

Verein für
Association pour le
Associazione per il



medizinische Qualitätskontrolle
contrôle de qualité médical
controllo di qualità medico

Survey Report

2015 - 2

Survey Specimens

The homogeneity and stability of all specimens were checked before and/or during shipment and no irregularities were noted. The suitability tests were performed by the laboratories of the Universitätsspital Zürich (University Hospital Zurich) (<http://www.uzl.usz.ch/>).

The following survey specimens were produced specifically for MQ by a sub-contractor:
B1 Strep A Test, B2 Uricult, H4 Parasitic Hematology, K14 tumor marker

Determination of target values

For each target value, the type of determination per ISO17043: 2010 B2.1 is indicated (column "type"):

- a Value known due to production.
- b Certified reference value for use with special specimens
- c Reference value determined by analysis
- d Consensus values of expert laboratories
- e Consensus values of the participants

For methods groups with more than 9 participants, consensus values of the participants ("e") are generally determined. In order to provide all participants with target values that are as meaningful as possible, other methods may also be applied for smaller method groups.

Uncertainty of the determined target values

The standard uncertainty (u_x) is calculated using the following formula (ISO13528):

$$u_x = (\text{target value}/100) * (1.25/\sqrt{\text{number of participants}}) * \%CV$$

- u_x has the same unit as the target value
- u_x can be compared with the standard deviation of the participants' collective ($SD = \text{target value} * \%CV/100$)
- For participant numbers >18, the standard uncertainty (u_x) is significantly lower than the scatter of the collective participants and can be neglected.

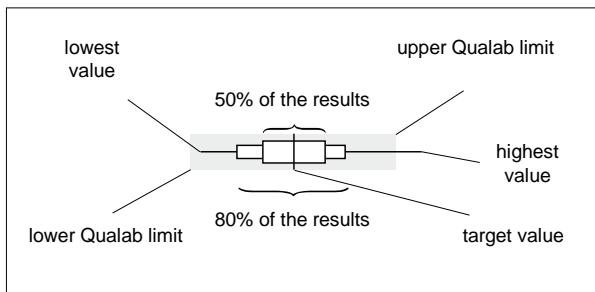
QUALAB and MQ tolerances

For all mandatory analyzes, QUALAB tolerances are used (www.qualab.ch, external quality control). For non-mandatory analyzes, the tolerances are specified by MQ's survey specimen leader.

If the determined uncertainty, u_x , of the target value is greater than 15% of the QUALAB or MQ tolerance, the letter indicating the type of target detection is marked with an additional star (example "e*"). Thereby, we are alerting the participants to the fact that the uncertainty of the target value can have an impact on the evaluation.

Graphics

The results are shown graphically as follows:



Comparison of Devices

The data in this report allows you to compare the performance of different devices. However, remember to consider the following:

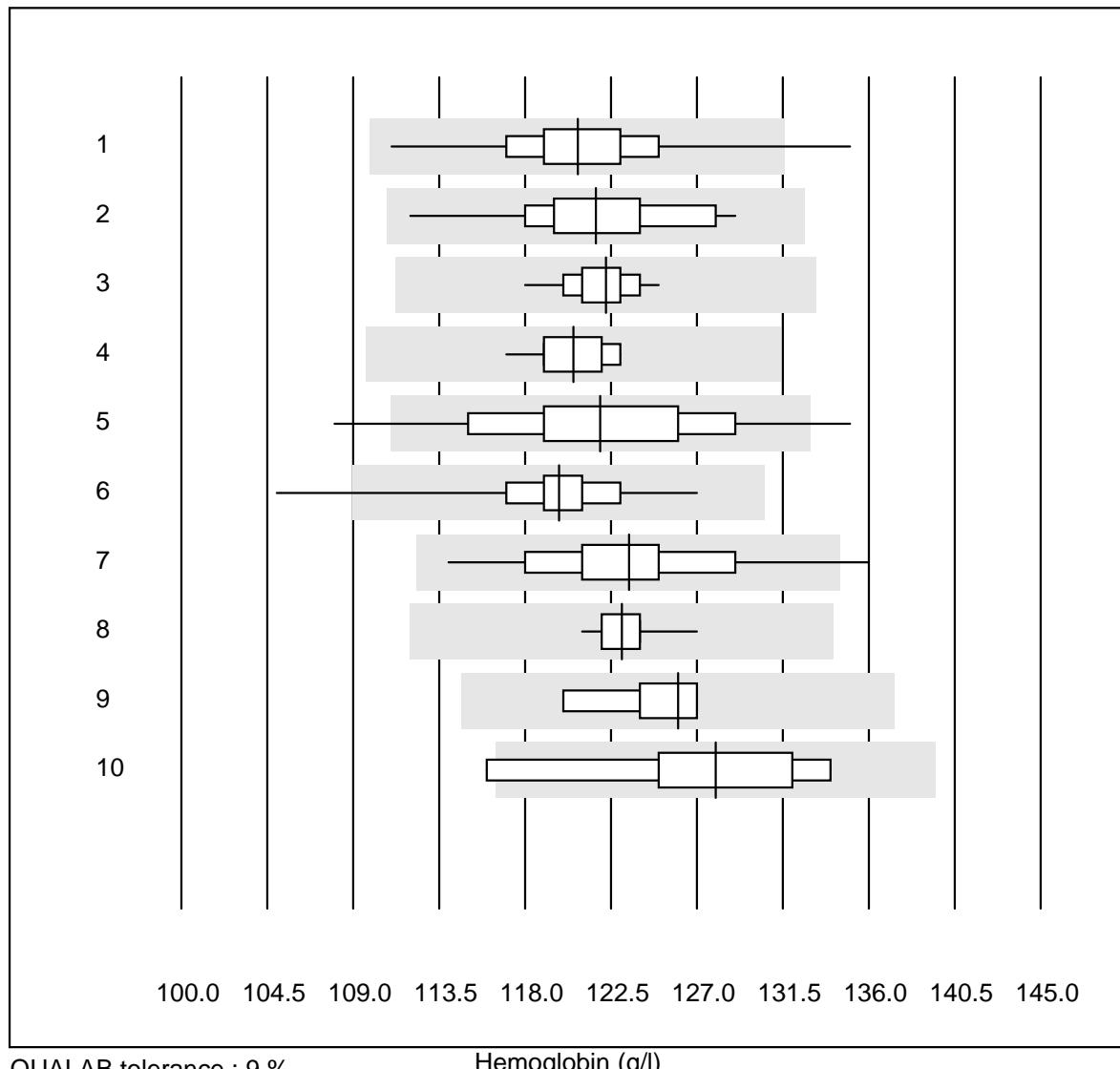
- The chemical control K1 is a ready-to-use commercial control serum. Even if the specimen is of human origin, it is possible that matrix effects occur. These are device-specific and result in different target values.
- Only one specimen was measured. Since the scatter of the results is dependent on the nature of the specimen (matrix effects) and on the signal strength, the determined coefficient of variations (CV in %) cannot be applied generally.
- A large number of runaways is due to administrative errors (wrong unit, results mixed up) or to operator errors (wrong sample, not correctly taken up in solution, not mixed well) and has nothing to do with the type of device.

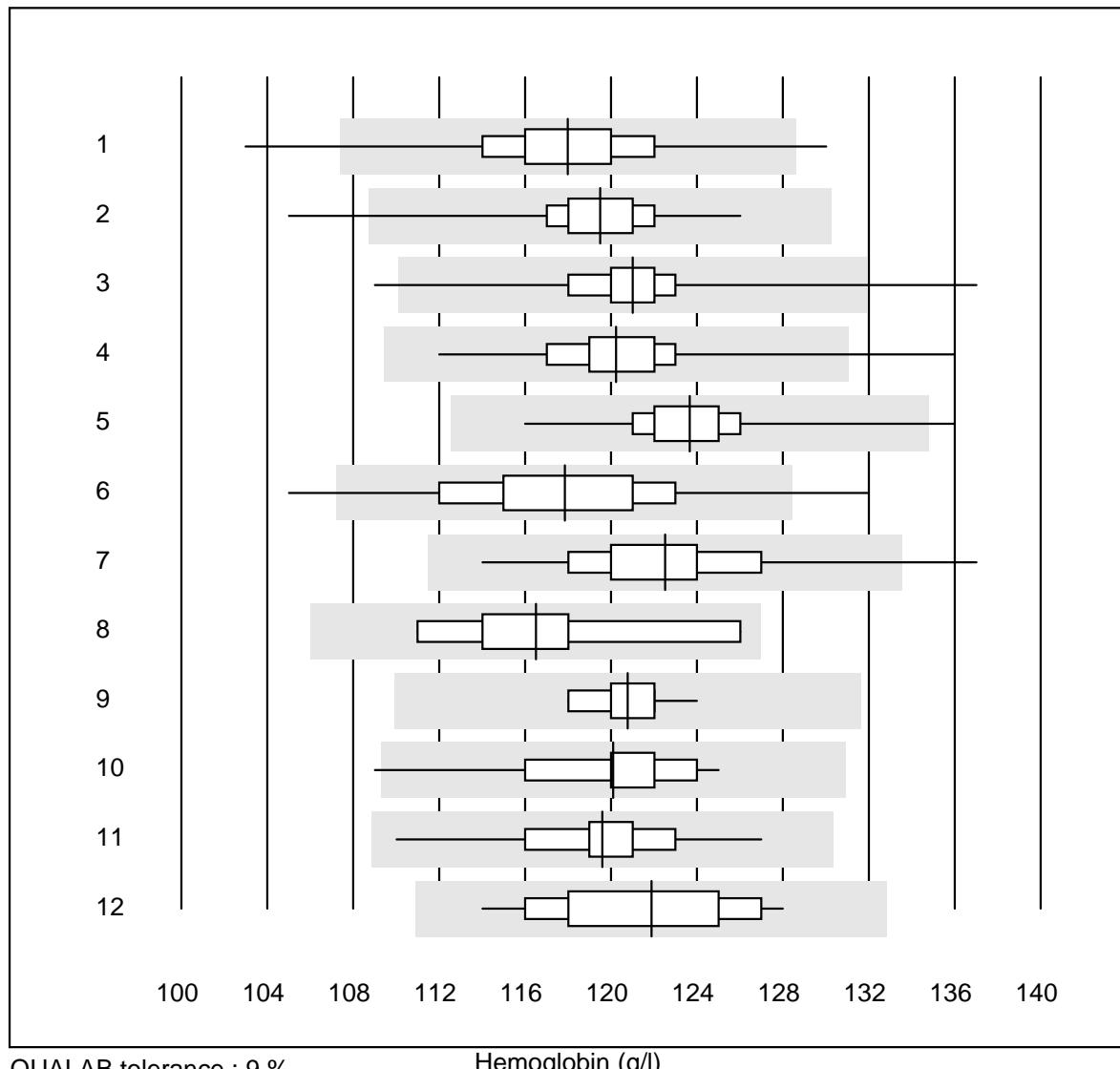
Zürich, 6.7.2015

Dr. R. Fried
Survey Director

Publication of this report or any portion thereof without our prior written consent is not permitted. The original is archived at www.mqzh.ch

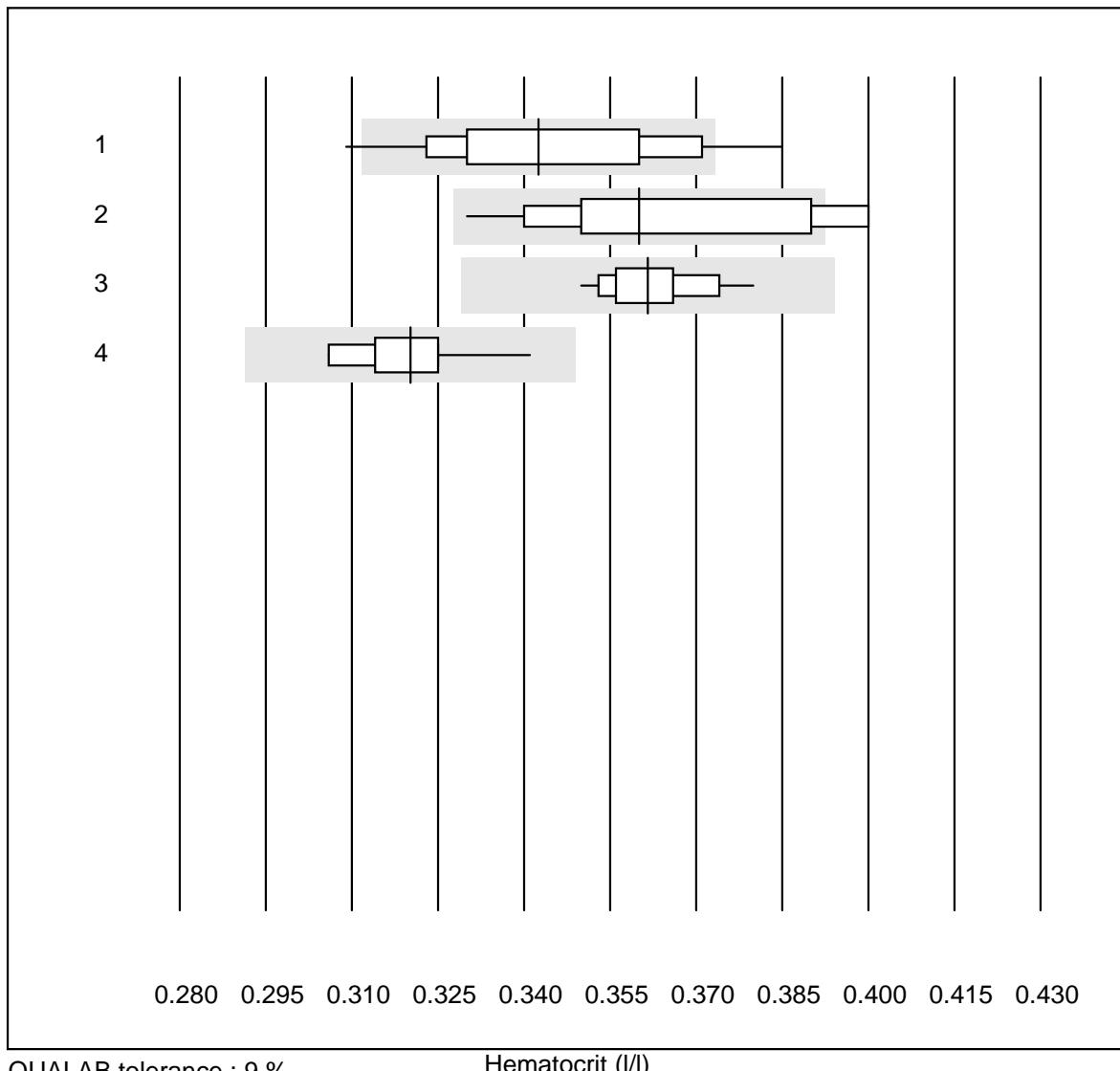
Hemoglobin



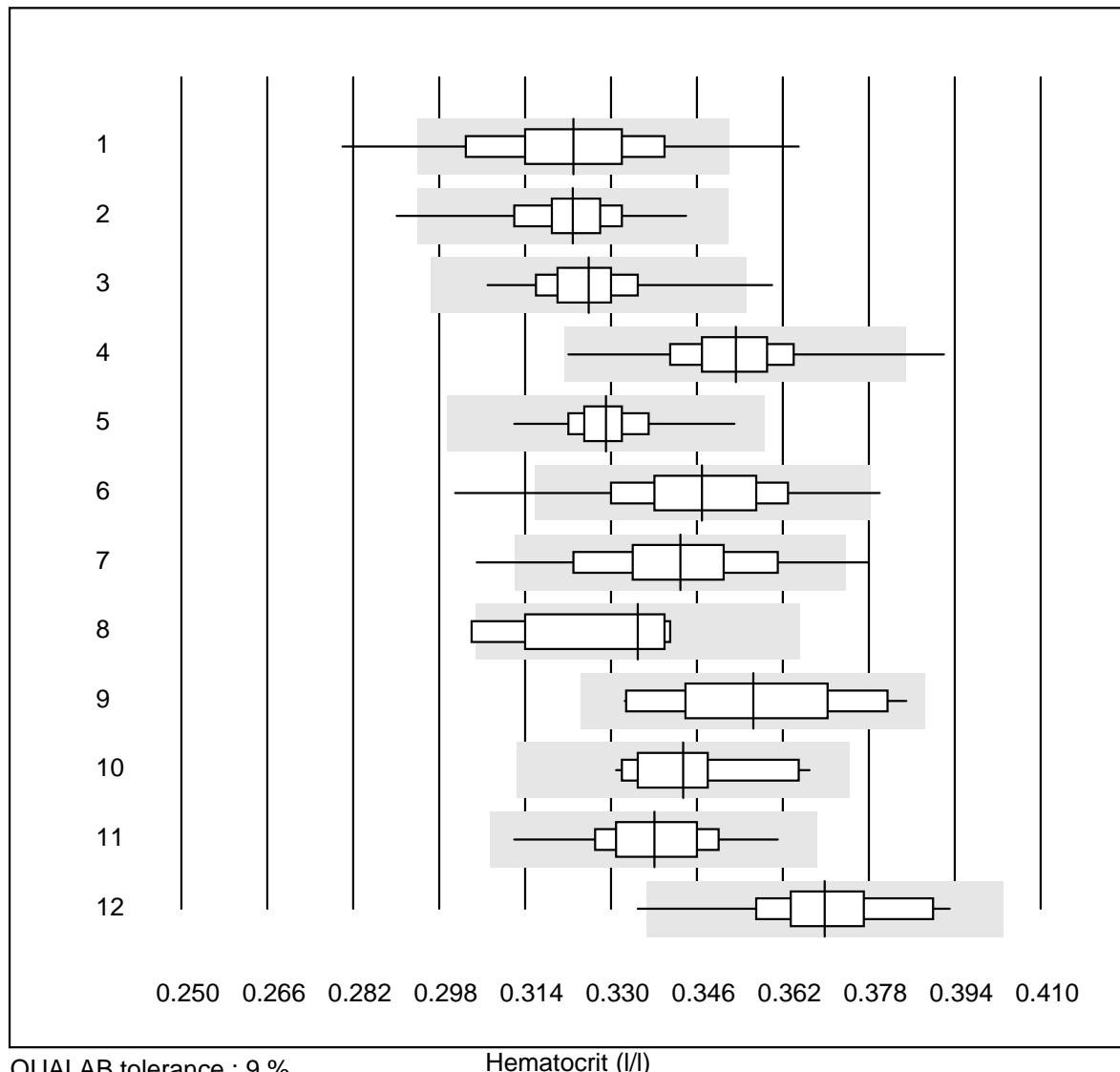
Hemoglobin

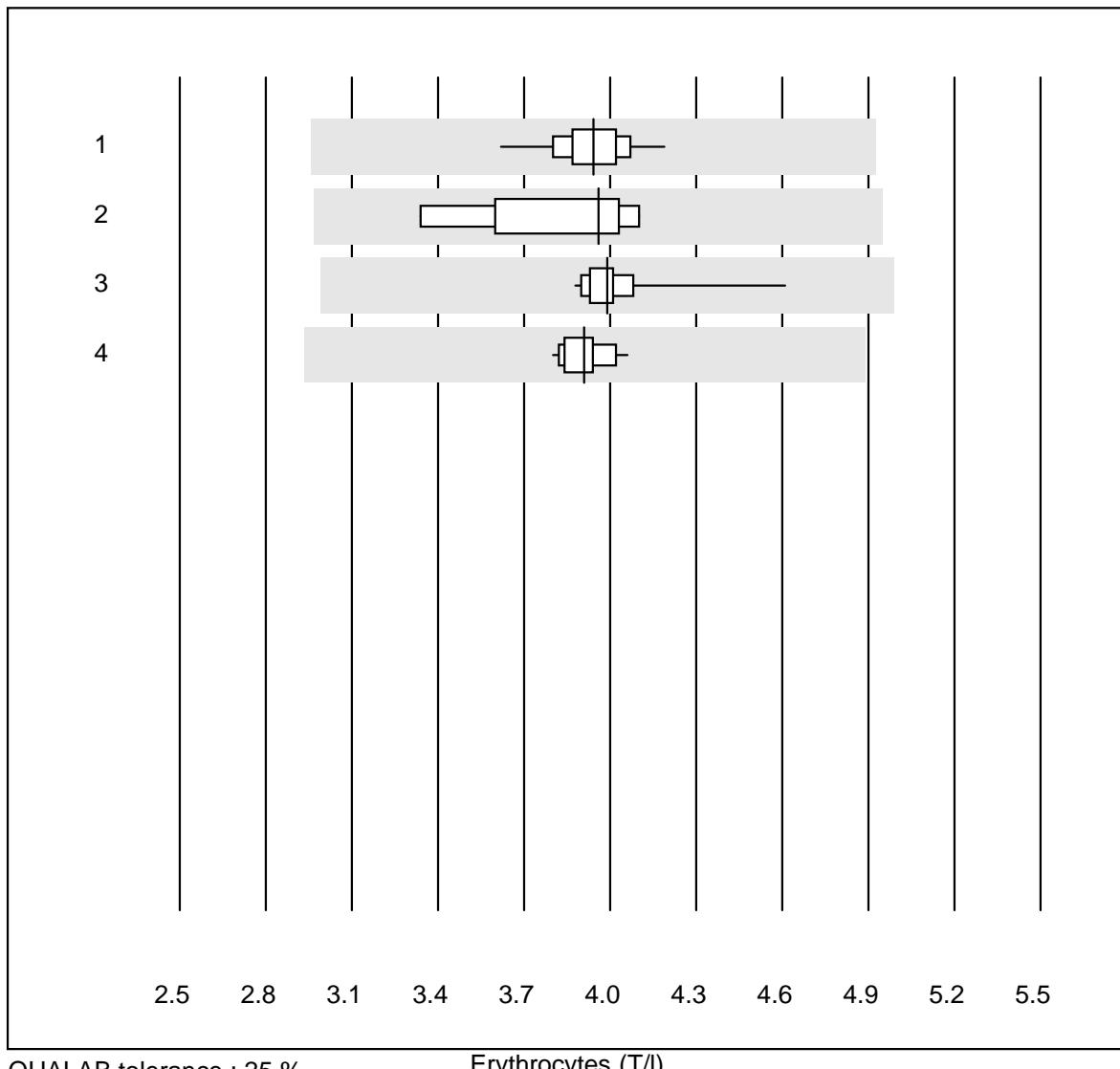
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Abx Micros	863	95.6	1.0	3.4	118.0	3.0	e
2 Microsemi	242	97.9	0.4	1.7	119.5	2.0	e
3 Sysmex KX21	436	95.8	1.4	2.8	121.0	2.2	e
4 Sysmex Poch - 100i	218	97.2	0.5	2.3	120.2	2.2	e
5 Sysmex XP 300	161	97.5	0.6	1.9	123.7	1.9	e
6 Mythic	242	96.2	2.1	1.7	117.8	3.7	e
7 Swelab	70	94.3	4.3	1.4	122.5	3.3	e
8 MS4	6	100.0	0.0	0.0	116.5	4.3	e*
9 Abacus Junior	13	100.0	0.0	0.0	120.8	1.4	e
10 Medonic	18	88.8	5.6	5.6	120.1	3.0	e
11 Nihon Kohden Celltac	38	89.5	0.0	10.5	119.6	2.8	e
12 Samsung HC10	45	100.0	0.0	0.0	121.9	3.3	e

Hematocrit

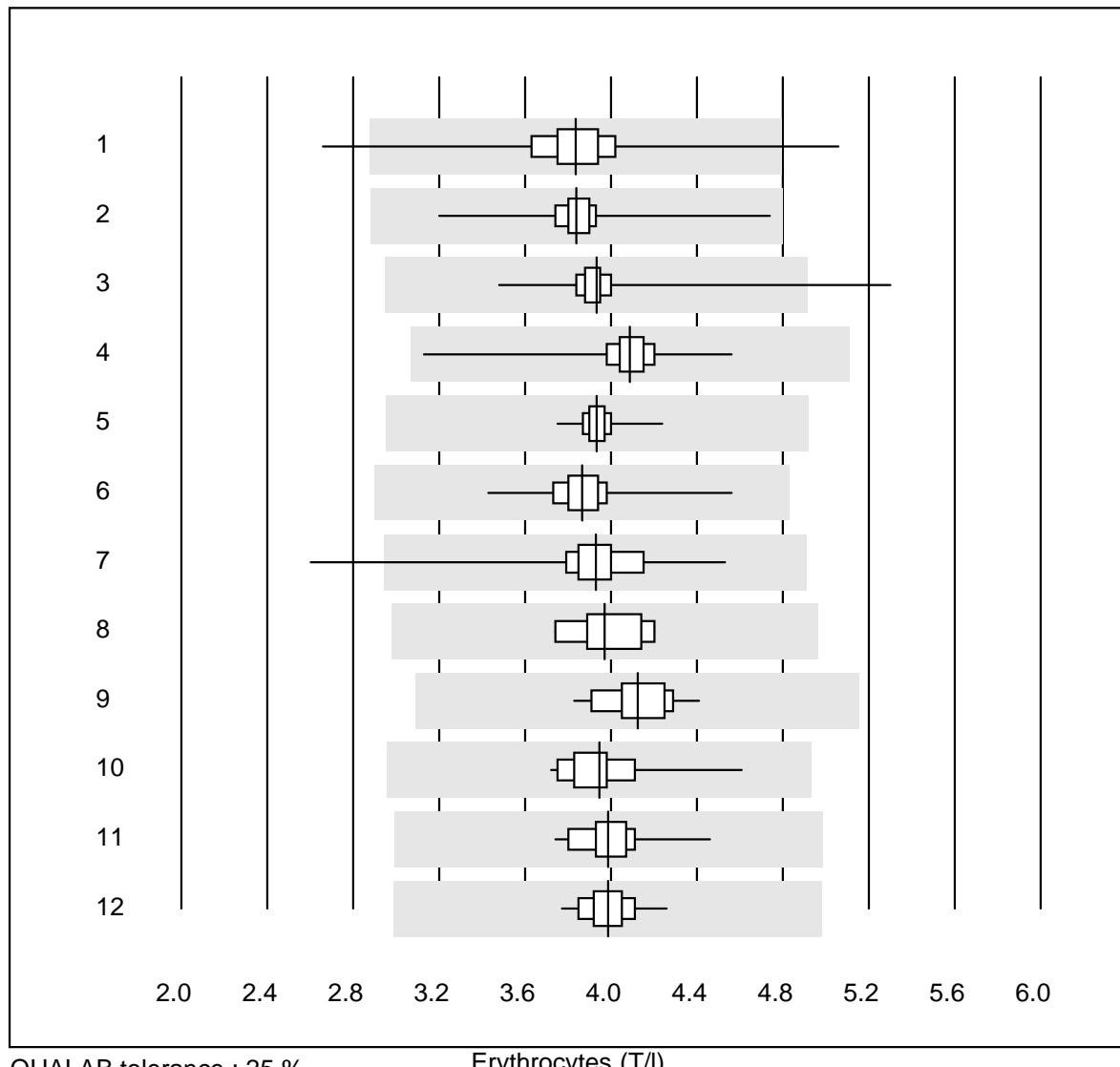


No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Automat	42	88.1	9.5	2.4	0.34	5.7	e
2 Centrifuge	15	80.0	13.3	6.7	0.36	6.3	e*
3 Sysmex XT/XE/XS	38	94.7	0.0	5.3	0.36	2.1	e
4 ABX Pentra	11	90.9	0.0	9.1	0.32	3.0	e

Hematocrit

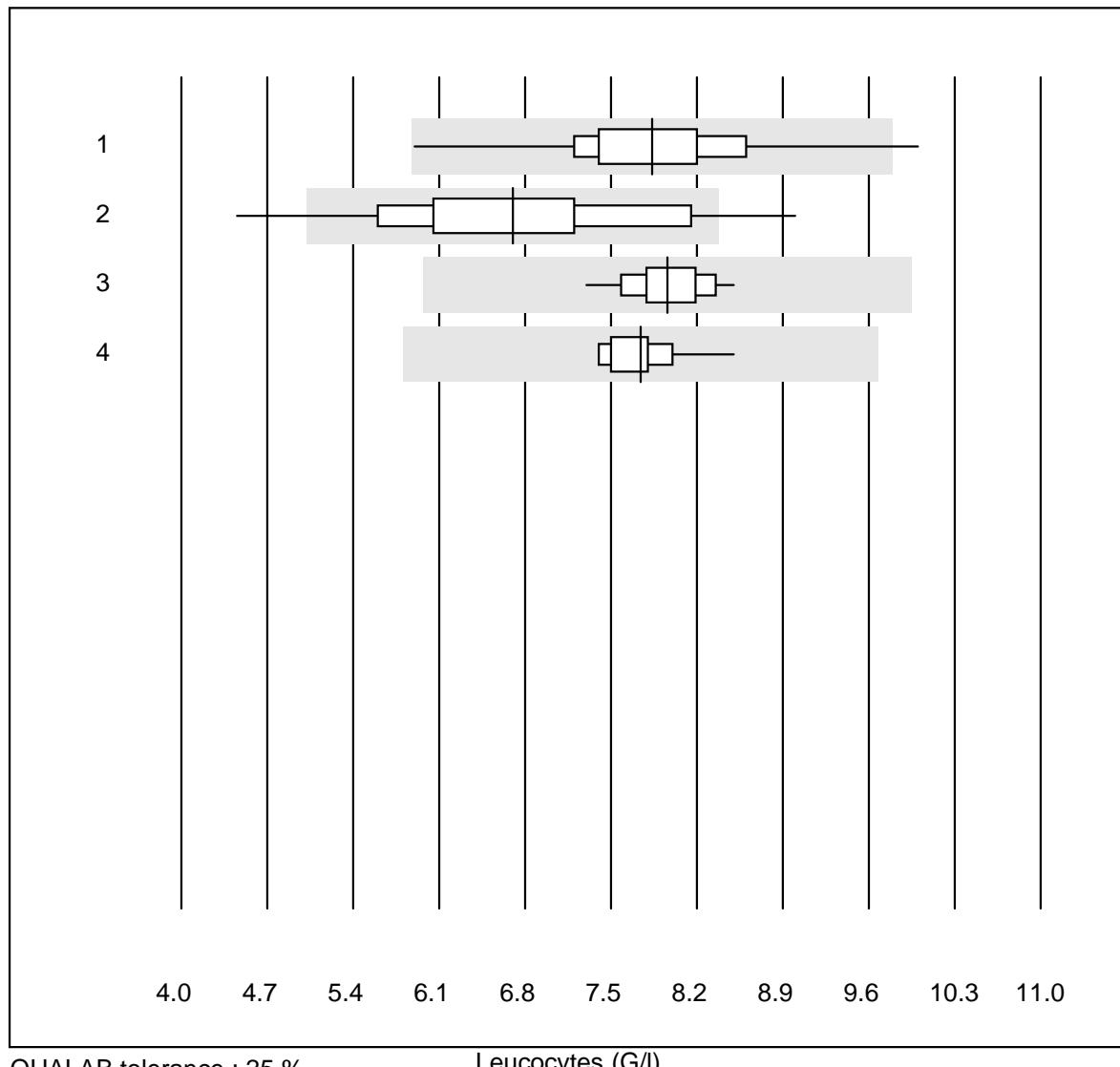
Erythrocytes

Erythrocytes



No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Abx Micros	863	97.7	0.7	1.6	3.84	5.2	e
2 Microsemi	244	100.0	0.0	0.0	3.84	3.2	e
3 Sysmex KX21	436	97.7	0.7	1.6	3.93	3.8	e
4 Sysmex Poch - 100i	218	98.2	0.0	1.8	4.09	3.0	e
5 Sysmex XP 300	160	98.7	0.0	1.3	3.93	1.5	e
6 Mythic	242	98.3	0.0	1.7	3.86	3.1	e
7 Swelab	70	97.2	1.4	1.4	3.93	5.7	e
8 MS4	6	100.0	0.0	0.0	3.97	4.3	e
9 Abacus Junior	13	100.0	0.0	0.0	4.12	3.8	e
10 Medonic	18	100.0	0.0	0.0	3.95	4.9	e
11 Samsung HC10	45	100.0	0.0	0.0	3.99	3.3	e
12 Nihon Kohden Celltac	38	92.1	0.0	7.9	3.99	2.7	e

Leucocytes

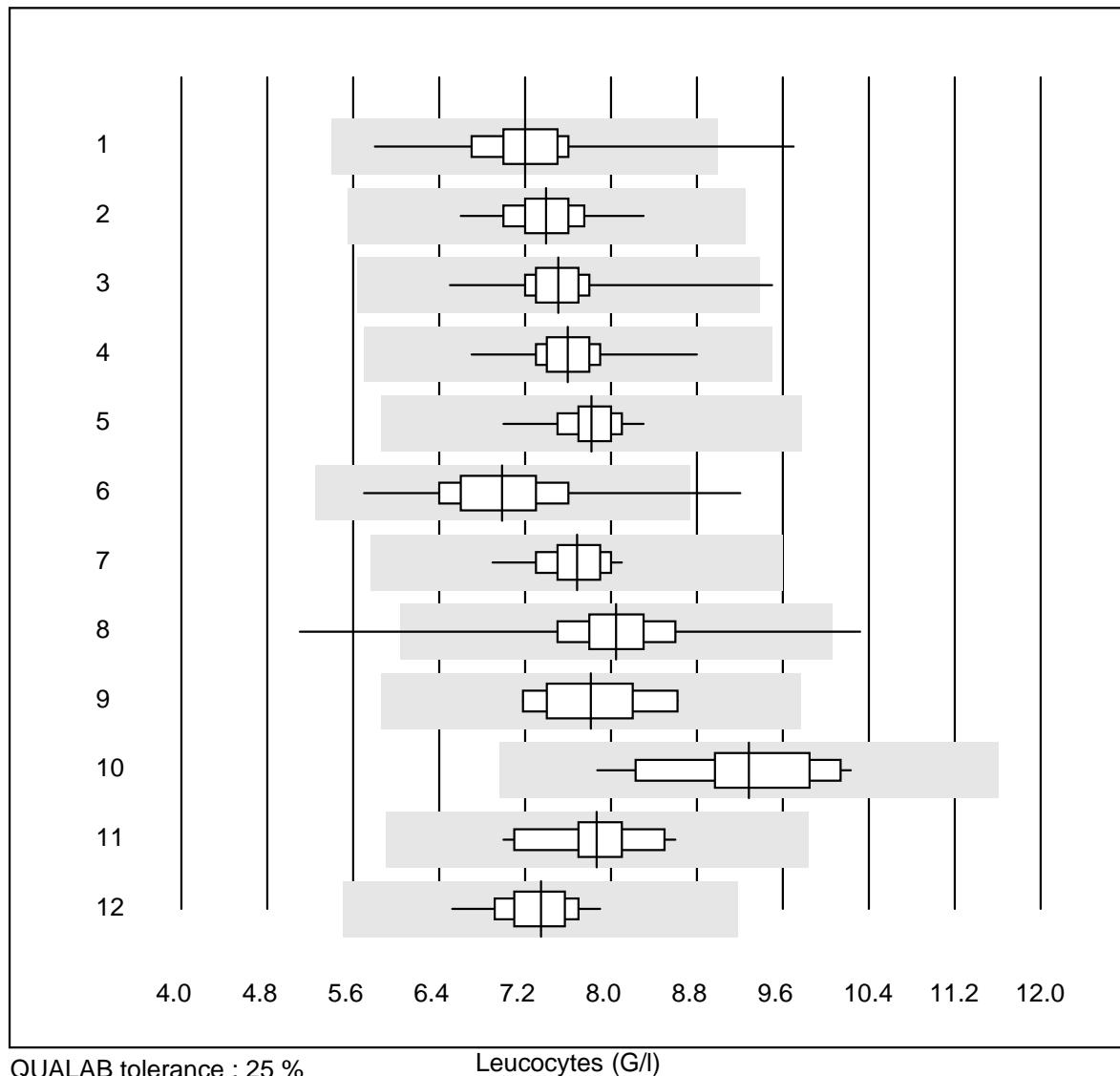


QUALAB tolerance : 25 %

Leucocytes (G/l)

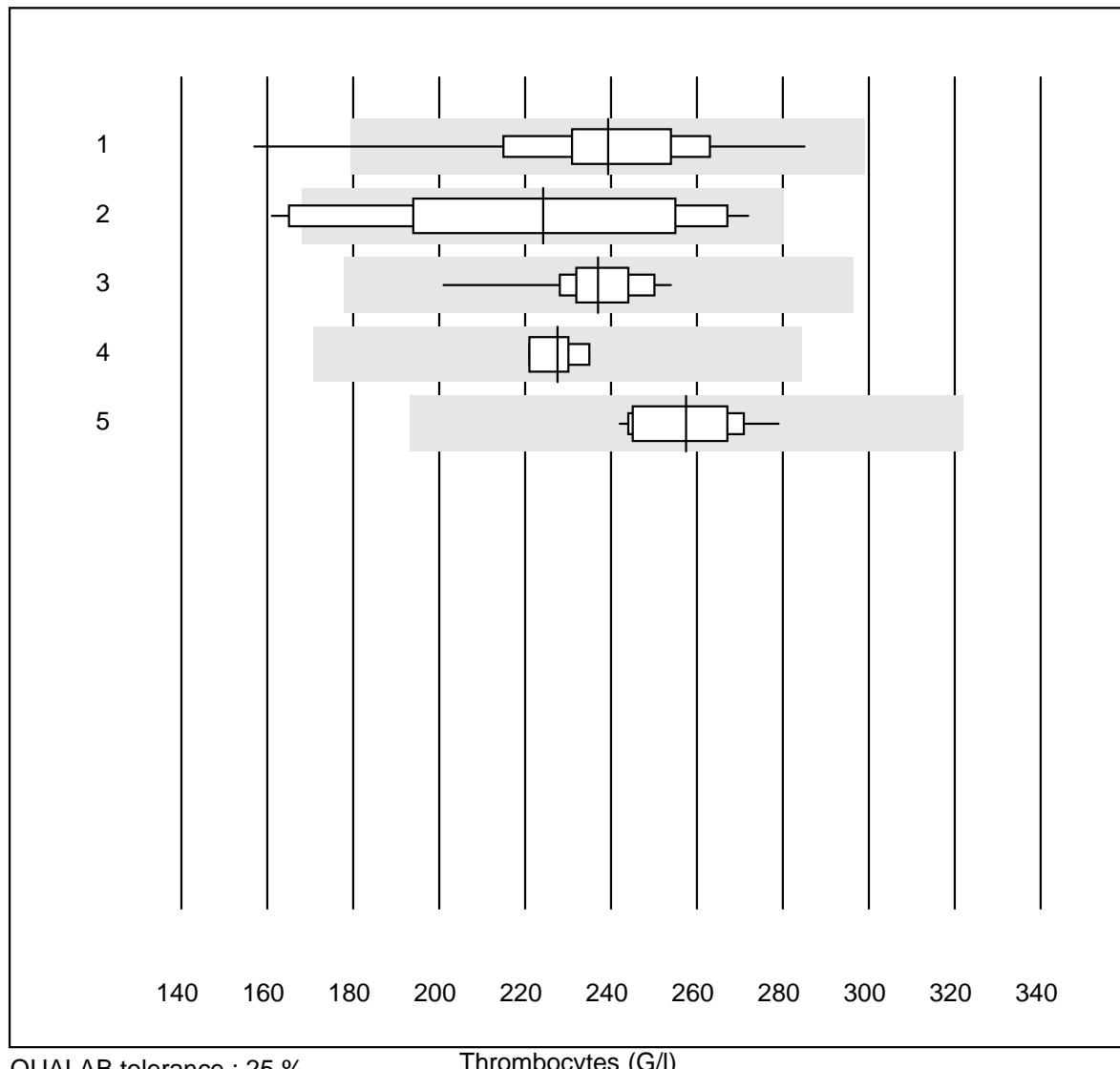
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Automat	37	97.3	2.7	0.0	7.84	8.9	e
2 Microscopic	58	91.4	8.6	0.0	6.70	14.2	e
3 Sysmex XT/XE/XS	39	100.0	0.0	0.0	7.96	3.7	e
4 ABX Pentra	11	100.0	0.0	0.0	7.74	4.0	e

Leucocytes

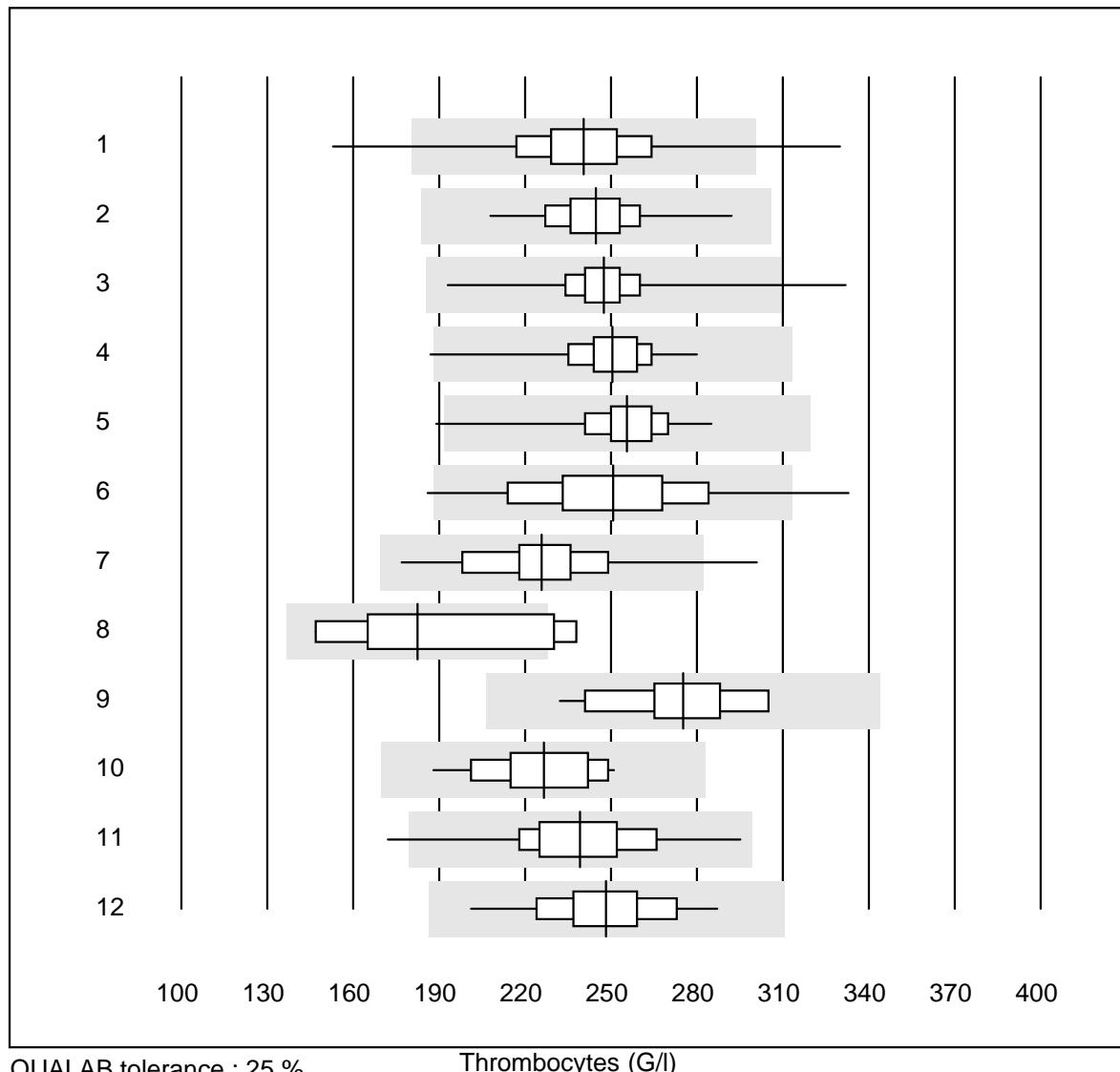


No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Abx Micros	861	99.2	0.2	0.6	7.20	5.3	e
2 Microsemi	245	100.0	0.0	0.0	7.40	3.9	e
3 Sysmex KX21	435	99.6	0.2	0.2	7.51	3.9	e
4 Sysmex Poch - 100i	218	98.6	0.0	1.4	7.60	3.6	e
5 Sysmex XP 300	161	100.0	0.0	0.0	7.82	3.0	e
6 Mythic	242	98.8	0.4	0.8	6.99	7.2	e
7 Nihon Kohden Celltac	38	97.4	0.0	2.6	7.68	3.5	e
8 Swelab	70	97.1	2.9	0.0	8.05	7.8	e
9 MS4	6	100.0	0.0	0.0	7.82	7.0	e
10 Abacus Junior	13	100.0	0.0	0.0	9.29	7.9	e
11 Medonic	18	100.0	0.0	0.0	7.87	5.4	e
12 Samsung HC10	45	100.0	0.0	0.0	7.35	4.2	e

