gamma-GT

units: U/I

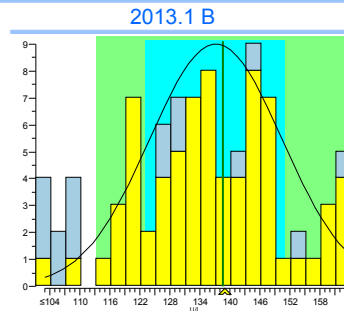


	2013.1	cumulative
Trueness	-1.2%	-1.2%
Precision	1.2%	1.2%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 <span style="border: 1px solid green; padding: 2px;">2</span>	6.0 <span style="border: 1px solid green; padding: 2px;">2</span>
Sigma-SA	6.0	6.0
Score pictogram		
Regression line	$-3.8 + 1.032 \cdot x$	$-3.8 + 1.032 \cdot x$

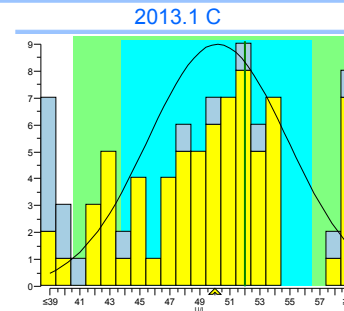
Consensus group IFCC traceerbaar  
Method IFCC traceable



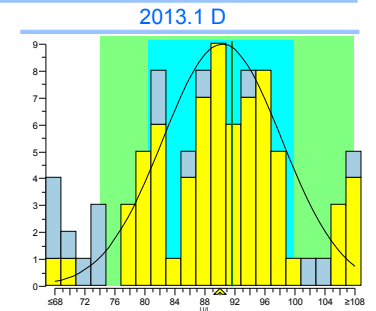
	cons.	meth.	ref.	lab
mean	74.2	74.2	75.6	73
SD	6.8	6.8		
n	72	72		
no	1	1		



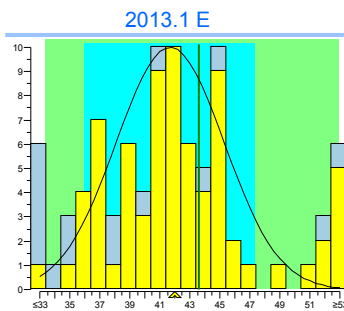
	cons.	meth.	ref.	lab
mean	138.2	138.2	139.7	140
SD	13.3	13.3		
n	72	72		
no	1	1		



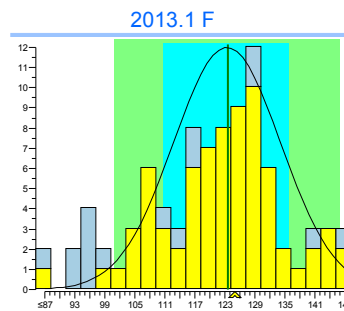
	cons.	meth.	exp.	lab
mean	50.2	50.2	52	50
SD	4.6	4.6		
n	72	72		
no	2	2		



	cons.	meth.	ref.	lab
mean	90.3	90.3	91.6	90
SD	7.9	7.9		
n	71	71		
no	1	1		



	cons.	meth.	ref.	lab
mean	41.7	41.7	43.6	42
SD	3.6	3.6		
n	73	73		
no	6	6		



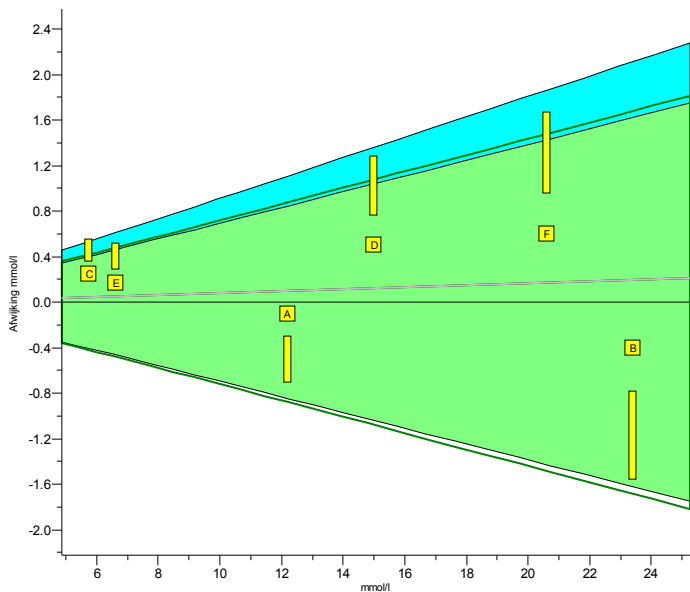
	cons.	meth.	ref.	lab
mean	123.4	123.4	123.6	125
SD	10.7	10.7		
n	73	73		
no	1	1		

Legend  
 IFCC traceable  
 IFCC non-traceable

# INPUTS 2013.1

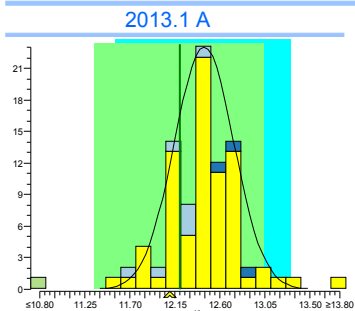
Glucose

units: mmol/l

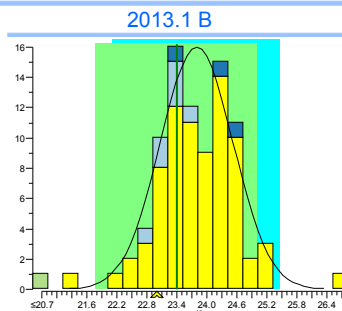


	2013.1	cumulative
Trueness	+1.2%	+1.2%
Precision	2.6%	2.6%
Number	6	6
Outliers	0	0
Sigma-TE	4.0	4.0
Sigma-SA	4.2	4.2
Score pictogram		
Regression line	<u>0.00 + 1.008.x</u>	<u>0.00 + 1.008.x</u>

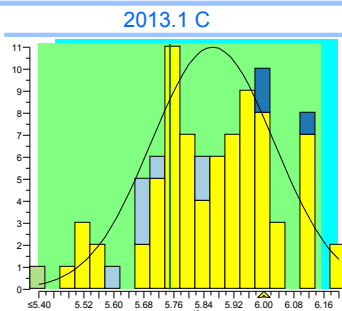
Consensus group: Natte chemie  
Method: Hexokinase, automatic