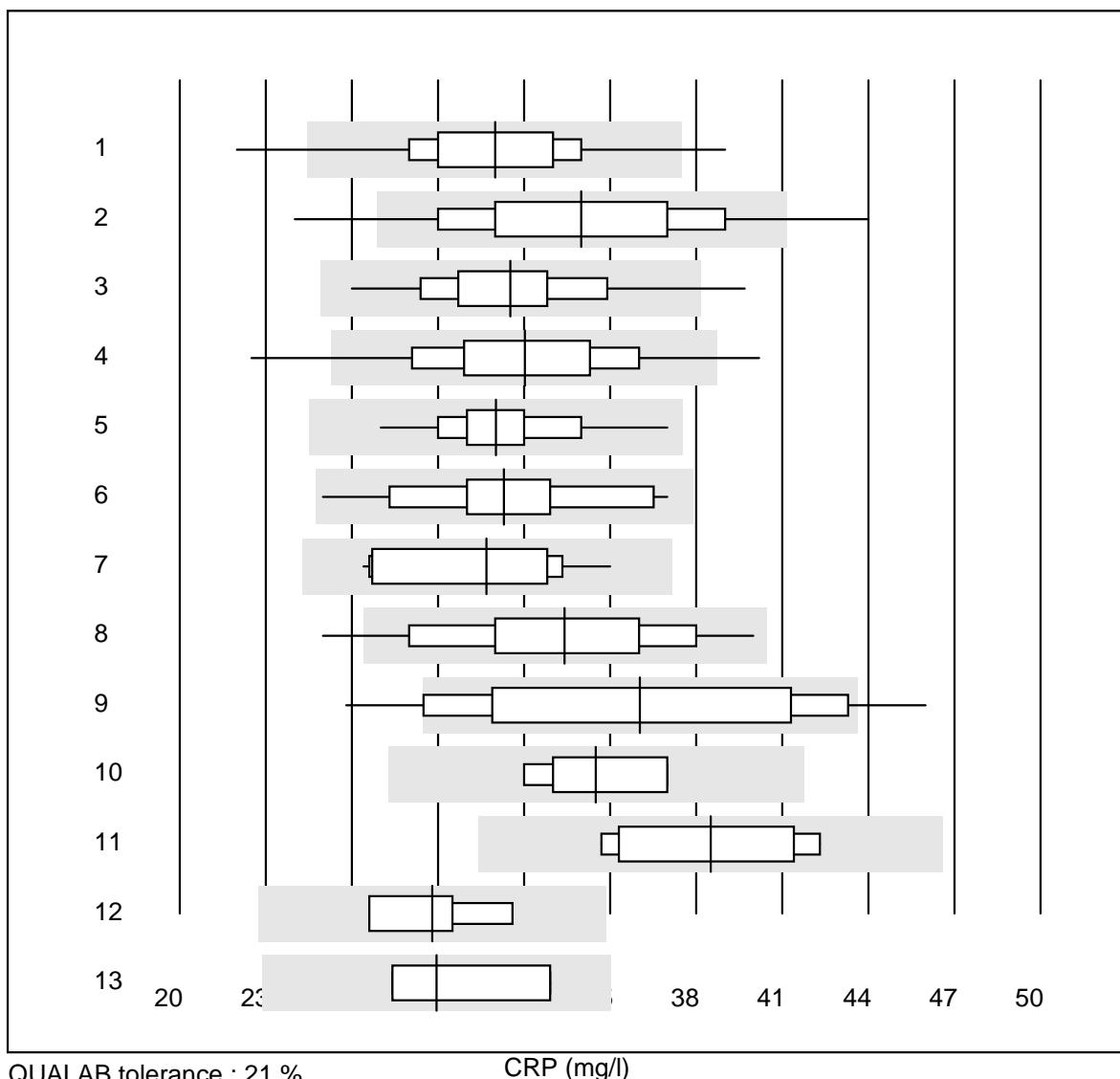
Thrombocytes



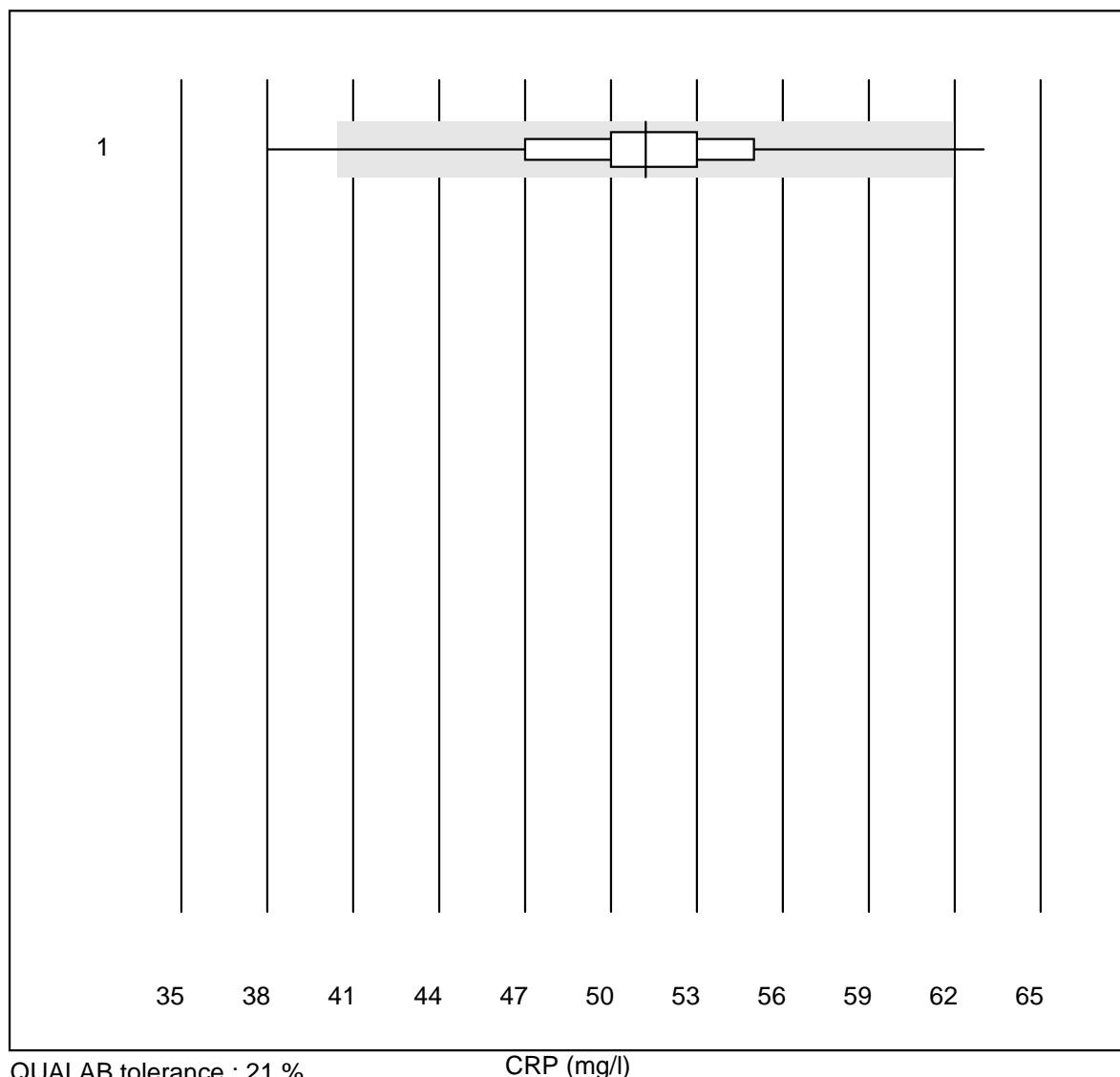
Thrombocytes



No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Abx Micros	862	96.9	2.1	1.0	240.4	8.8	e
2 Microsemi	244	100.0	0.0	0.0	244.8	5.3	e
3 Sysmex KX21	436	98.2	0.9	0.9	247.4	5.2	e
4 Sysmex Poch - 100i	216	99.0	0.5	0.5	250.6	5.1	e
5 Sysmex XP 300	161	99.4	0.6	0.0	255.6	4.9	e
6 Mythic	242	98.4	1.2	0.4	250.7	10.7	e
7 Swelab	70	98.6	1.4	0.0	225.7	8.9	e
8 MS4	6	66.7	33.3	0.0	182.5	19.7	e*
9 Abacus Junior	13	100.0	0.0	0.0	275.2	8.4	e
10 Medonic	18	100.0	0.0	0.0	226.5	7.7	e
11 Nihon Kohden Celltac	38	97.4	2.6	0.0	239.3	9.5	e
12 Samsung HC10	45	97.8	0.0	2.2	248.3	7.5	e

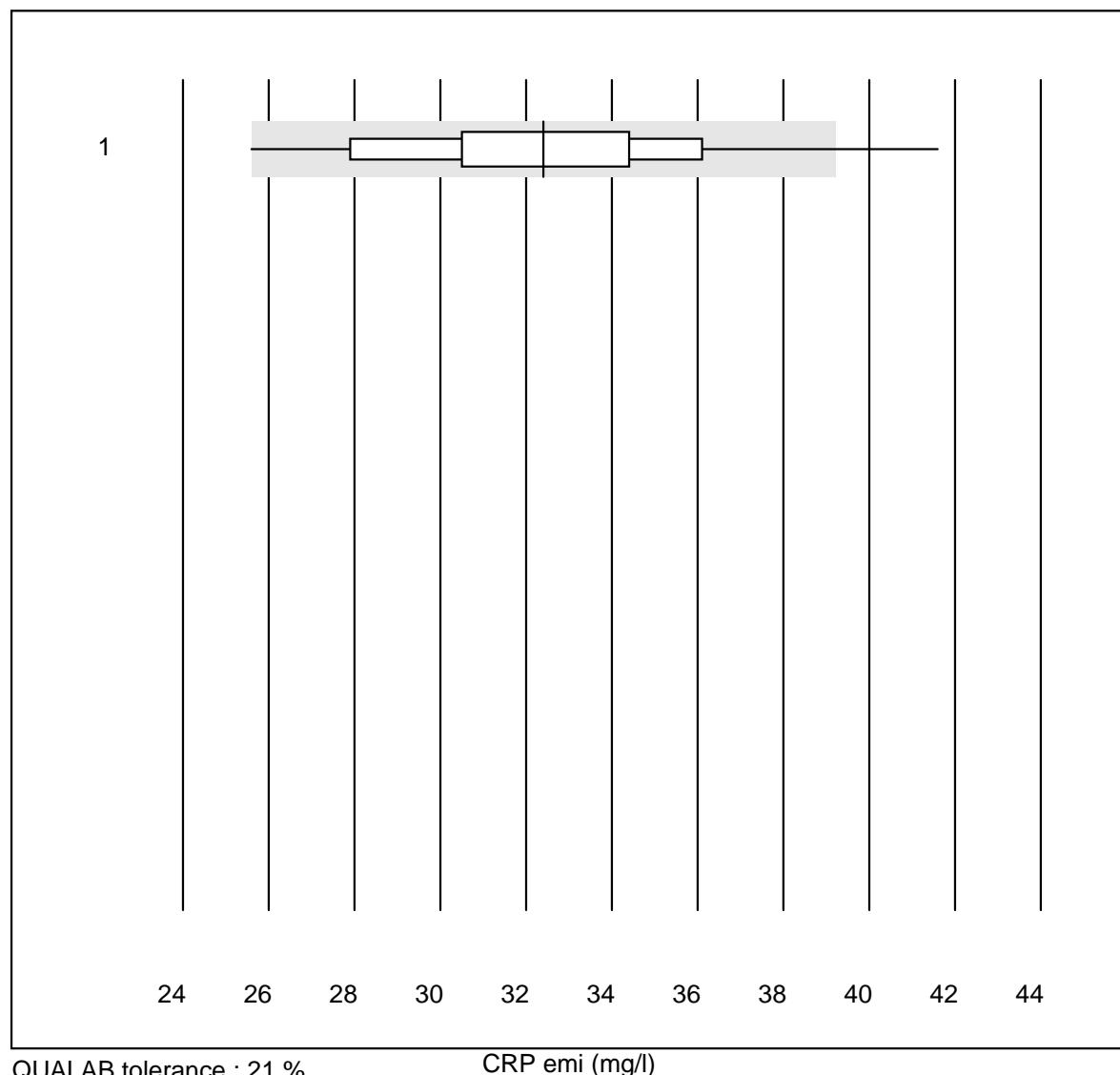
CRP

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Afinion	1147	98.6	1.1	0.3	31.0	8.2	e
2 NycoCard SingleTest-	460	81.7	6.3	12.0	34.0	11.8	e
3 Abx Micros	135	97.8	1.5	0.7	31.5	8.0	e
4 ABX Micros CRP200	332	95.2	3.9	0.9	32.0	10.0	e
5 Quick Read go	87	98.9	0.0	1.1	31.0	6.2	e
6 Turbidimetry	40	97.5	0.0	2.5	31.3	8.7	e
7 Cobas	11	100.0	0.0	0.0	30.7	9.6	e*
8 Fuji Dri-Chem	23	78.3	4.3	17.4	33.4	10.8	e
9 Eurolyser	126	70.6	15.9	13.5	36.0	15.9	e
10 AQT 90 FLEX	6	100.0	0.0	0.0	34.5	6.5	e*
11 Spotchem D-Concept	7	100.0	0.0	0.0	38.5	7.5	e*
12 Spotchem SI-3510	4	100.0	0.0	0.0	28.8	7.3	e*
13 Other methods	4	75.0	0.0	25.0	29.0	9.1	e*

CRP

No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	QuikRead (Vollblut)	179	96.7	1.1	2.2	51.2	6.5	e

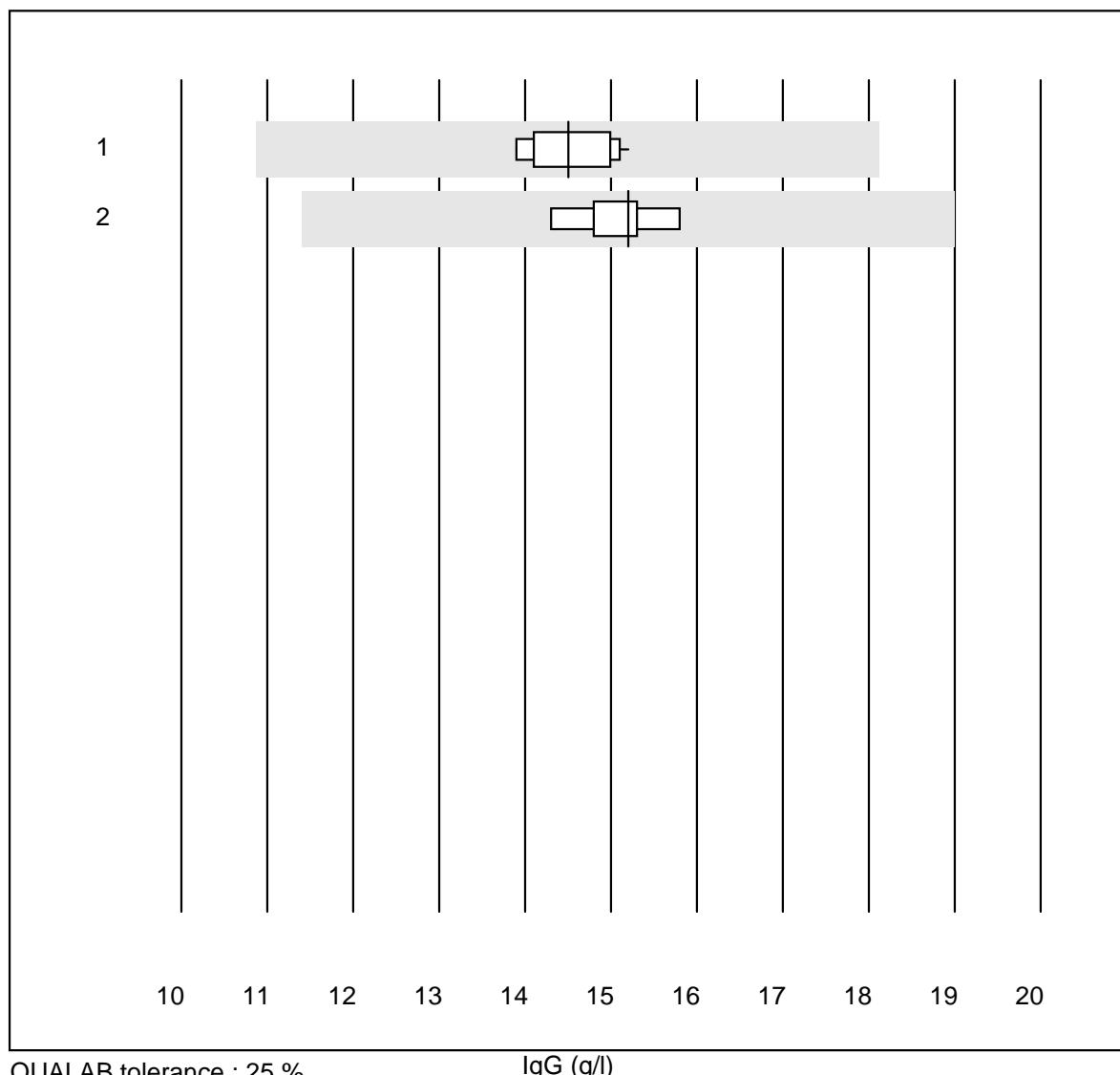
CRP emi



No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Microsemi	243	97.6	1.6	0.8	32.4	9.5	e

I2 Plasmaproteins

IgG

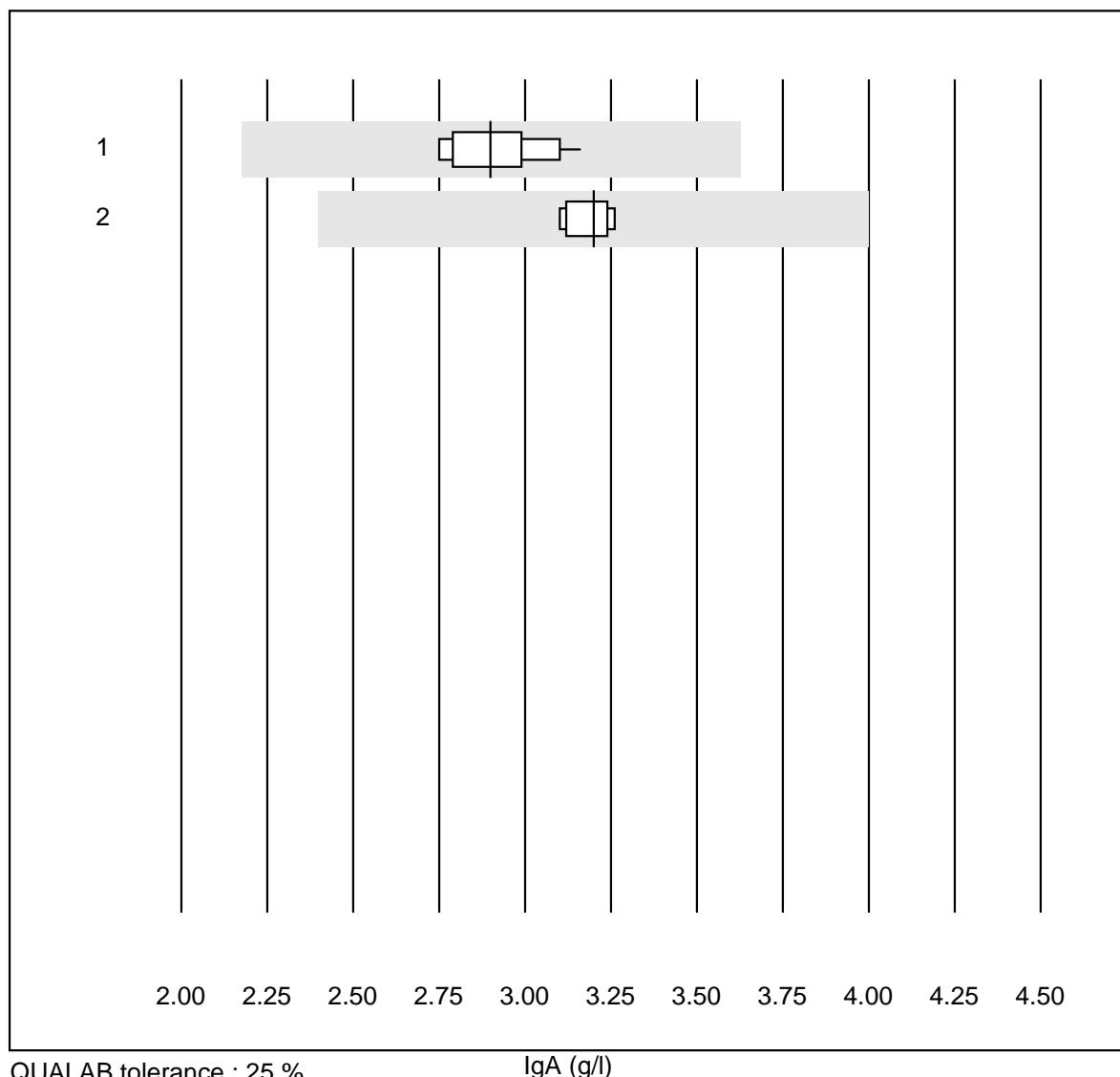


QUALAB tolerance : 25 %

IgG (g/l)

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Turbidimetry	10	100.0	0.0	0.0	14.5	3.3	e
2 Nephelometry	7	85.7	0.0	14.3	15.2	3.4	e

I2 Plasmaproteins

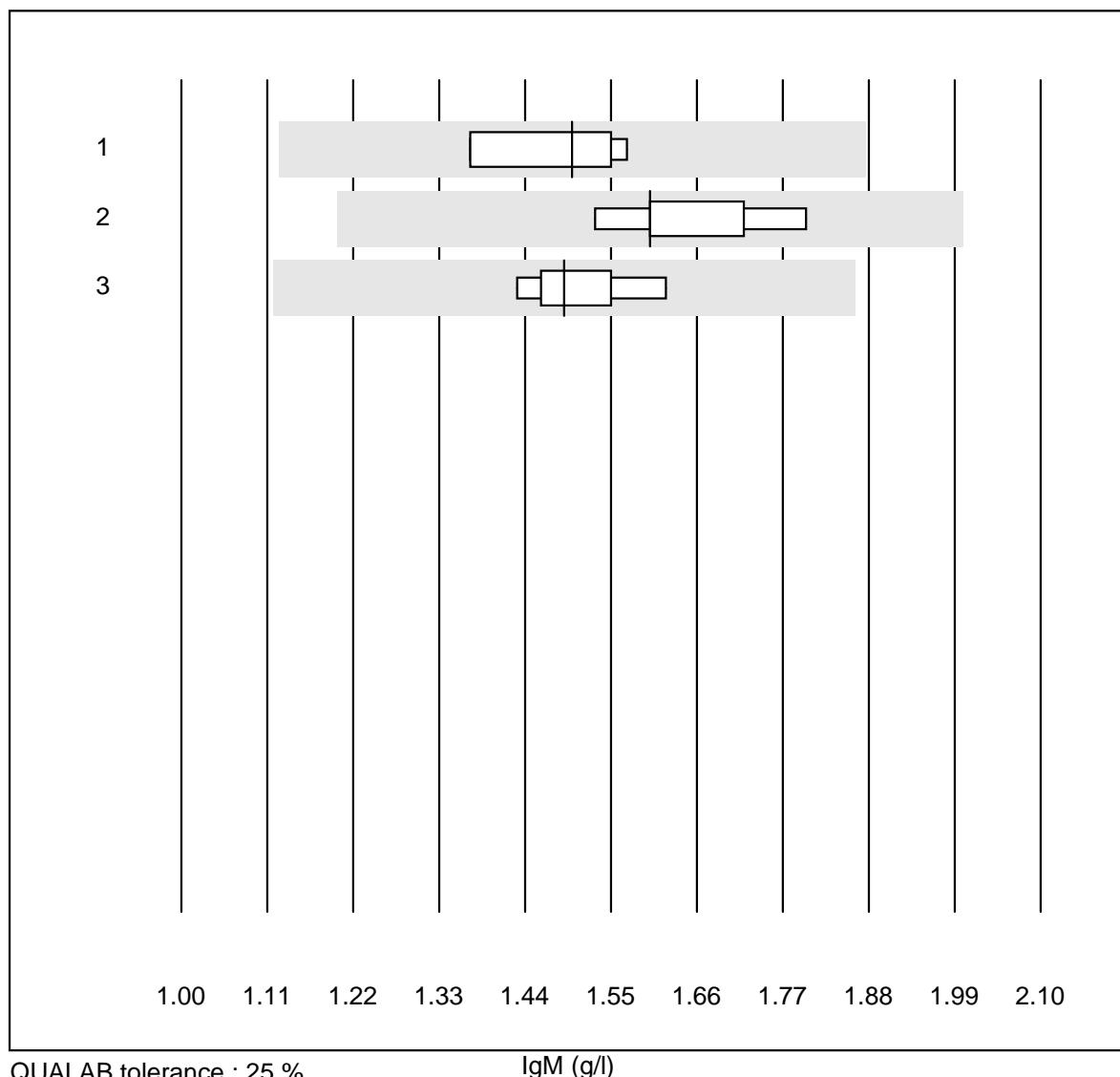
IgA

QUALAB tolerance : 25 %

IgA (g/l)

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Turbidimetry	10	100.0	0.0	0.0	2.9	4.9	e
2 Nephelometry	7	85.7	0.0	14.3	3.2	2.0	e

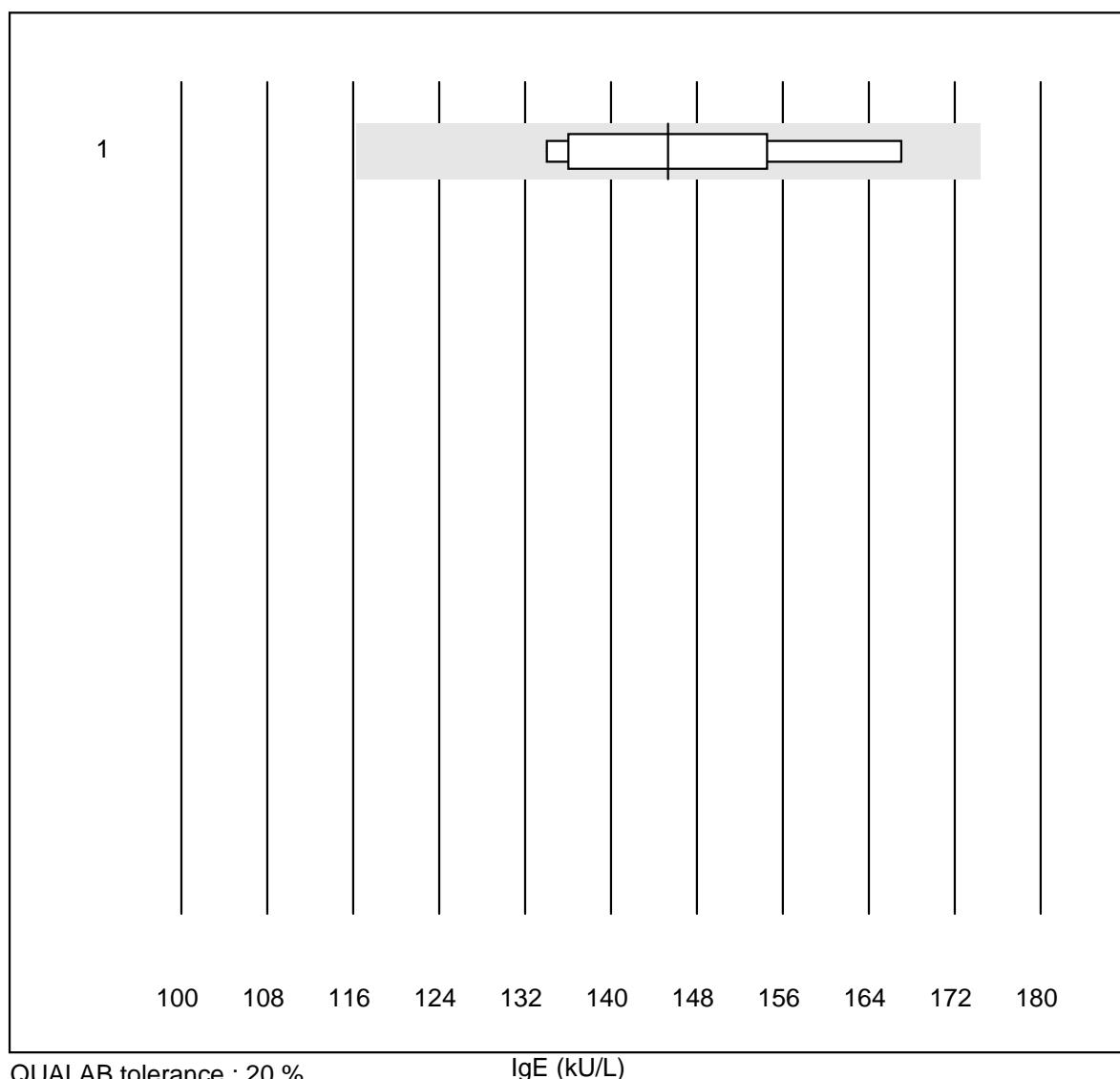
I2 Plasmaproteins

IgM

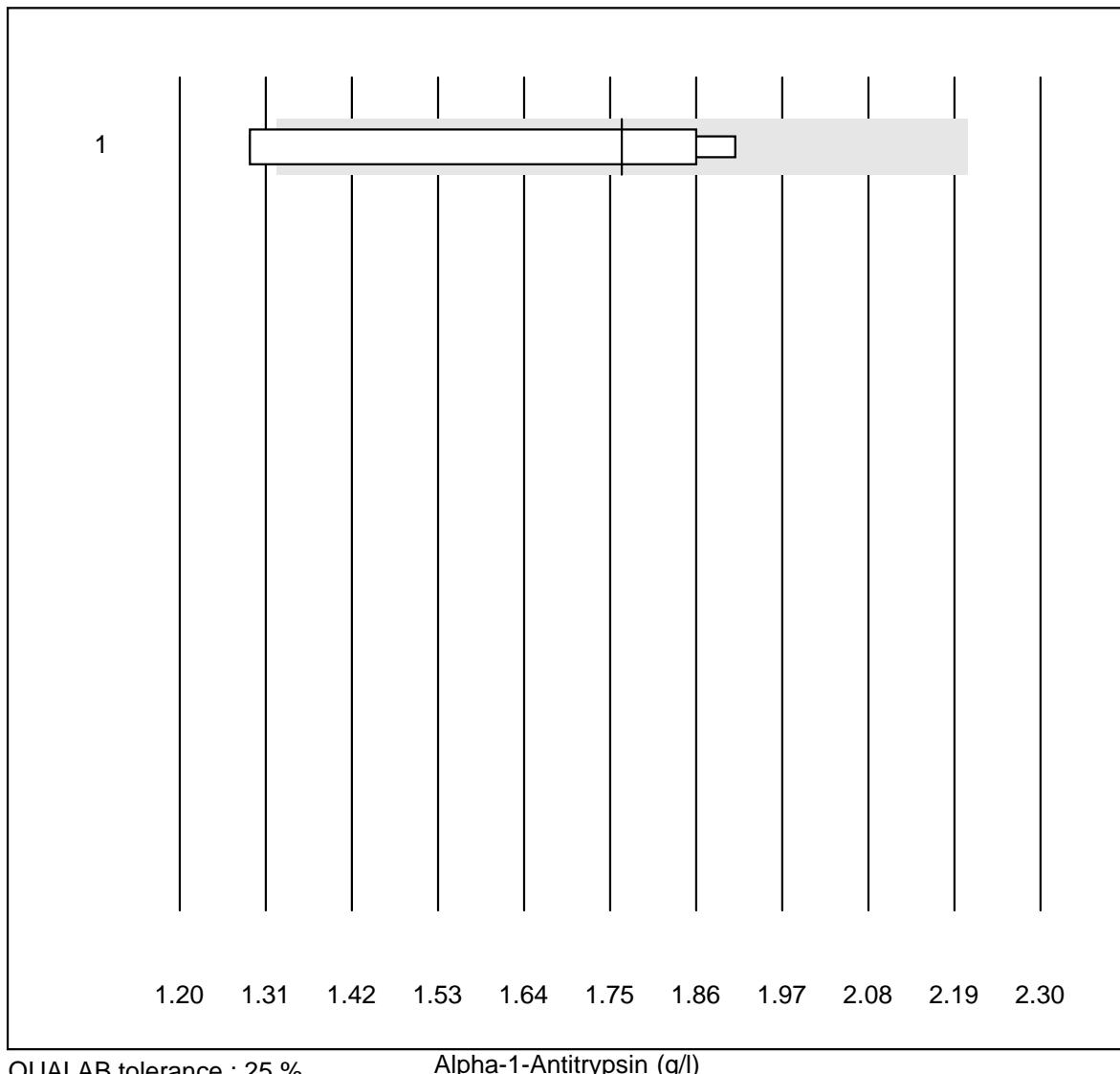
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Turbidimetry	4	100.0	0.0	0.0	1.5	6.3	e*
2 Nephelometry	7	85.7	0.0	14.3	1.6	5.9	e
3 Cobas Integra 800/40	6	100.0	0.0	0.0	1.5	4.5	e

I2 Plasmaproteins

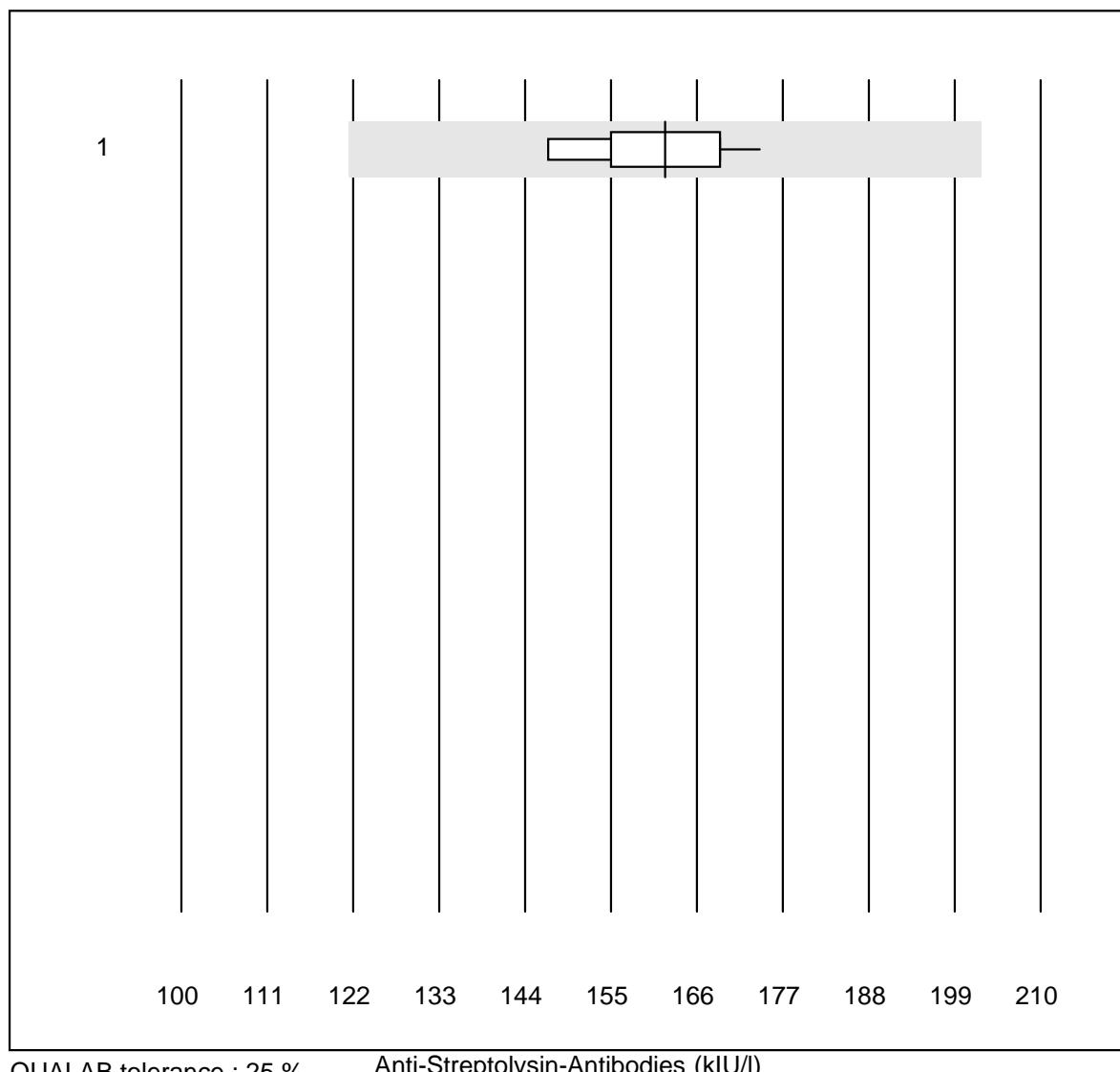
IgE



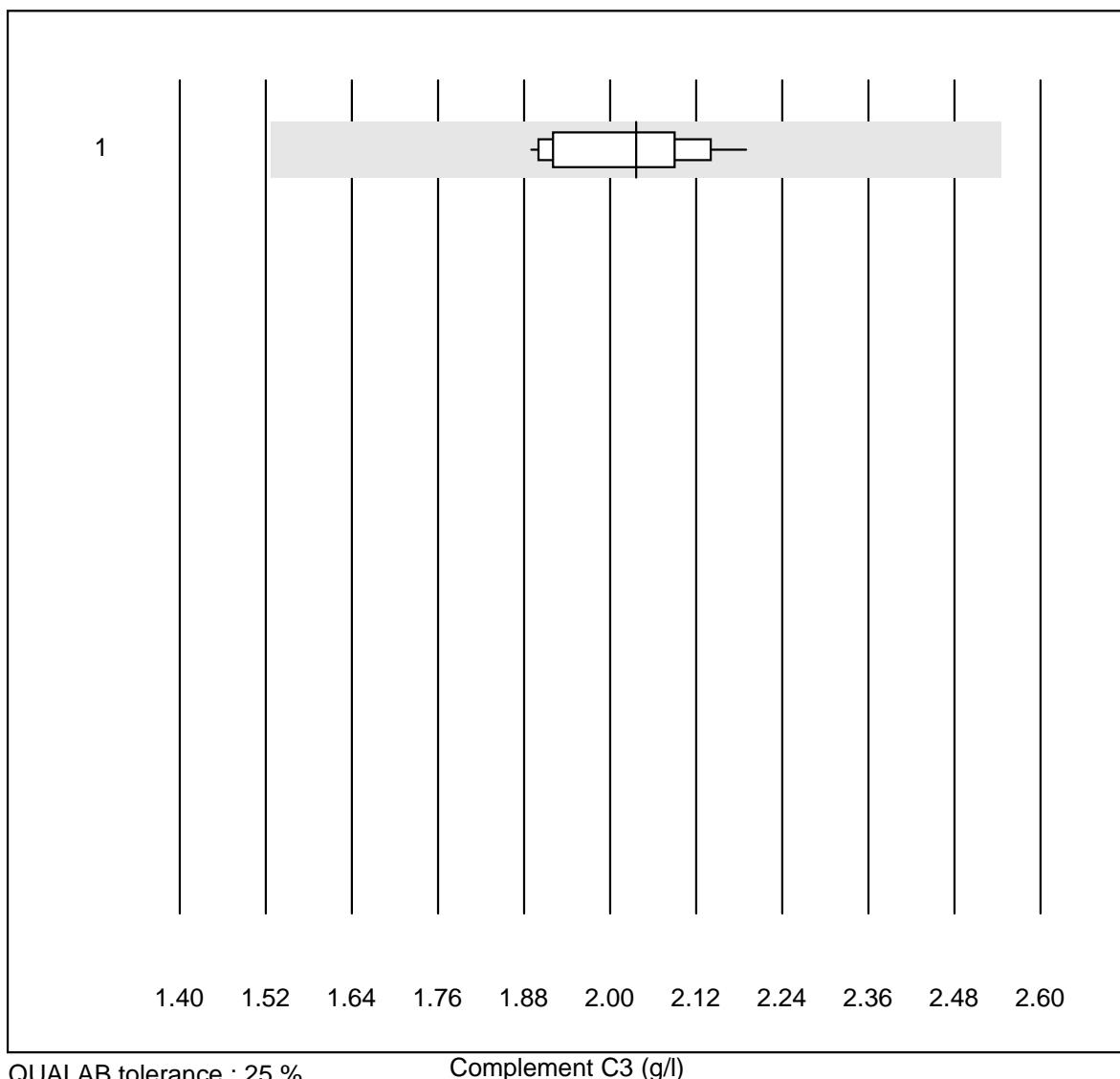
Alpha-1-Antitrypsin



Anti-Streptolysin-Antibodies

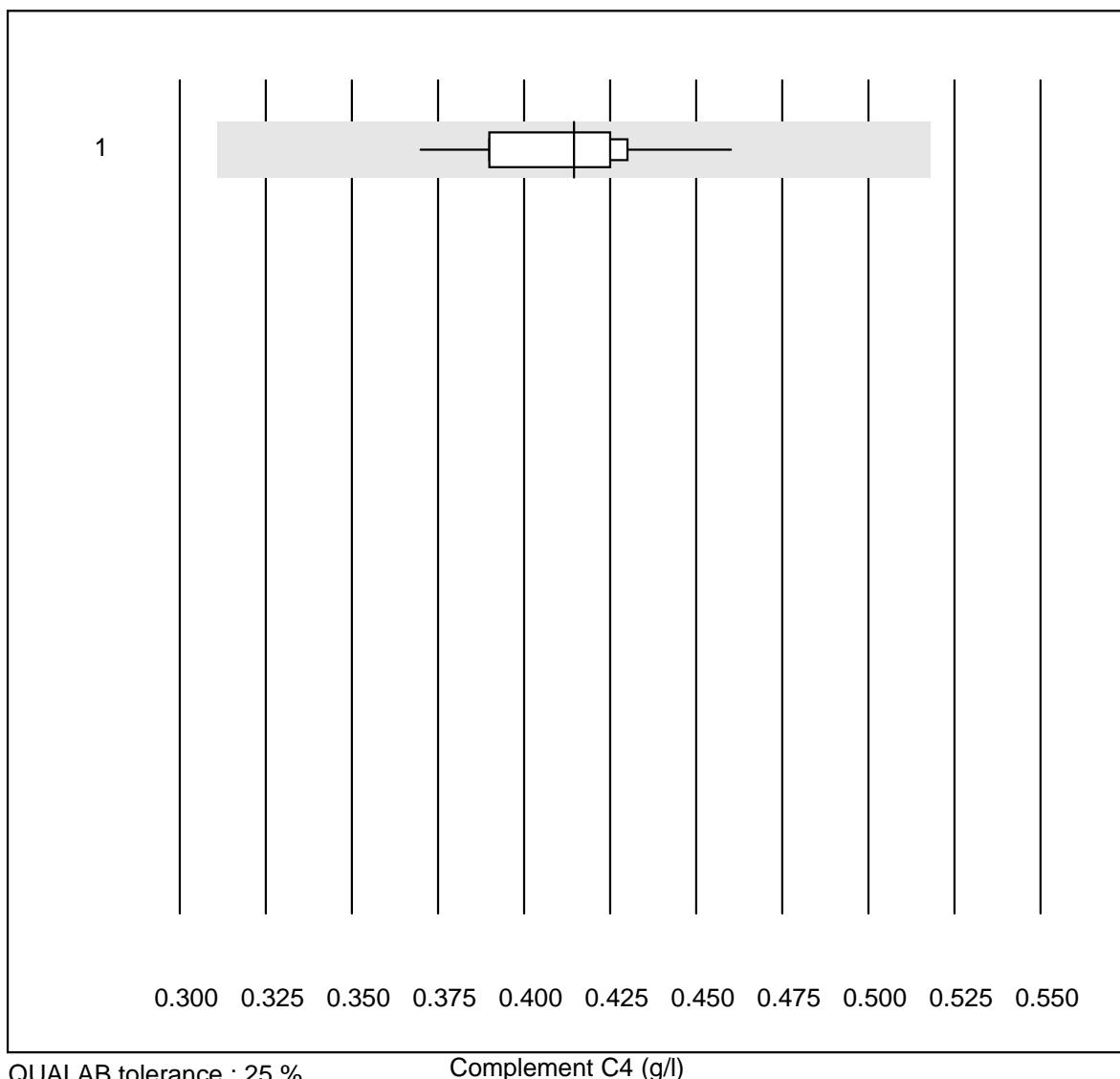


No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 all Participants	10	100.0	0.0	0.0	162	5.8	e