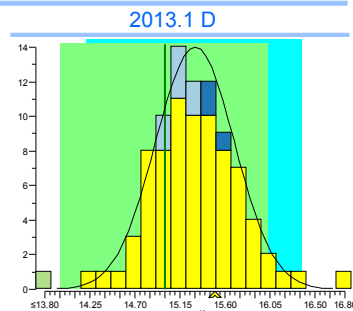
	cons.	meth.	ref.	lab
mean	12.44	12.44	12.2	12.1
SD	0.30	0.31		
n	88	77		
no	3	2		



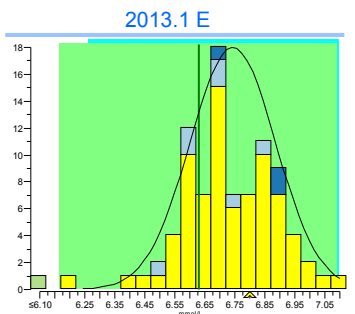
	cons.	meth.	ref.	lab
mean	23.8	23.8	23.4	23
SD	0.7	0.7		
n	88	77		
no	3	2		



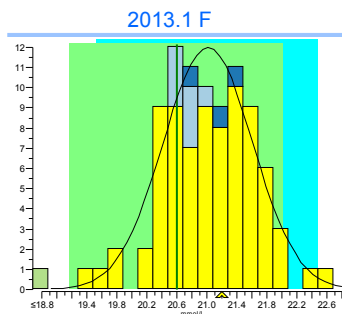
	cons.	meth.	exp.	lab
mean	5.86	5.87	5,75	6
SD	0.16	0.16		
n	88	77		
no	1	0		