Complement C3

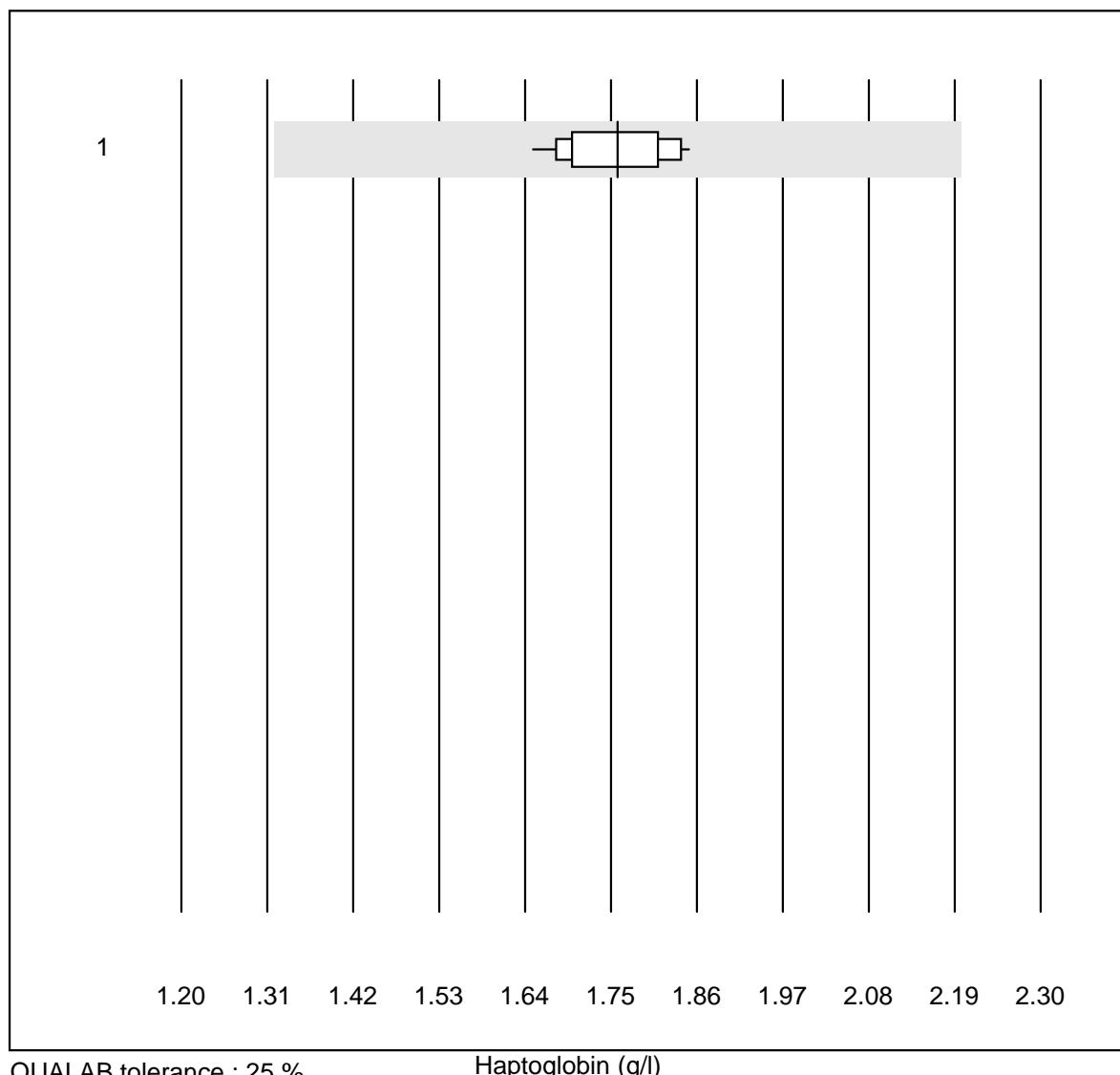
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	all Participants	12	100.0	0.0	0.0	2.04	4.8	e

Complement C4



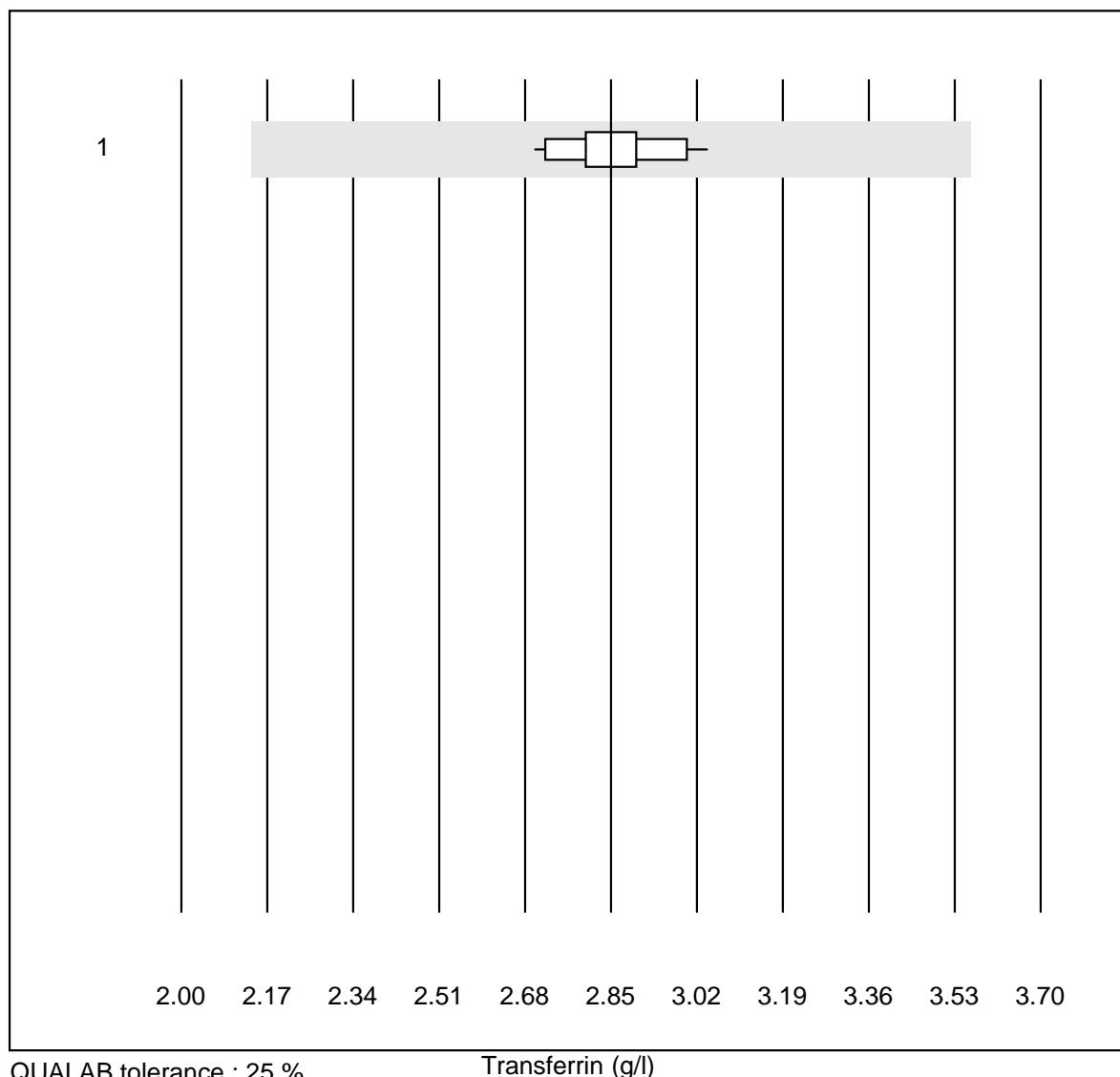
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 all Participants	11	100.0	0.0	0.0	0.41	5.8	e

Haptoglobin



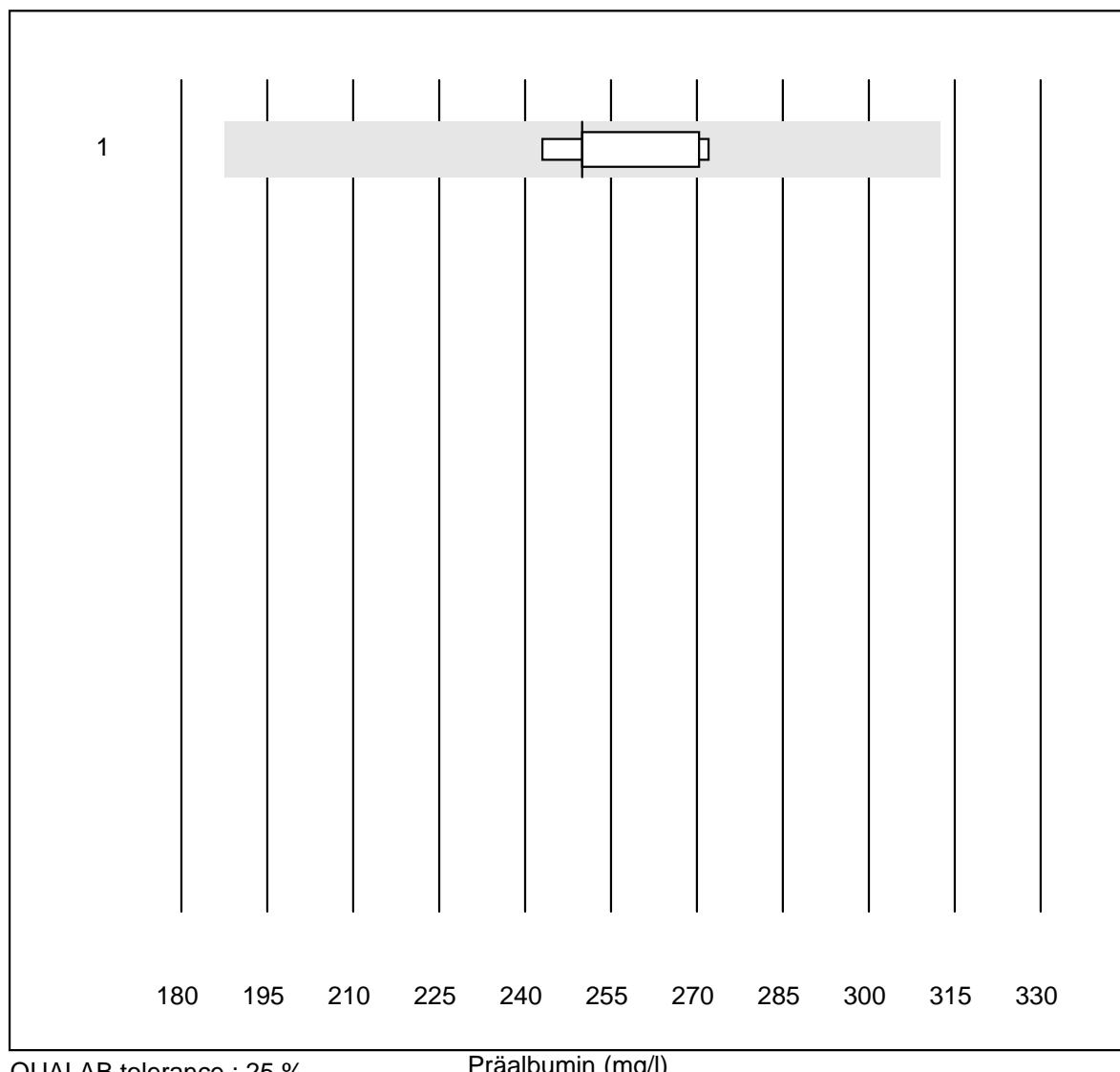
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	all Participants	13	100.0	0.0	0.0	1.76	3.7	e

Transferrin



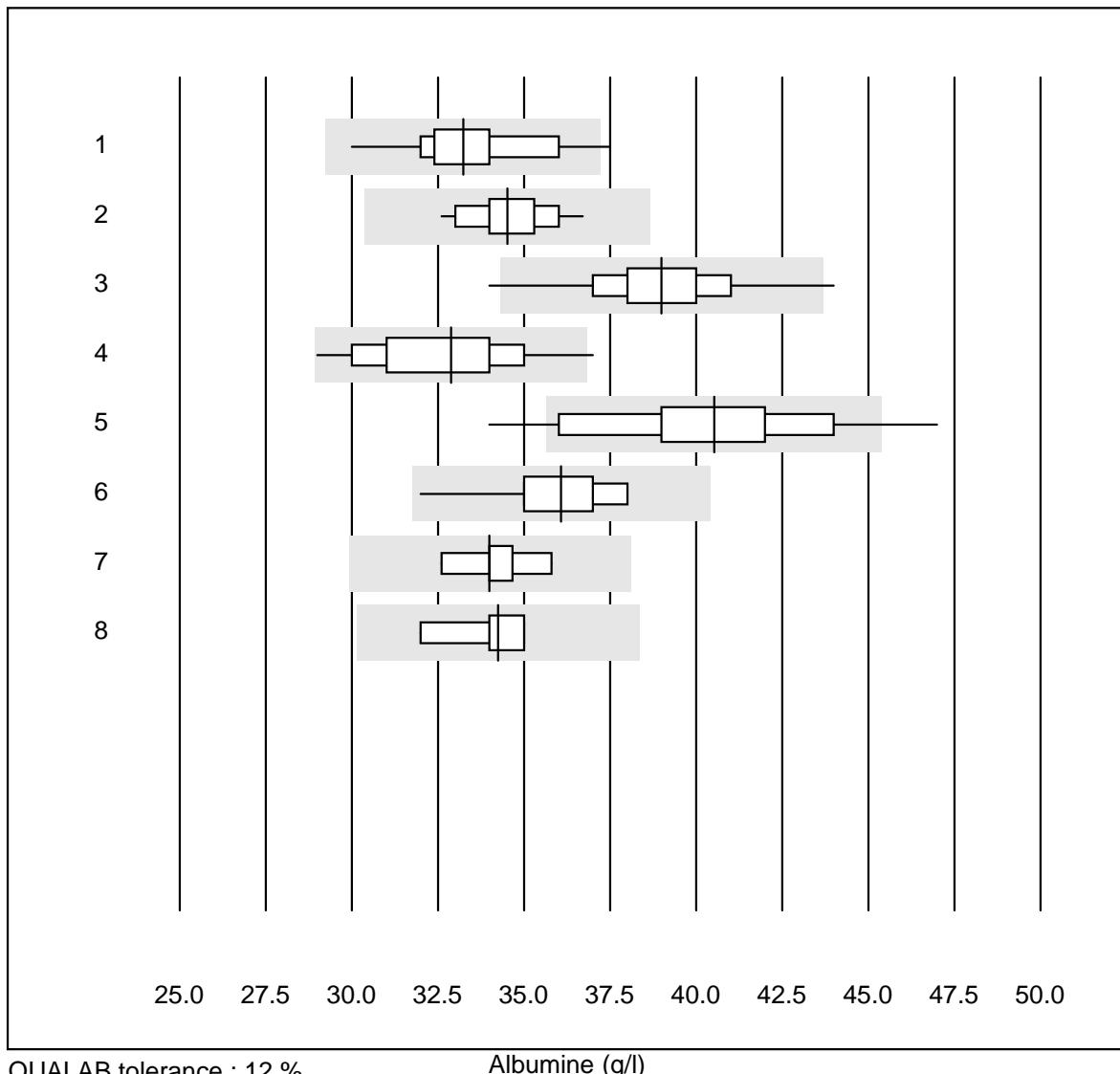
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	all Participants	14	100.0	0.0	0.0	2.85	3.6	e

Präalbumin



No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 all Participants	6	100.0	0.0	0.0	250.0	4.7	e

Albumine

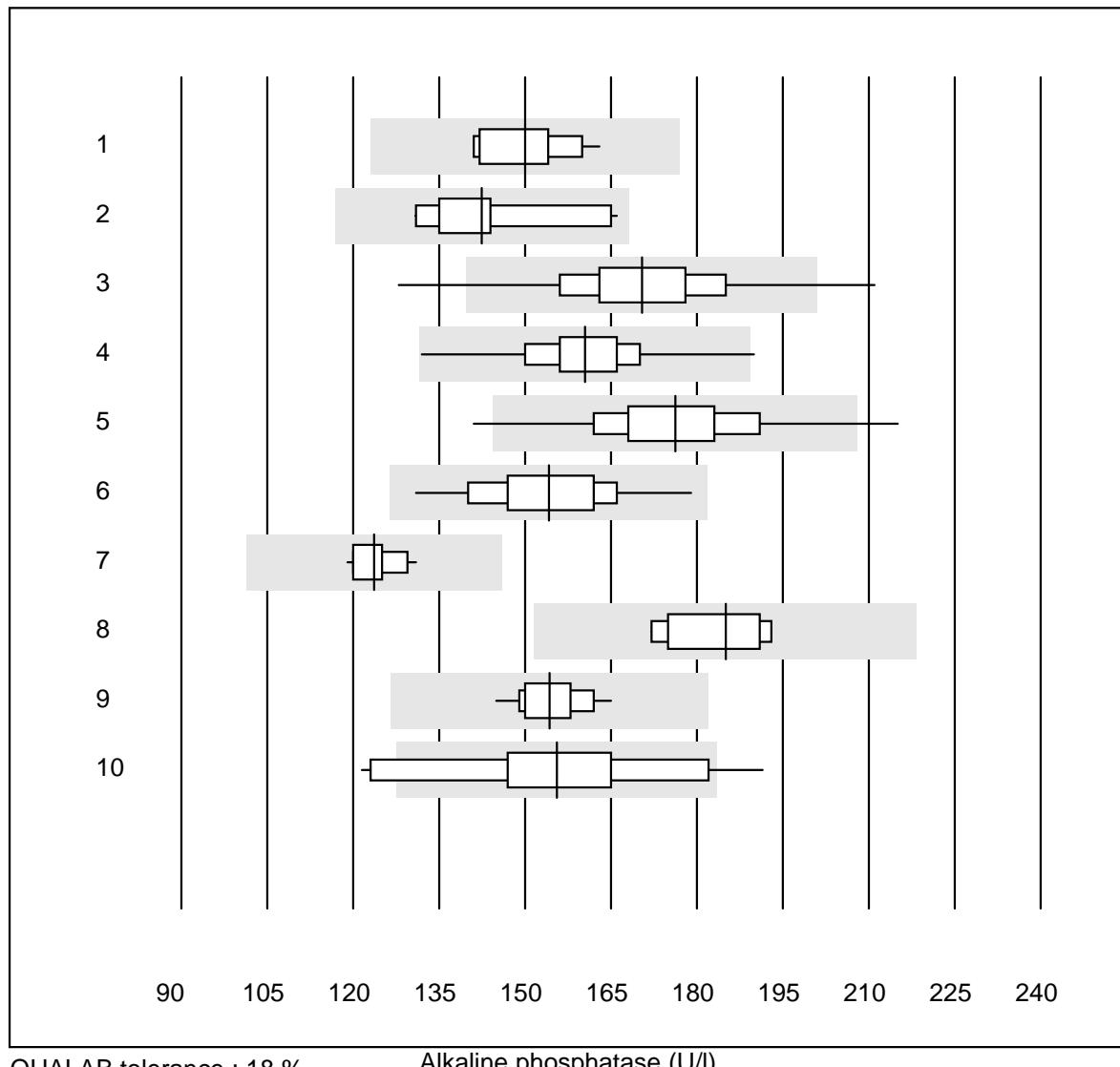


QUALAB tolerance : 12 %

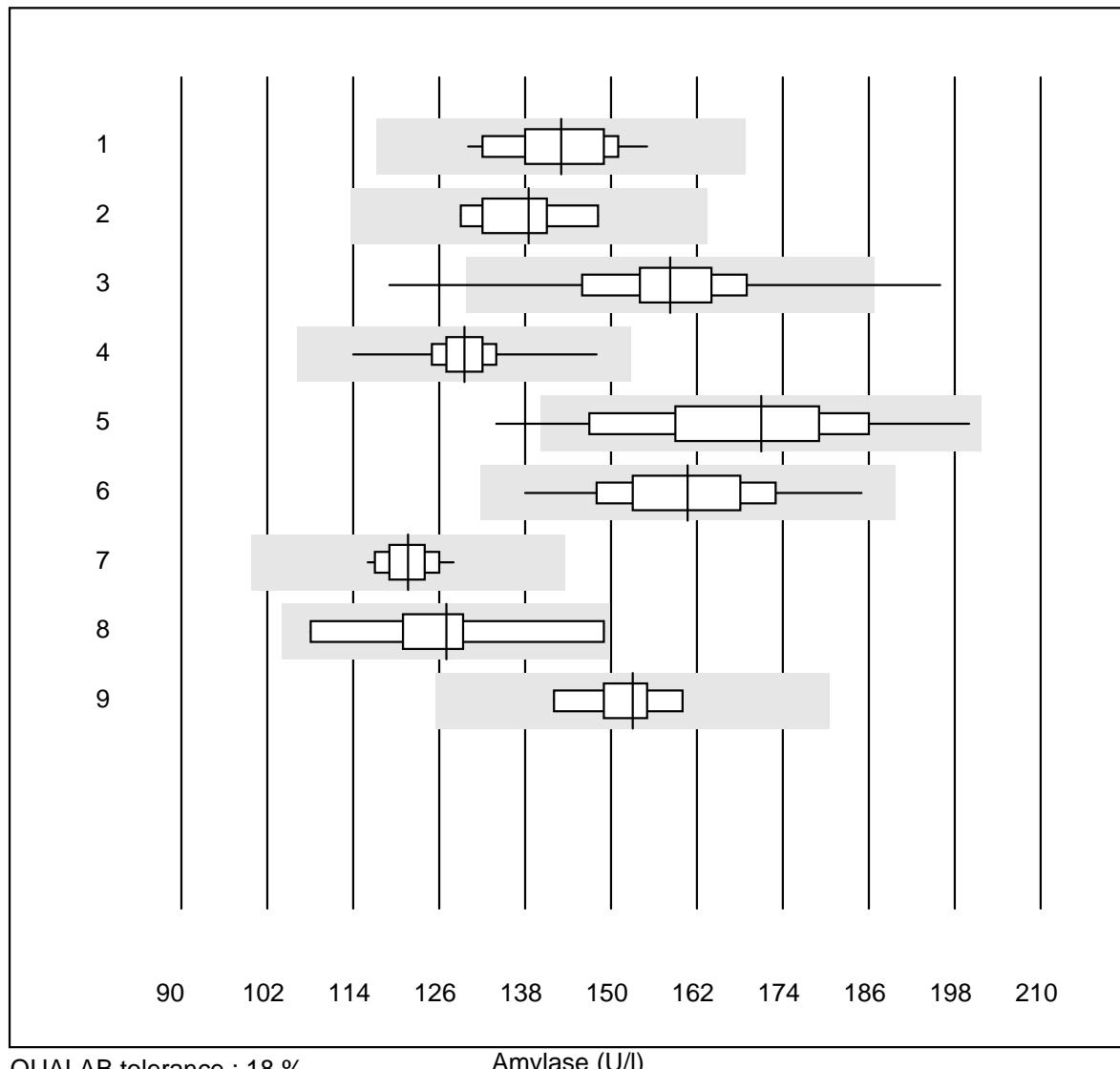
Albumine (g/l)

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Standard chemistry	17	94.1	5.9	0.0	33	5.0	e
2 Cobas	13	100.0	0.0	0.0	35	3.5	e
3 Fuji Dri-Chem	172	98.8	1.2	0.0	39	3.8	e
4 Spotchem/Ready	41	97.6	2.4	0.0	33	6.1	e
5 Spotchem D-Concept	70	90.0	8.6	1.4	41	6.8	e
6 Piccolo	26	100.0	0.0	0.0	36	3.8	e
7 Abx Mira	8	100.0	0.0	0.0	34	2.9	e
8 Hitachi S40/M40	8	87.5	0.0	12.5	34	3.0	e

Alkaline phosphatase

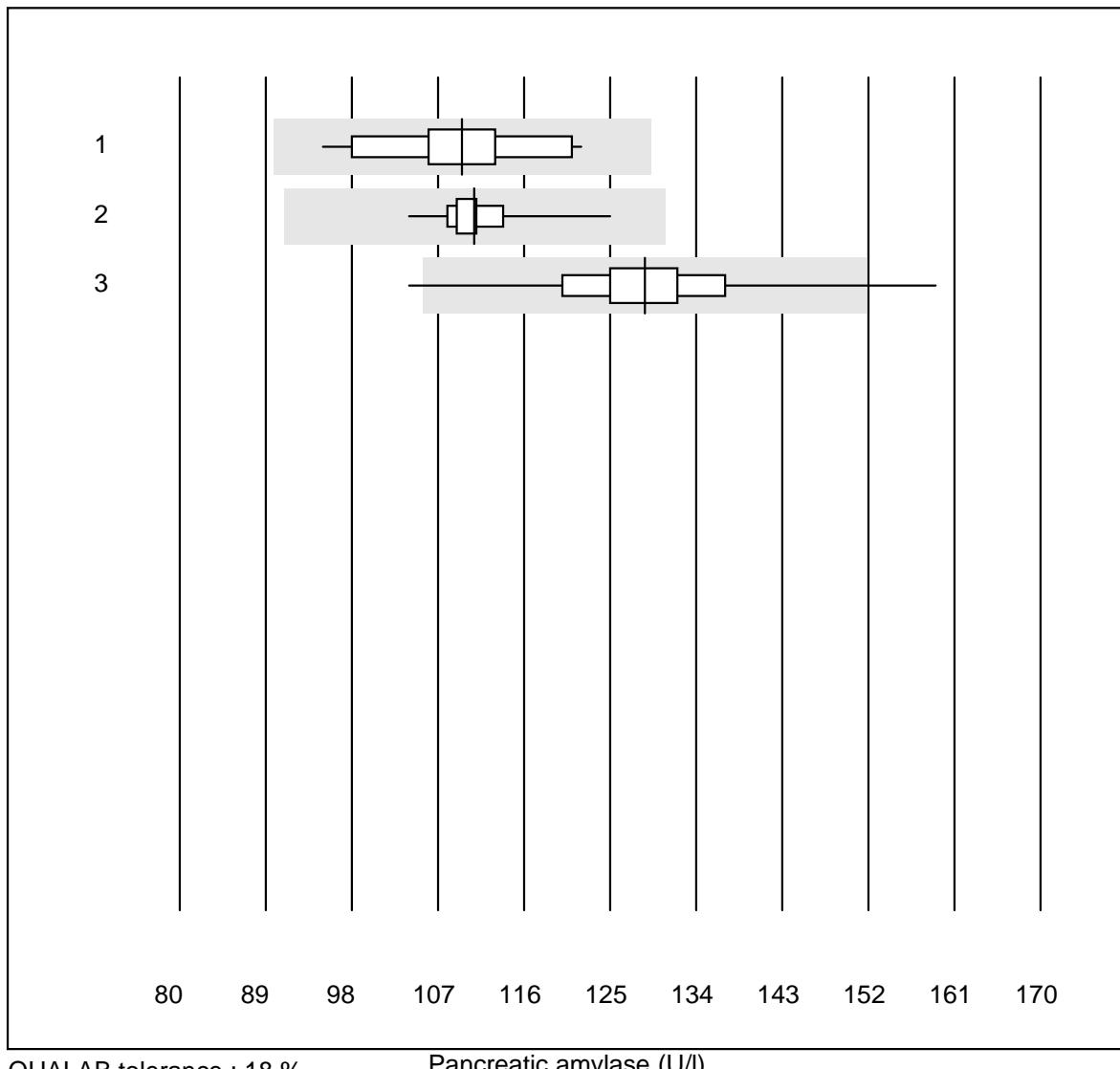


Amylase



No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 IFCC EPS liquid 37°C	11	100.0	0.0	0.0	143	5.6	e
2 Cobas	6	100.0	0.0	0.0	139	5.0	e
3 Reflotron	184	96.7	3.3	0.0	158	6.5	e
4 Fuji Dri-Chem	490	99.8	0.0	0.2	130	3.1	e
5 Spotchem/Ready	78	96.2	3.8	0.0	171	8.9	a
6 Spotchem D-Concept	95	98.9	0.0	1.1	161	6.1	e
7 Piccolo	25	96.0	0.0	4.0	122	2.8	e
8 Abx Mira	9	100.0	0.0	0.0	127	8.6	e*
9 Hitachi S40/M40	6	100.0	0.0	0.0	153	4.1	e

Pancreatic amylase

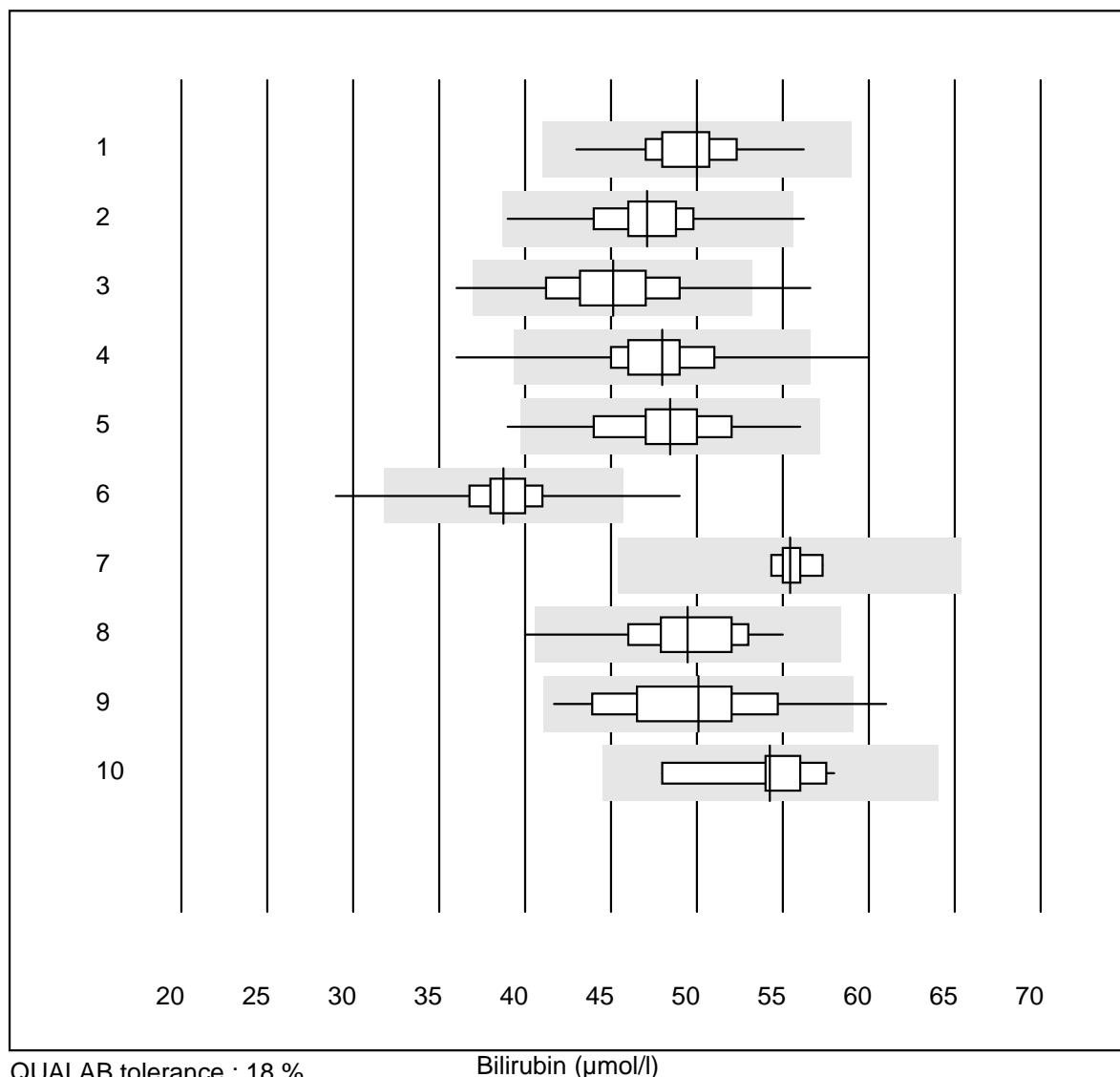


QUALAB tolerance : 18 %

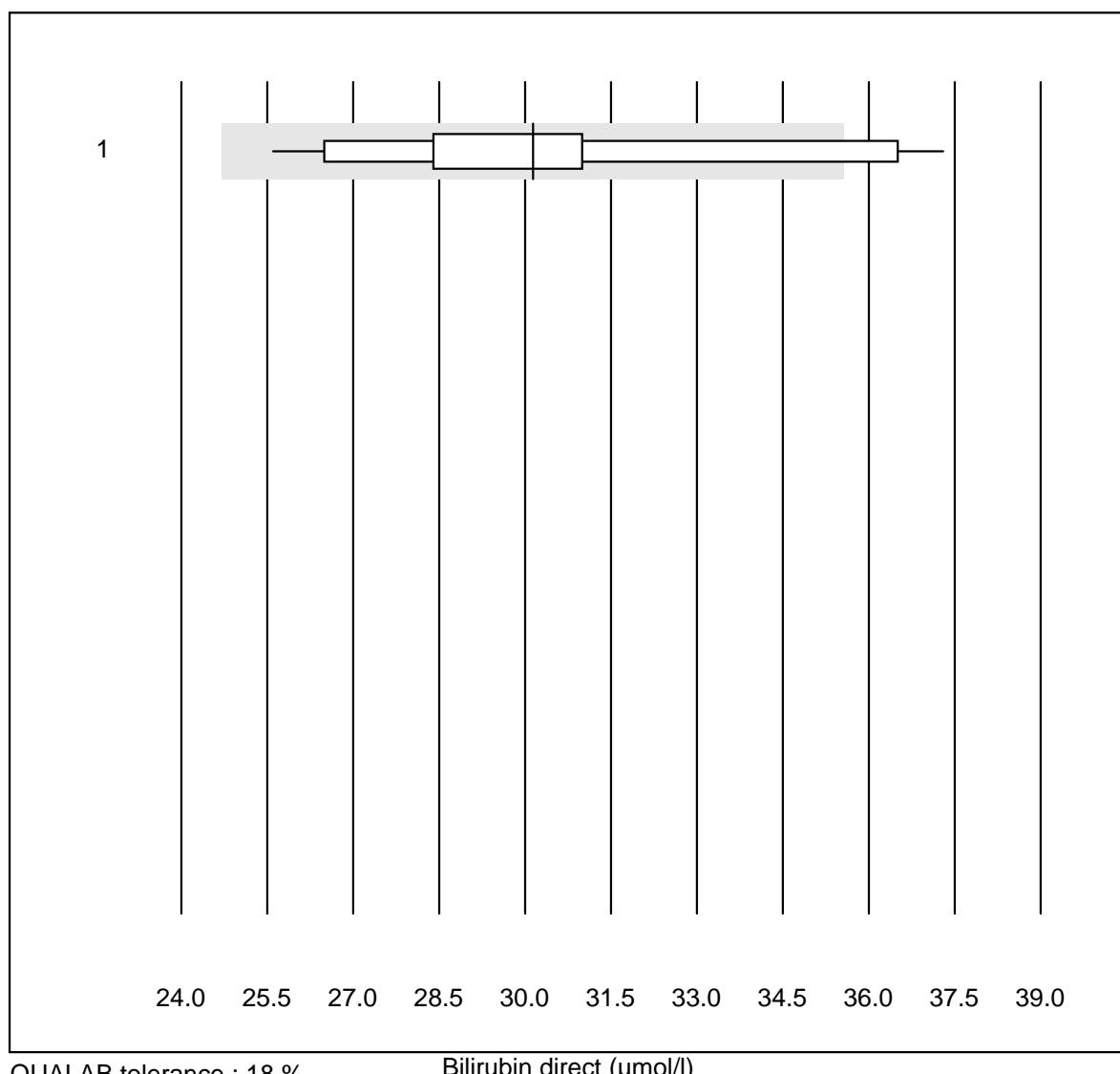
Pancreatic amylase (U/l)

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 IFCC EPS liquid 37°C	18	94.4	0.0	5.6	110	6.8	e
2 Cobas	13	100.0	0.0	0.0	111	4.4	e
3 Reflotron	444	97.5	1.8	0.7	129	5.6	e

Bilirubin

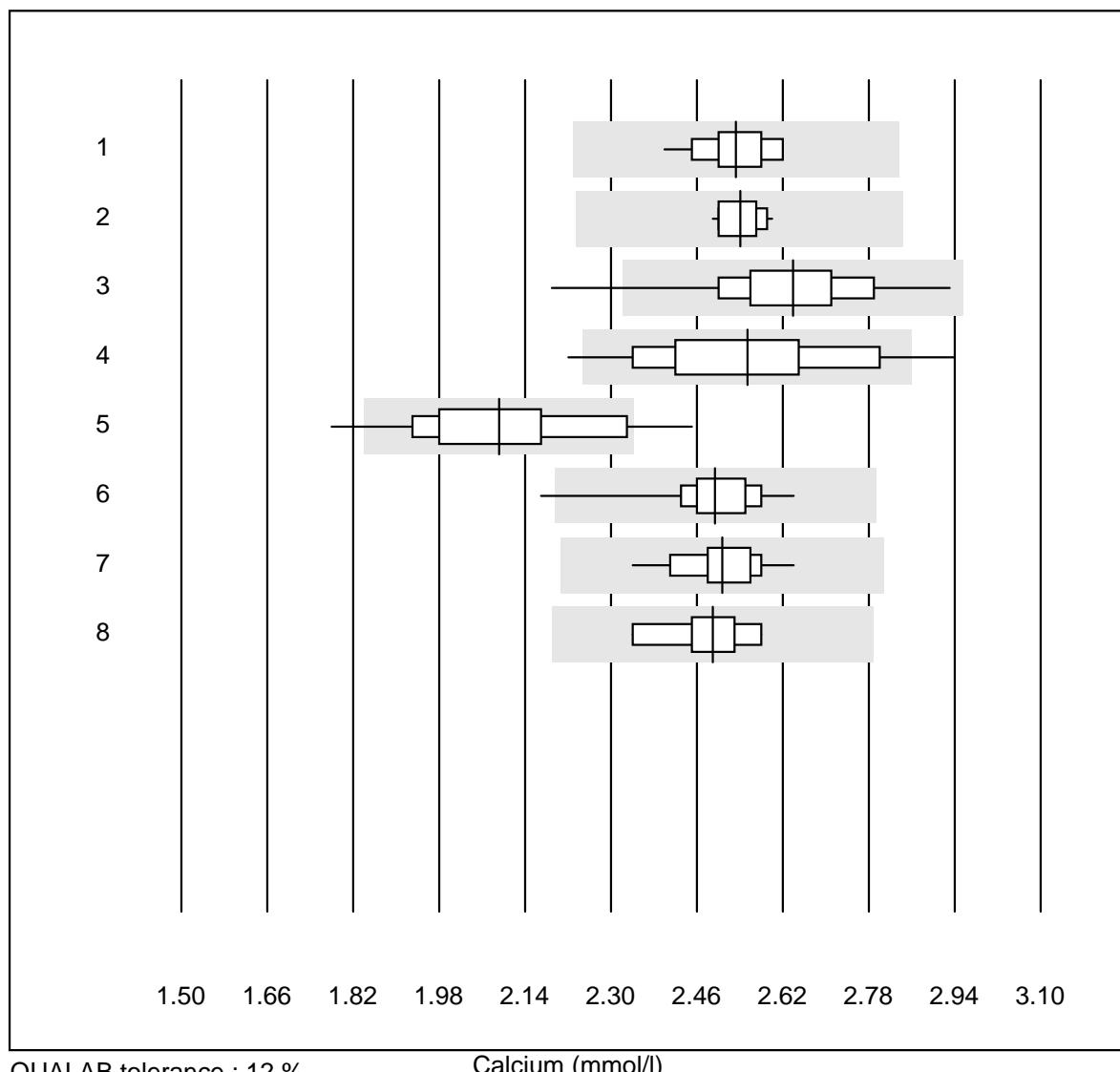


Bilirubin direct



No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Fuji Dri-Chem	29	86.3	10.3	3.4	30.1	10.4	e

Calcium

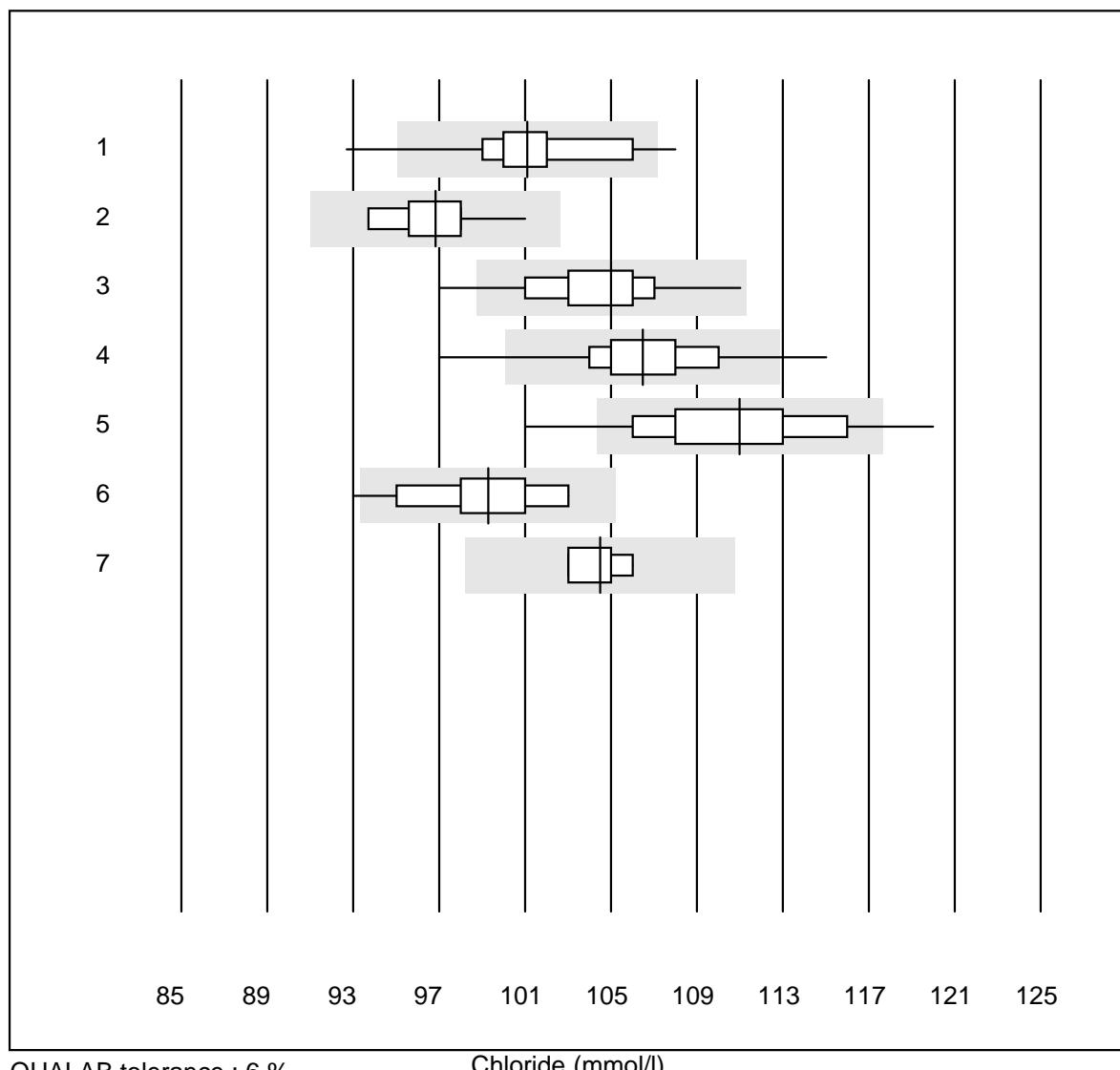


QUALAB tolerance : 12 %

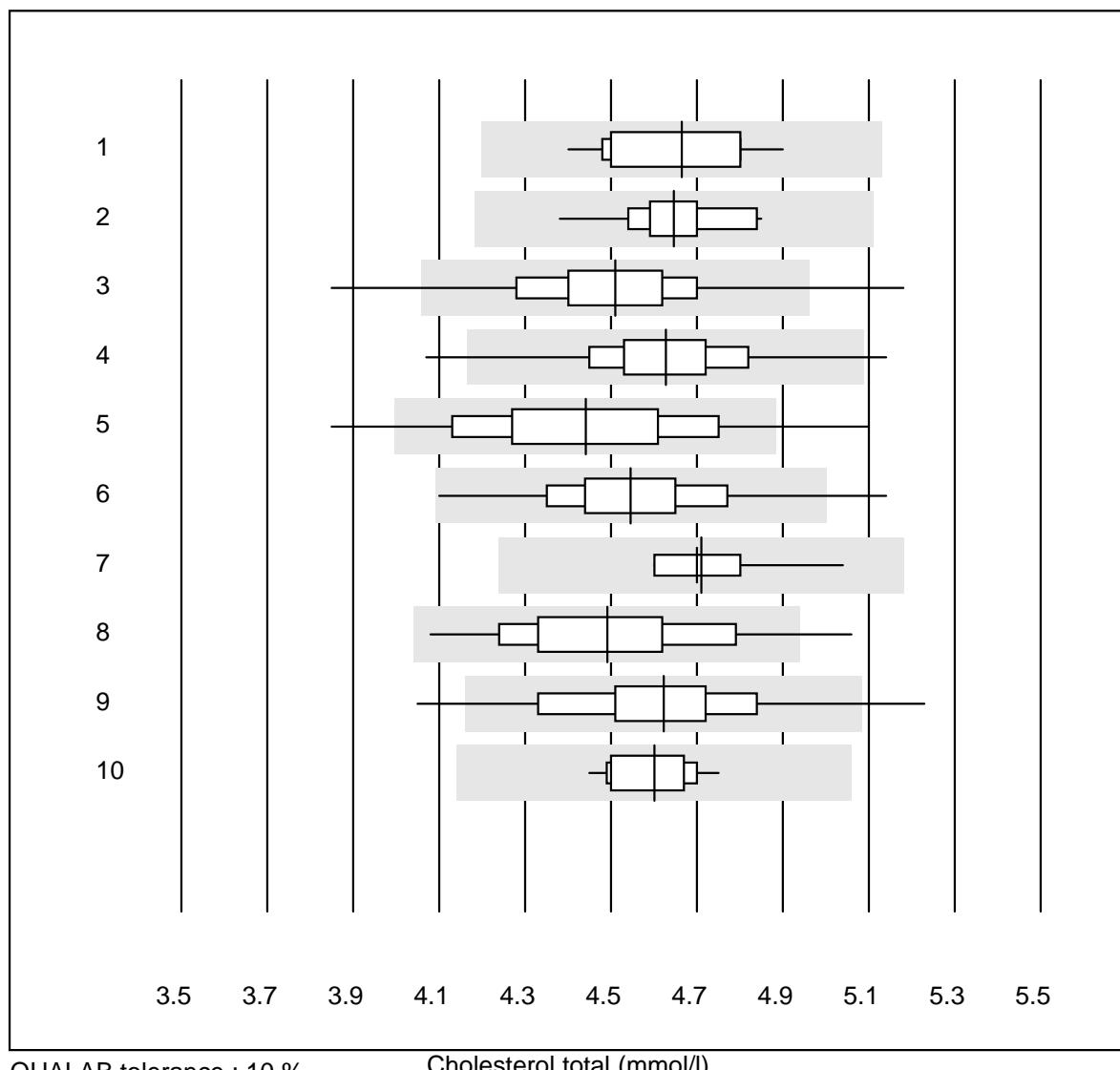
Calcium (mmol/l)

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Standard chemistry	19	100.0	0.0	0.0	2.53	2.3	e
2 Cobas	13	100.0	0.0	0.0	2.54	1.6	e
3 Fuji Dri-Chem	339	99.4	0.3	0.3	2.64	4.3	e
4 Spotchem/Ready	44	95.5	4.5	0.0	2.55	6.5	e
5 Spotchem D-Concept	66	84.8	15.2	0.0	2.09	7.5	e
6 Piccolo	26	96.2	3.8	0.0	2.49	3.5	e
7 Abx Mira	13	100.0	0.0	0.0	2.51	3.1	e
8 Hitachi S40/M40	10	90.0	0.0	10.0	2.49	2.8	e

Chloride

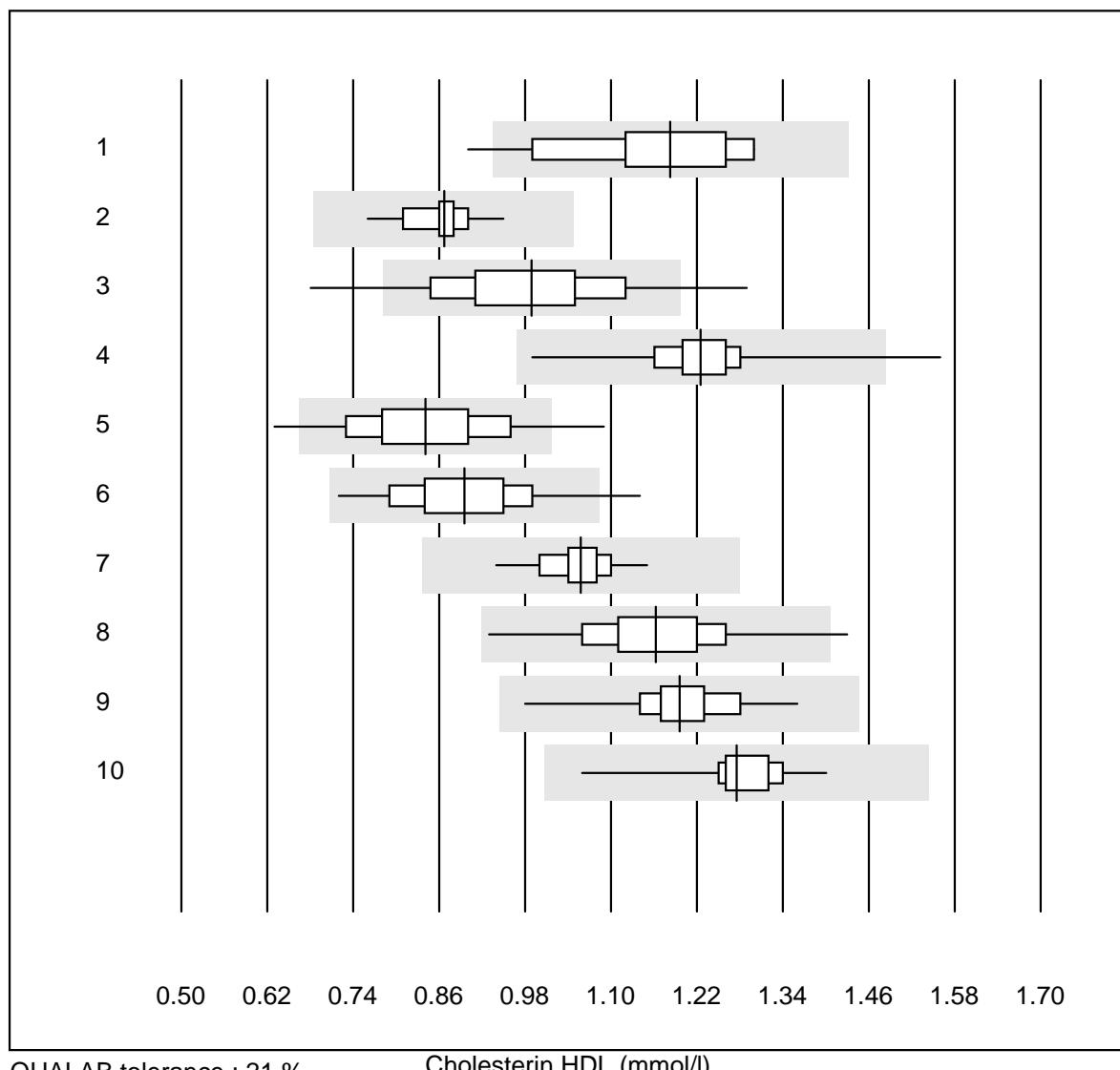


No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 ISE	15	86.7	13.3	0.0	101	3.4	e*
2 Cobas	10	100.0	0.0	0.0	97	2.1	e
3 Fuji Dri-Chem	578	96.7	2.1	1.2	105	2.3	e
4 Spotchem D-Concept	114	96.4	1.8	1.8	106	2.5	e
5 Spotchem EL-SE 1520	111	81.1	10.8	8.1	111	3.6	e
6 Piccolo	17	94.1	5.9	0.0	99	2.7	e
7 iStat Chem8	4	100.0	0.0	0.0	105	1.2	e

Cholesterol total

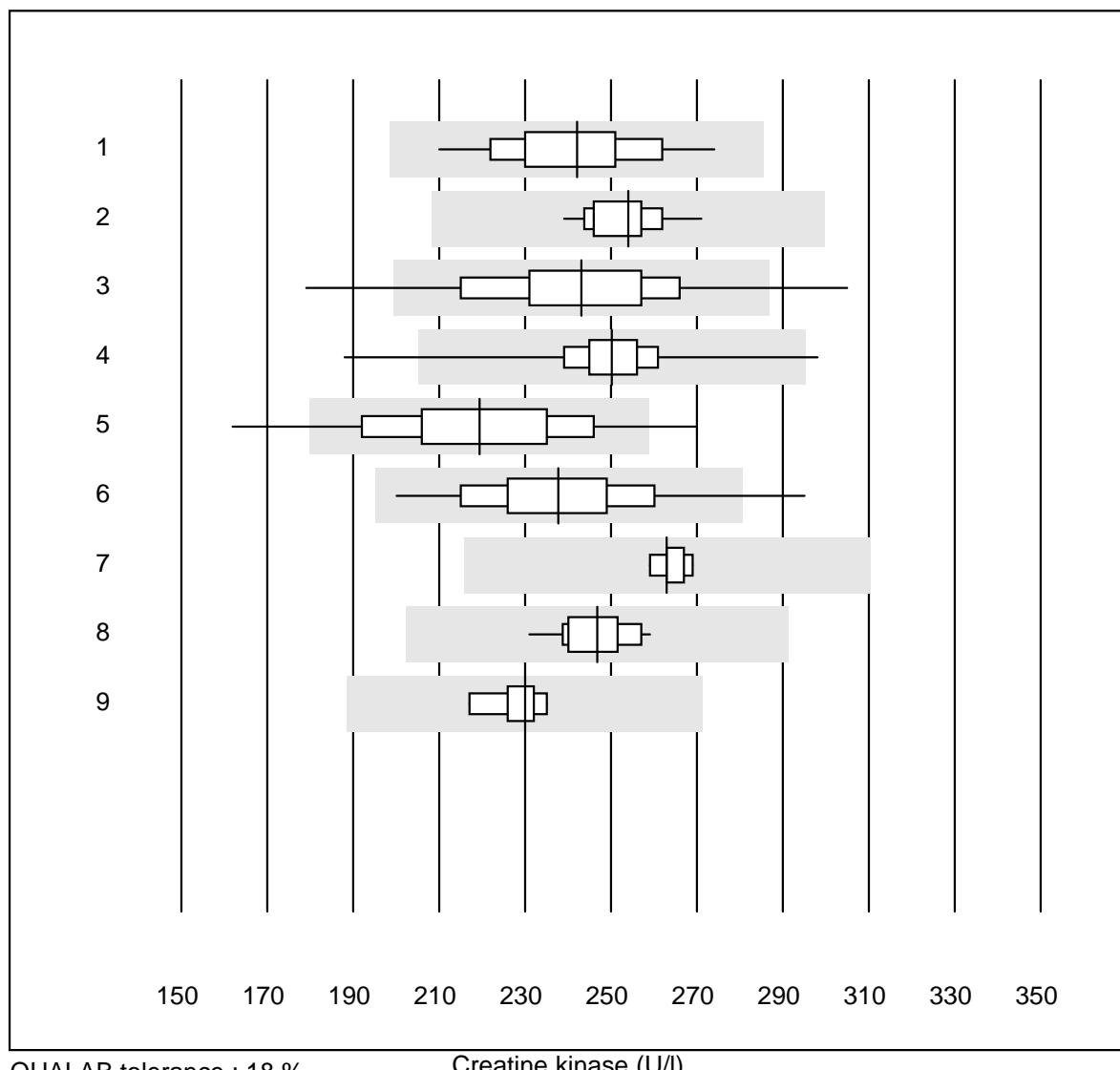
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Standard chemistry	19	100.0	0.0	0.0	4.7	3.0	e
2	Cobas	16	100.0	0.0	0.0	4.6	2.5	e
3	Reflotron	790	97.7	1.5	0.8	4.5	3.8	e
4	Fuji Dri-Chem	677	99.2	0.4	0.4	4.6	3.3	e
5	Spotchem/Ready	144	88.2	6.9	4.9	4.4	5.5	e
6	Spotchem D-Concept	131	98.5	1.5	0.0	4.5	3.7	e
7	Piccolo	22	100.0	0.0	0.0	4.7	2.1	e
8	Cholestech LDX	192	96.9	2.1	1.0	4.5	4.7	e
9	Abx Mira	19	89.5	10.5	0.0	4.6	5.2	e
10	Hitachi S40/M40	12	100.0	0.0	0.0	4.6	2.0	e

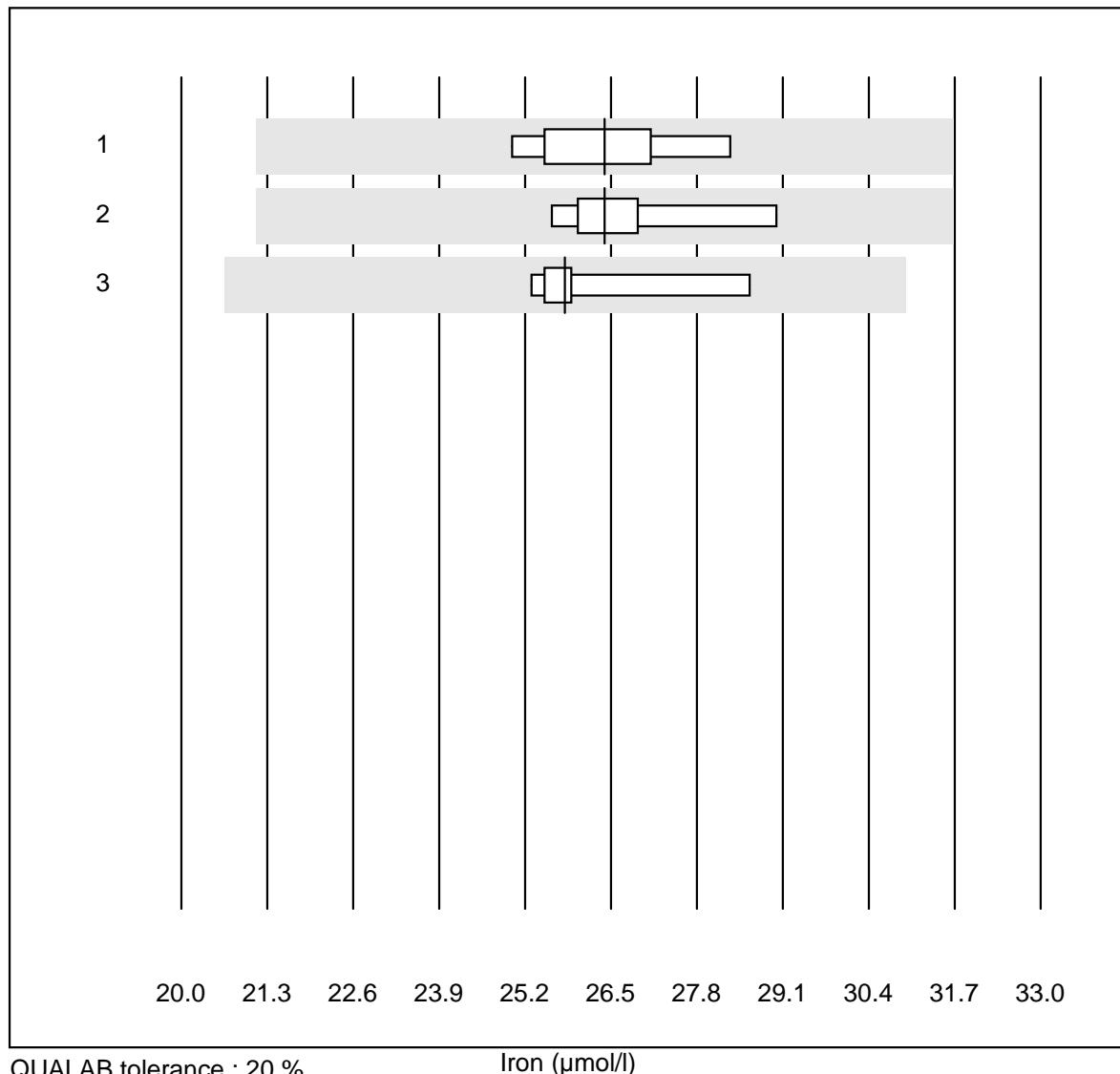
Cholesterin HDL



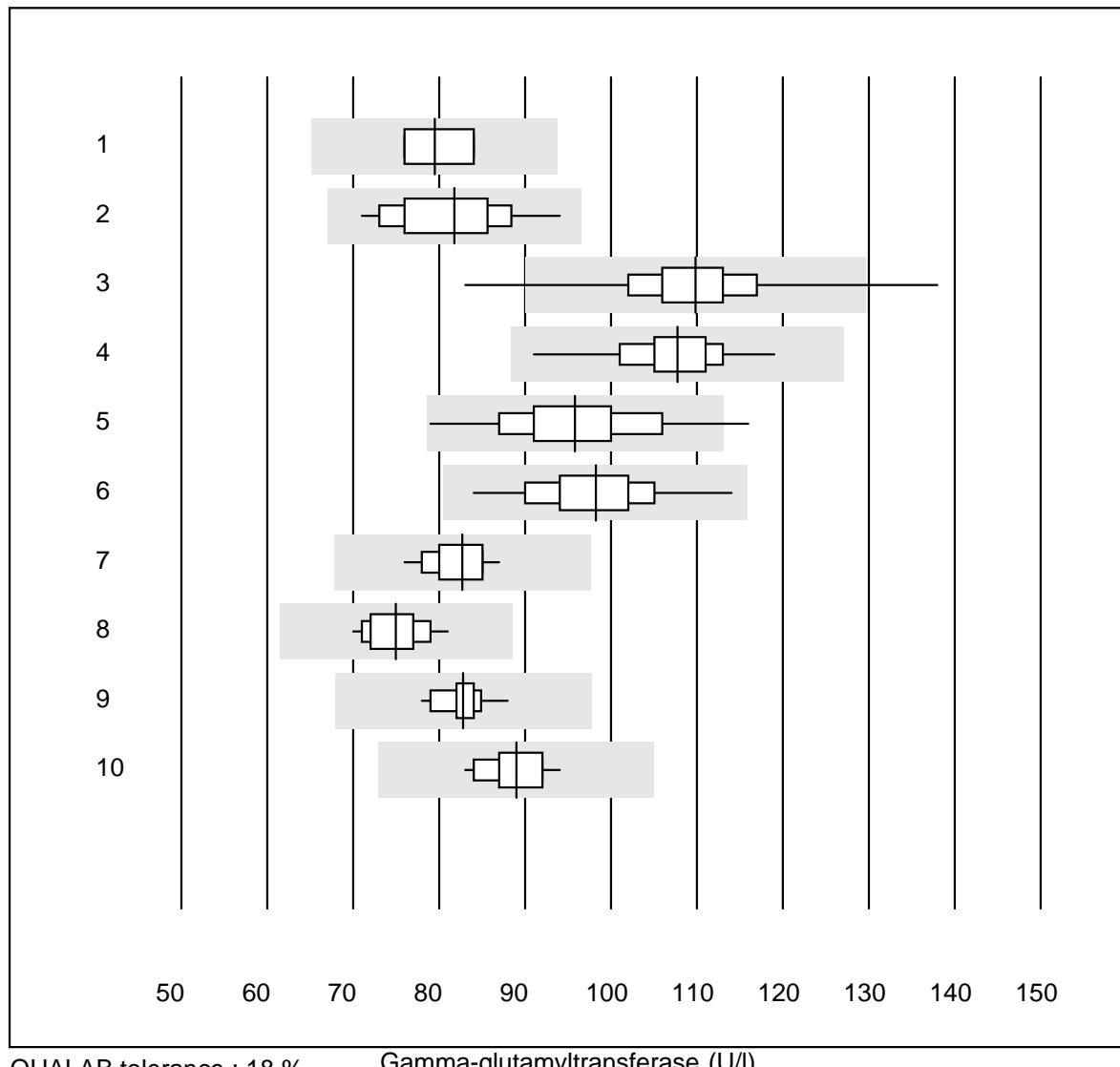
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Wet chemistry, direc	17	88.2	5.9	5.9	1.18	10.5	e*
2	Cobas	14	92.9	0.0	7.1	0.87	5.2	e
3	Reflotron	594	88.7	6.4	4.9	0.99	11.0	e
4	Fuji Dri-Chem	639	98.9	0.2	0.9	1.23	4.1	e
5	Spotchem/Ready	131	93.1	3.8	3.1	0.84	10.4	e
6	Spotchem D-Concept	129	98.4	1.6	0.0	0.90	8.8	e
7	Piccolo	22	100.0	0.0	0.0	1.06	4.2	e
8	Cholestech LDX	192	97.9	0.5	1.6	1.16	7.5	e
9	Abx Mira	18	100.0	0.0	0.0	1.20	6.2	e
10	Hitachi S40/M40	11	100.0	0.0	0.0	1.28	6.6	e

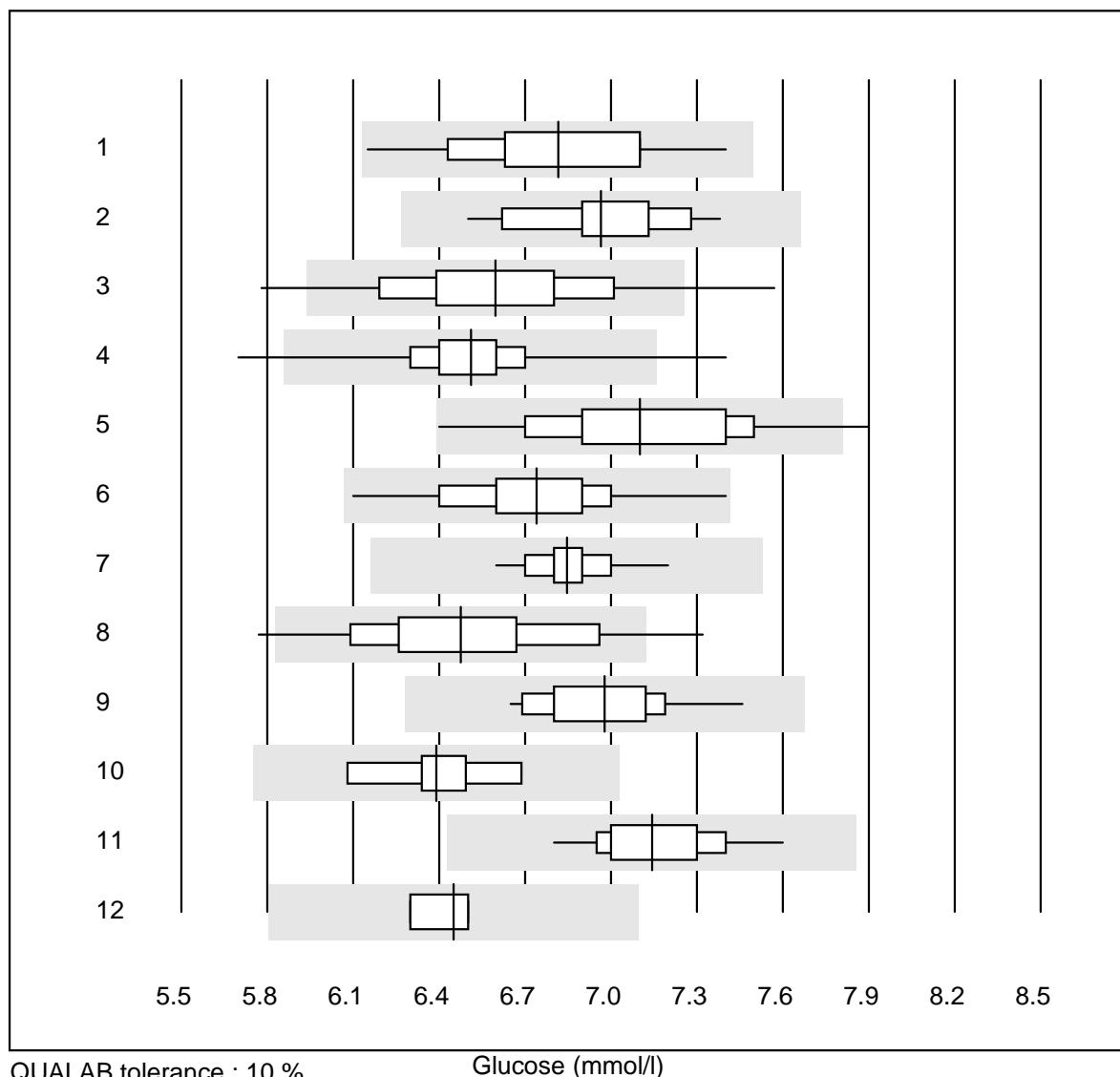
Creatine kinase



Iron

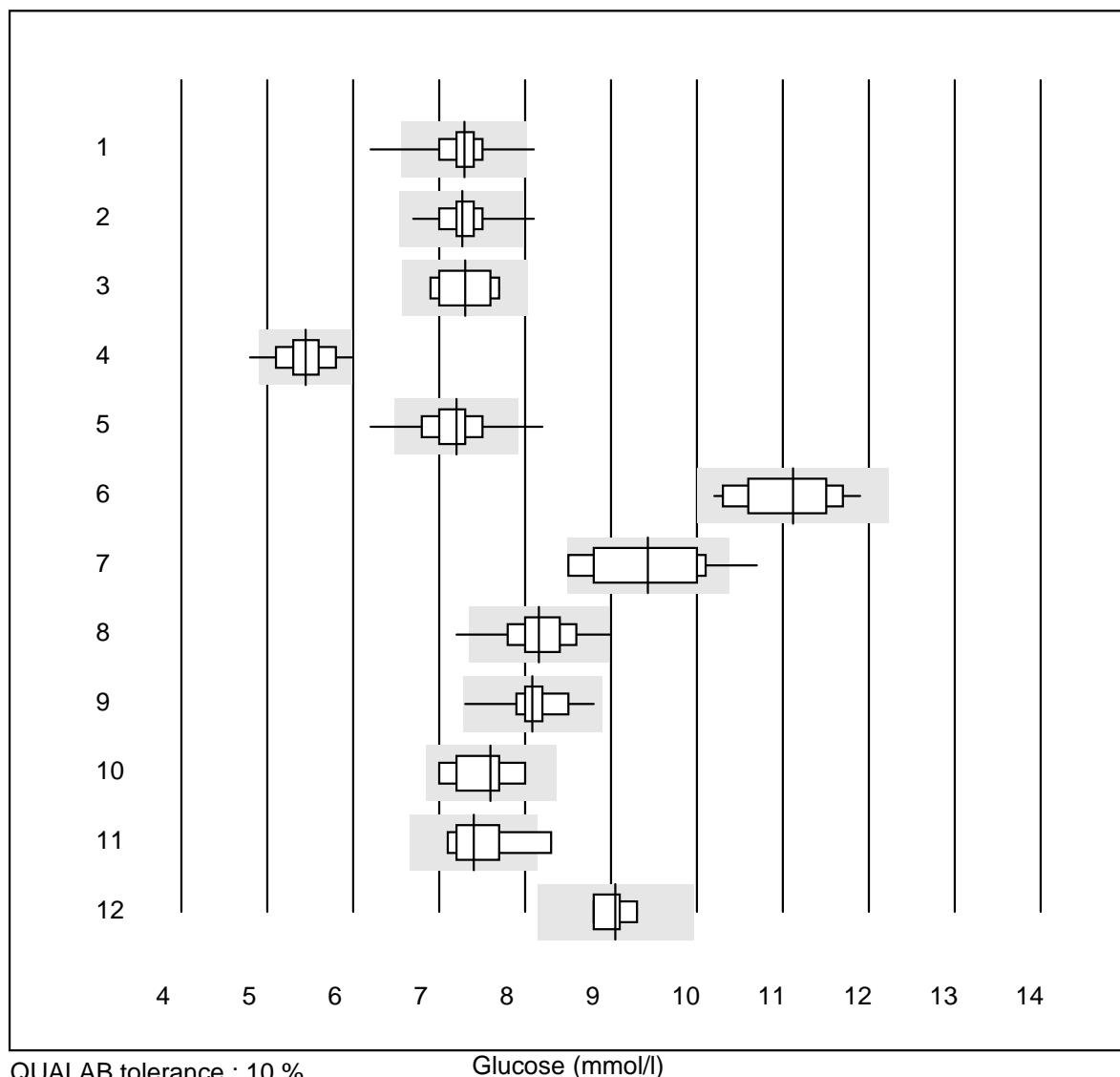
Gamma-glutamyltransferase



Glucose

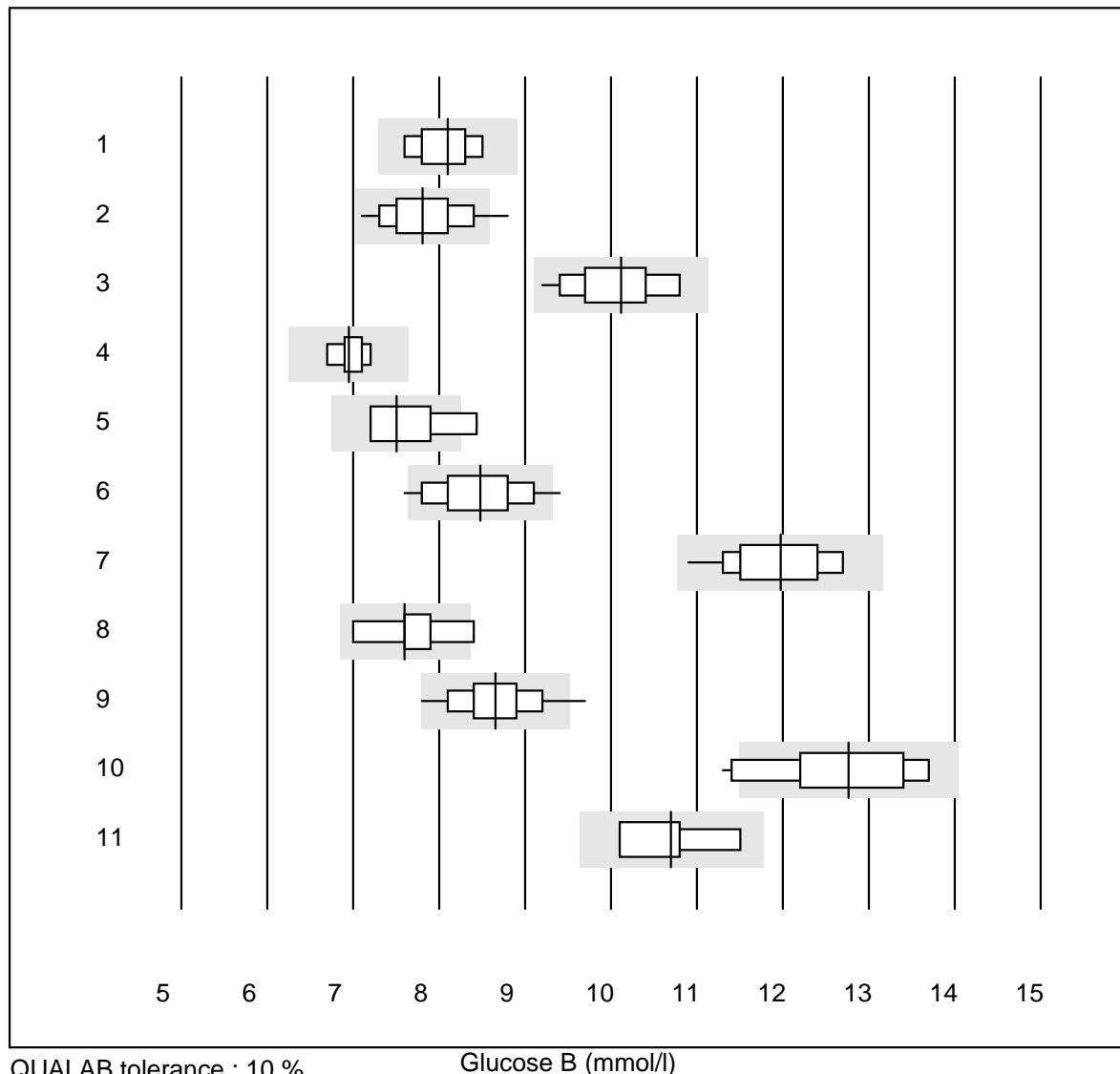
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Standard chemistry	25	100.0	0.0	0.0	6.8	4.4	e
2	Cobas	17	100.0	0.0	0.0	7.0	3.2	e
3	Reflotron	908	94.3	3.7	2.0	6.6	4.8	e
4	Fuji Dri-Chem	677	99.5	0.4	0.1	6.5	2.3	e
5	Spotchem/Ready	134	98.5	1.5	0.0	7.1	4.4	e
6	Spotchem D-Concept	132	100.0	0.0	0.0	6.7	3.9	e
7	Piccolo	33	93.9	0.0	6.1	6.8	1.9	e
8	Cholestech LDX	154	94.2	5.2	0.6	6.5	5.0	e
9	Abx Mira	20	100.0	0.0	0.0	7.0	3.0	e
10	Lange	6	83.3	0.0	16.7	6.4	3.5	e*
11	Hitachi S40/M40	15	100.0	0.0	0.0	7.1	2.8	e
12	iStat Chem8	4	100.0	0.0	0.0	6.5	1.5	e

Glucose

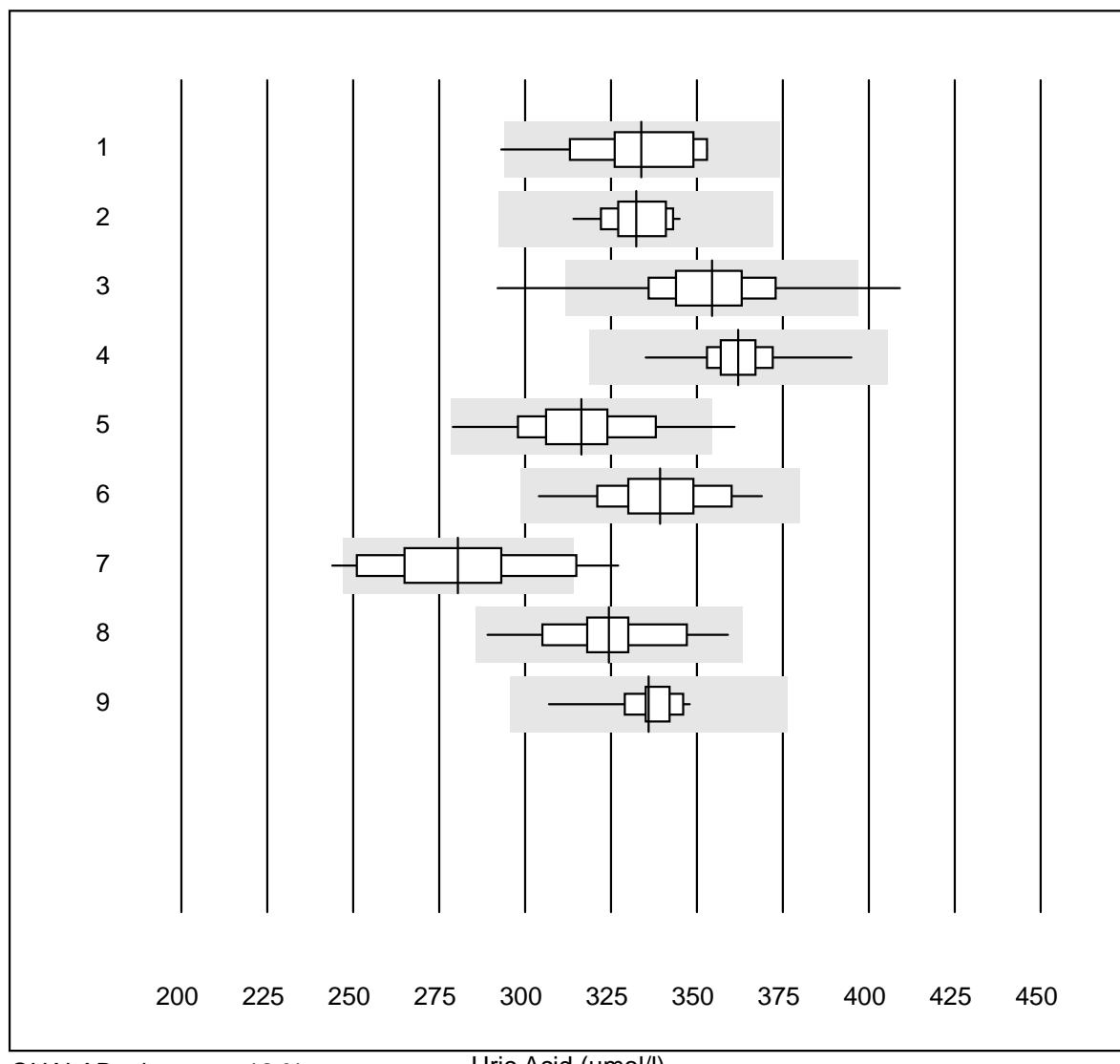


No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Accu-Chek Aviva	371	98.7	0.8	0.5	7.3	3.0	e
2 Accu-Chek Inform 2	232	99.6	0.4	0.0	7.3	2.7	e
3 Accu-Chek Mobile	5	100.0	0.0	0.0	7.3	4.8	e*
4 Bayer Contour 2 (5s)	61	90.1	6.6	3.3	5.4	4.8	e
5 Bayer Contour XT/NEX	1212	98.3	0.9	0.8	7.2	3.8	e
6 Bayer Breeze 2	17	88.2	0.0	11.8	11.1	4.6	e
7 Glucocard	11	81.8	9.1	9.1	9.4	7.4	e*
8 Hemocue 201+ P-equiv	83	91.6	2.4	6.0	8.2	4.1	e
9 Hemocue 201RT P-equiv	33	100.0	0.0	0.0	8.1	3.6	e
10 FreeStyle Precision	5	100.0	0.0	0.0	7.6	5.3	e*
11 Freestyle Freedom li	9	88.9	11.1	0.0	7.4	5.4	e*
12 Sanofi BG Star	6	66.7	0.0	33.3	9.1	2.3	e

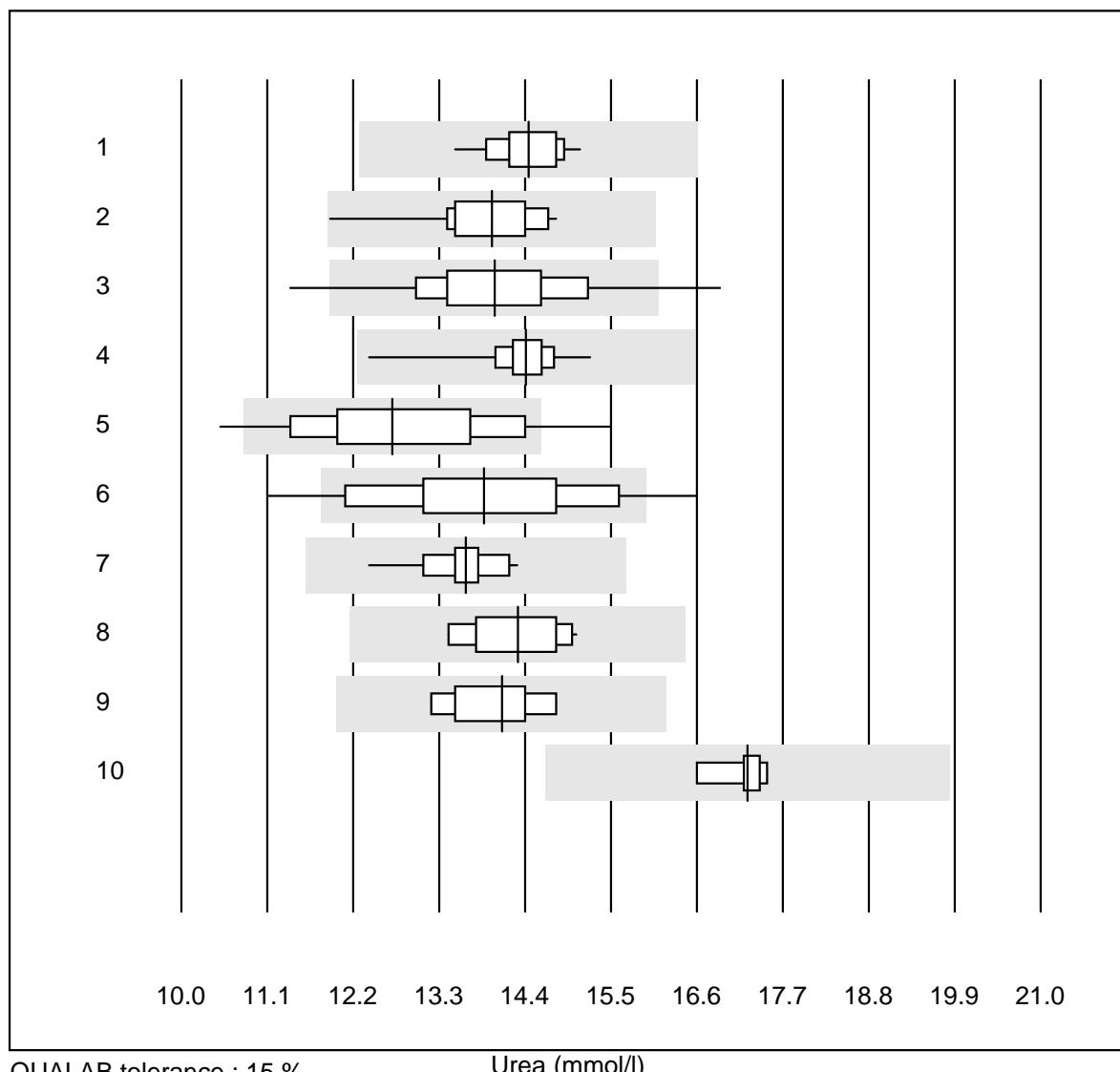
Glucose B



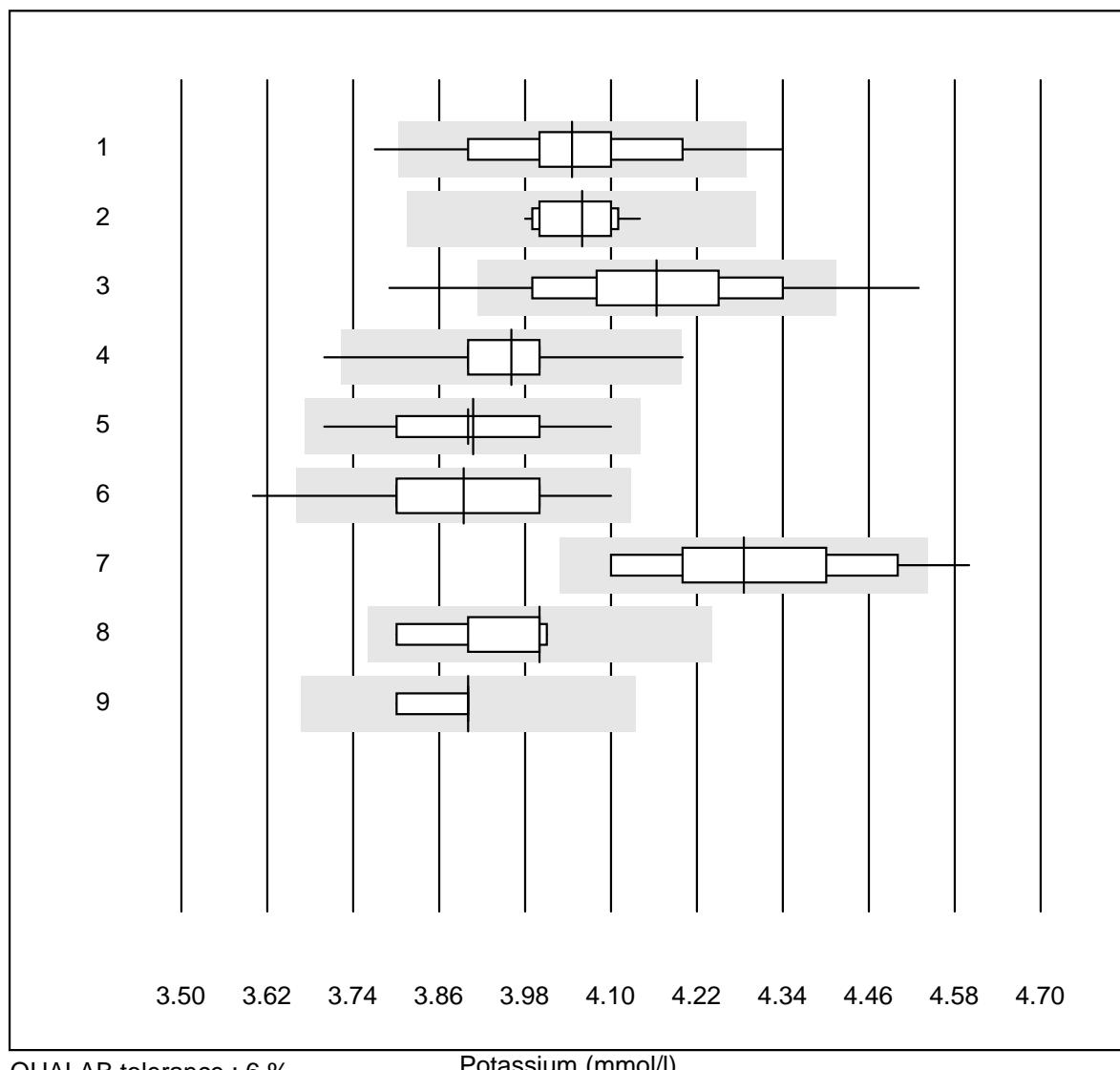
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Bayer Elite	9	88.9	0.0	11.1	8.1	3.8	e*
2 Hemocue 201+ (alt)	57	91.2	5.3	3.5	7.8	5.4	e
3 OneTouch Ultra	28	100.0	0.0	0.0	10.1	4.5	e
4 OneTouch Verio	6	100.0	0.0	0.0	7.0	2.5	e
5 AccuChek Compact	4	75.0	25.0	0.0	7.5	6.9	e*
6 Bayer Contour (15s)	46	87.0	6.5	6.5	8.5	5.4	e
7 Healthpro	14	100.0	0.0	0.0	12.0	4.5	e
8 Mylife UNIO	5	80.0	20.0	0.0	7.6	6.6	e*
9 mylife Pura	57	96.5	3.5	0.0	8.7	5.0	e
10 Omnitest	16	81.2	12.5	6.3	12.8	5.9	e*
11 Alpha Check	4	100.0	0.0	0.0	10.7	5.4	e*