	cons.	meth.	ref.	lab
mean	15.30	15.31	15.0	15.5
SD	0.39	0.41		
n	88	77		
no	2	1		



	cons.	meth.	ref.	lab
mean	6.74	6.75	6.63	6.8
SD	0.14	0.14		
n	89	78		
no	2	1		



	cons.	meth.	ref.	lab
mean	21.01	21.03	20.6	21.2
SD	0.60	0.63		
n	89	78		
no	1	0		

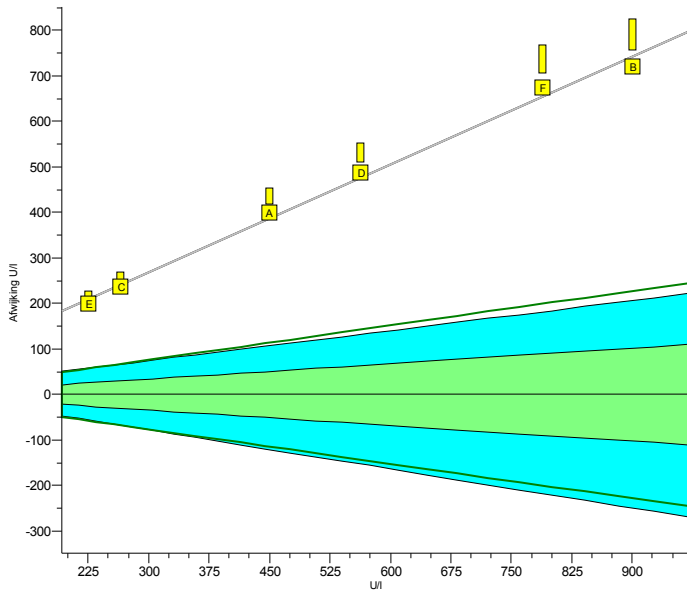
Legend

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

# INPUTs 2013.1

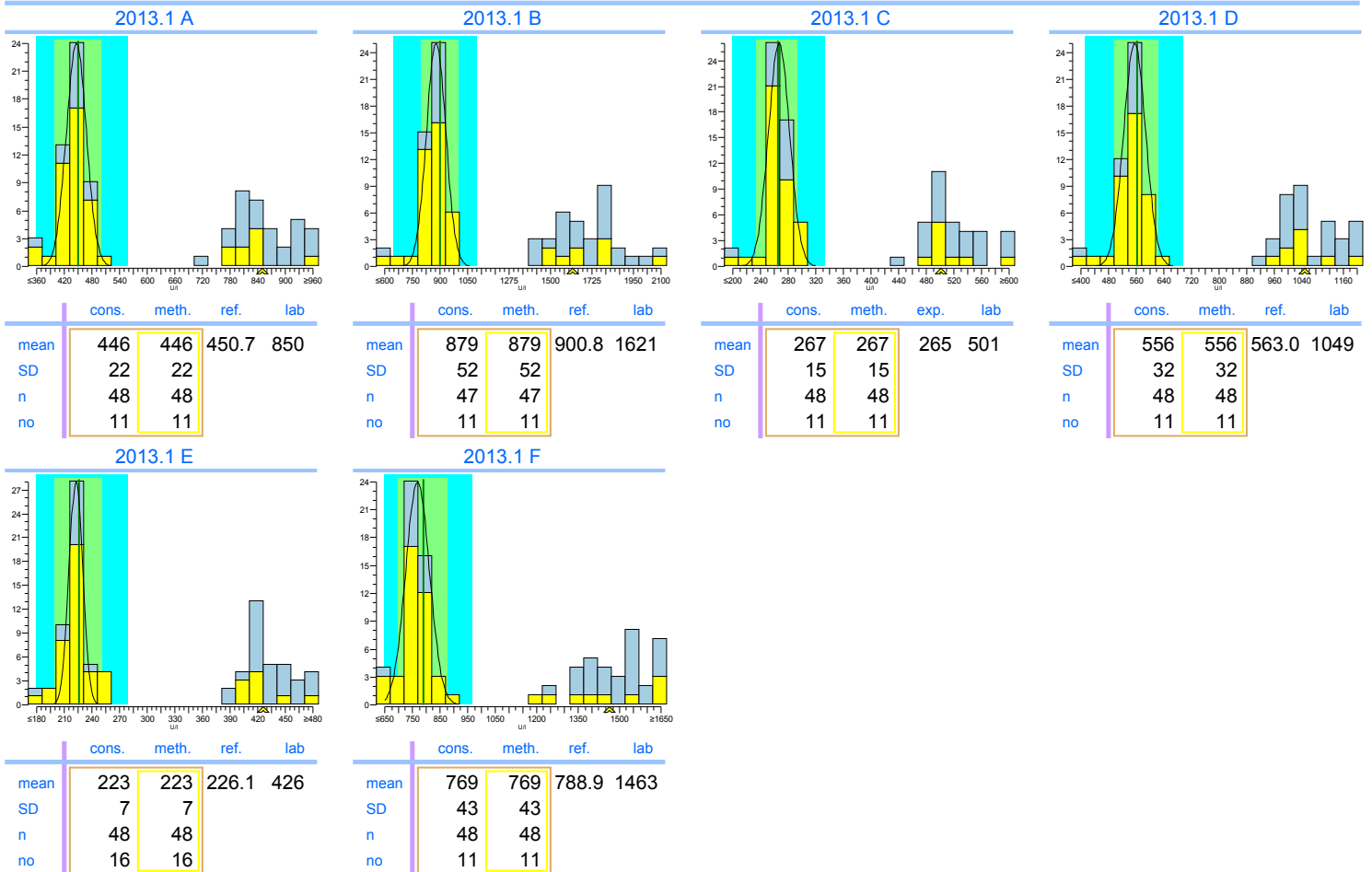
LD

units: U/I



	2013.1	cumulative
Trueness	+85%	+85%
Precision	3.3%	3.3%
Number	6	6
Outliers	0	0
Sigma-TE	-3.0	-3.0
Sigma-SA	-3.0 <span style="border: 1px solid red; padding: 0 2px;">0</span>	-3.0 <span style="border: 1px solid red; padding: 0 2px;">0</span>
Score pictogram	<span style="border: 1px solid cyan; padding: 0 2px;"> </span> <span style="border: 1px solid yellow; padding: 0 2px;"> </span>	<span style="border: 1px solid cyan; padding: 0 2px;"> </span> <span style="border: 1px solid yellow; padding: 0 2px;"> </span>
Regression line	$32 + 1.791 \cdot x$	$32 + 1.791 \cdot x$

Consensus group IFCC traceerbaar  
Method IFCC traceable

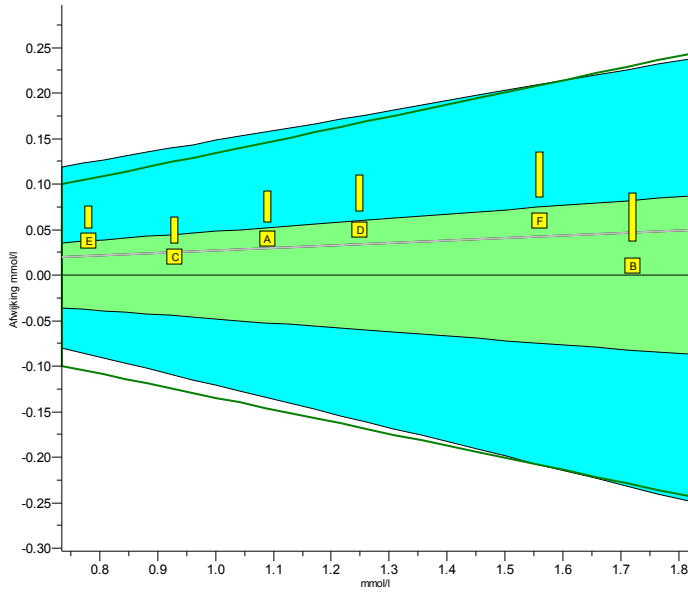


Legend  
  IFCC traceable      IFCC non-traceable

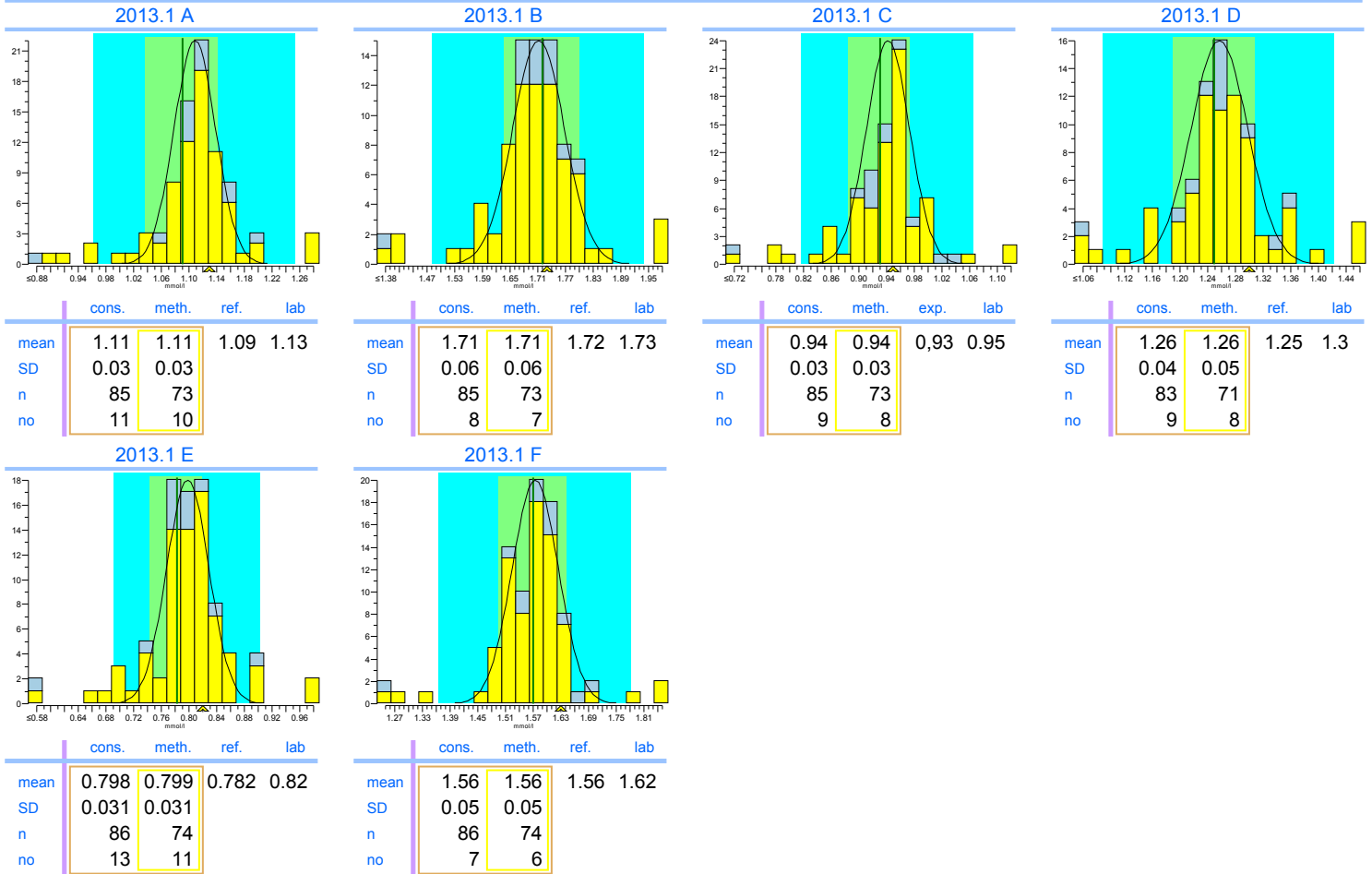
# INPUTS 2013.1

## Magnesium

units: mmol/l



	2013.1	cumulative
Trueness	+3.0%	+3.0%
Precision	1.3%	1.3%
Number	6	6
Outliers	0	0
Sigma-TE	2.1	2.1
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	<u>0.00 + 1.028.x</u>	<u>0.00 + 1.028.x</u>
Consensus group	Overall	
Method	Colorimetric	