Uric Acid

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Standard chemistry	19	89.4	5.3	5.3	334	4.7	e
2 Cobas	12	100.0	0.0	0.0	332	2.8	e
3 Reflotron	788	98.1	0.8	1.1	354	4.2	e
4 Fuji Dri-Chem	676	99.7	0.0	0.3	362	2.1	e
5 Spotchem/Ready	121	99.2	0.8	0.0	316	4.9	e
6 Spotchem D-Concept	127	100.0	0.0	0.0	339	4.2	e
7 Piccolo	23	78.3	17.4	4.3	280	8.5	e*
8 Abx Mira	18	100.0	0.0	0.0	324	5.0	e
9 Hitachi S40/M40	13	100.0	0.0	0.0	336	3.0	e

Urea

Potassium

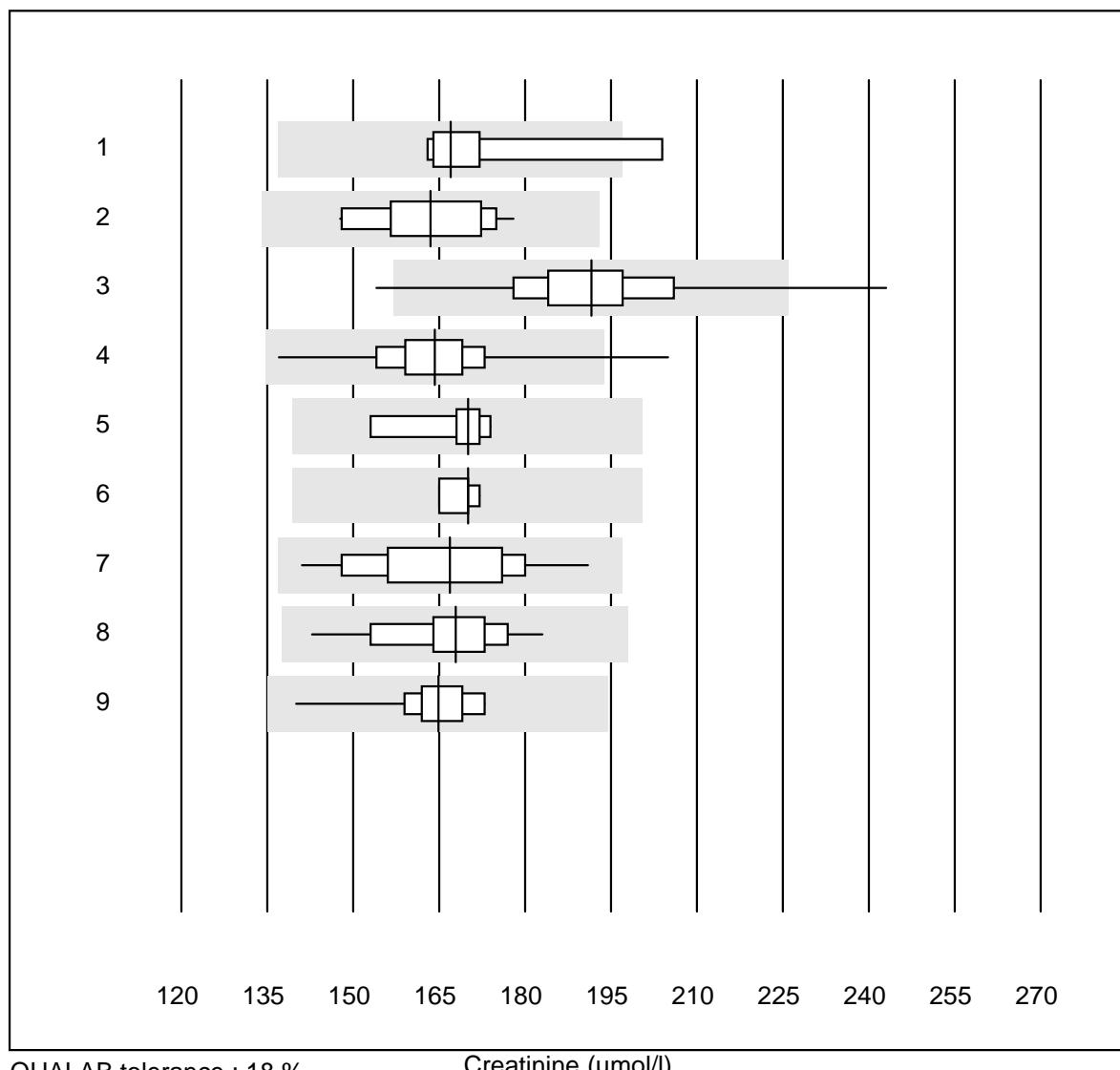


QUALAB tolerance : 6 %

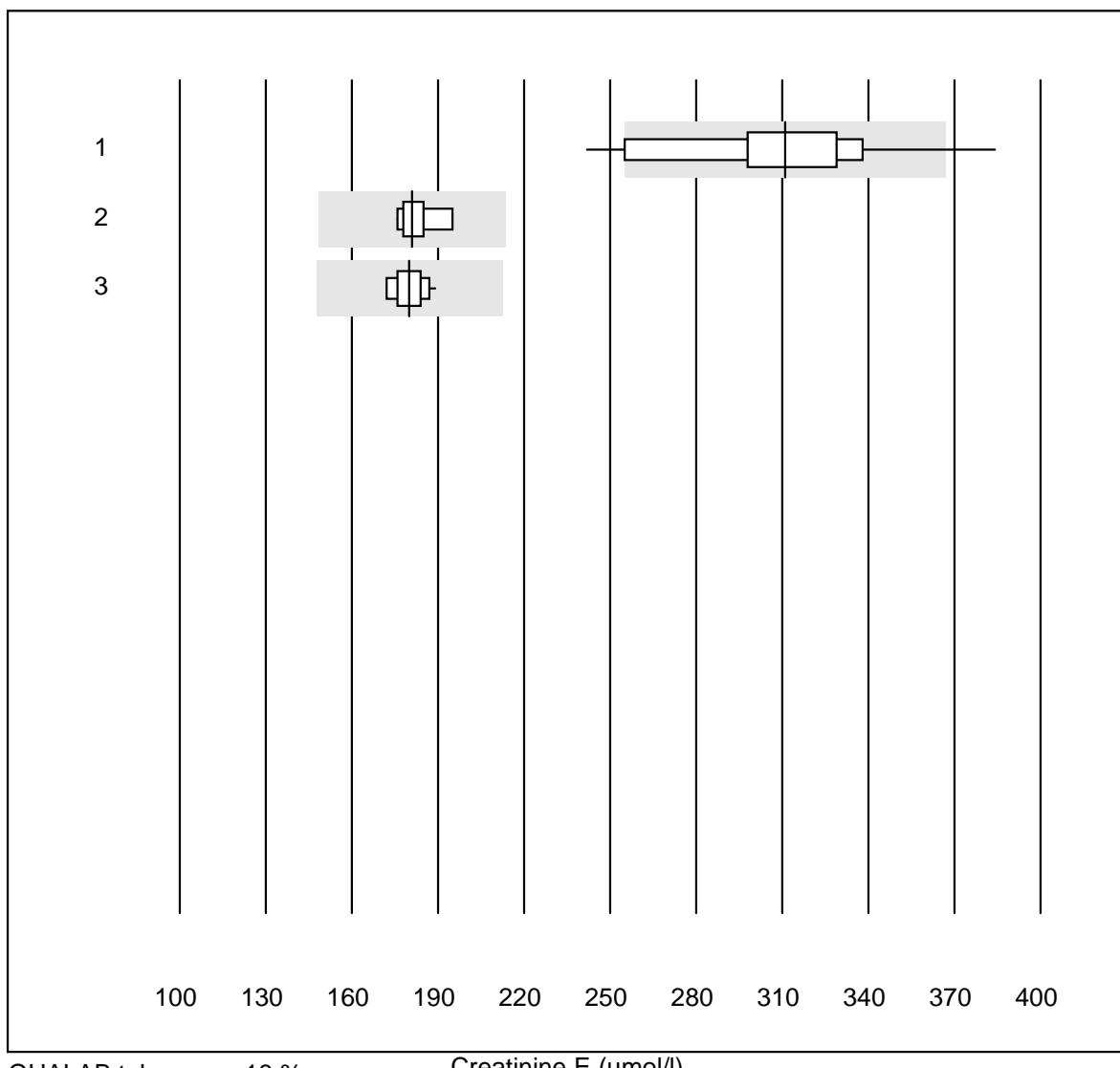
Potassium (mmol/l)

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 ISE	27	92.6	7.4	0.0	4.05	2.8	e
2 Cobas	17	100.0	0.0	0.0	4.06	1.3	e
3 Reflotron	816	89.0	7.1	3.9	4.16	3.3	e
4 Fuji Dri-Chem	709	97.2	1.4	1.4	3.96	1.9	e
5 Spotchem D-Concept	128	100.0	0.0	0.0	3.91	1.7	e
6 Spotchem EL-SE 1520	117	93.1	2.6	4.3	3.89	2.4	e
7 Piccolo	18	72.2	5.6	22.2	4.29	3.8	e*
8 Abx Mira	5	100.0	0.0	0.0	4.00	2.3	e*
9 iStat Chem8	6	100.0	0.0	0.0	3.90	1.1	e

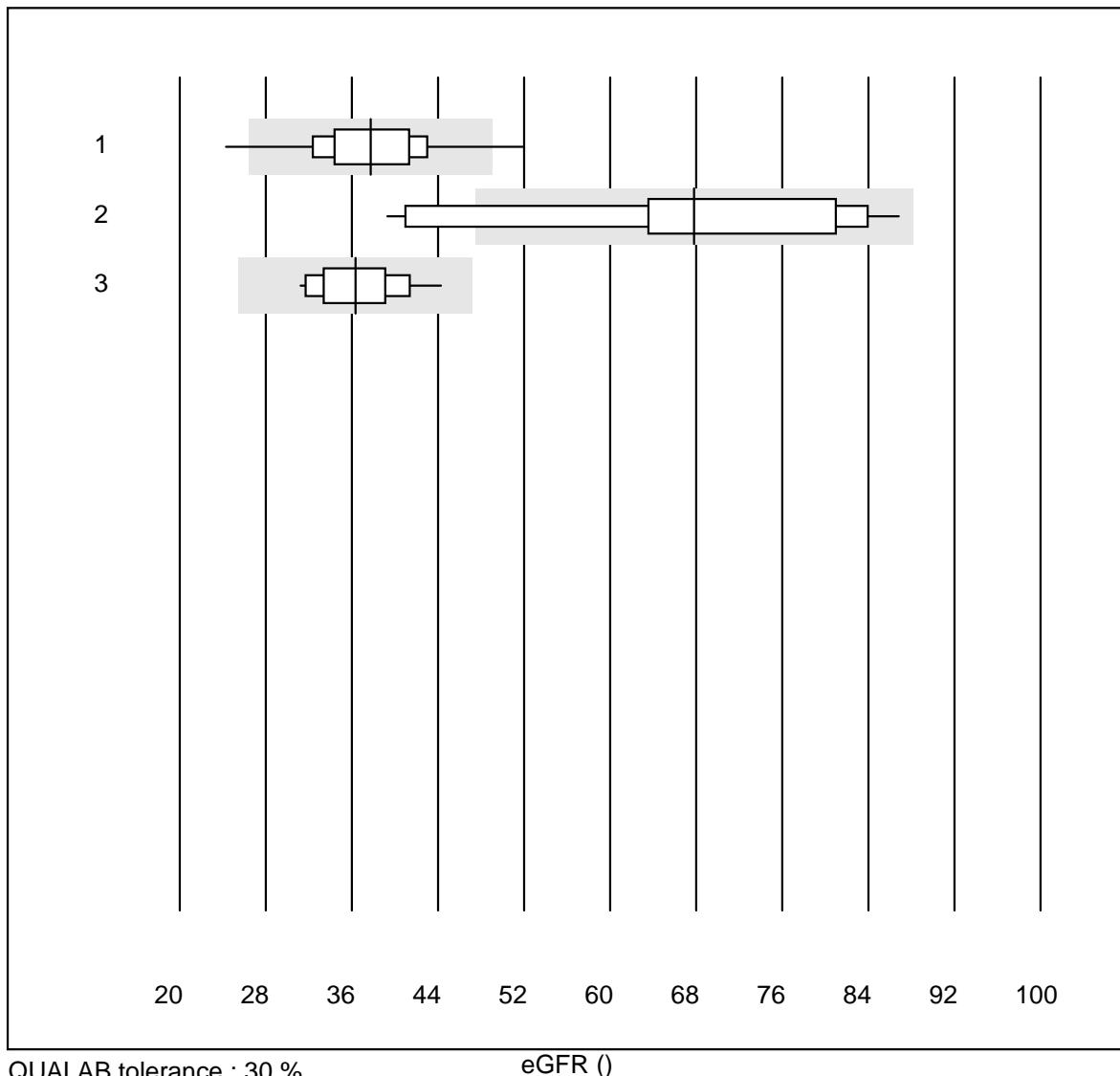
Creatinine

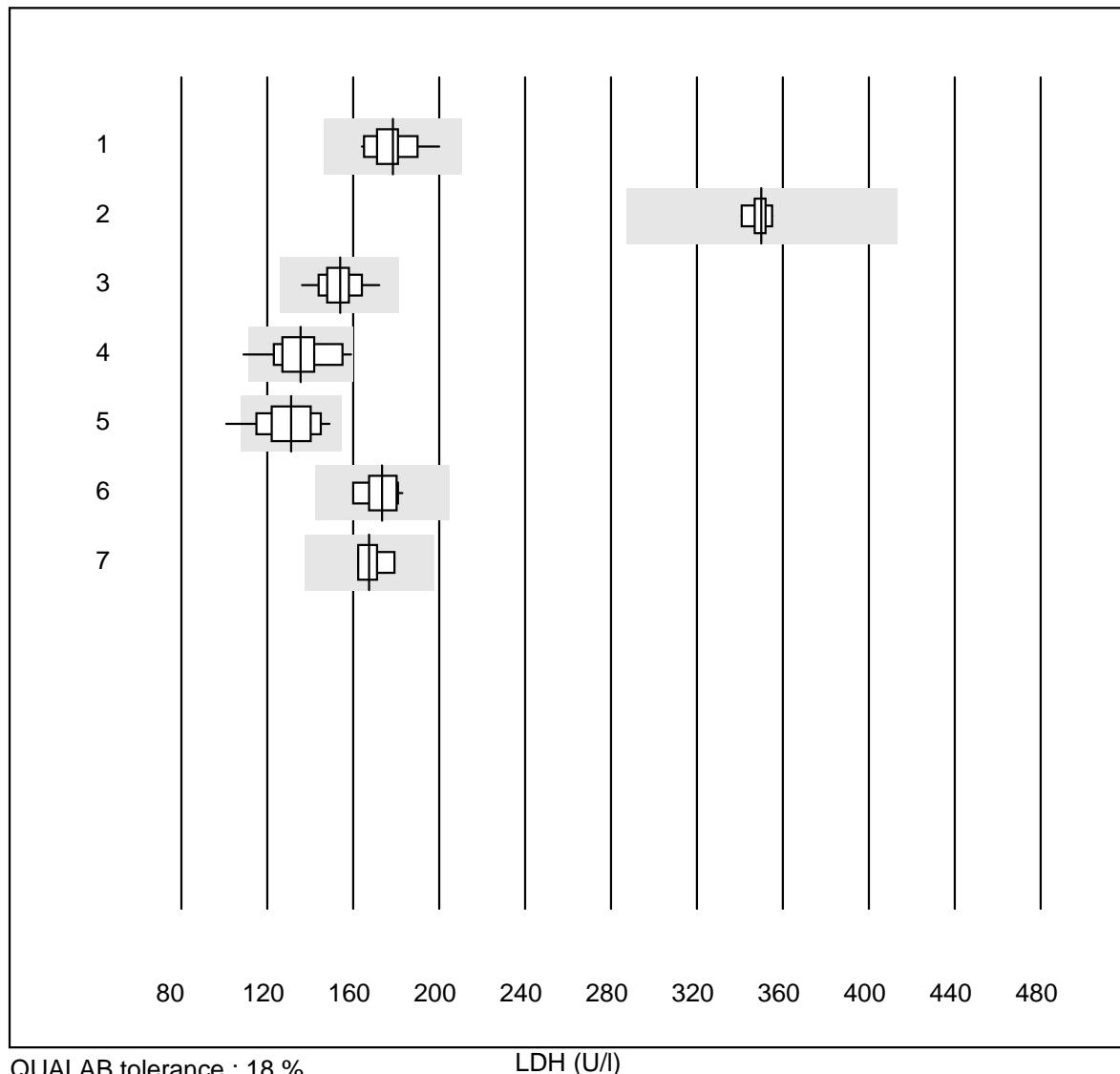


No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Standard chemistry	8	87.5	12.5	0.0	167	7.9	e*
2	Cobas	18	100.0	0.0	0.0	164	5.8	e
3	Reflotron	995	97.6	1.6	0.8	192	6.2	e
4	Fuji Dri-Chem	743	98.8	0.5	0.7	164	5.1	e
5	Jaffé	9	100.0	0.0	0.0	170	3.6	e
6	Enzymatic	4	100.0	0.0	0.0	170	1.8	e
7	Piccolo	31	96.8	0.0	3.2	167	7.8	e
8	Abx Mira	20	100.0	0.0	0.0	168	5.3	e
9	Hitachi S40/M40	14	100.0	0.0	0.0	165	5.1	e

Creatinine E

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Statsensor i / Nova	26	84.6	15.4	0.0	311	9.8	e
2 iStat Chem8	8	100.0	0.0	0.0	181	3.2	e
3 ABL700/800 Radiomete	10	100.0	0.0	0.0	180	3.1	e

eGFR

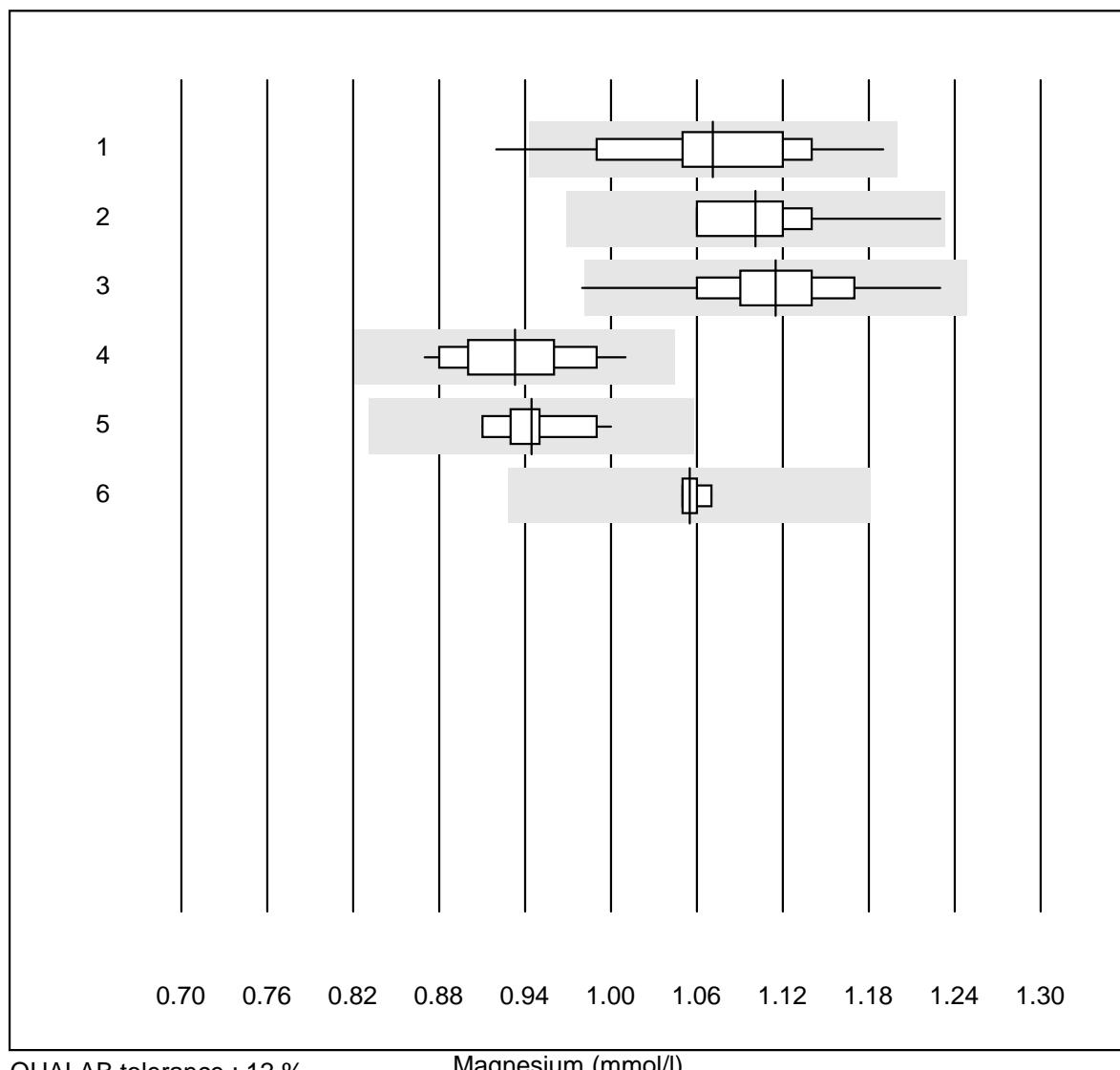
LDH

QUALAB tolerance : 18 %

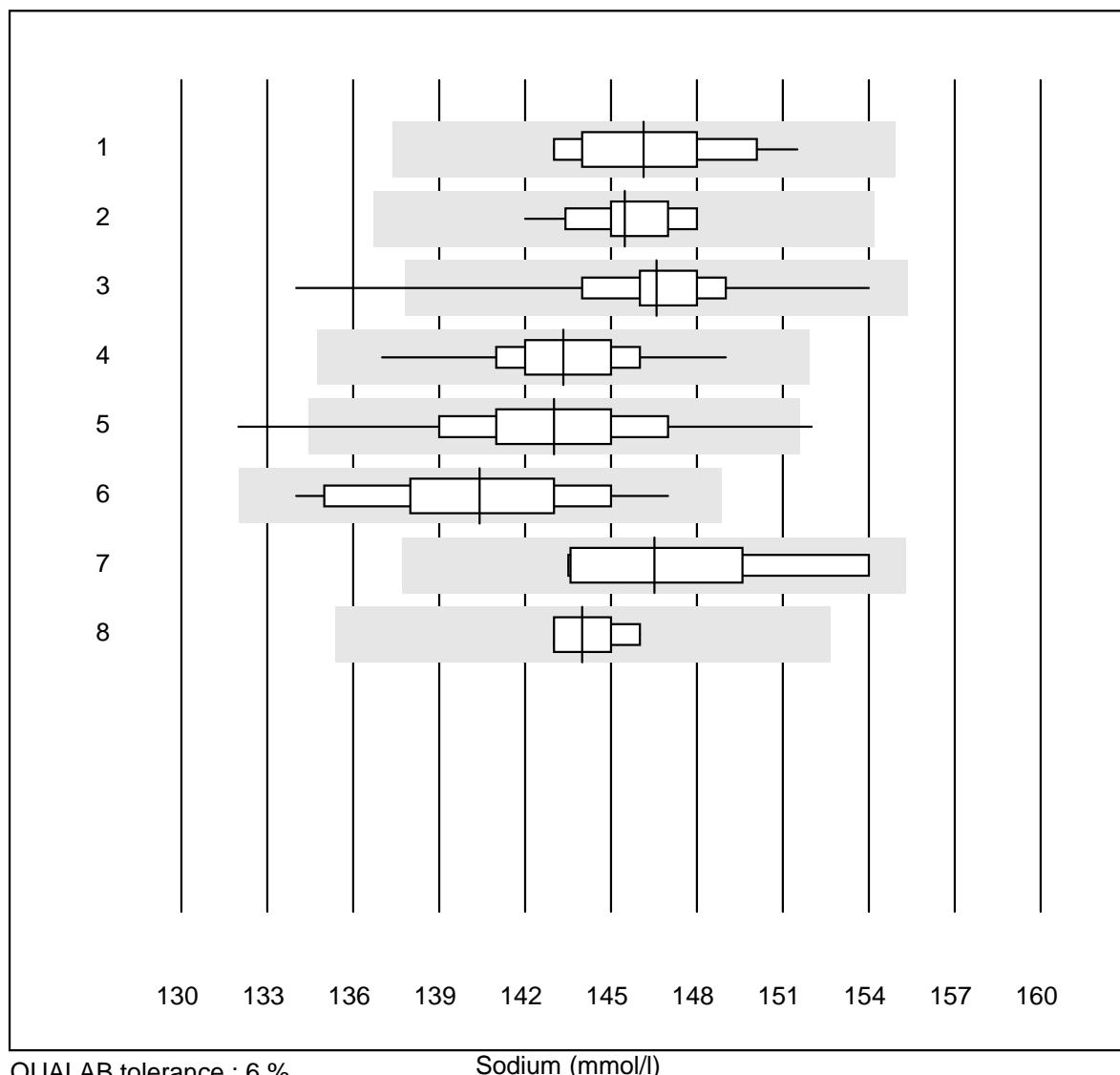
LDH (U/l)

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 IFCC	20	100.0	0.0	0.0	178	5.4	e
2 Cobas	9	100.0	0.0	0.0	350	1.2	e
3 Fuji Dri-Chem	140	99.3	0.0	0.7	154	4.7	e
4 Spotchem/Ready	38	92.1	2.6	5.3	136	8.6	e
5 Spotchem D-Concept	35	94.2	2.9	2.9	131	9.1	e
6 Abx Mira	11	90.9	0.0	9.1	173	4.4	e
7 Hitachi S40/M40	4	100.0	0.0	0.0	168	4.5	e*

Magnesium

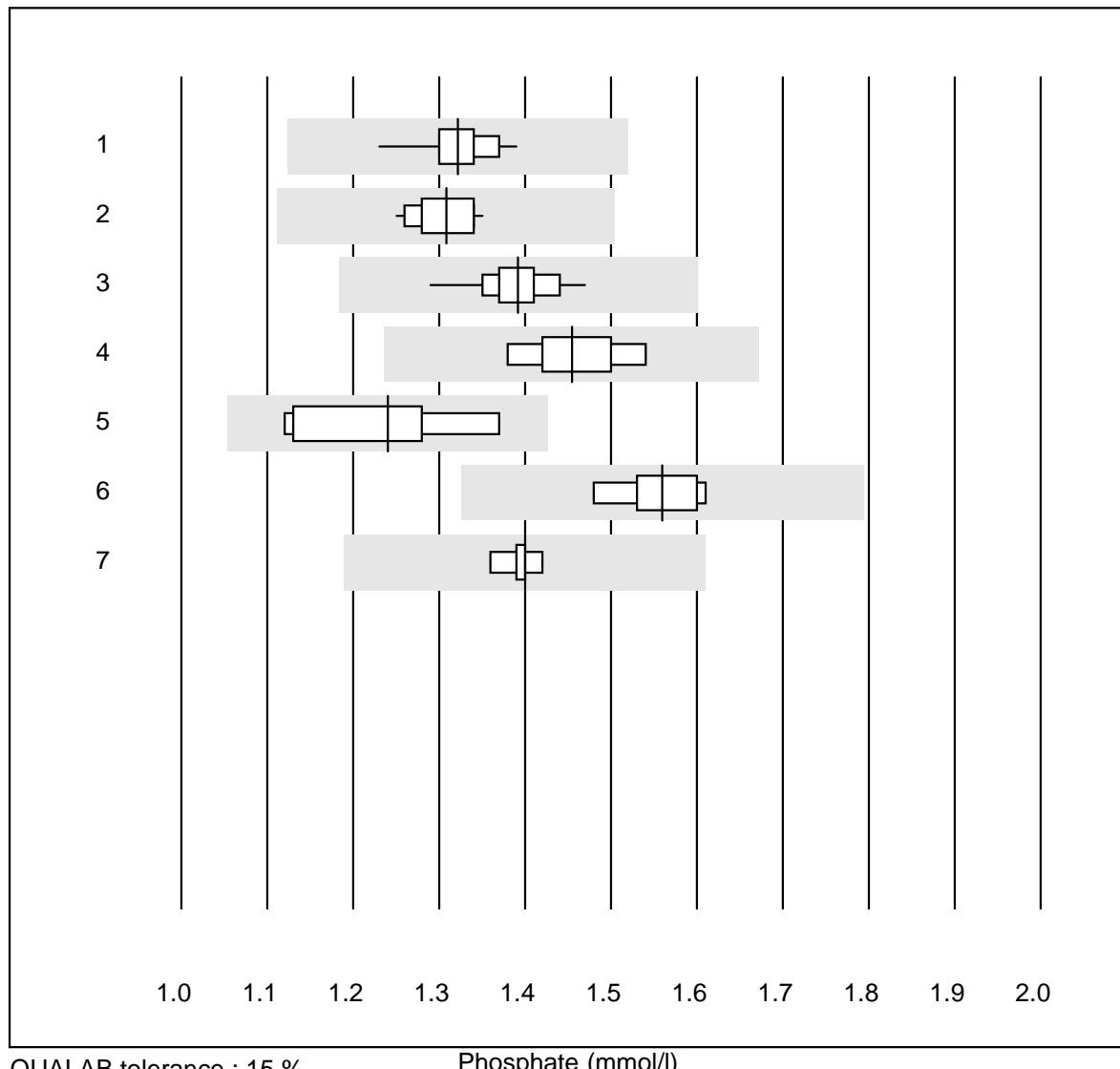


Sodium



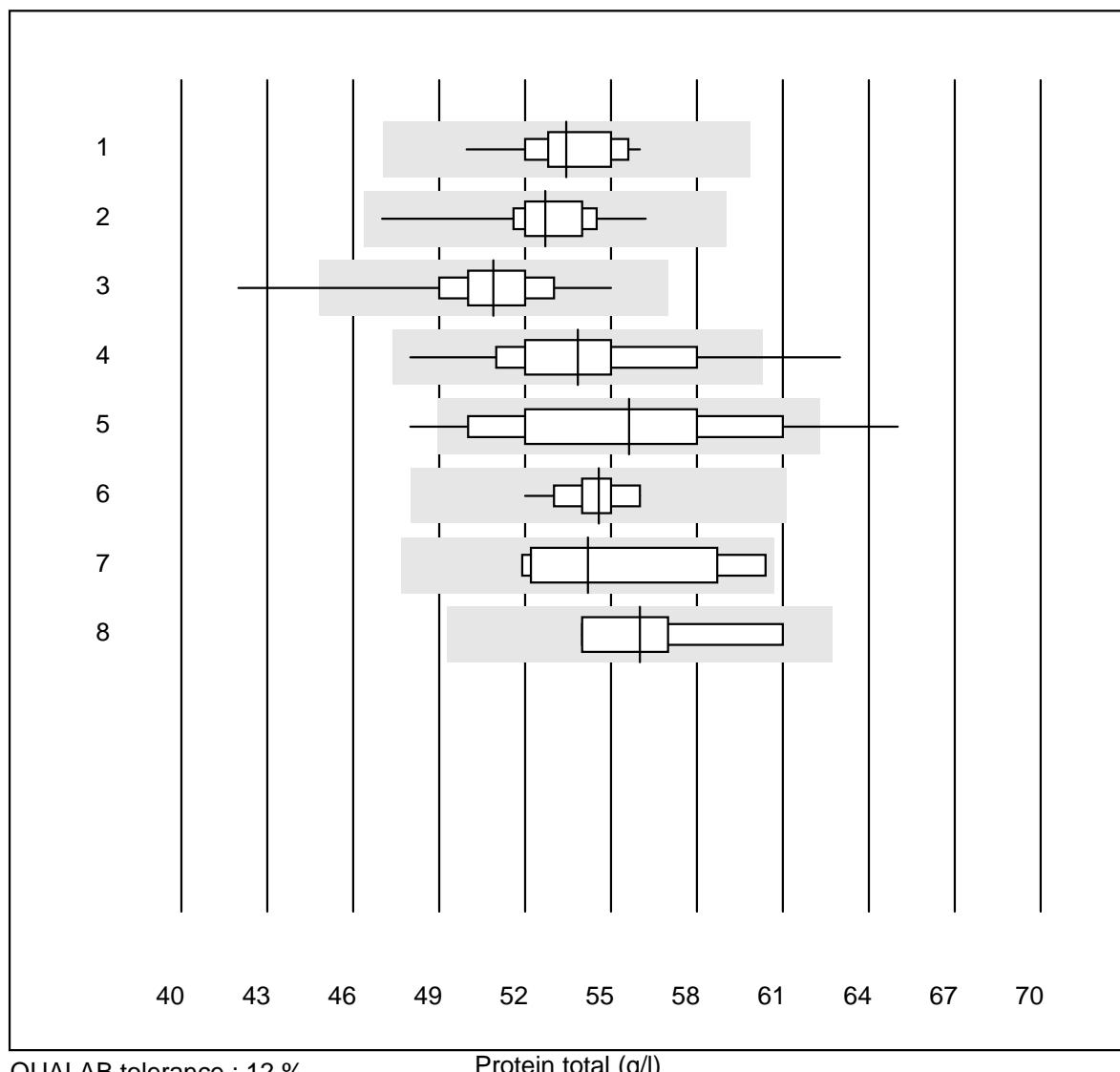
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 ISE	26	100.0	0.0	0.0	146	1.7	e
2 Cobas	16	100.0	0.0	0.0	145	1.1	e
3 Fuji Dri-Chem	655	98.6	0.8	0.6	147	1.5	e
4 Spotchem D-Concept	123	100.0	0.0	0.0	143	1.3	e
5 Spotchem EL-SE 1520	117	95.7	2.6	1.7	143	2.2	e
6 Piccolo	19	100.0	0.0	0.0	140	2.5	e
7 Abx Mira	6	100.0	0.0	0.0	147	2.8	e*
8 iStat Chem8	6	100.0	0.0	0.0	144	0.8	e

Phosphate

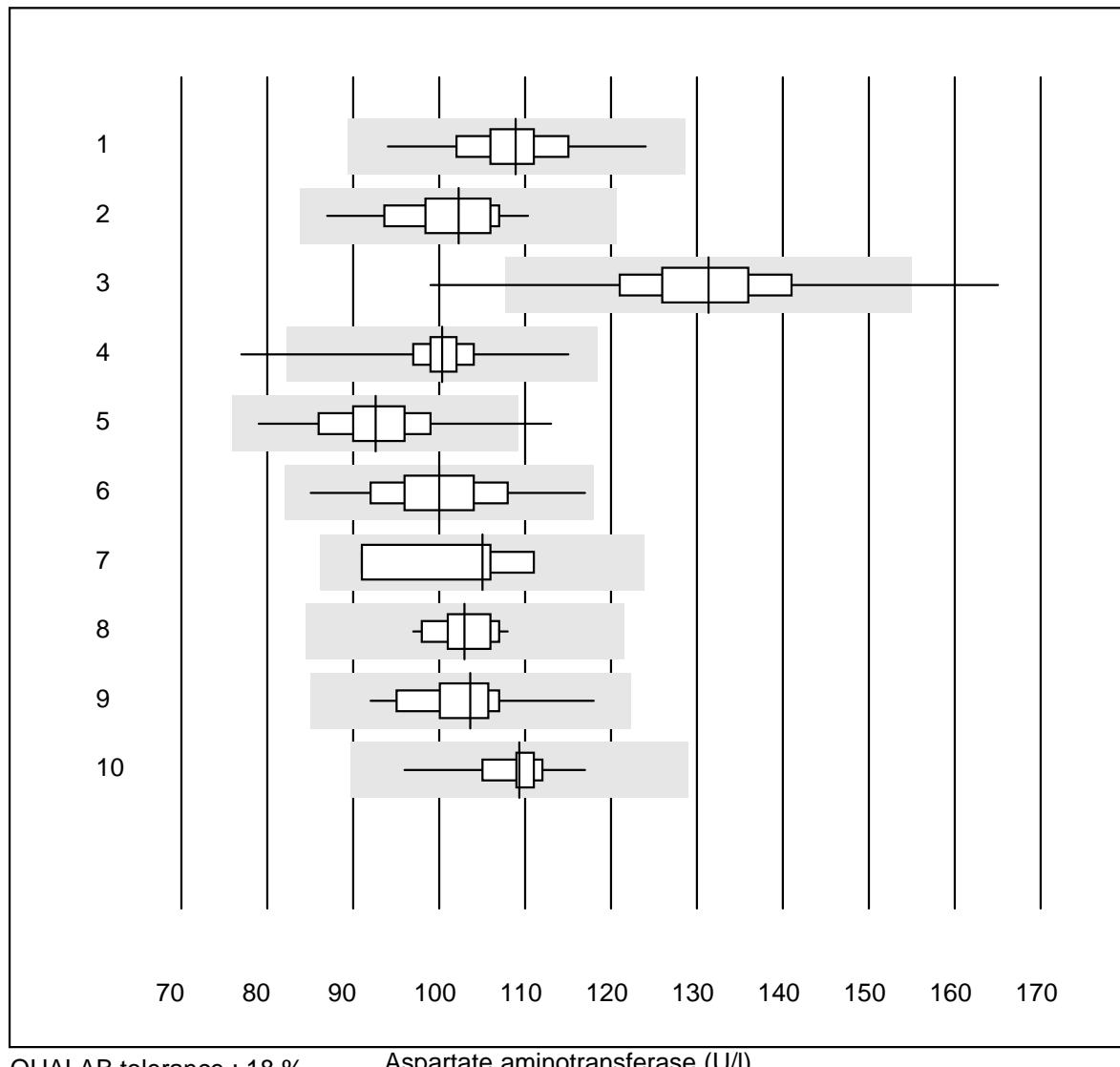


No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Standard chemistry	12	100.0	0.0	0.0	1.3	3.1	e
2	Cobas	11	100.0	0.0	0.0	1.3	2.5	e
3	Fuji Dri-Chem	73	100.0	0.0	0.0	1.4	2.6	e
4	Spotchem D-Concept	15	100.0	0.0	0.0	1.5	3.7	e
5	Spotchem/Ready	7	100.0	0.0	0.0	1.2	7.2	e*
6	Piccolo	5	100.0	0.0	0.0	1.6	3.4	e
7	Abx Mira	5	100.0	0.0	0.0	1.4	1.6	e

Protein total

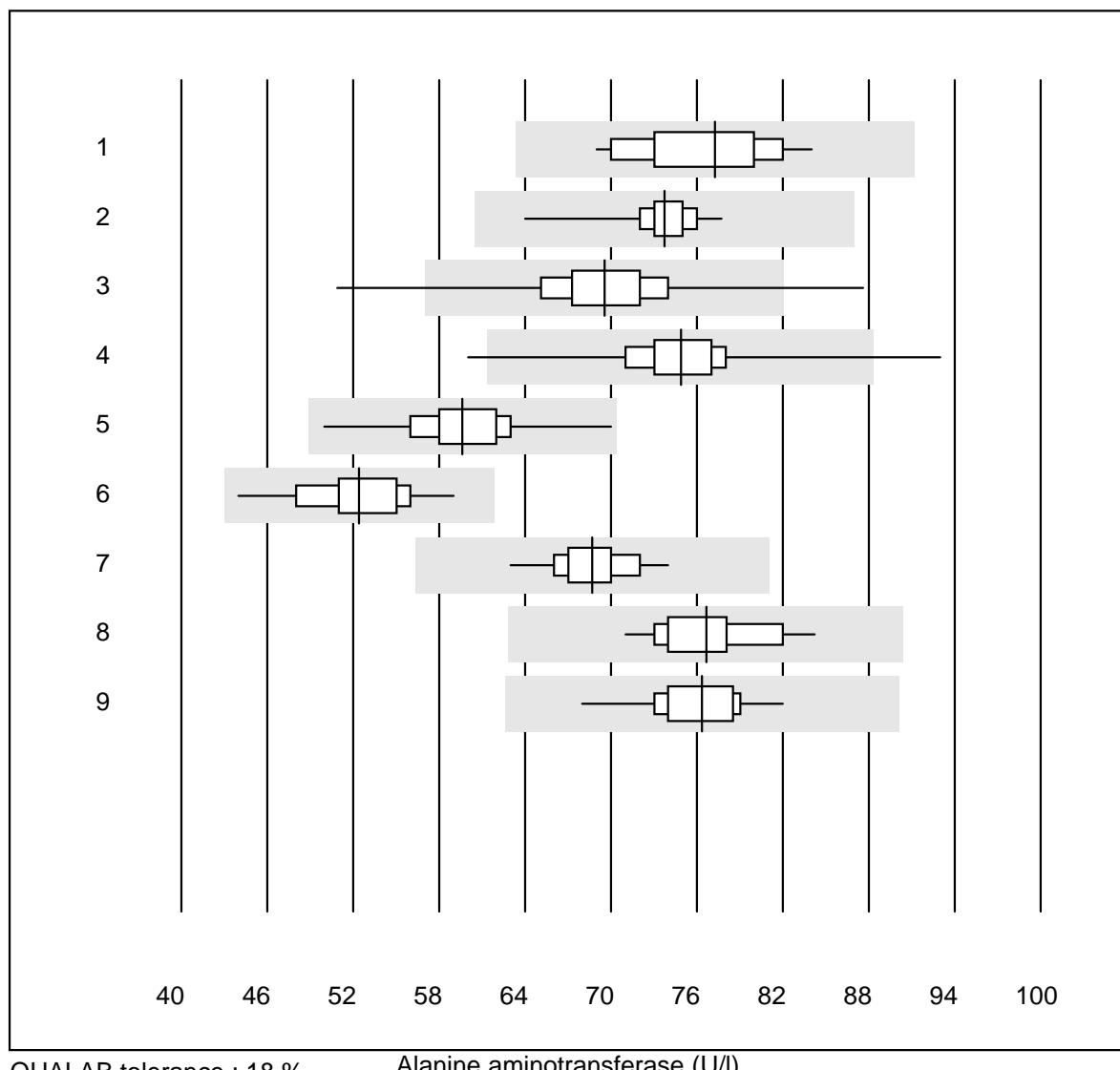


Aspartate aminotransferase



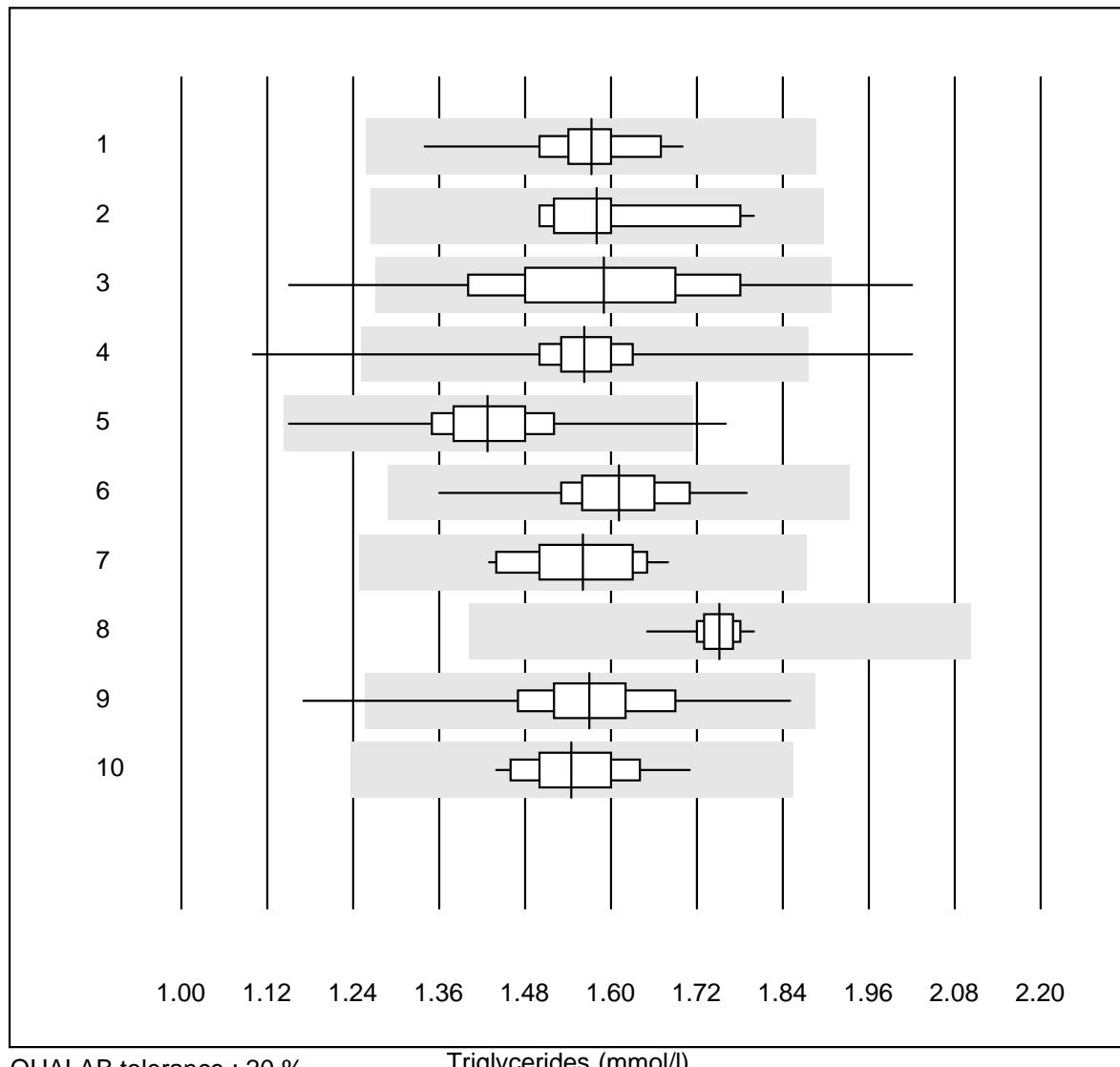
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 IFCC with Pyridox 37	16	100.0	0.0	0.0	109	5.8	e
2 Cobas	17	100.0	0.0	0.0	102	5.7	e
3 Reflotron	900	98.3	1.1	0.6	131	6.2	e
4 Fuji Dri-Chem	716	98.8	0.6	0.6	100	3.4	e
5 Spotchem/Ready	156	98.8	0.6	0.6	93	5.8	e
6 Spotchem D-Concept	136	99.3	0.0	0.7	100	6.0	e
7 IFCC with Pyridox 37	4	100.0	0.0	0.0	105	8.3	e*
8 Piccolo	30	96.7	0.0	3.3	103	3.0	e
9 Abx Mira	20	100.0	0.0	0.0	104	5.5	e
10 Hitachi S40/M40	16	93.7	0.0	6.3	109	4.1	e