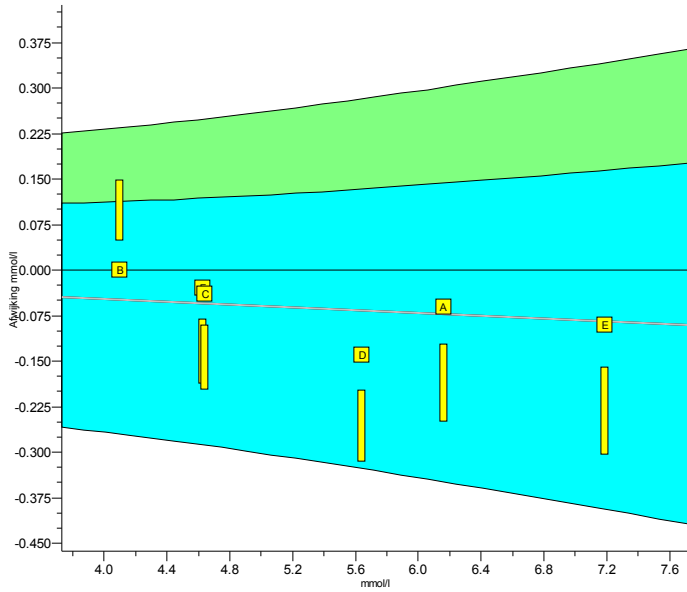
Legend



# INPUTS 2013.1

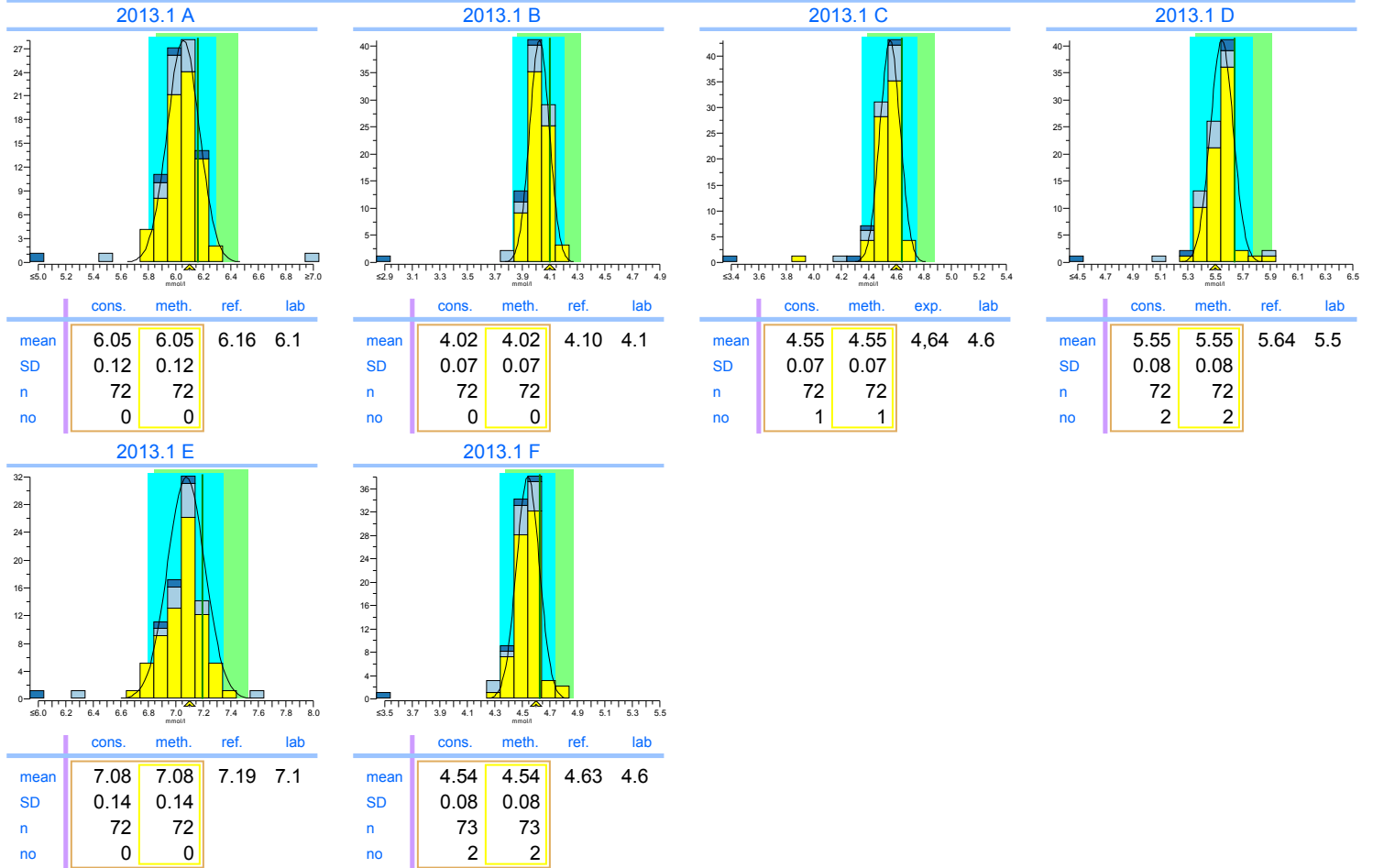
## Potassium

units: mmol/l



	2013.1	cumulative
Trueness	-1.1%	-1.1%
Precision	0.88%	0.88%
Number	6	6
Outliers	0	0
Sigma-TE	5.4 <span style="border: 1px solid green; padding: 2px;">2</span>	5.4 <span style="border: 1px solid green; padding: 2px;">2</span>
Sigma-SA	4.4	4.4
Score pictogram		
Regression line	<u>0.00 + 0.988.x</u>	<u>0.00 + 0.988.x</u>

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



Legend

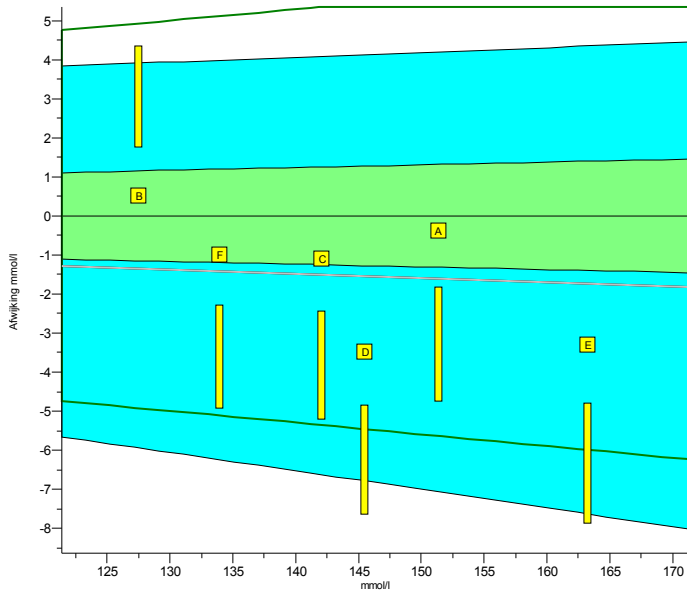
- ISE indirect (with predilution)
- ISE direct (no predilution)
- Other methods



# INPUTS 2013.1

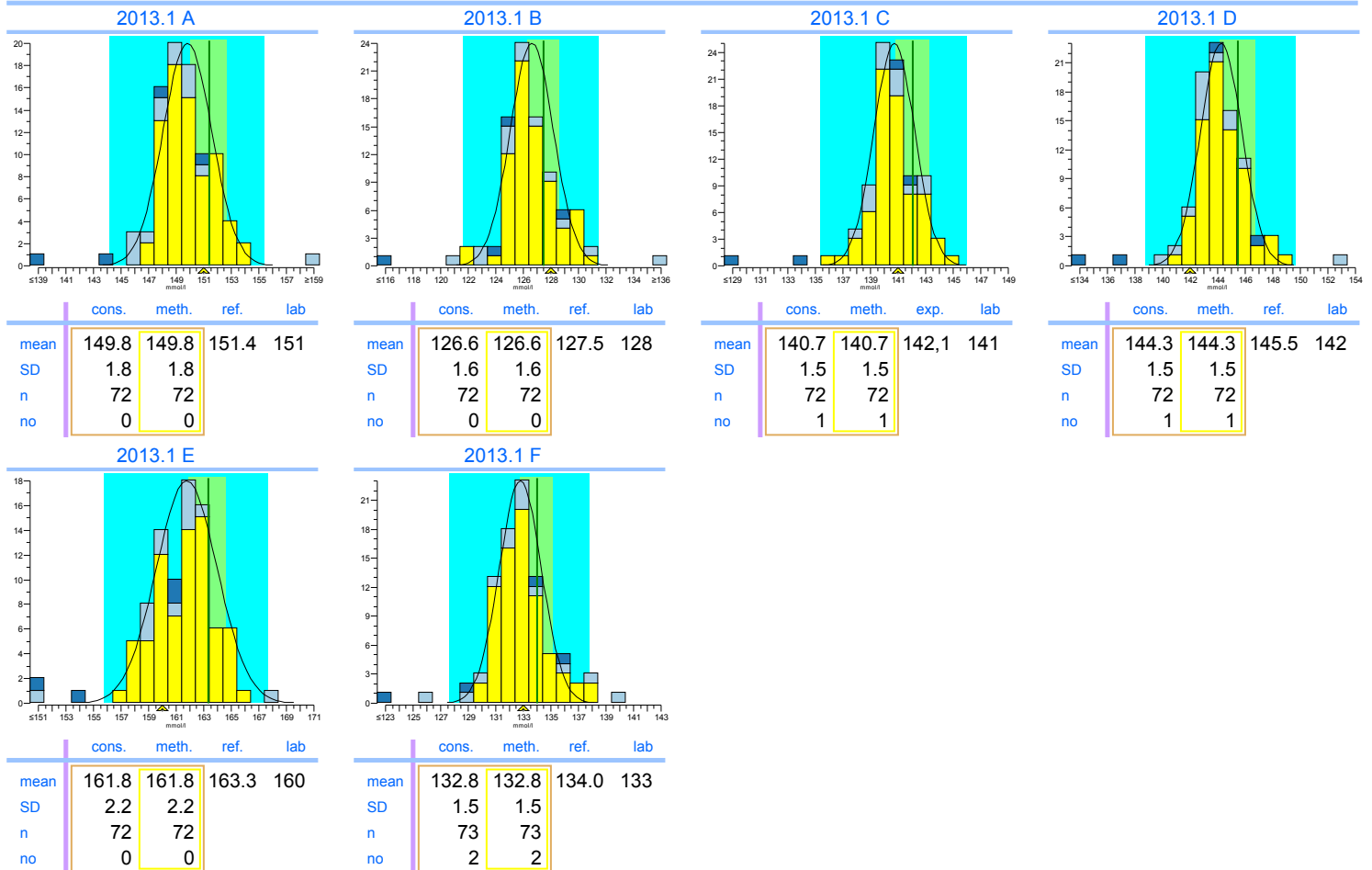
## Sodium

units: mmol/l



	2013.1	cumulative
Trueness	-1.0%	-1.0%
Precision	1.1%	1.1%
Number	6	6
Outliers	0	0
Sigma-TE	0.5	0.5
Sigma-SA	4.2 <span style="background-color: green; color: white;">1</span>	4.2 <span style="background-color: green; color: white;">1</span>
Score pictogram		
Regression line	$0.0 + 0.989.x$	$0.0 + 0.989.x$

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



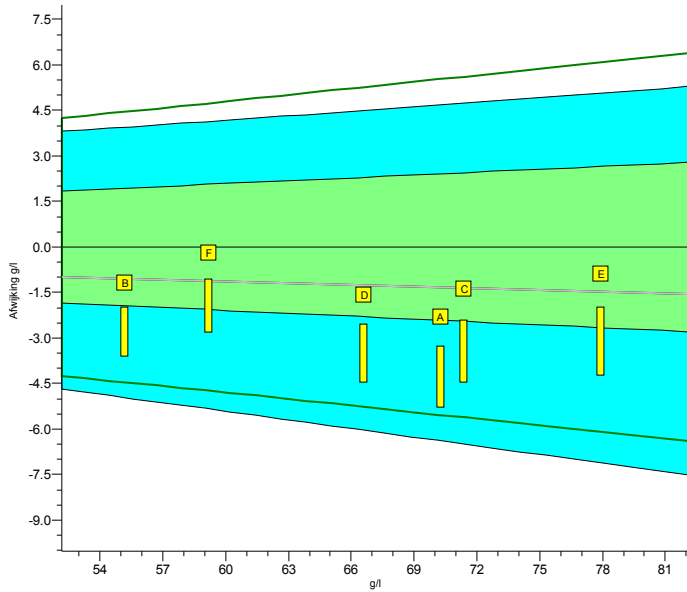
Legend

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

# INPUTS 2013.1

## Total Protein

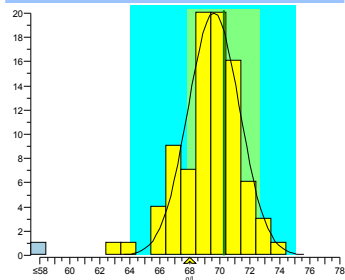
units: g/l



	2013.1	cumulative
Trueness	-1.9%	-1.9%
Precision	1.2%	1.2%
Number	6	6
Outliers	0	0
Sigma-TE	2.1	2.1
Sigma-SA	5.9 <span style="background-color: green; color: white; padding: 2px;">2</span>	5.9 <span style="background-color: green; color: white; padding: 2px;">2</span>
Score pictogram		
Regression line	$0.0 + 0.981.x$	$0.0 + 0.981.x$