Alanine aminotransferase

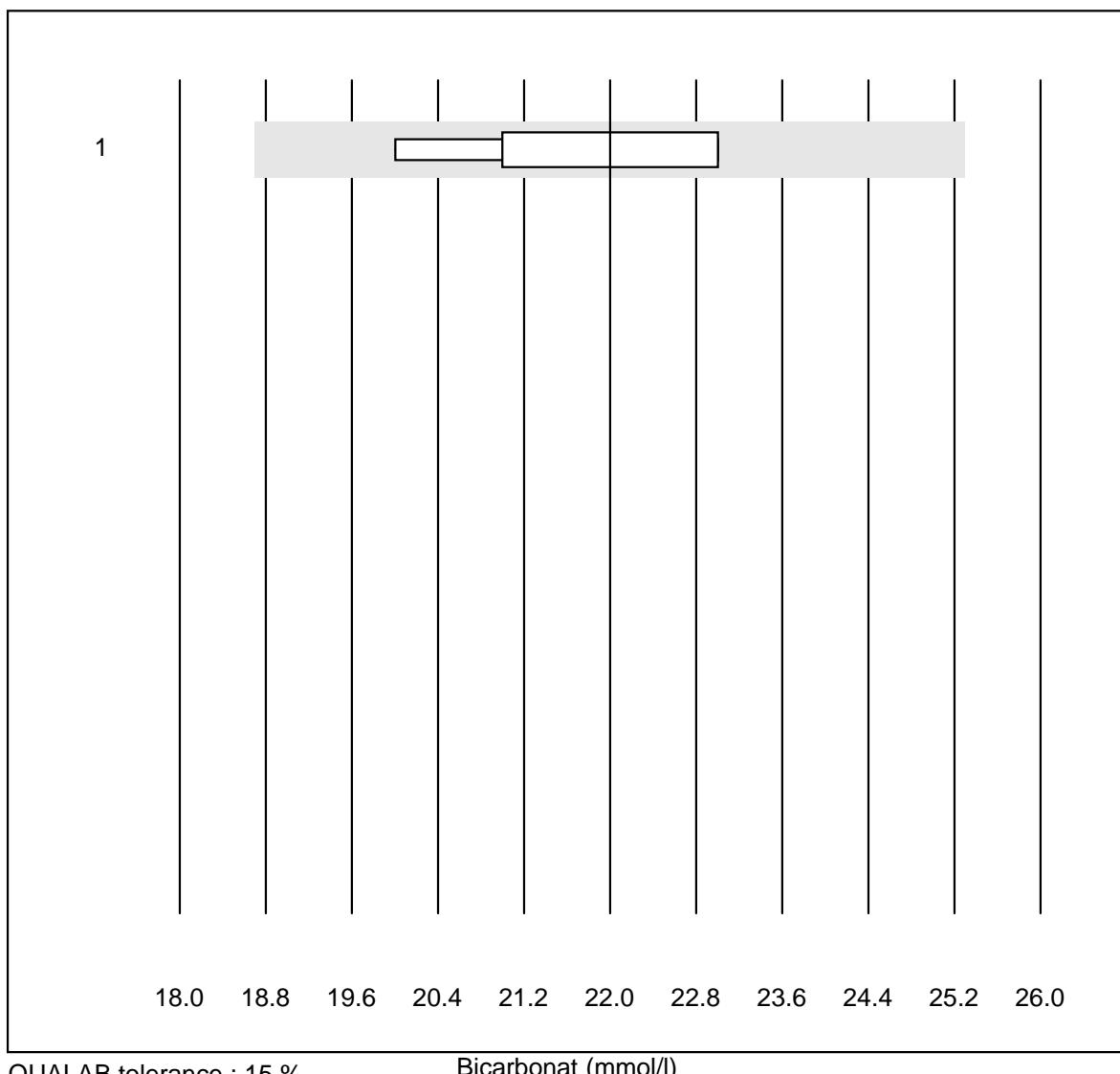


No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 IFCC with Pyridox 37	17	94.1	0.0	5.9	77	5.7	e
2 Cobas	18	100.0	0.0	0.0	74	3.8	e
3 Reflotron	935	97.8	1.0	1.2	70	5.5	e
4 Fuji Dri-Chem	730	98.9	0.3	0.8	75	4.0	e
5 Spotchem/Ready	159	98.7	0.0	1.3	60	5.7	e
6 Spotchem D-Concept	141	99.3	0.0	0.7	52	5.8	e
7 Piccolo	31	96.8	0.0	3.2	69	3.4	e
8 Abx Mira	20	100.0	0.0	0.0	77	4.6	e
9 Hitachi S40/M40	15	100.0	0.0	0.0	76	4.3	e

Triglycerides

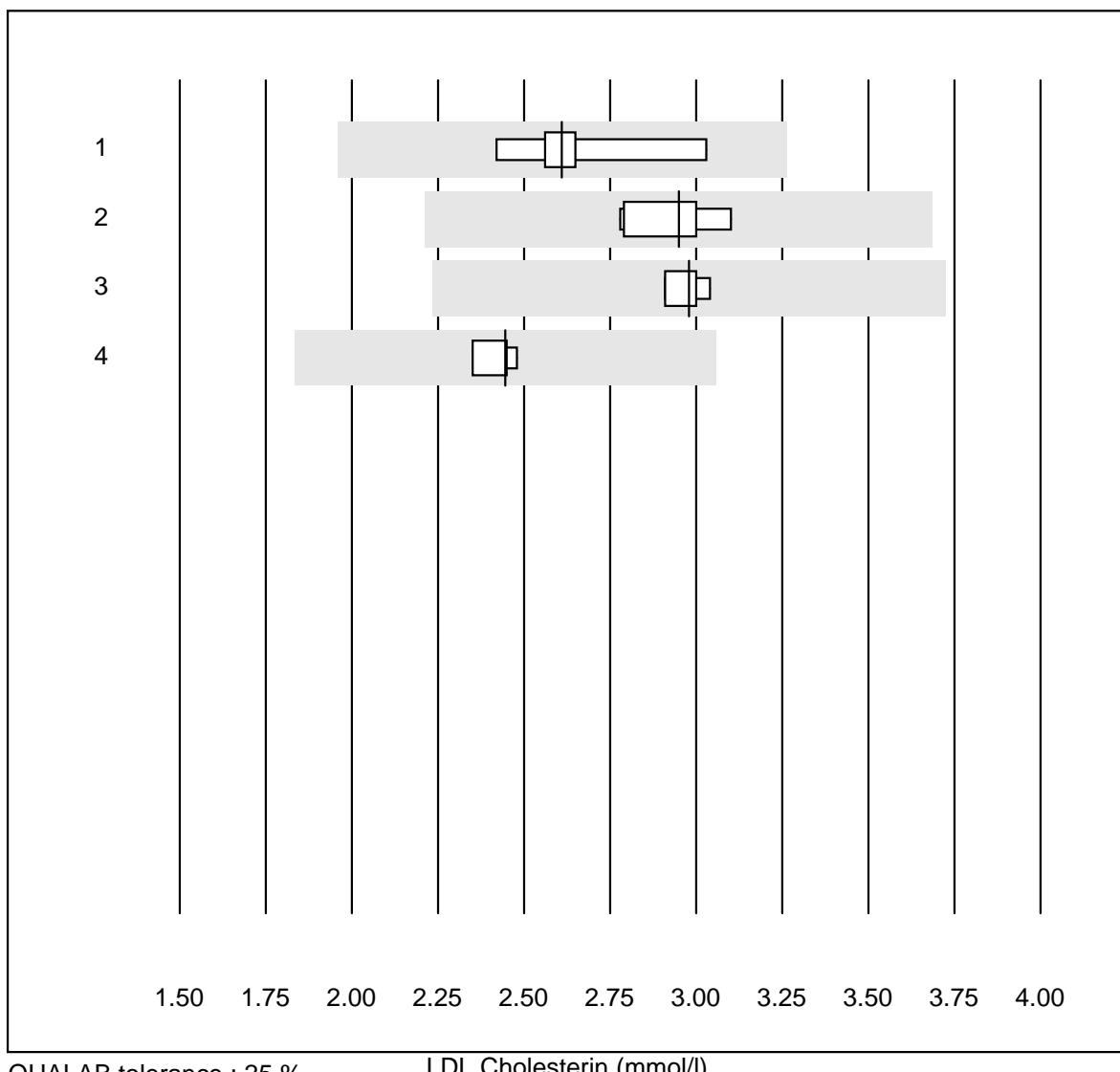


Bicarbonat



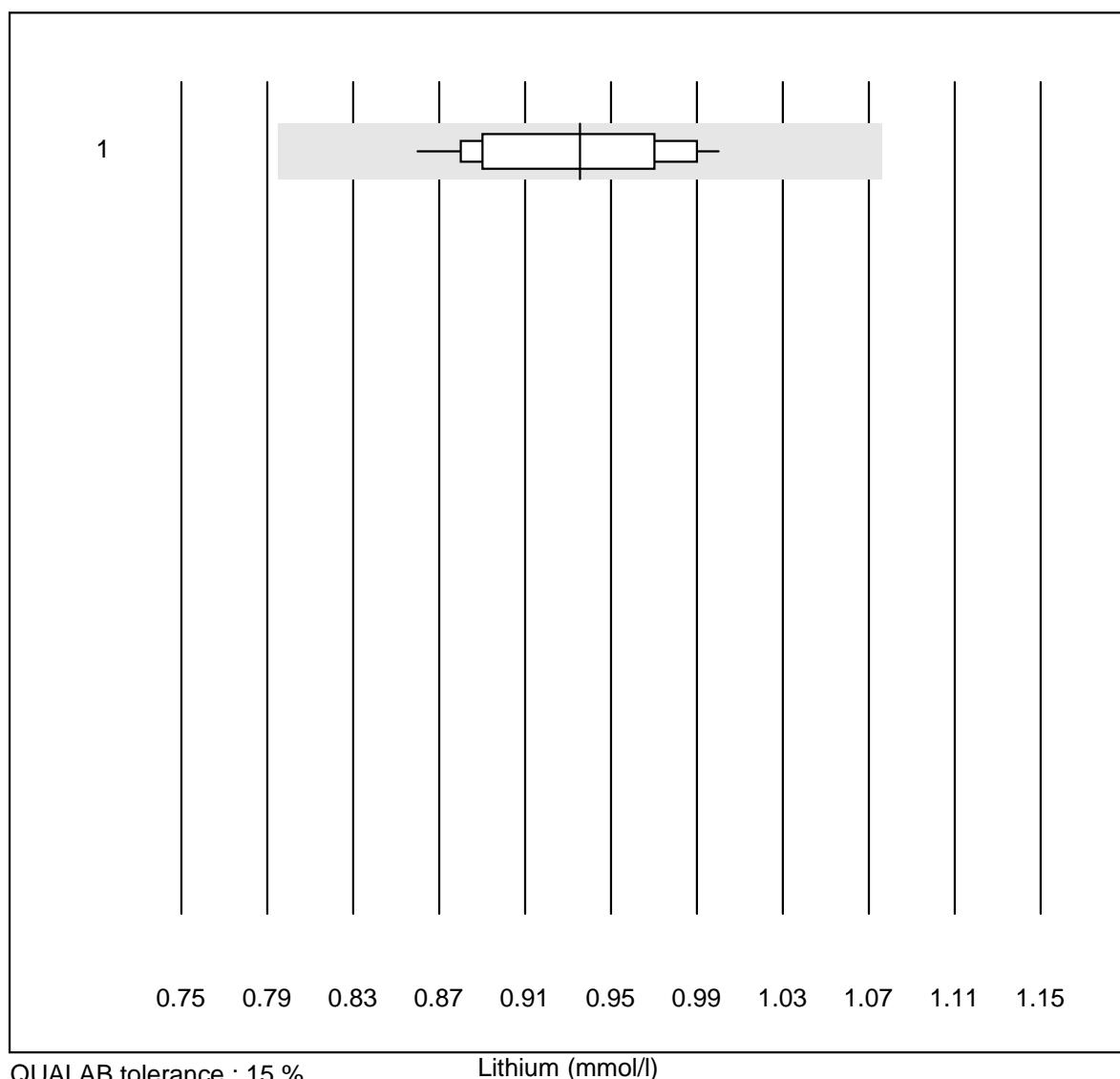
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Piccolo	7	100.0	0.0	0.0	22	5.6	e*

LDL Cholesterin



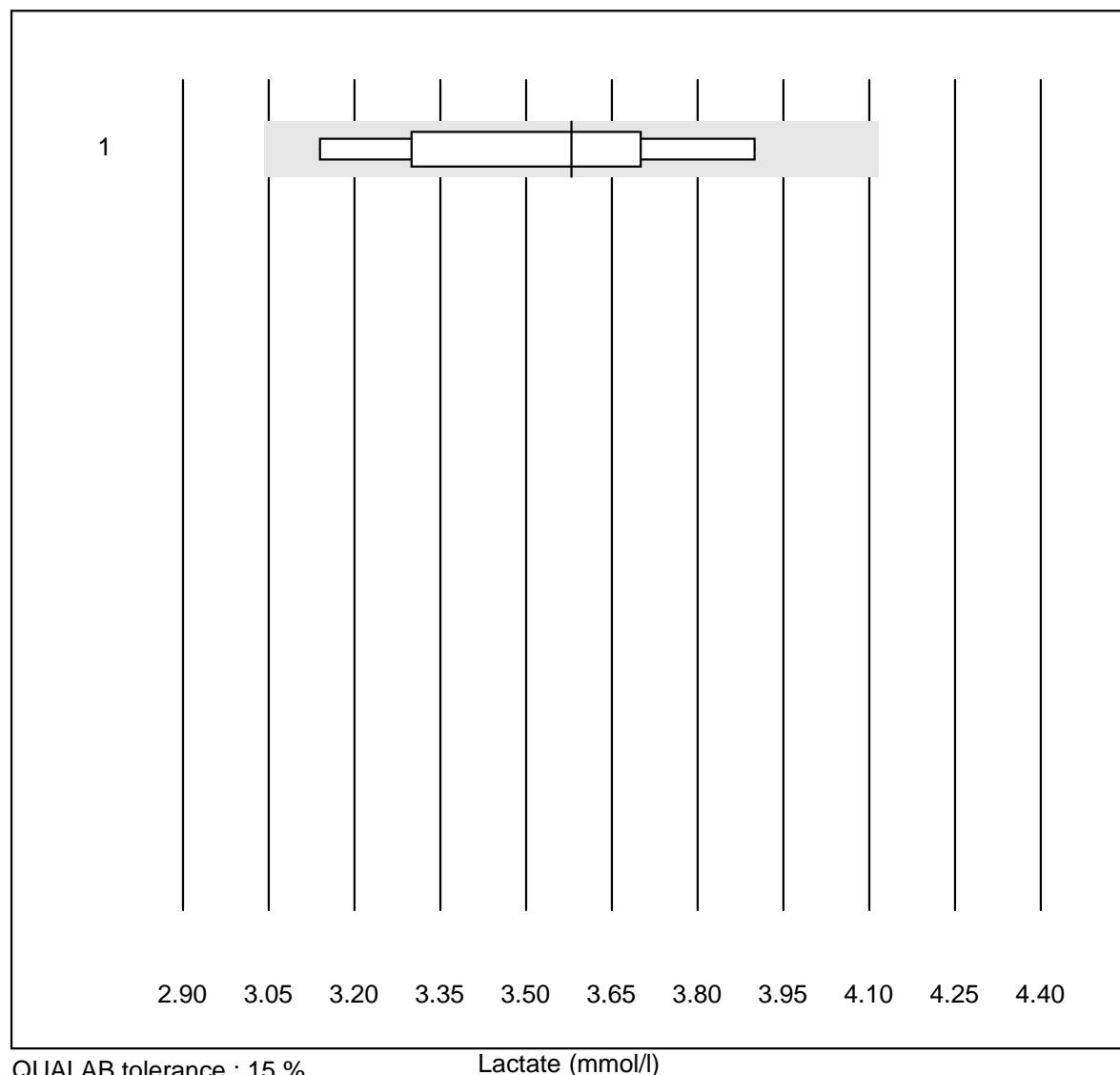
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Abx Mira	9	100.0	0.0	0.0	2.6	6.7	e
2 Standard chemistry	8	100.0	0.0	0.0	3.0	4.4	e
3 Roche, Cobas	4	100.0	0.0	0.0	3.0	1.9	e
4 Hitachi S40/M40	4	100.0	0.0	0.0	2.4	2.3	e

Lithium

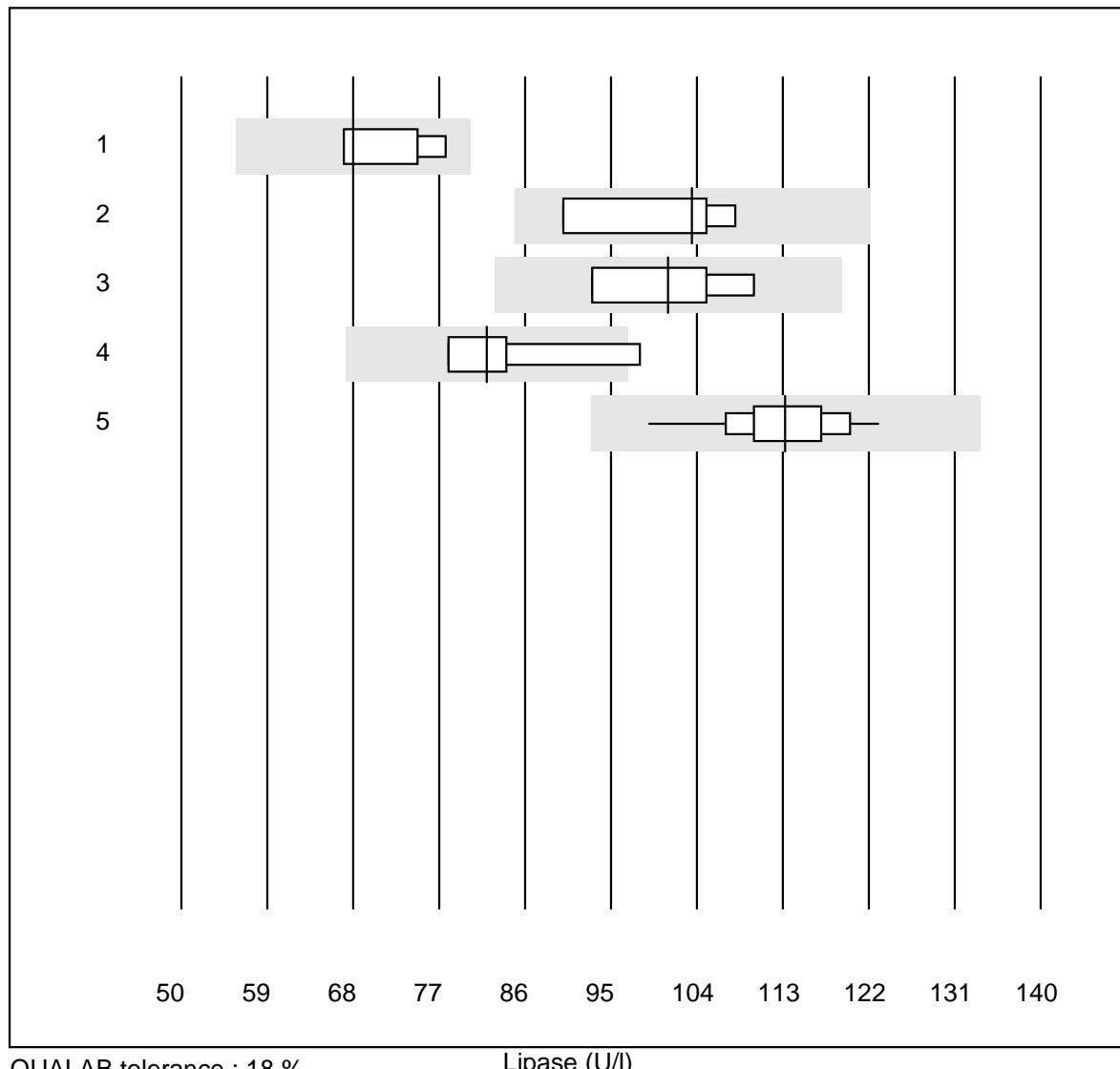


No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	all Participants	14	100.0	0.0	0.0	0.94	4.6	e

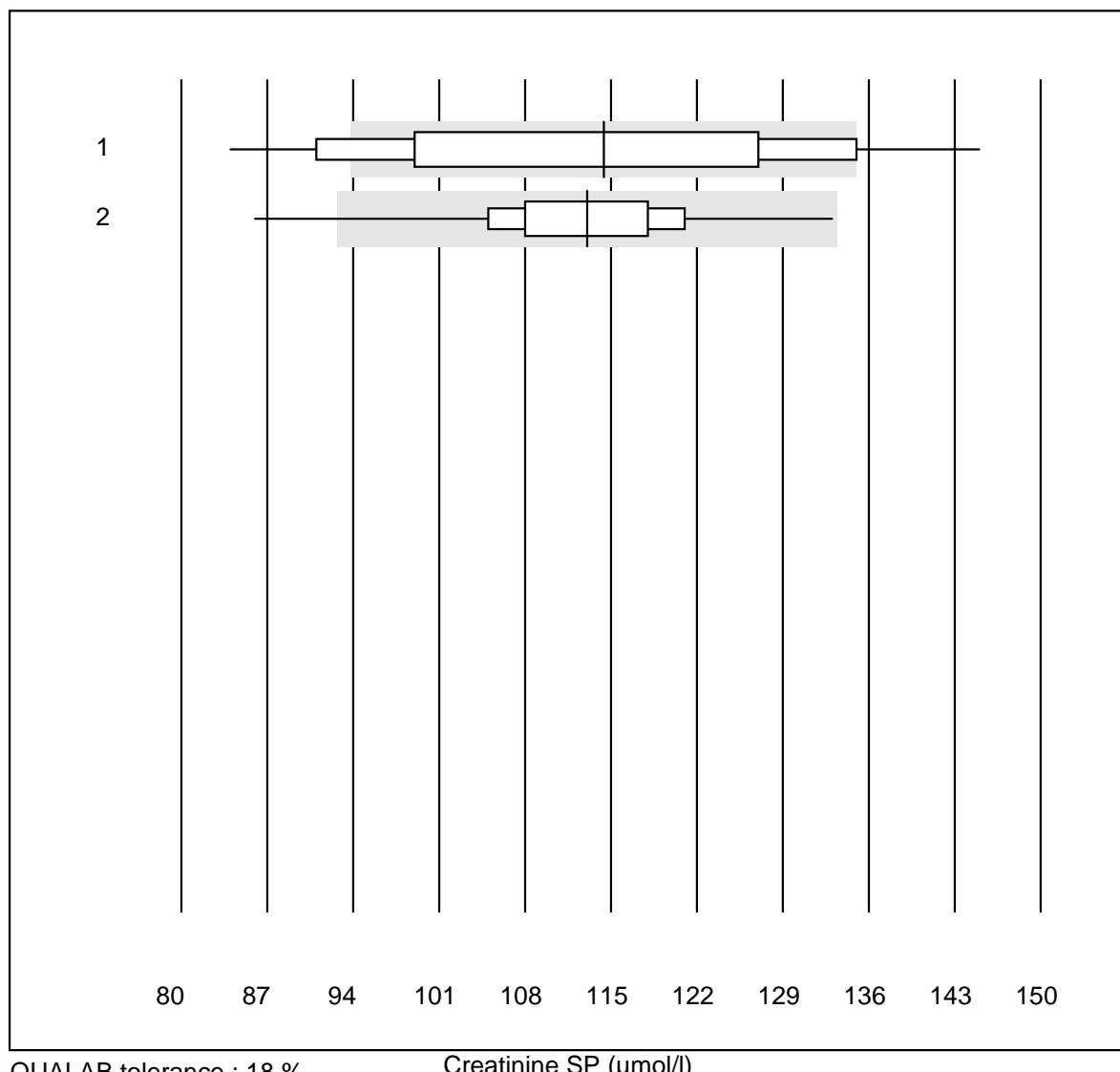
Lactate



No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	all Participants	7	100.0	0.0	0.0	3.58	7.2	e*

Lipase

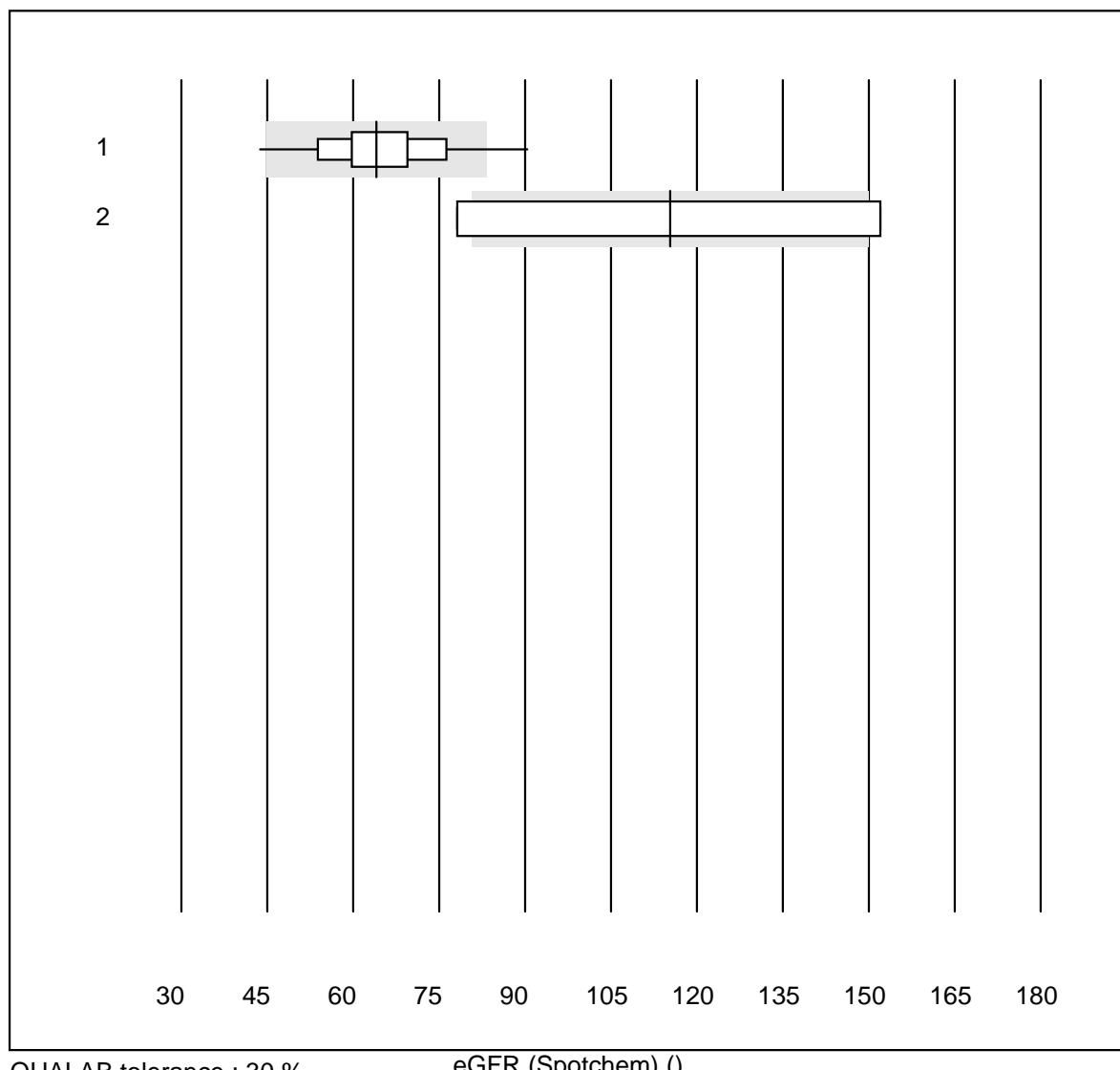
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Abx Mira	4	100.0	0.0	0.0	68.0	7.2	e*
2 Architect	4	100.0	0.0	0.0	103.5	7.8	e*
3 Beckman/Olympus	4	100.0	0.0	0.0	101.0	7.6	e*
4 Cobas	8	87.5	12.5	0.0	82.0	7.5	e*
5 Fuji Dri-Chem	58	98.3	0.0	1.7	113.3	4.8	e

Creatinine SP

QUALAB tolerance : 18 %

Creatinine SP ($\mu\text{mol/l}$)

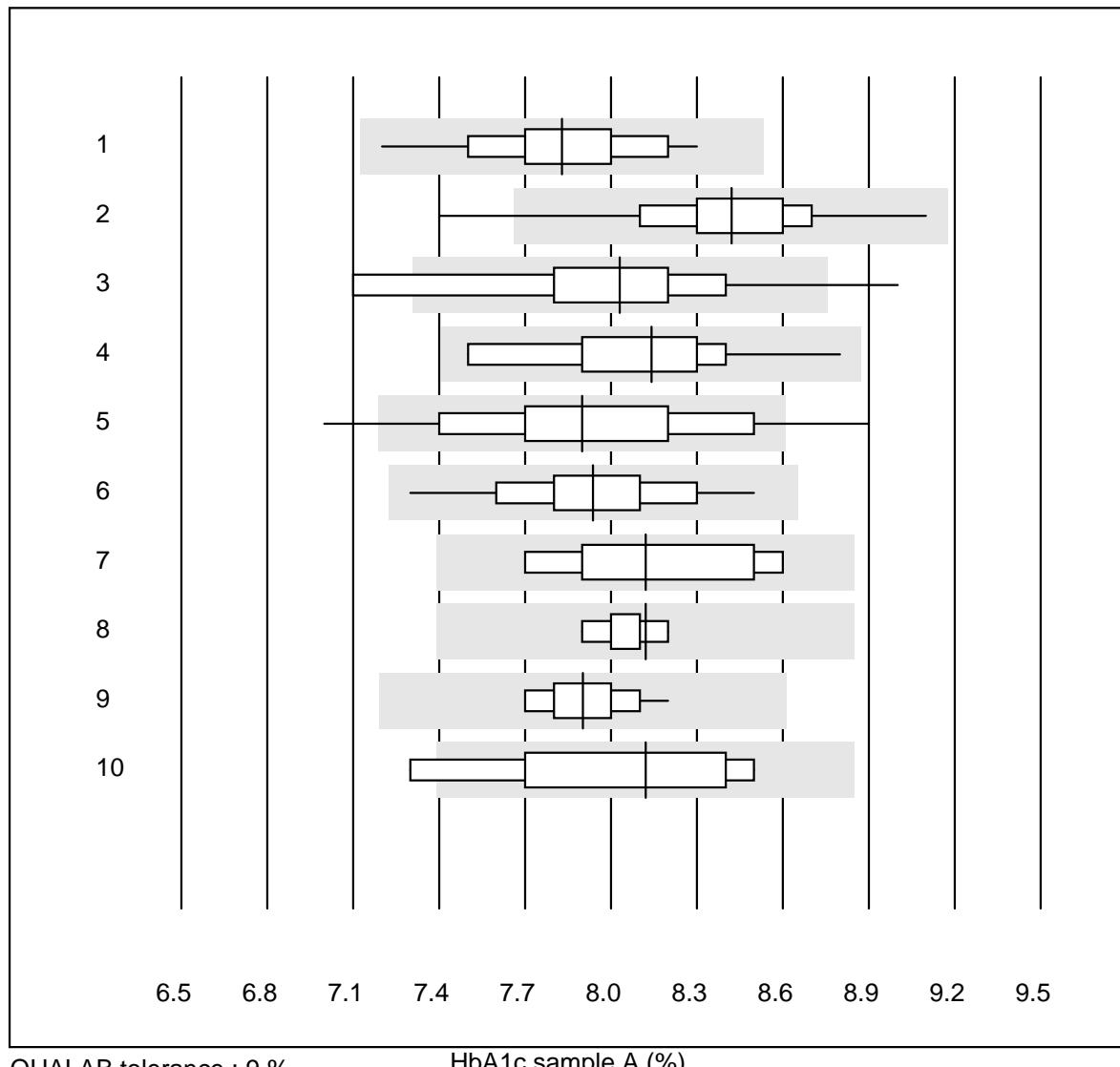
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Spotchem/Ready	156	67.9	23.1	9.0	114	14.2	e
2 Spotchem D-Concept	135	98.5	1.5	0.0	113	6.3	e

eGFR (Spotchem)

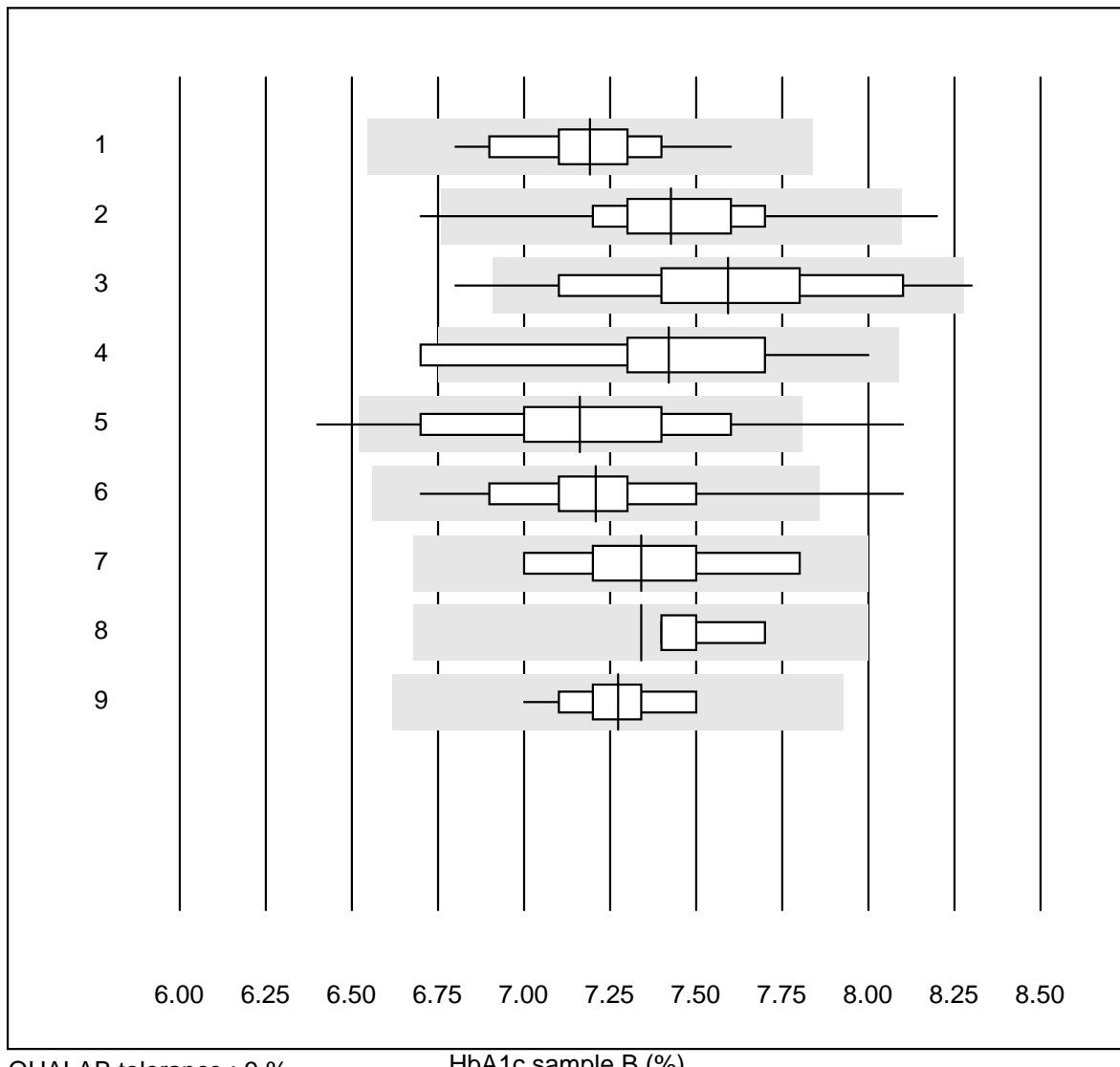
QUALAB tolerance : 30 %

eGFR (Spotchem) ()

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 CKD-EPI	94	88.3	5.3	6.4	64	13.9	e
2 Cockcroft-Gault	5	20.0	40.0	40.0	115	32.3	a

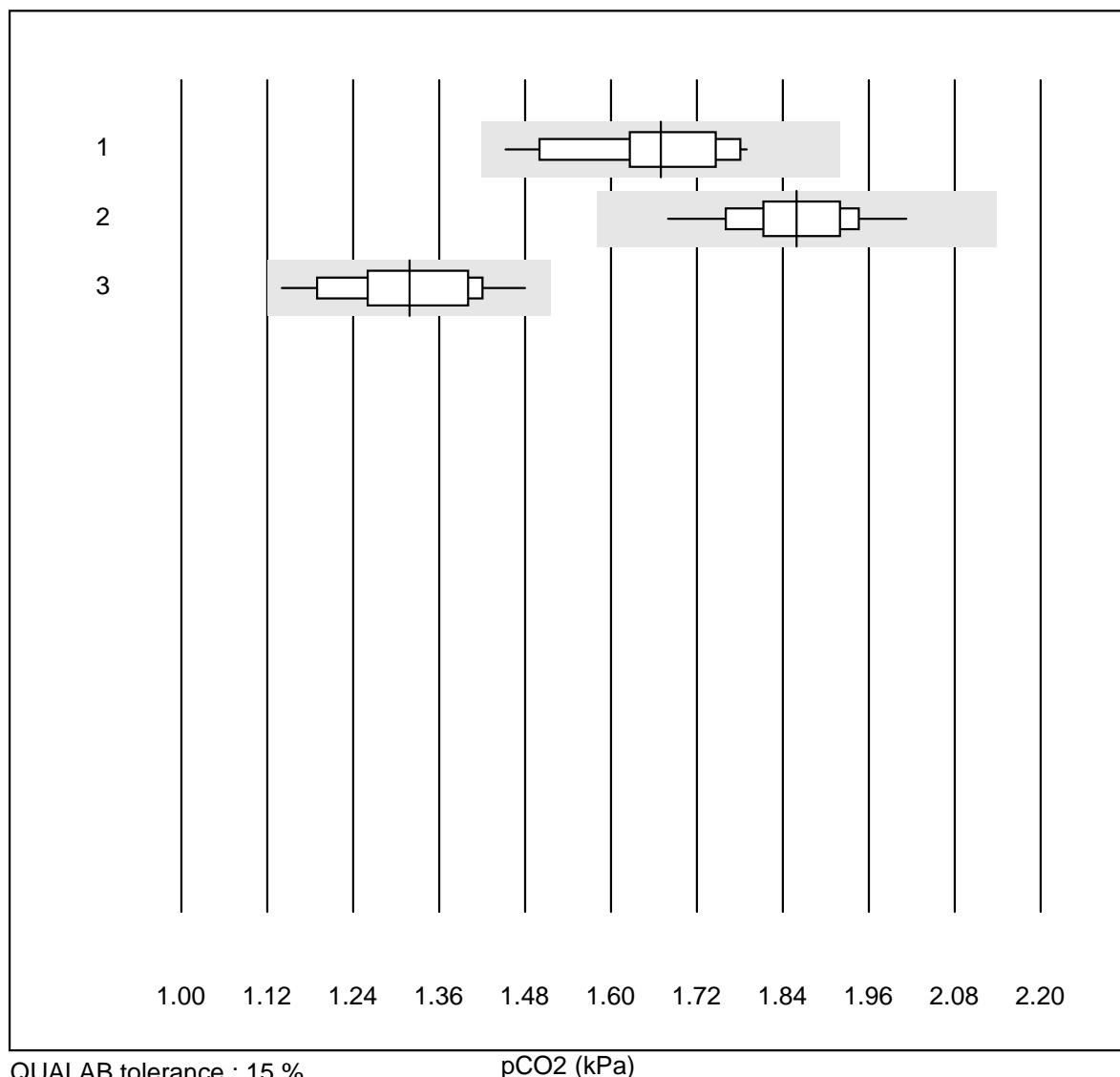
HbA1c sample A

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Cobas b101	26	100.0	0.0	0.0	7.8	3.5	e
2 Afinion	627	99.1	0.6	0.3	8.4	2.7	e
3 Eurolyser	11	72.7	18.2	9.1	8.0	6.5	e*
4 Hemocue HbA1c 501	10	100.0	0.0	0.0	8.1	4.3	e*
5 NycoCard	107	83.2	8.4	8.4	7.9	5.2	e
6 DCA2000/Vantage	200	100.0	0.0	0.0	7.9	3.1	e
7 Others	7	100.0	0.0	0.0	8.1	3.9	a
8 HPLC	6	100.0	0.0	0.0	8.1	1.3	a
9 Roche, Cobas	17	100.0	0.0	0.0	7.9	1.8	e
10 A1c Now	7	85.7	14.3	0.0	8.1	5.2	a