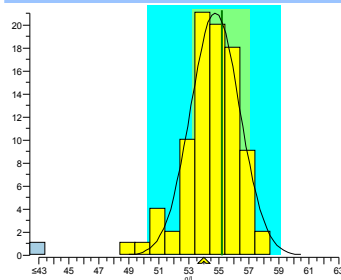
Consensus group Biureet  
Method Biurete, automatic

2013.1 A



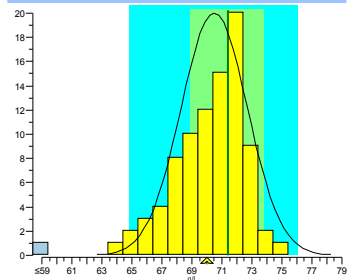
	cons.	meth.	ref.	lab
mean	69.6	69.6	70.3	68
SD	1.7	1.7		
n	88	88		
no	2	2		

2013.1 B



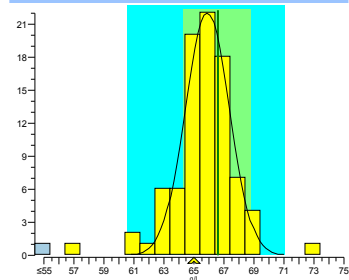
	cons.	meth.	ref.	lab
mean	54.8	54.8	55.2	54
SD	1.6	1.6		
n	88	88		
no	1	1		

2013.1 C



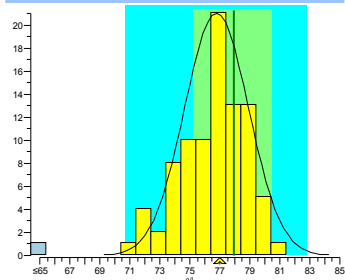
	cons.	meth.	exp.	lab
mean	70.5	70.5	71.4	70
SD	2.2	2.2		
n	87	87		
no	0	0		

2013.1 D



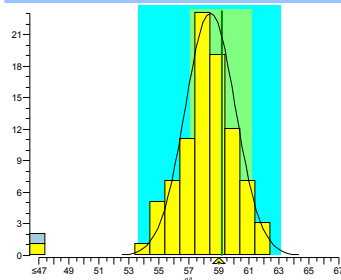
	cons.	meth.	ref.	lab
mean	65.9	65.9	66.6	65
SD	1.5	1.5		
n	88	88		
no	4	4		

2013.1 E



	cons.	meth.	ref.	lab
mean	76.8	76.8	77.9	77
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		

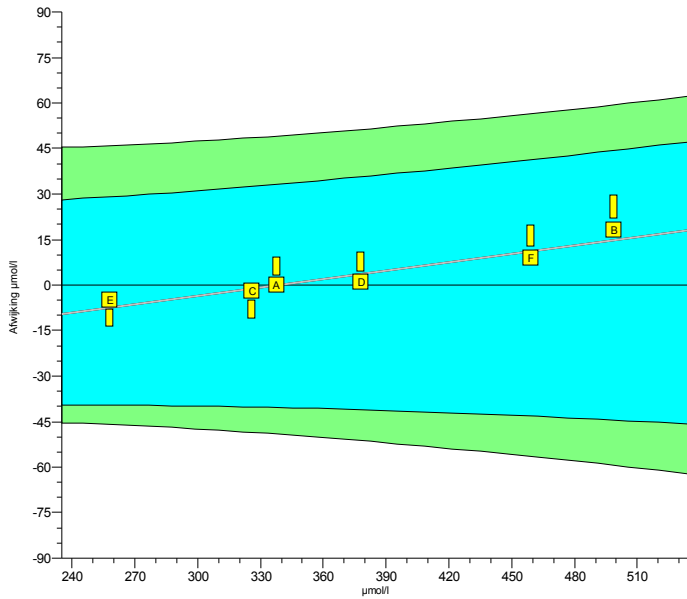
Legend

Biurete, automatic     Other methods

# INPUTS 2013.1

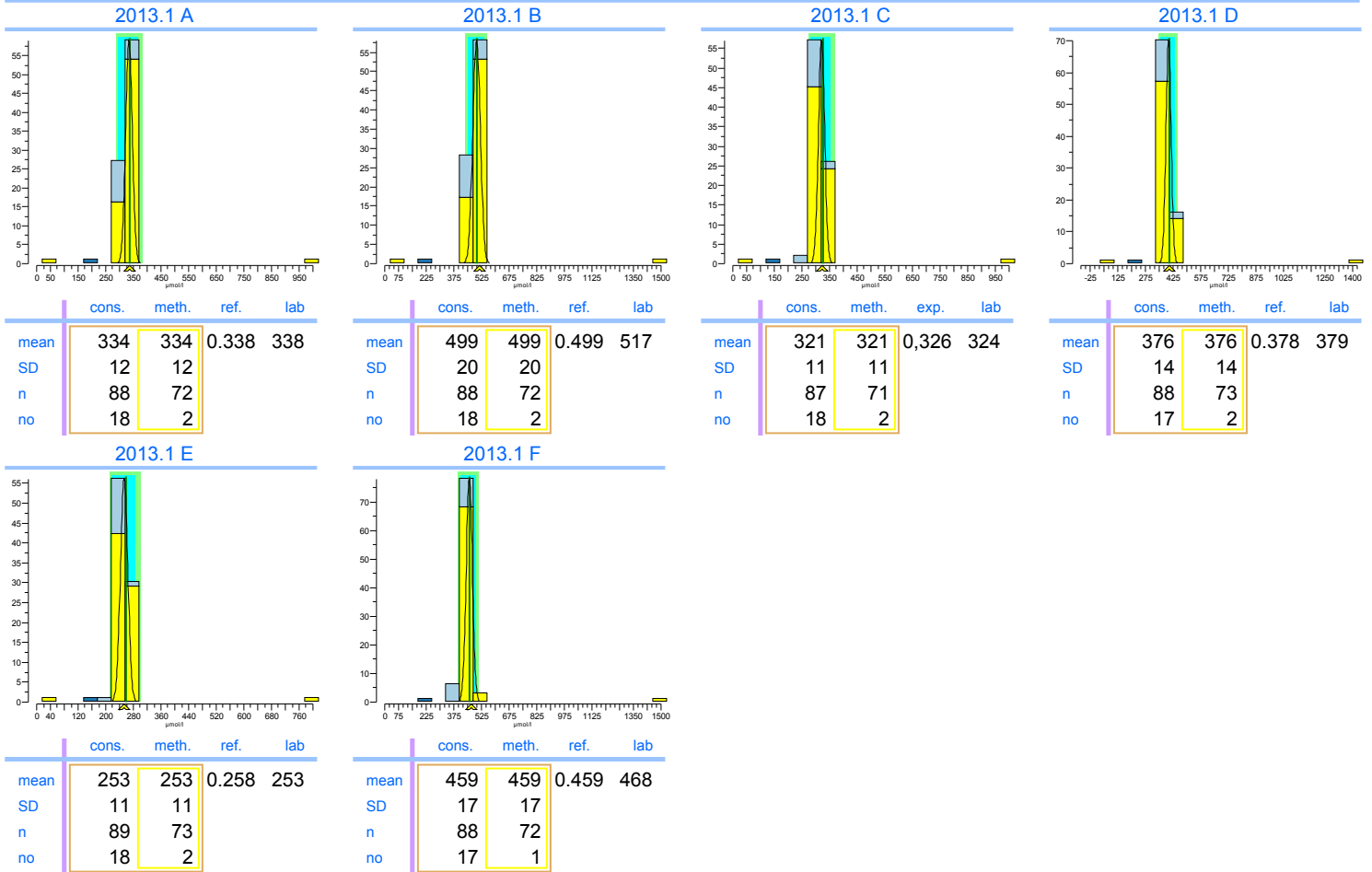
## Urate

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



	2013.1	cumulative
Trueness	+0.93%	+0.93%
Precision	0.71%	0.71%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 <span style="border: 1px solid black; padding: 2px;">2</span>	6.0 <span style="border: 1px solid black; padding: 2px;">2</span>
Sigma-SA	6.0	6.0
Score pictogram		
Regression line	$0 + 1.091 \cdot x$	$0 + 1.091 \cdot x$