HbA1c sample B

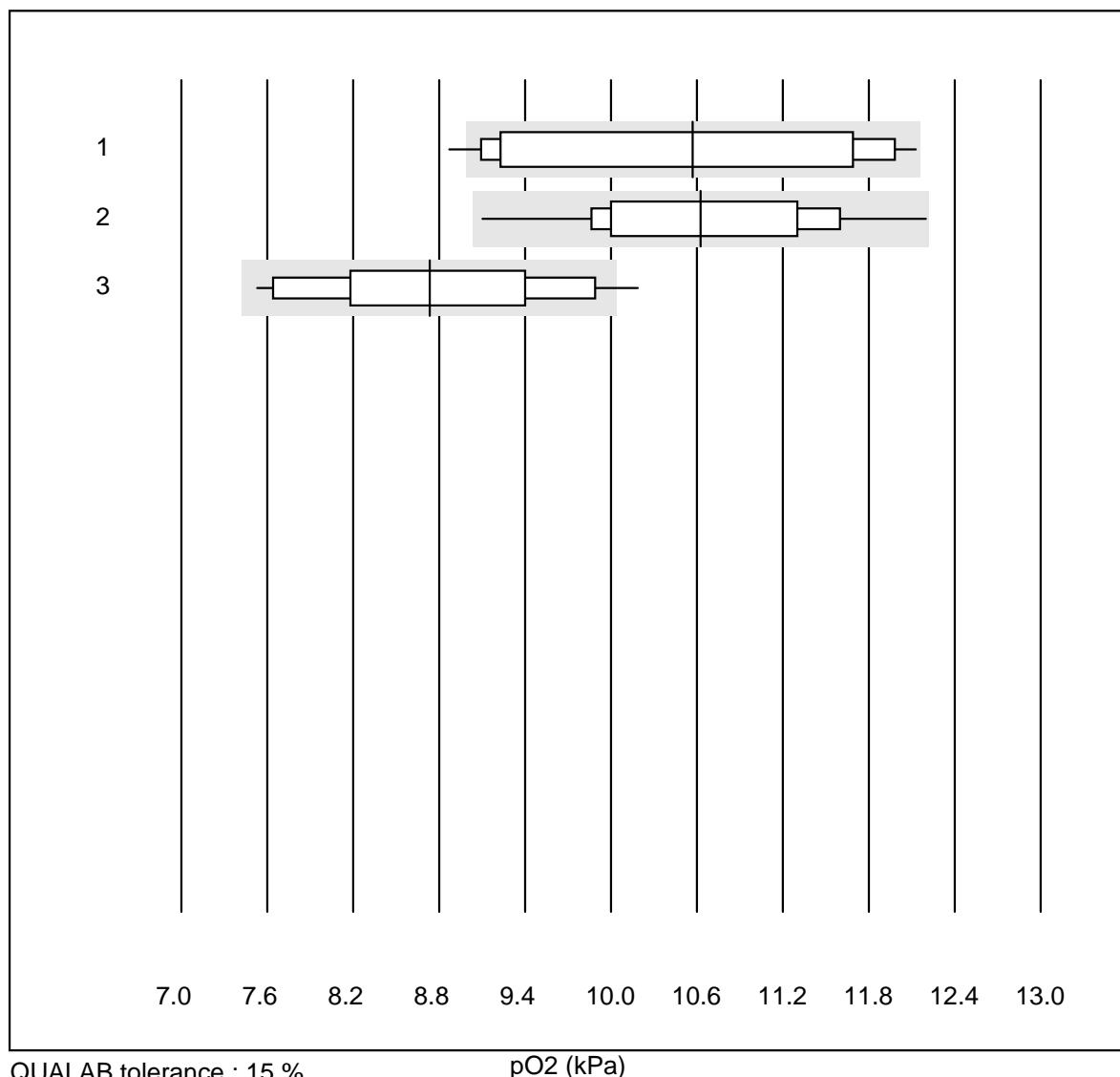
K4 Blood gases

pCO₂



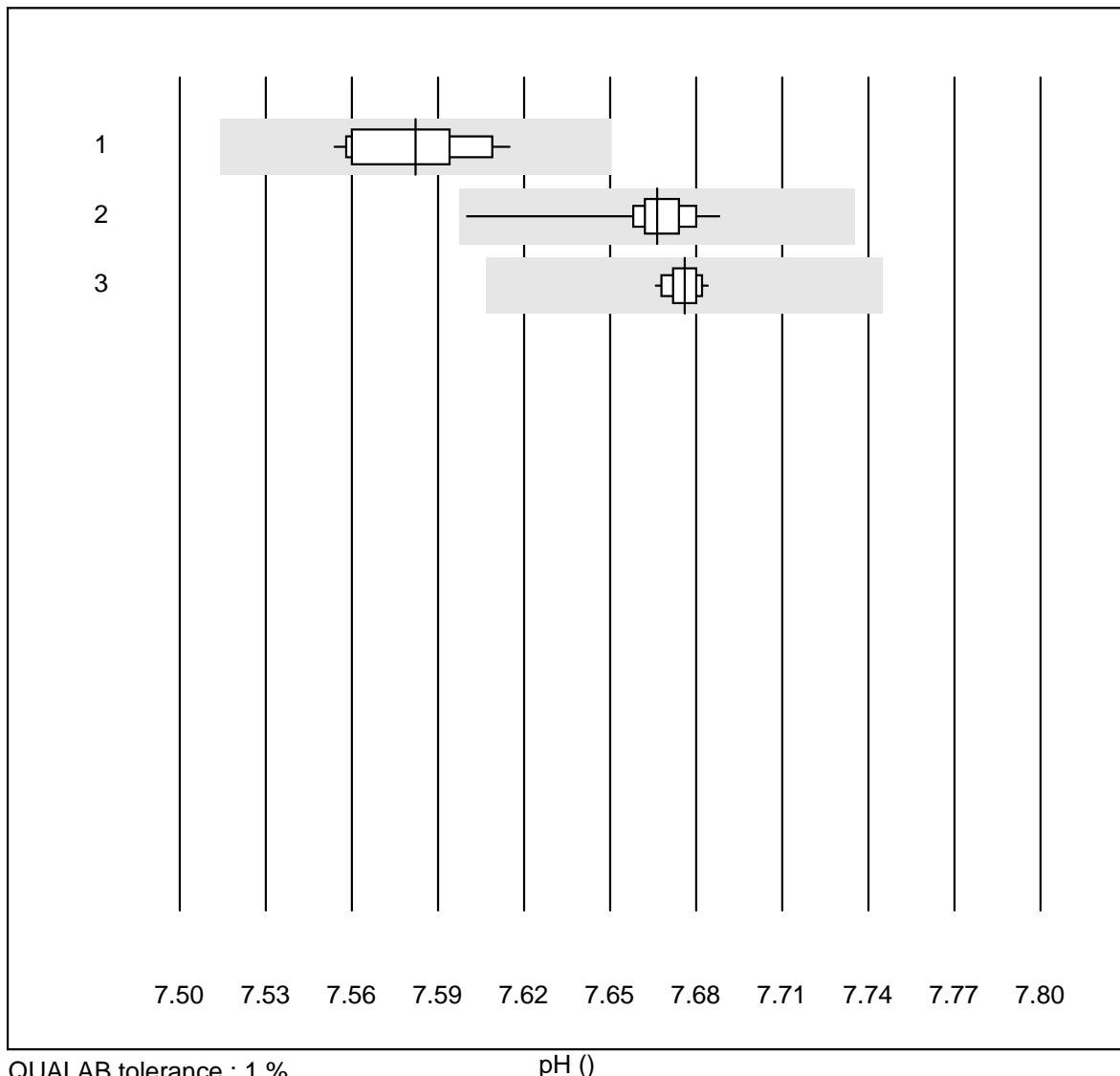
K4 Blood gases

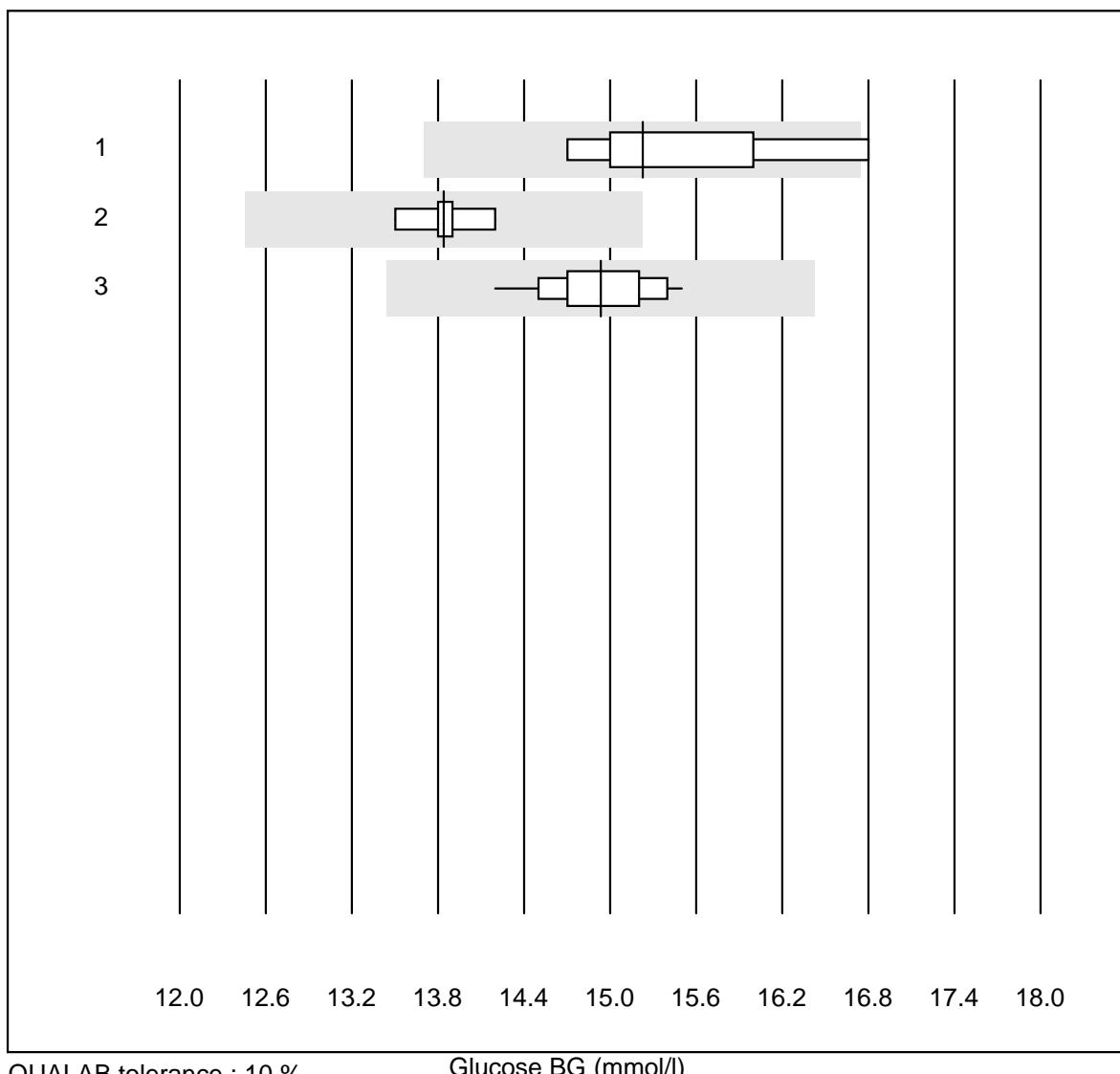
pO₂



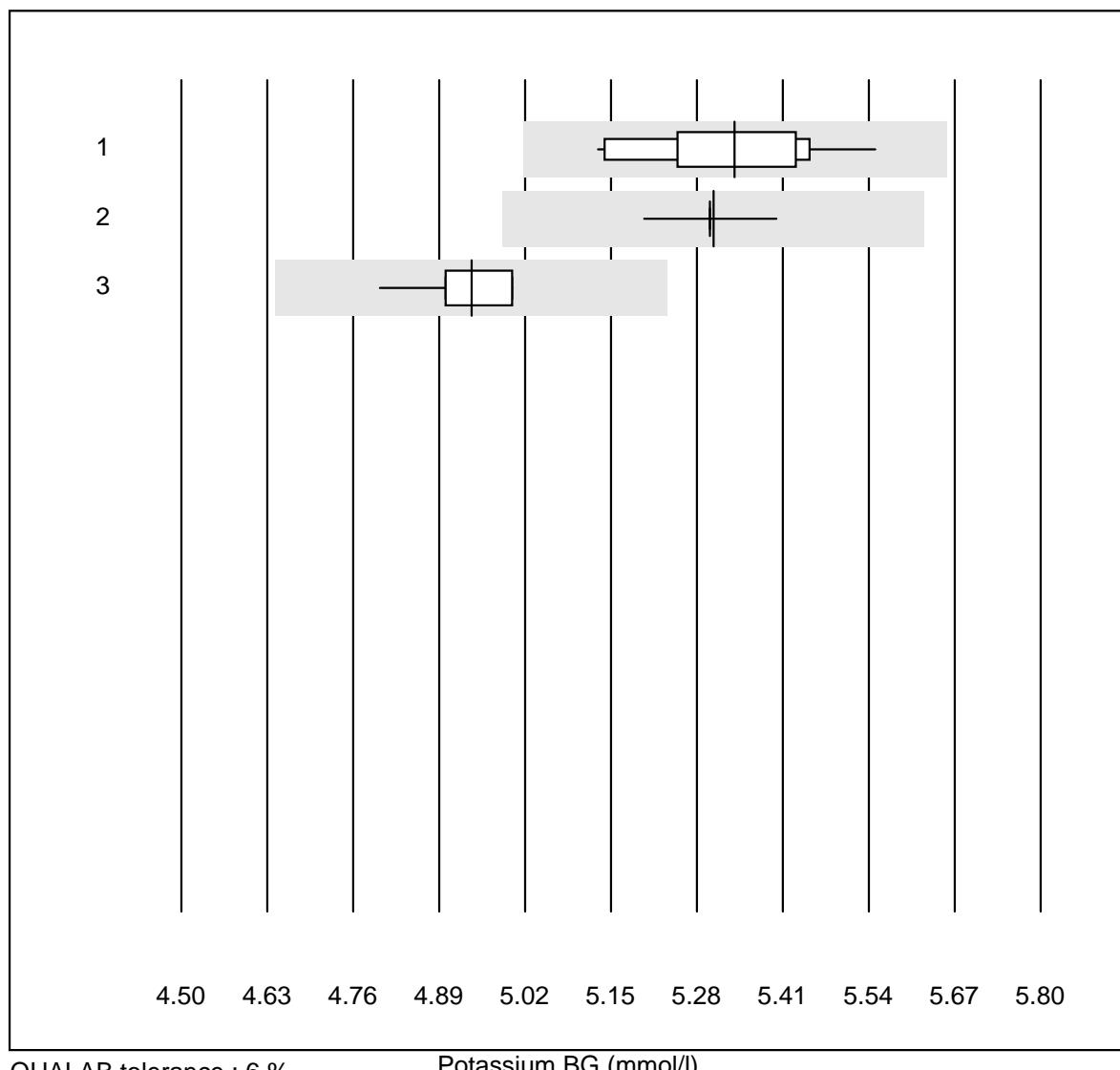
K4 Blood gases

pH



Glucose BG

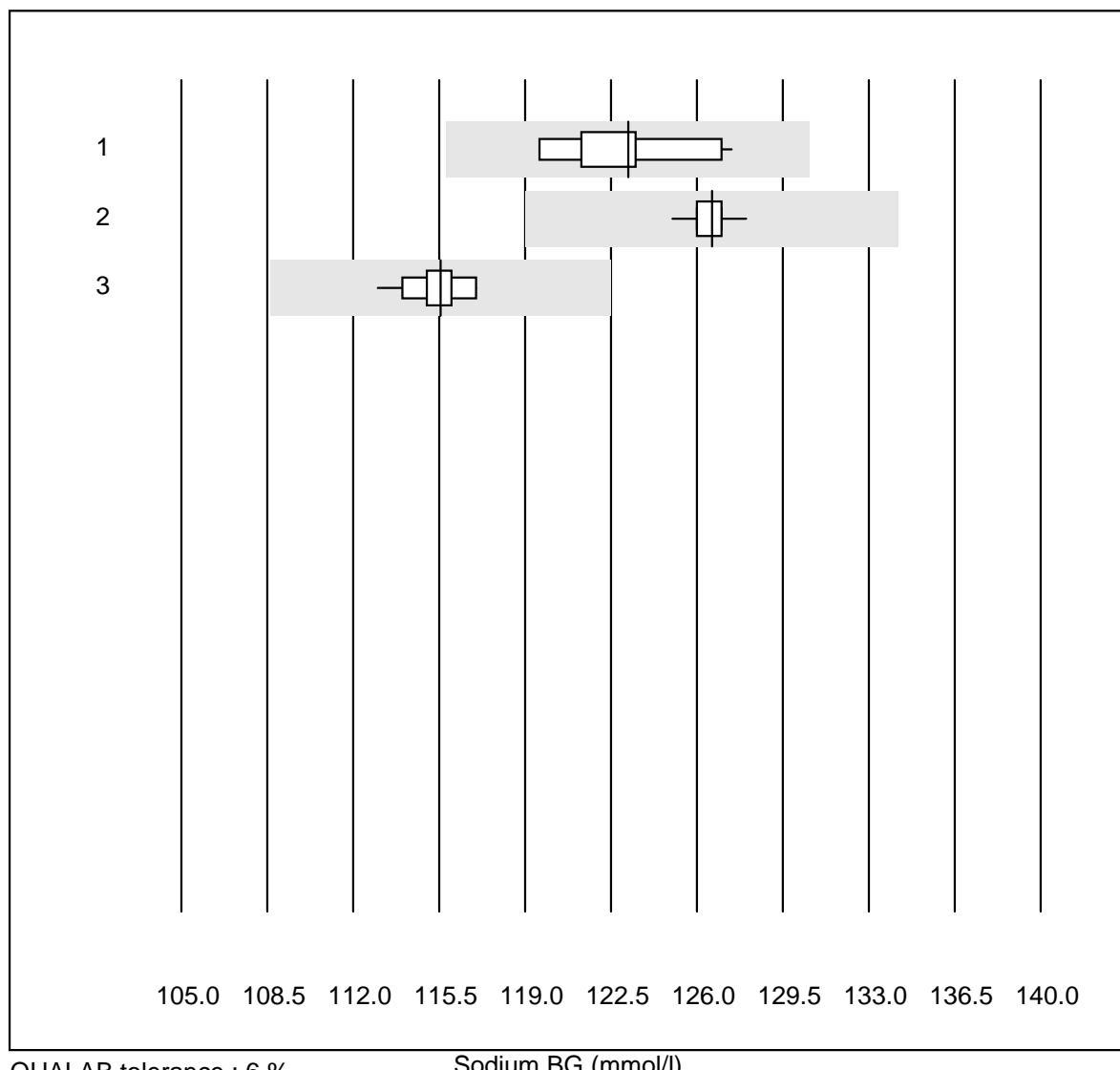
Potassium BG

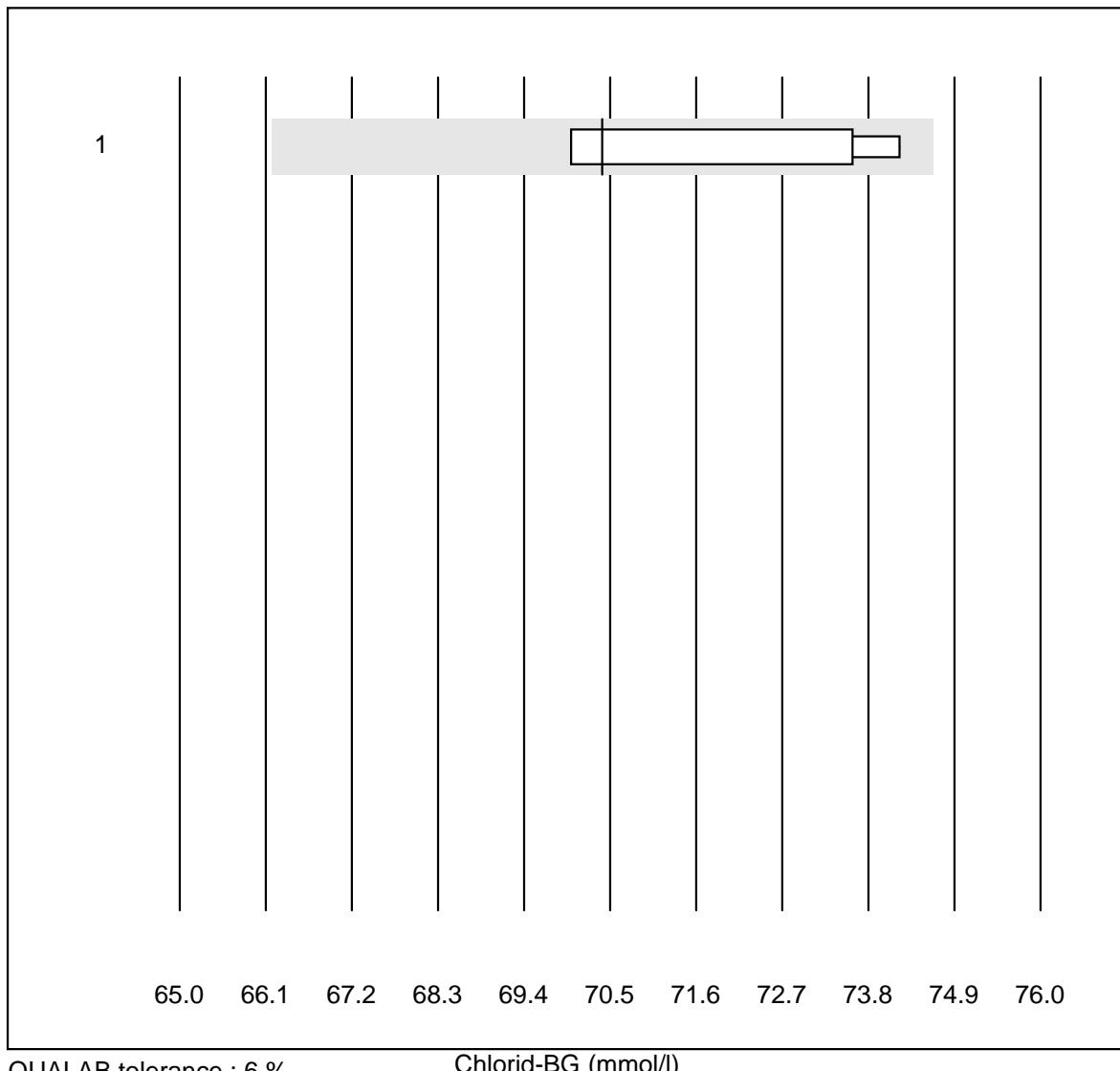


QUALAB tolerance : 6 %

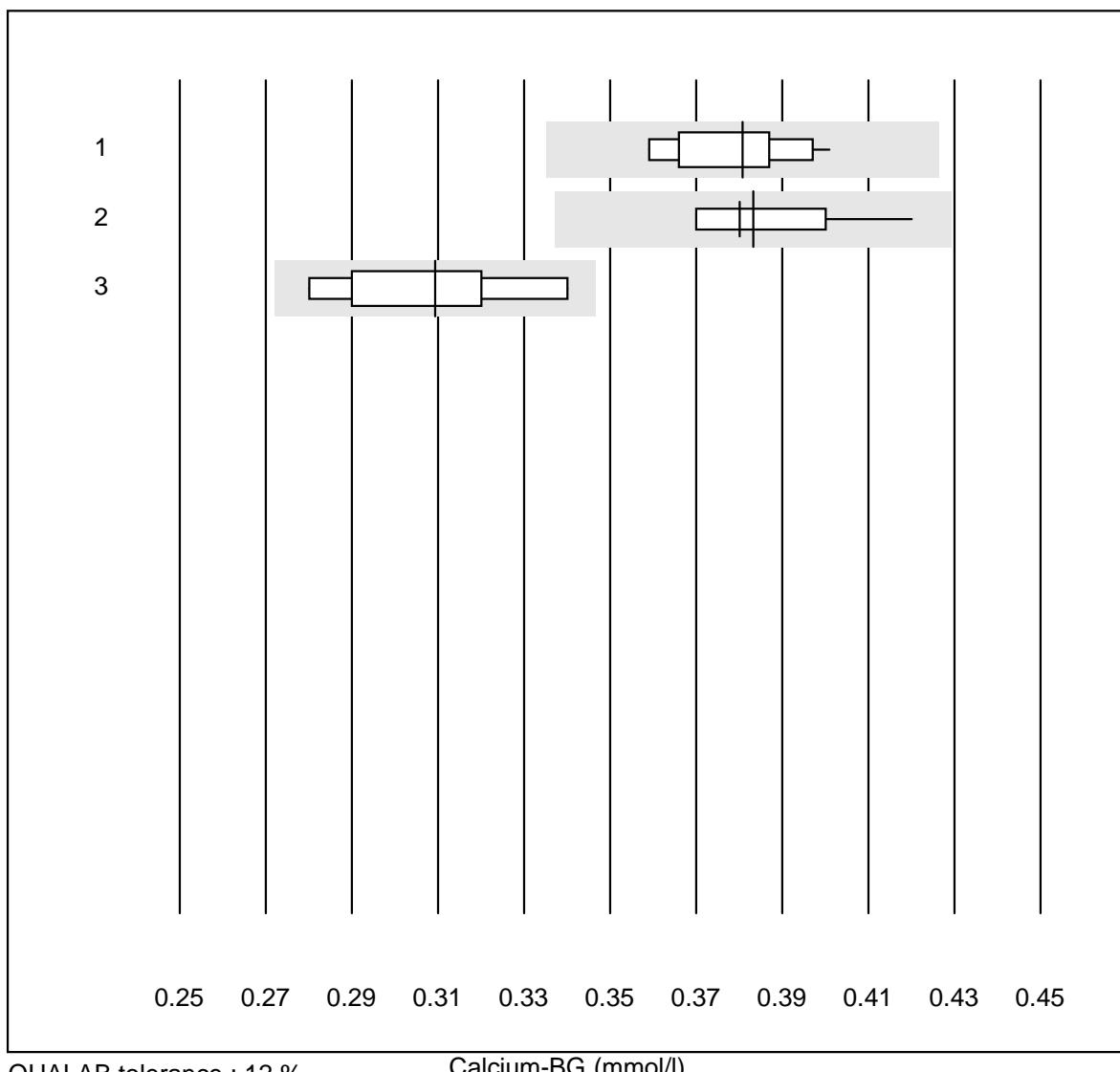
Potassium BG (mmol/l)

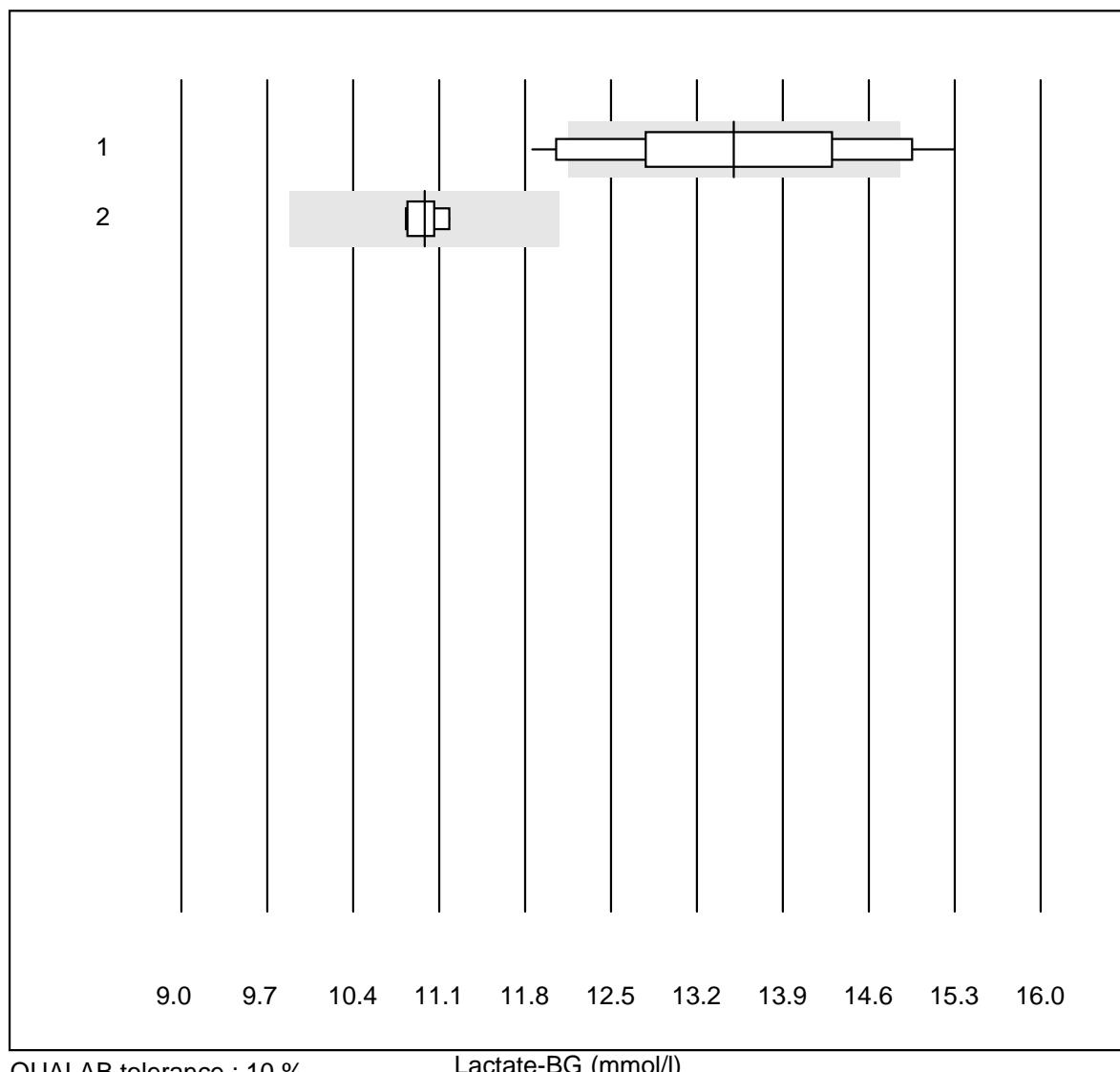
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Cobas b121/123/221	15	100.0	0.0	0.0	5.3	2.2	e
2 iStat	21	100.0	0.0	0.0	5.3	0.7	e
3 EPOC	19	100.0	0.0	0.0	4.9	1.2	e

Sodium BG

Chlorid-BG

No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Cobas b121/123/221	6	100.0	0.0	0.0	70.4	2.7	e*

Calcium-BG

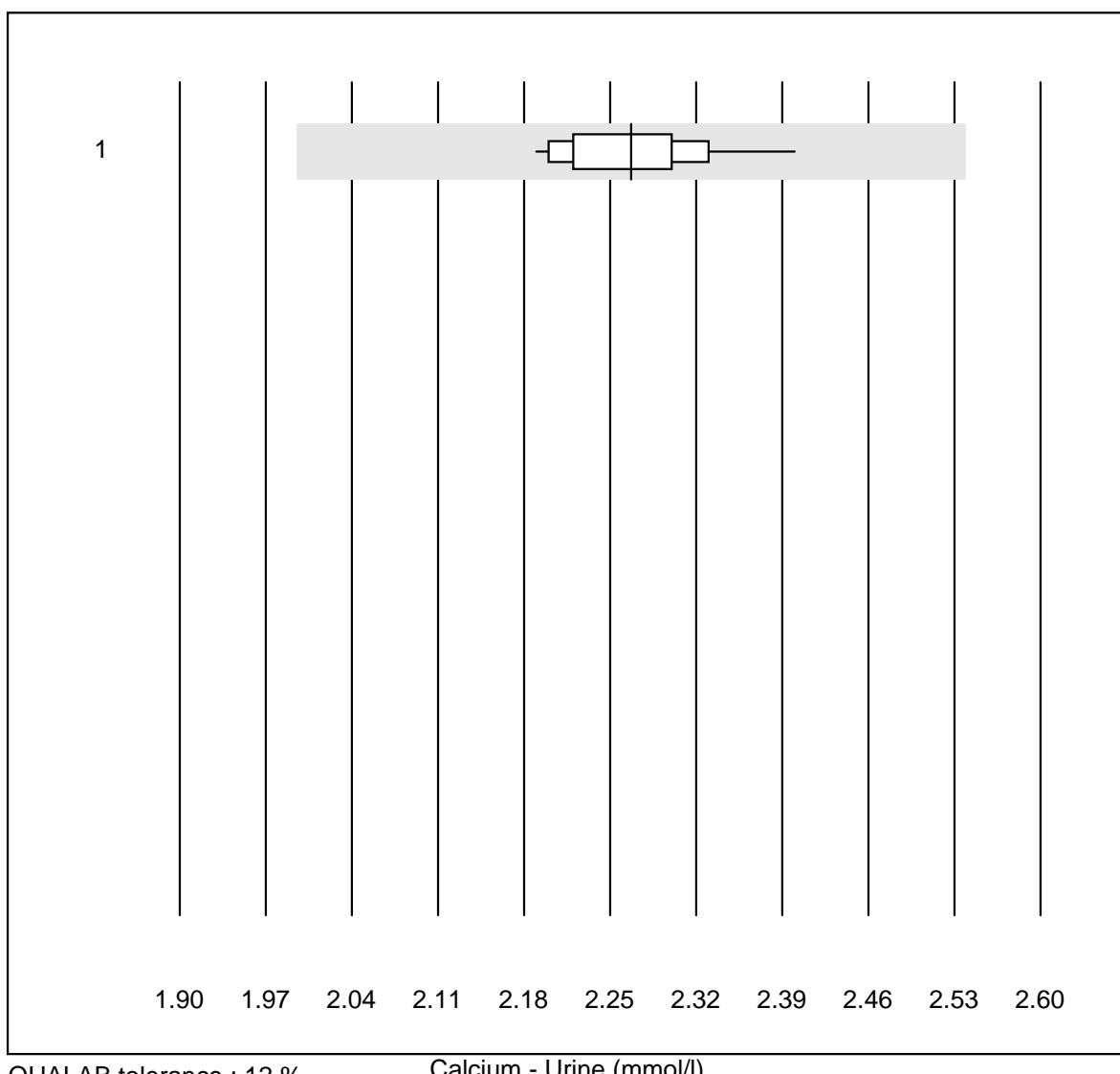
Lactate-BG

QUALAB tolerance : 10 %

Lactate-BG (mmol/l)

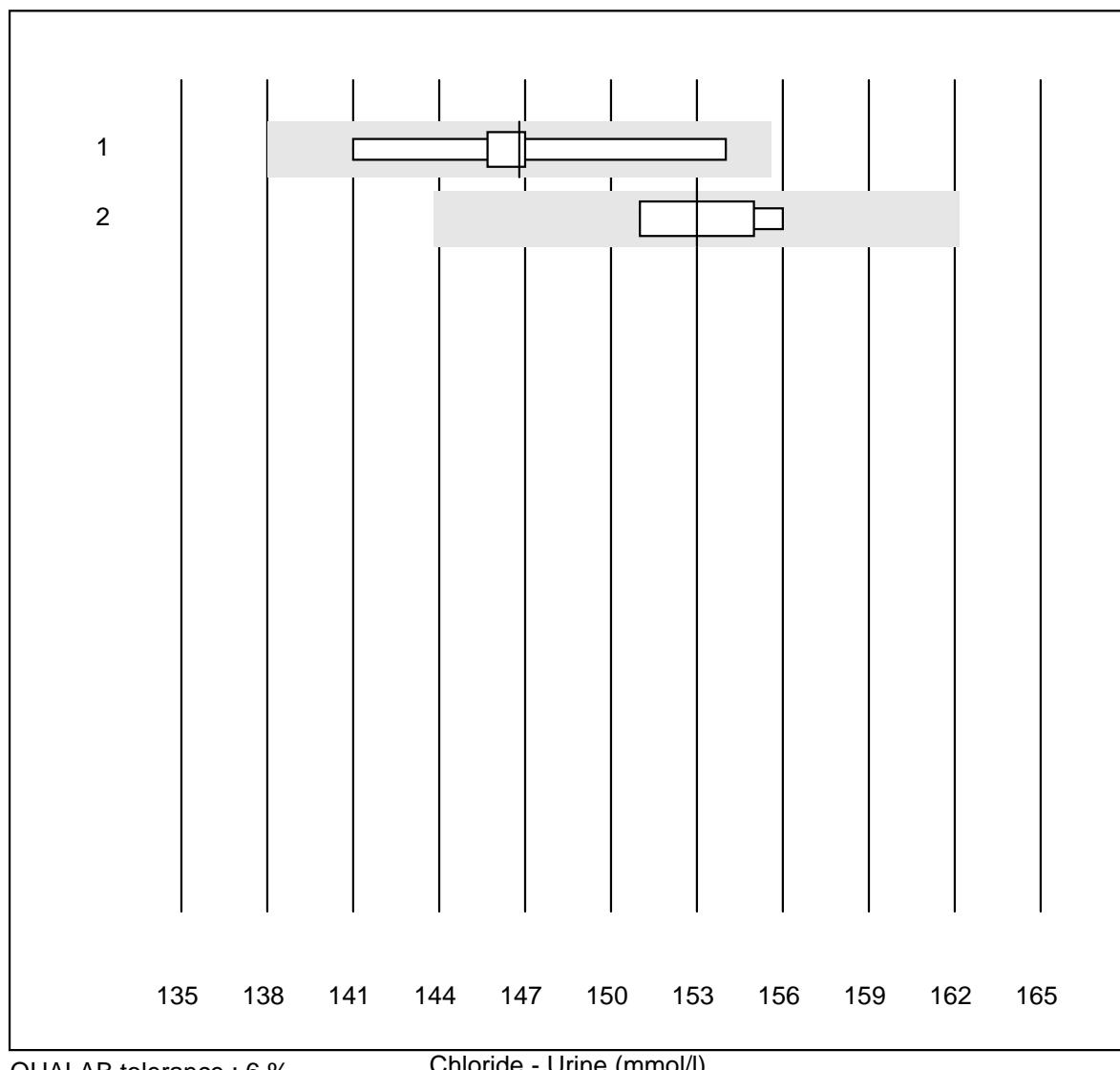
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 EPOC	20	70.0	20.0	10.0	13.50	7.5	e*
2 iStat	7	100.0	0.0	0.0	10.98	1.2	e

Calcium - Urine



No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Standard chemistry	13	100.0	0.0	0.0	2.27	2.6	e

Chloride - Urine

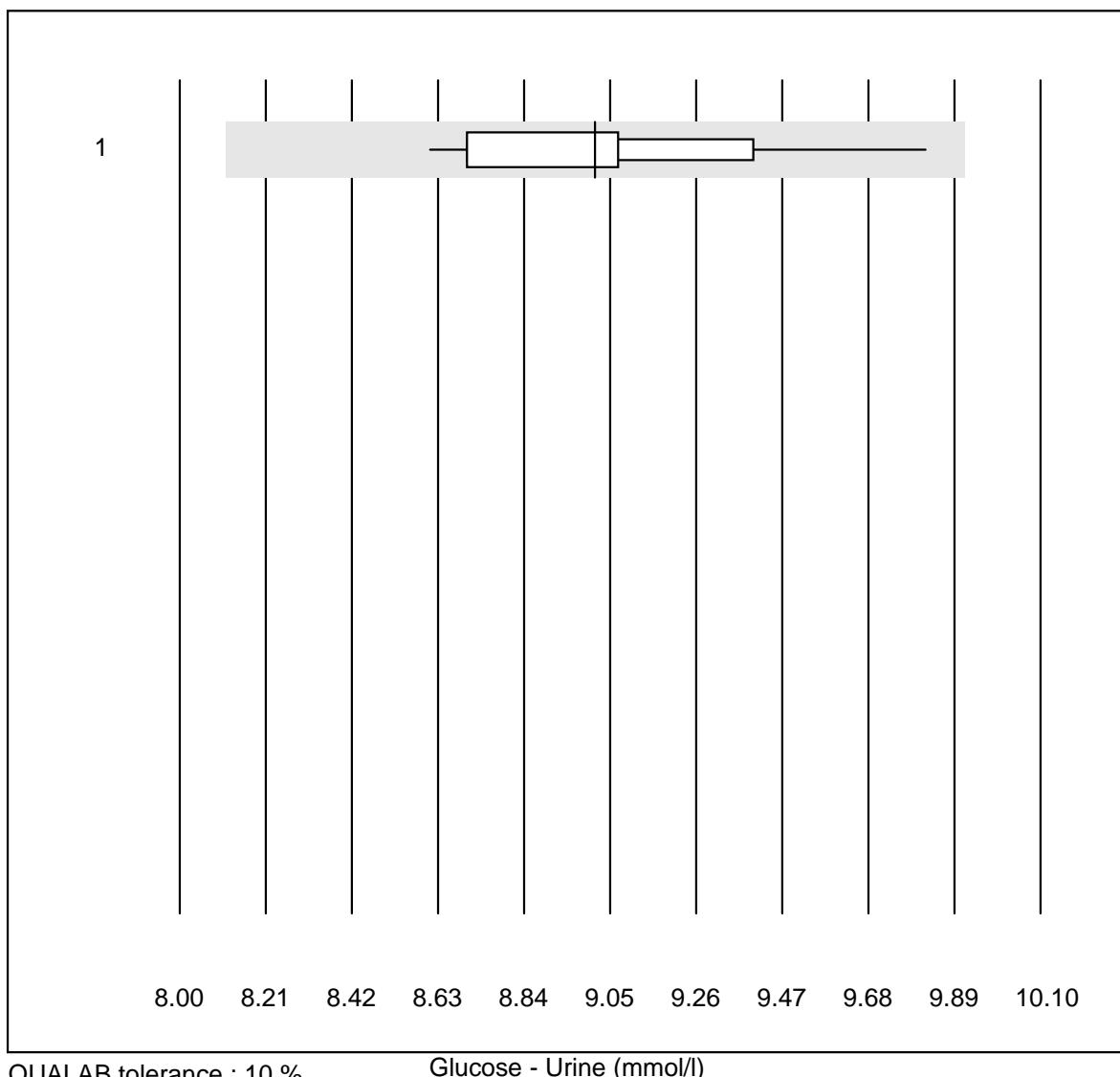


QUALAB tolerance : 6 %

Chloride - Urine (mmol/l)

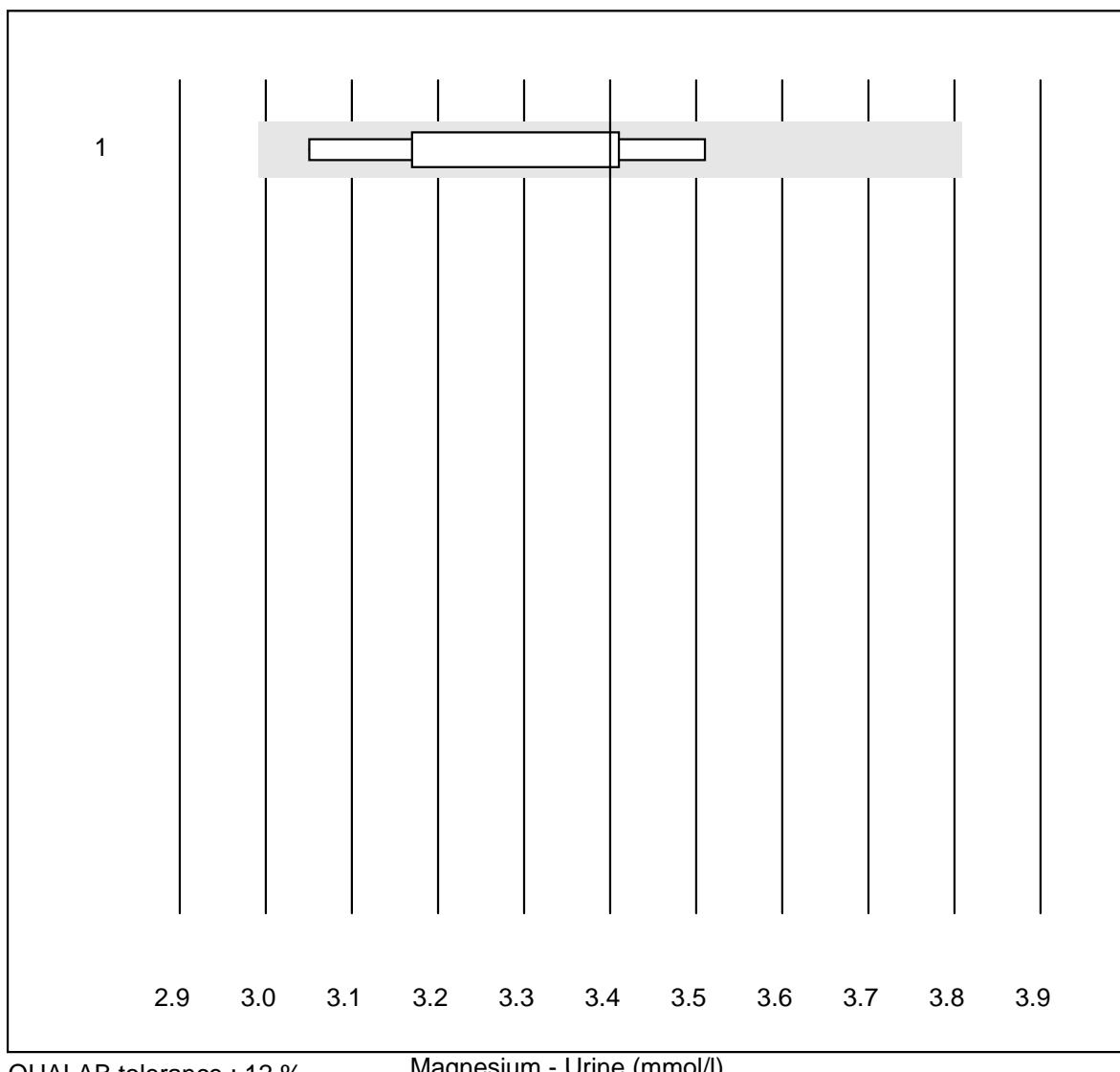
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Standard chemistry	5	100.0	0.0	0.0	147	3.2	e*
2 ISE	4	100.0	0.0	0.0	153	1.7	e*