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



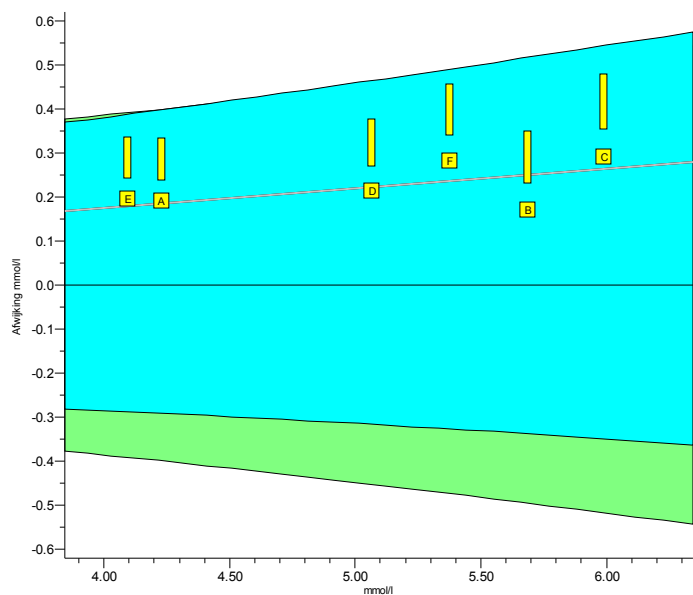
Legend

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

# INPUTS 2013.1

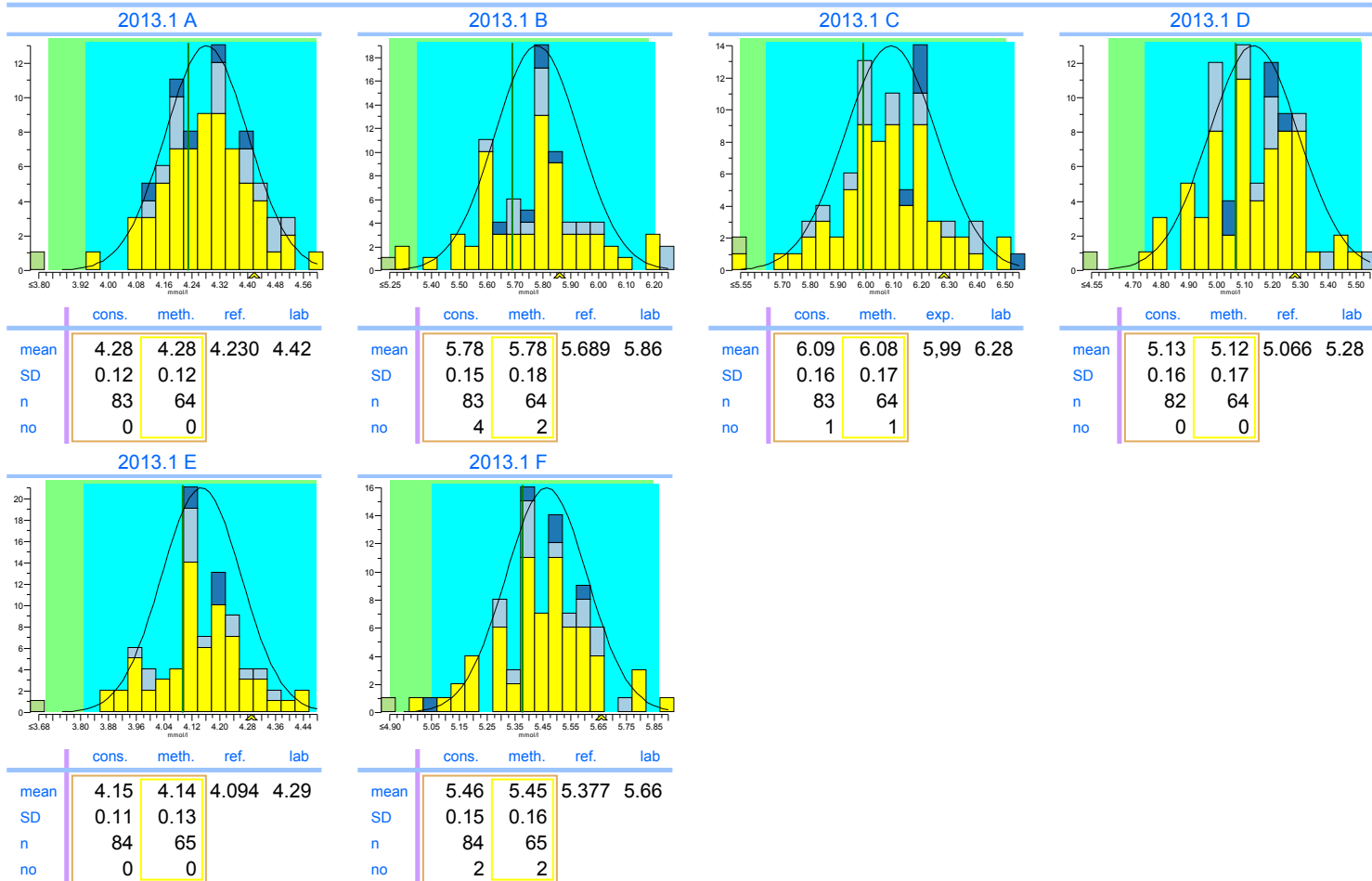
## Cholesterol

units: mmol/l



	2013.1	cumulative
Trueness	+4.4%	+4.4%
Precision	0.86%	0.86%
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	4.7	4.7
Score pictogram		
Regression line	<u>0.00 + 1.044.x</u>	<u>0.00 + 1.044.x</u>

Consensus group: Enzymatisch  
Method: Enzymatic, automatic, discrete



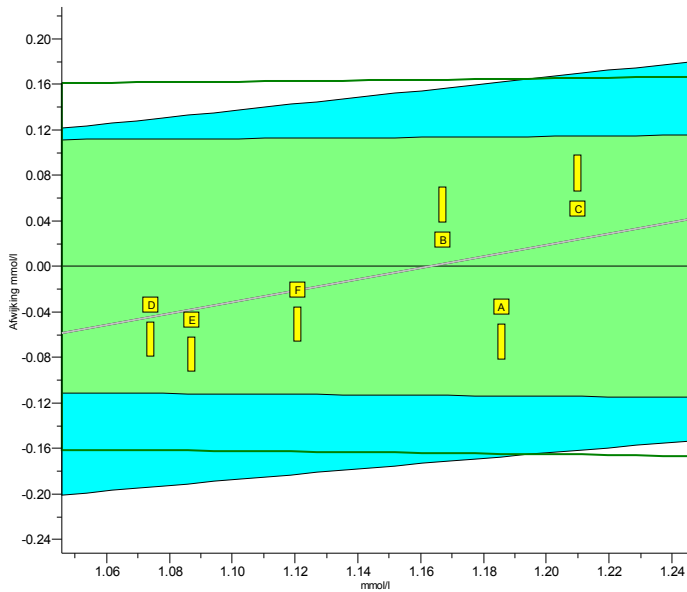
Legend

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

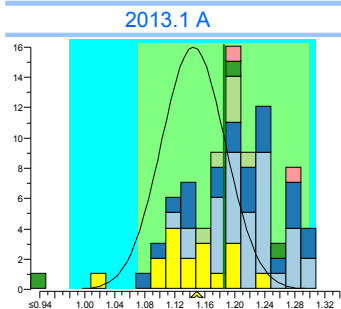
# INPUTS 2013.1

## HDL-Cholesterol

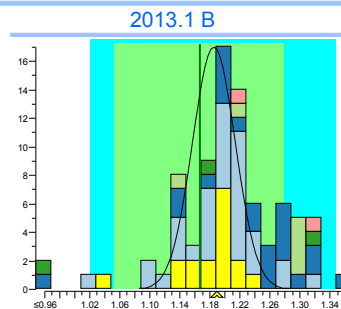
units: mmol/l



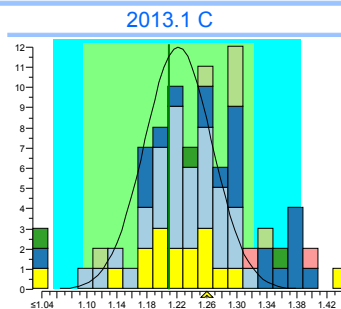
	2013.1	cumulative
Trueness	-0.95%	-0.95%
Precision	1.1%	1.1%
Number	6	6
Outliers	0	0
Sigma-TE	6.0	6.0
Sigma-SA	6.0 <span style="background-color: green; color: white; padding: 2px;">2</span>	6.0 <span style="background-color: green; color: white; padding: 2px;">2</span>
Score pictogram		
Regression line	$-0.58 + 1.501 \cdot x$	$-0.58 + 1.501 \cdot x$
Consensus group	Overall	
Method	Immunoinhibition	



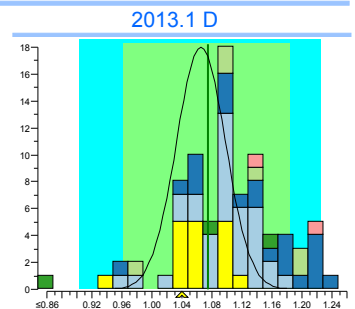
	cons.	meth.	ref.	lab
mean	1.14	1.14	1.186	1.15
SD	0.04	0.04		
n	17	17		
no	1	1		



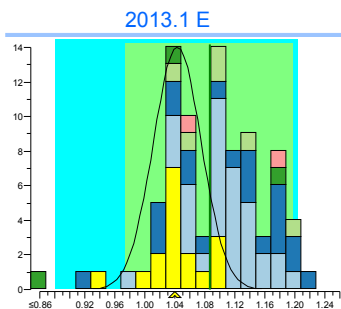
	cons.	meth.	ref.	lab
mean	1.19	1.19	1.167	1.19
SD	0.03	0.03		
n	17	17		
no	1	1		



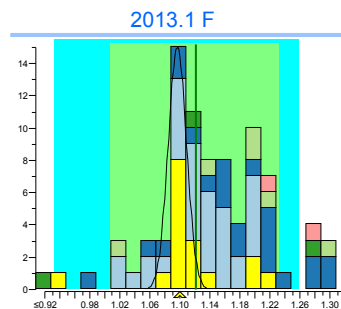
	cons.	meth.	exp.	lab
mean	1.22	1.22	1.21	1.26
SD	0.04	0.04		
n	17	17		
no	2	2		



	cons.	meth.	ref.	lab
mean	1.07	1.07	1.074	1.04
SD	0.03	0.03		
n	17	17		
no	1	1		



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



	cons.	meth.	ref.	lab
mean	1.10	1.10	1.121	1.1
SD	0.01	0.01		
n	17	17		
no	5	5		

Legend

- Immunoinhibition
- PEG modified enzyme, PEGME (Kyowa Medex)
- Accelerator Selective Detergent (=Ultra HDL<sup>+</sup>)
- Catalase method (Denka Seiken)
- Overige methoden
- Precipitation Technique