Glucose - Urine



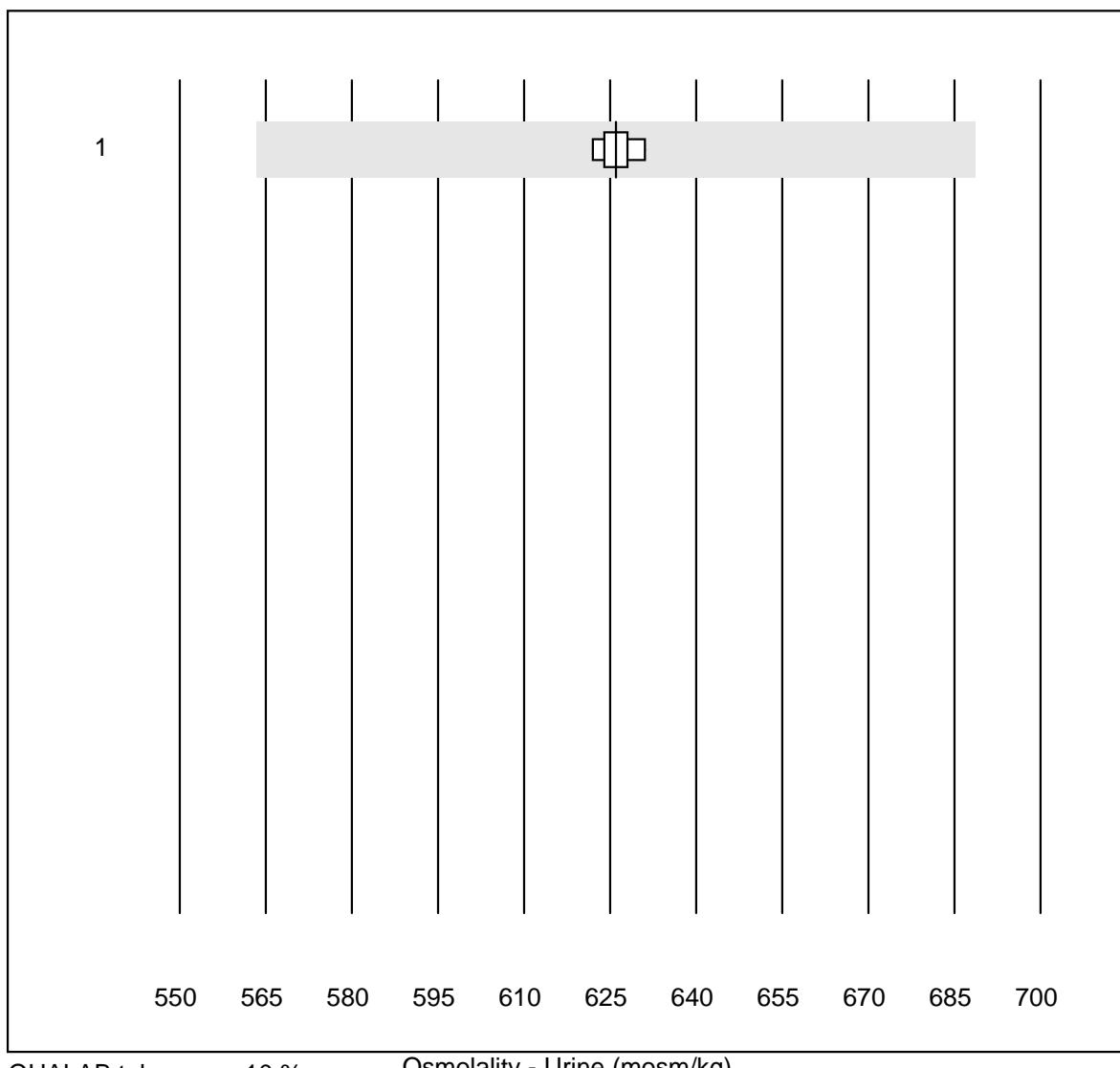
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Standard chemistry	12	100.0	0.0	0.0	9.0	3.9	e

Magnesium - Urine



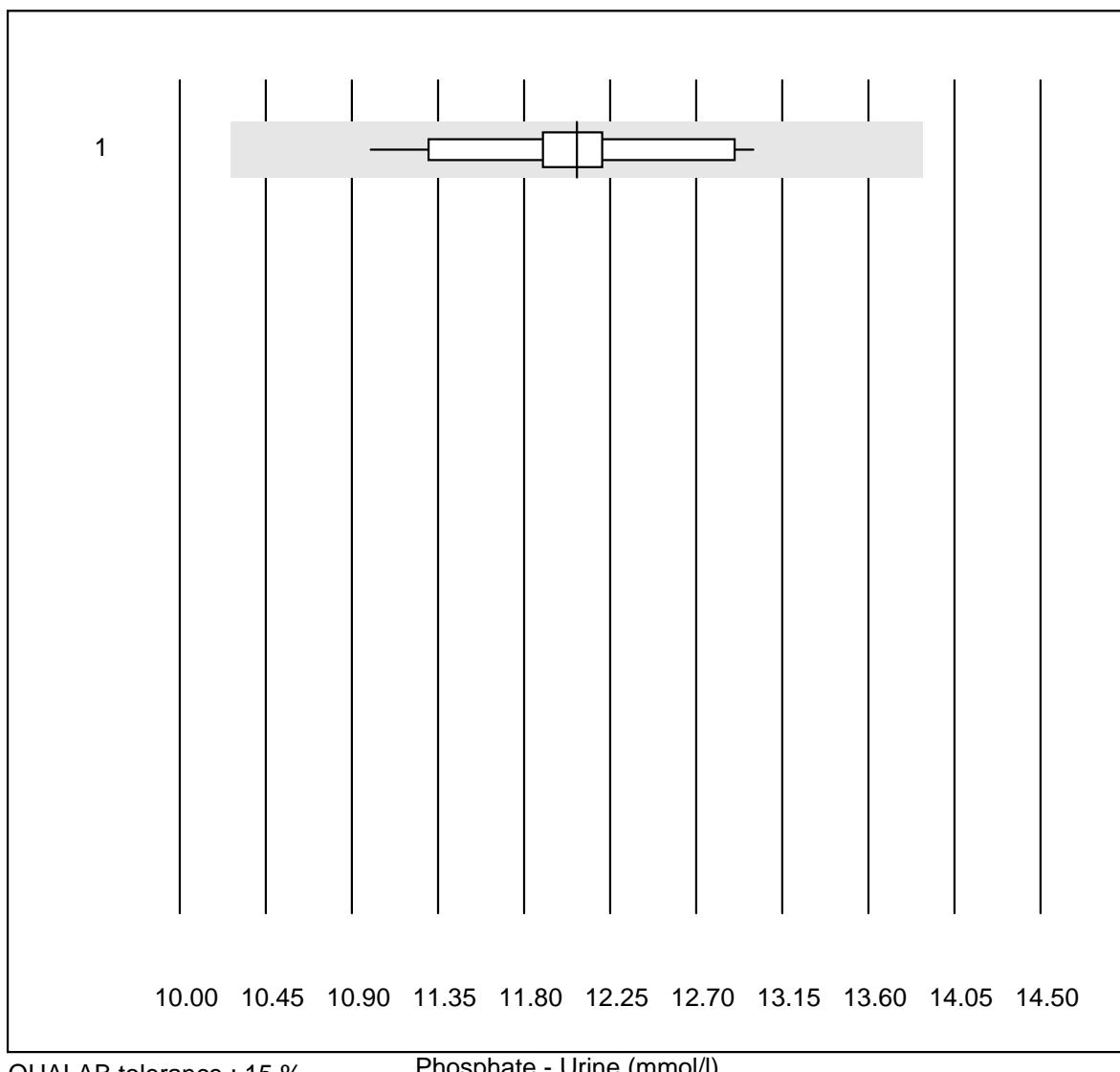
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Standard chemistry	8	100.0	0.0	0.0	3.4	5.1	e*

Osmolality - Urine



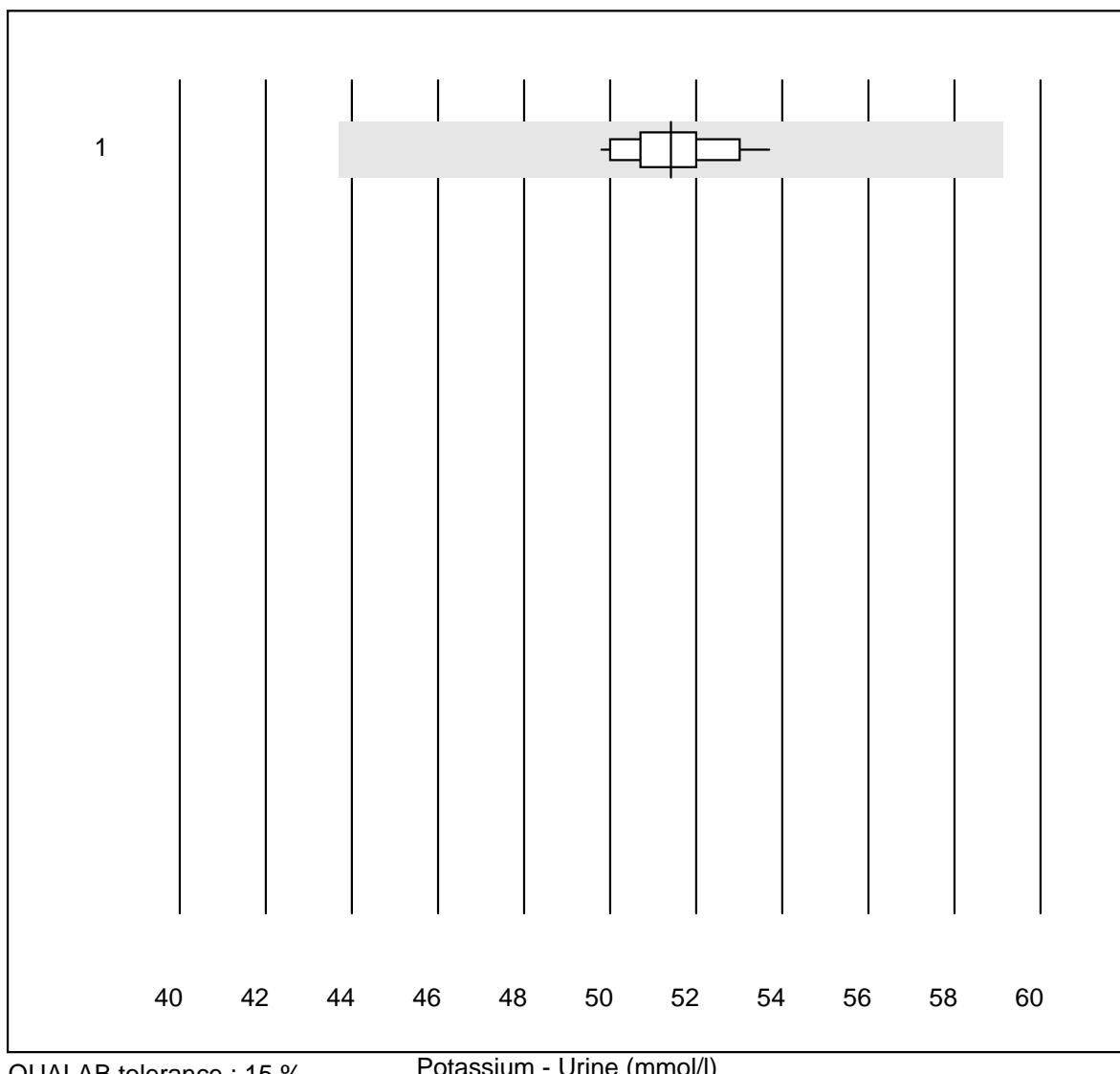
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Cryoskopy	6	100.0	0.0	0.0	626	0.5	e

Phosphate - Urine

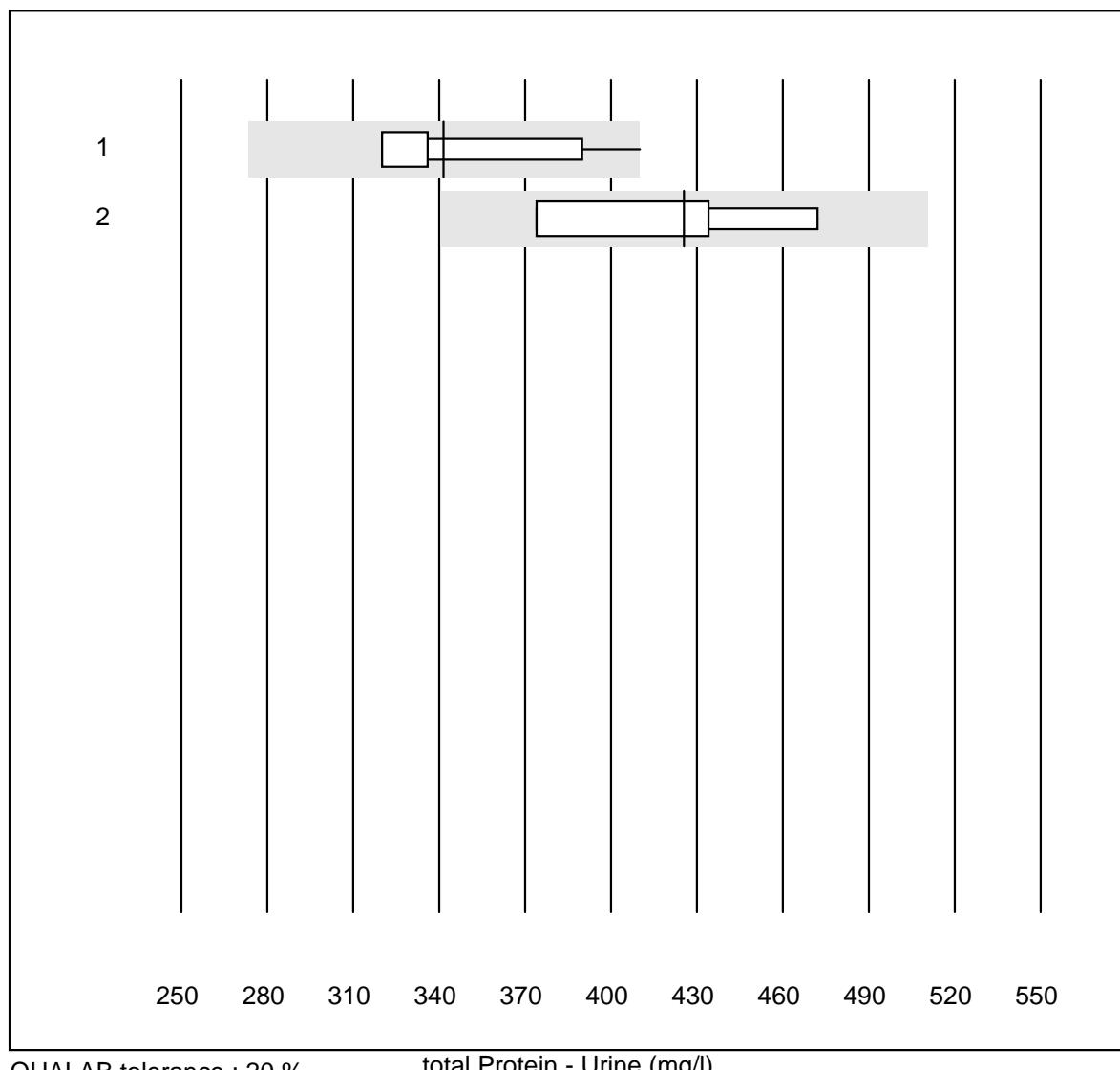


No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Standard chemistry	14	100.0	0.0	0.0	12.1	4.4	e

Potassium - Urine



No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 all Participants	18	100.0	0.0	0.0	51	2.2	e

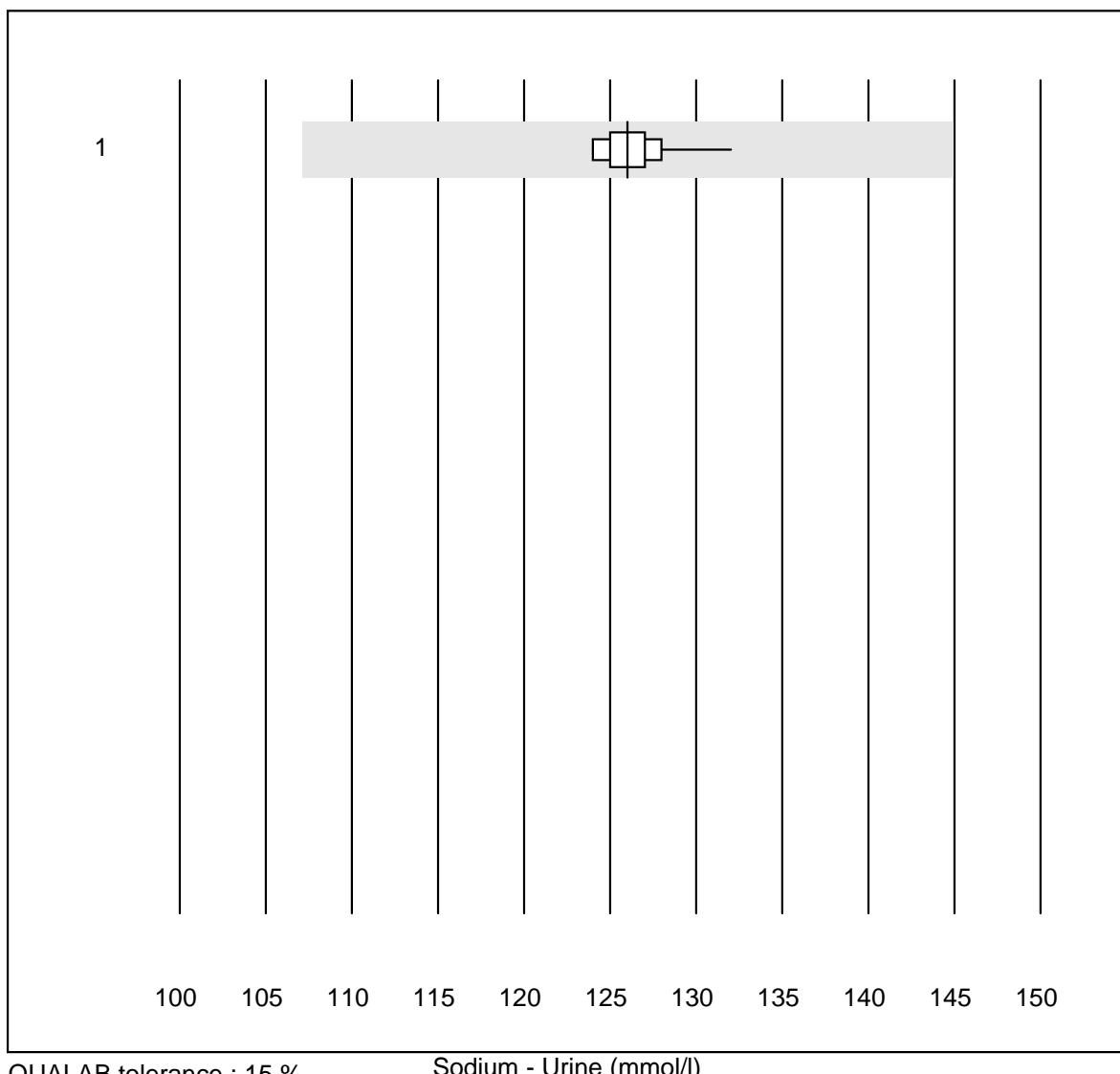
total Protein - Urine

QUALAB tolerance : 20 %

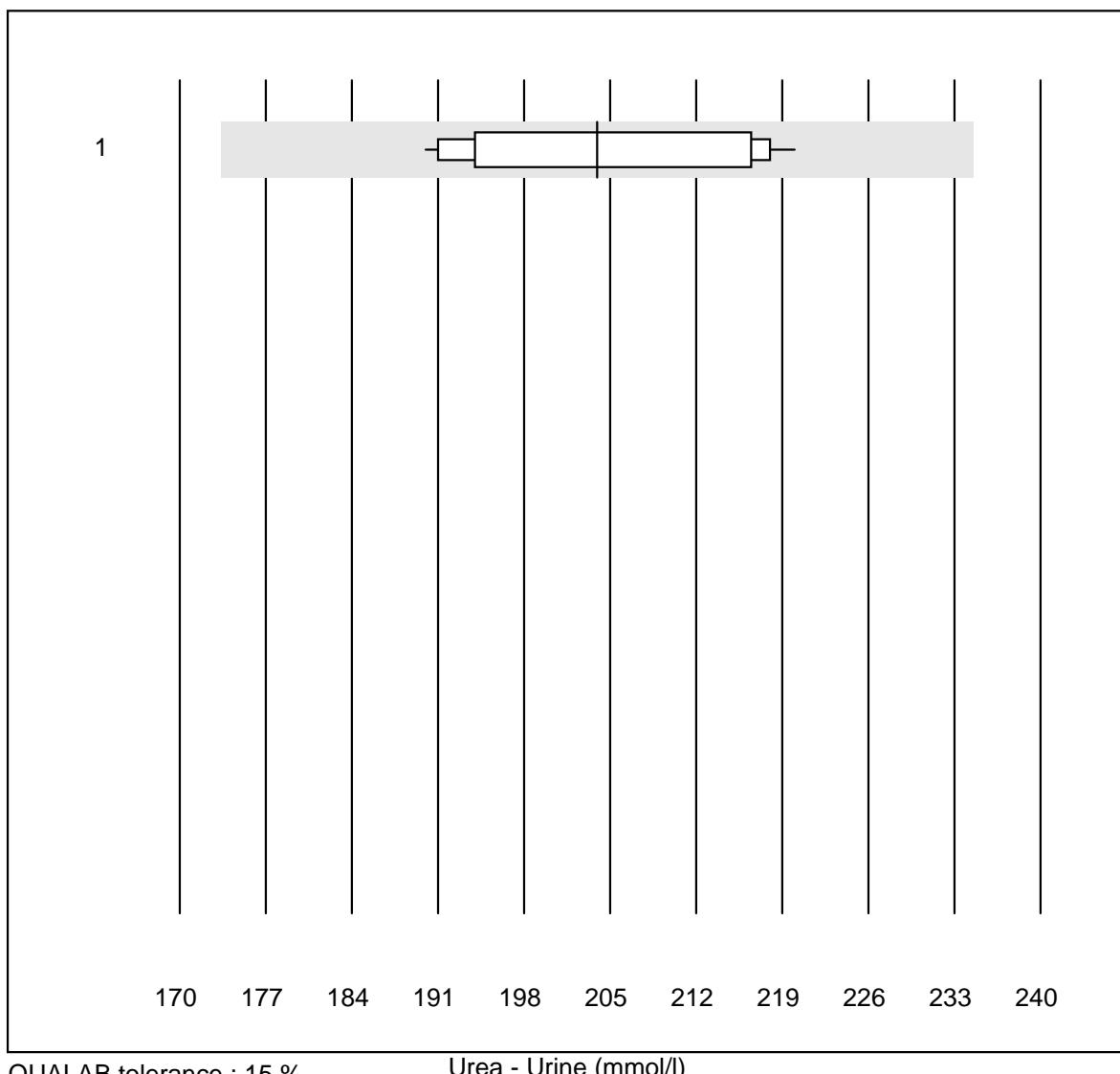
total Protein - Urine (mg/l)

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Cobas/Roche	12	91.7	8.3	0.0	341.6	8.5	e*
2 Other methods	4	100.0	0.0	0.0	425.5	9.6	e*

Sodium - Urine

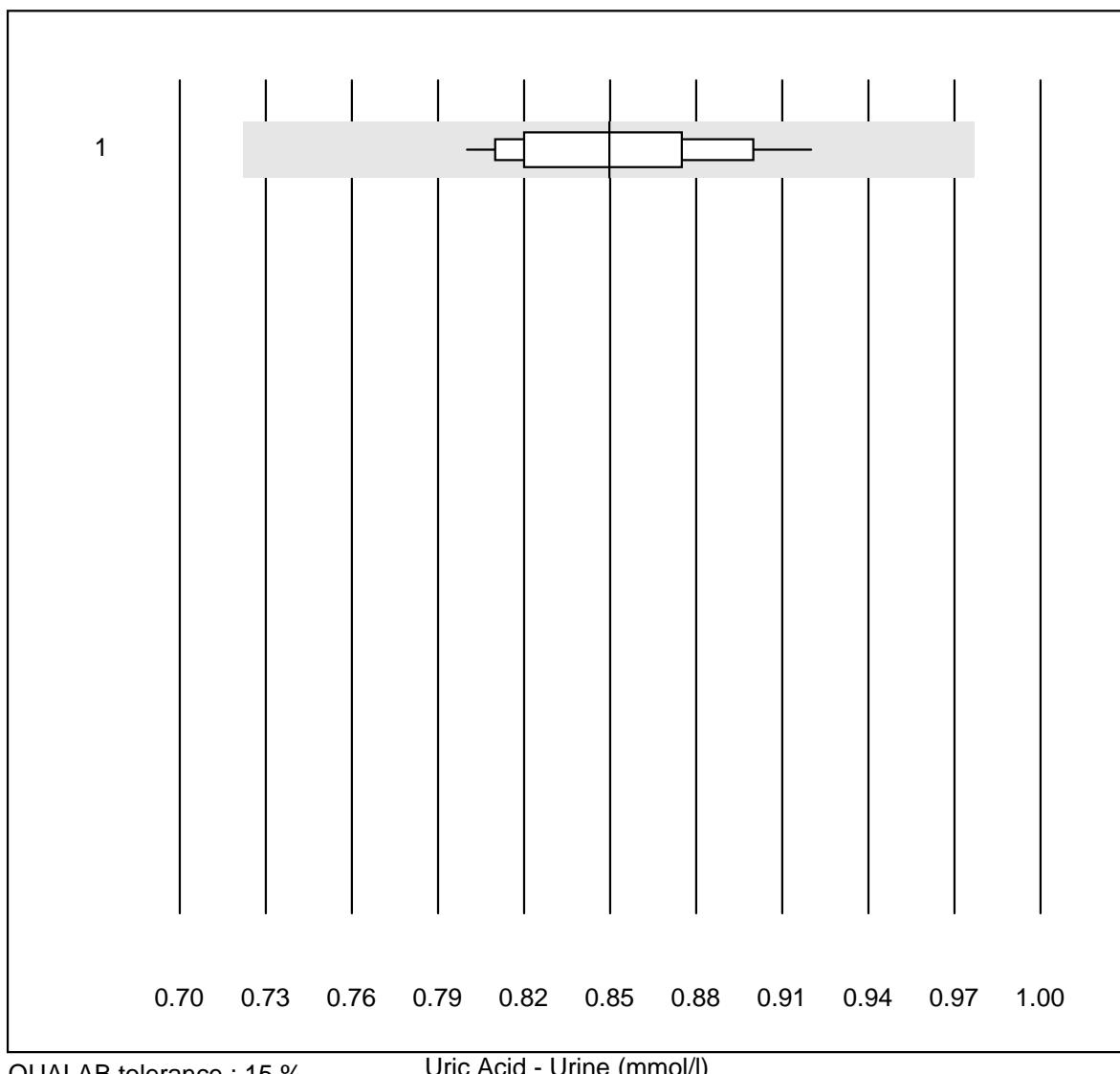


No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 all Participants	18	100.0	0.0	0.0	126	1.5	e

Urea - Urine

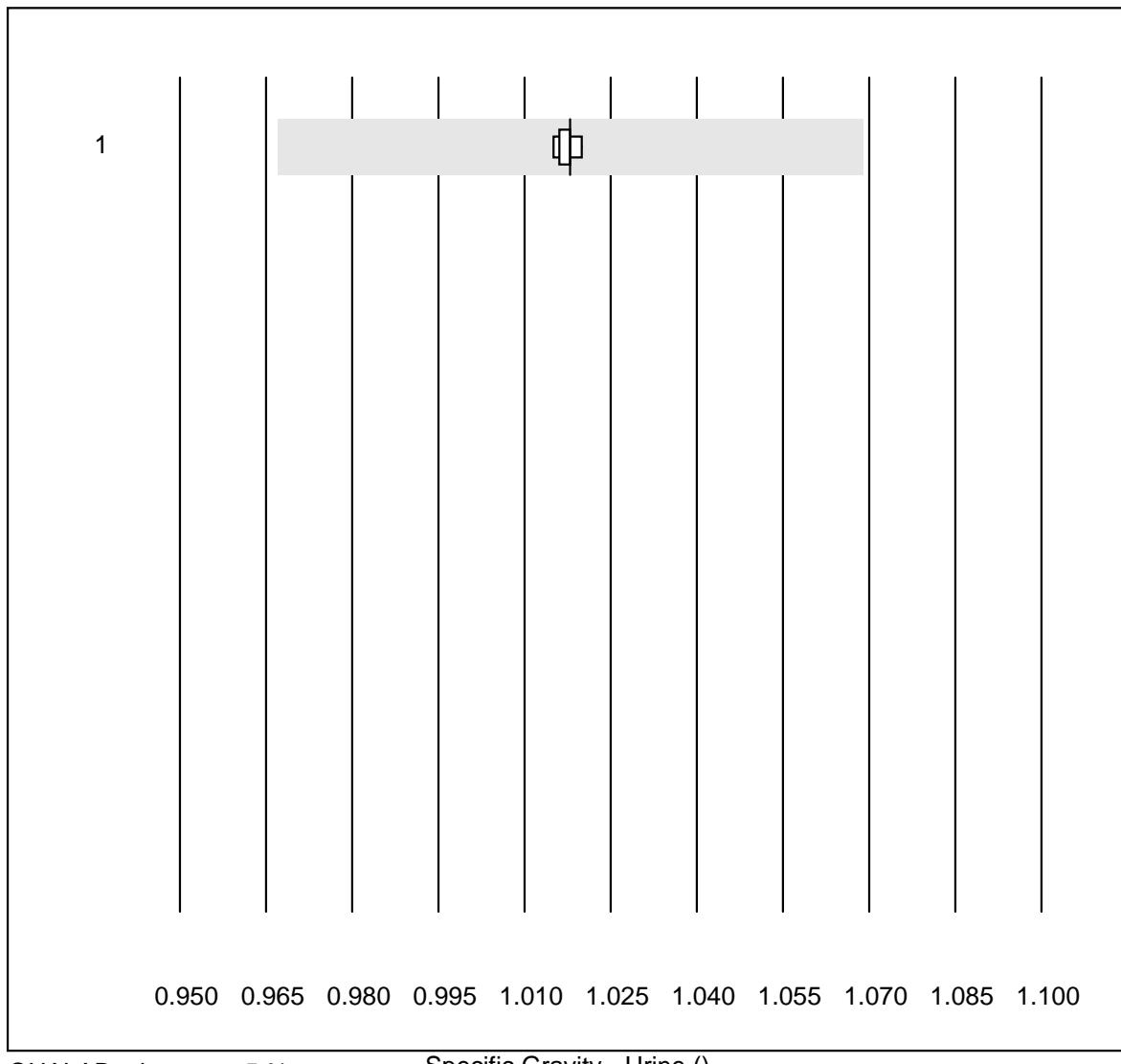
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Standard chemistry	14	100.0	0.0	0.0	204	5.3	e

Uric Acid - Urine



No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Standard chemistry	13	100.0	0.0	0.0	0.85	4.4	e

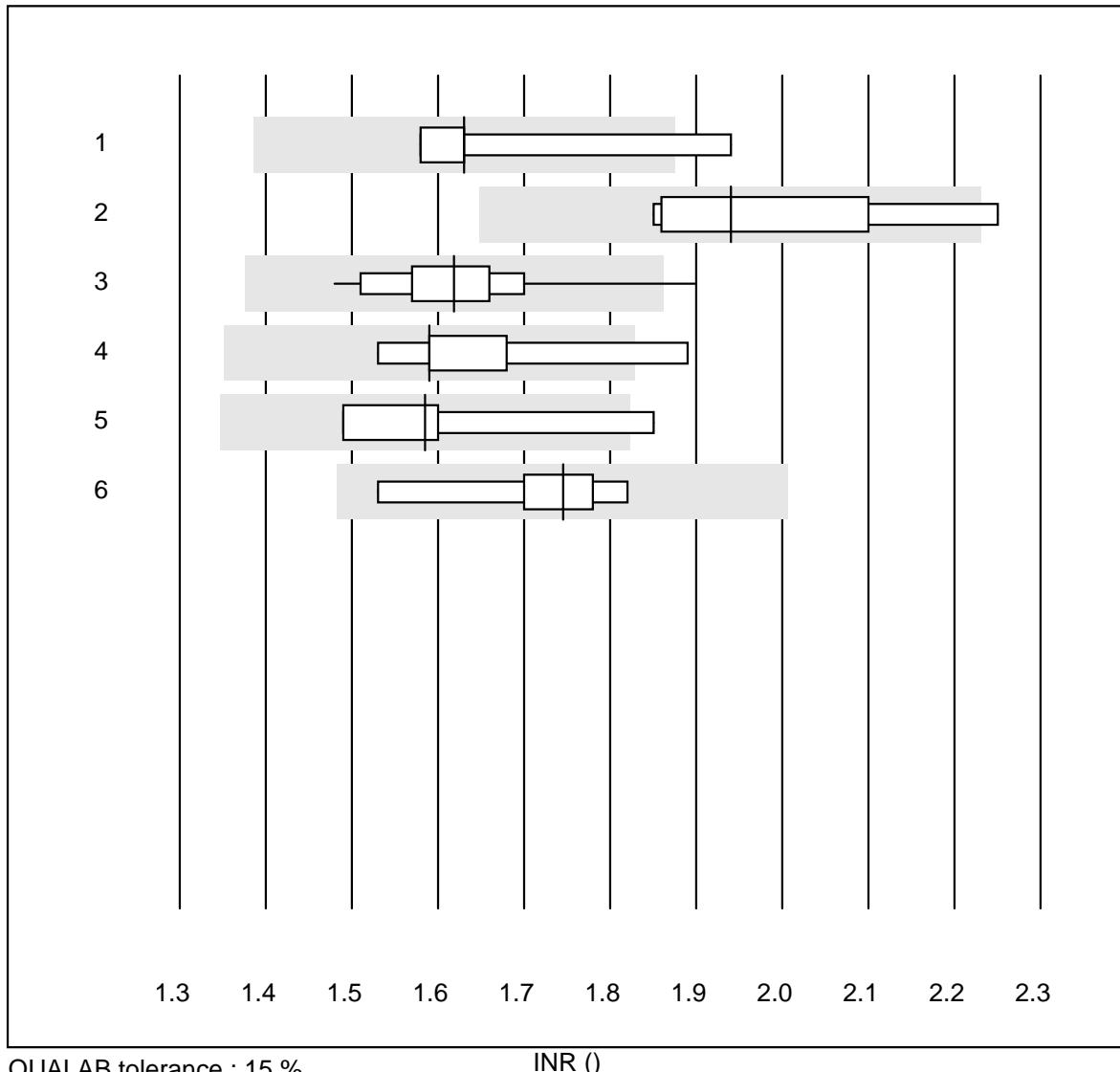
Specific Gravity - Urine



No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Refractometer	7	100.0	0.0	0.0	1.018	0.2	e

G1 Coagulation INR

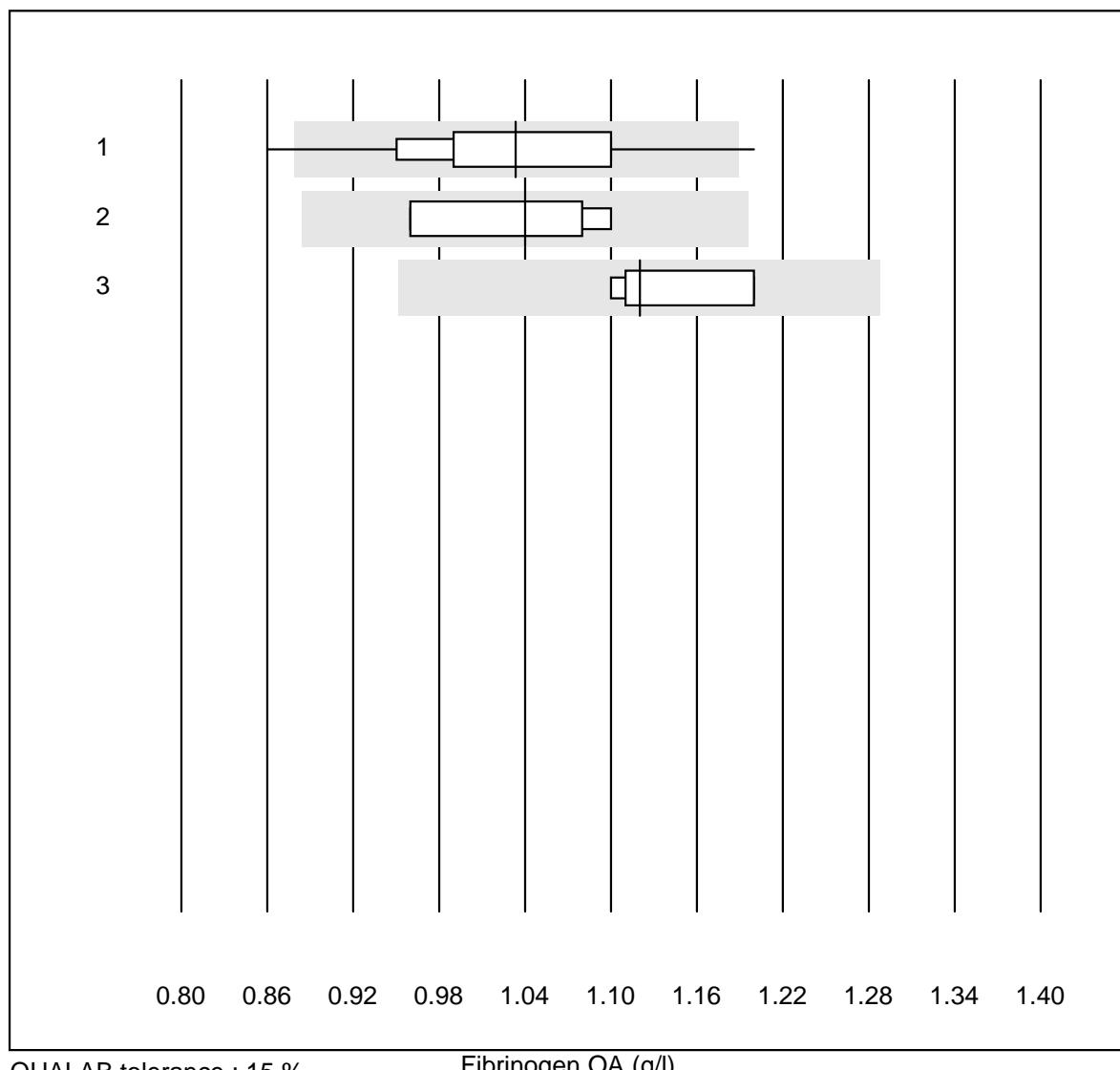
INR



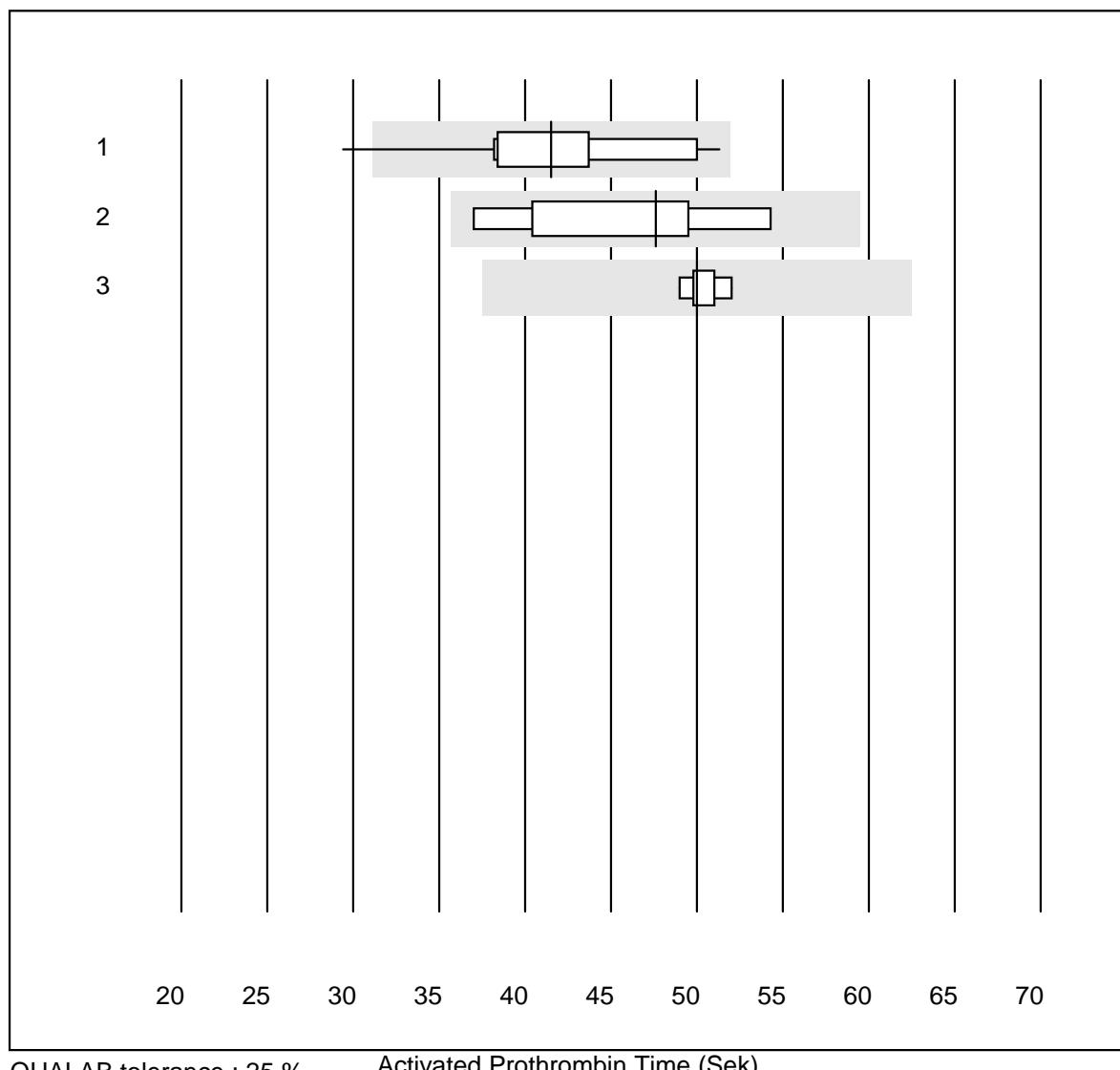
QUALAB tolerance : 15 %

INR ()

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Thromborel S	4	75.0	25.0	0.0	1.63	9.7	e*
2 Neoplastin Plus	5	80.0	20.0	0.0	1.94	8.6	e*
3 Innovin	17	94.1	5.9	0.0	1.62	6.0	e
4 Recombiplastin IL	5	80.0	20.0	0.0	1.59	8.5	e*
5 Other methods	4	75.0	25.0	0.0	1.59	9.6	e*
6 Neoplastin R	8	100.0	0.0	0.0	1.75	5.2	e*

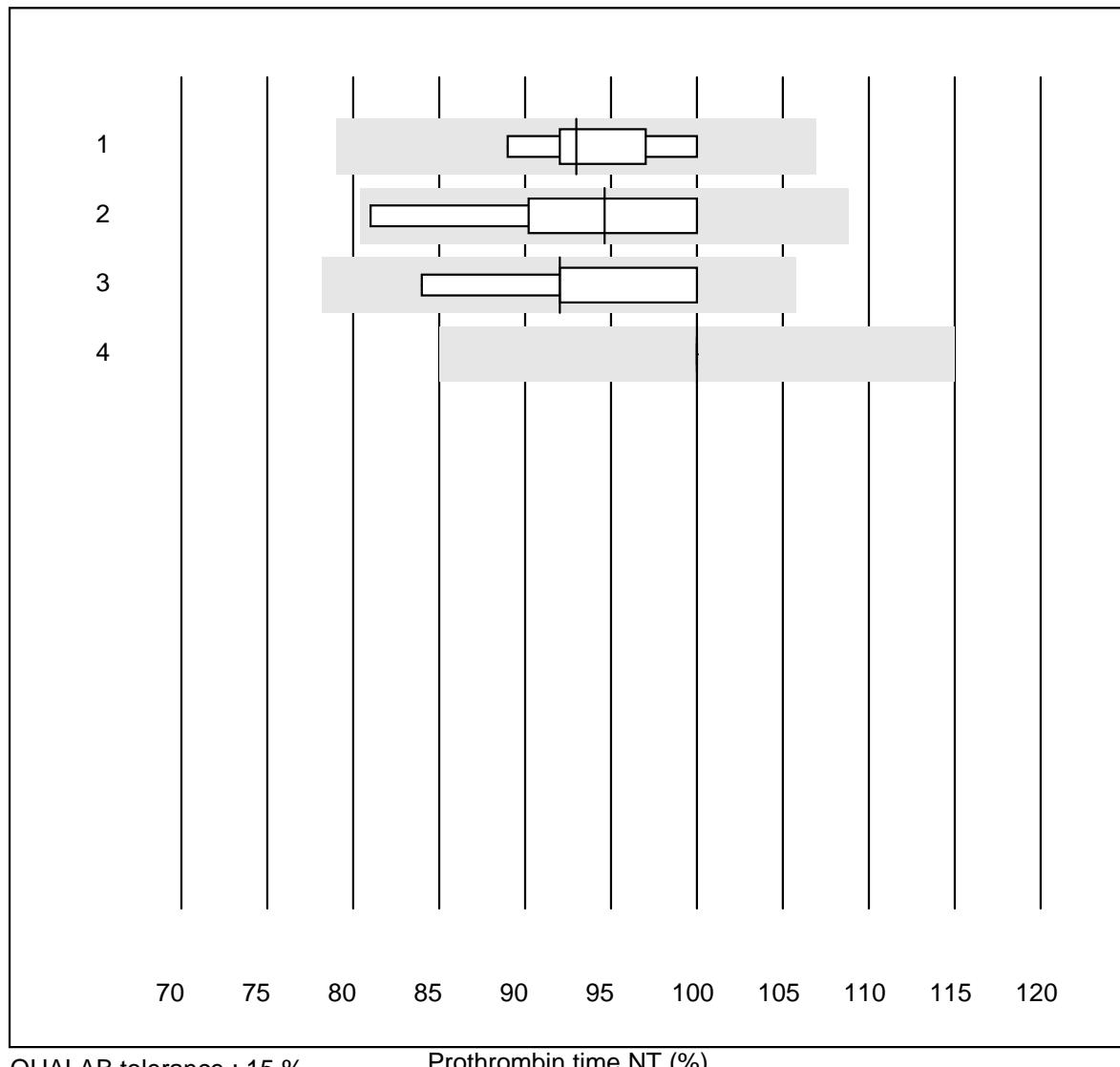
Fibrinogen OA

Activated Prothrombin Time

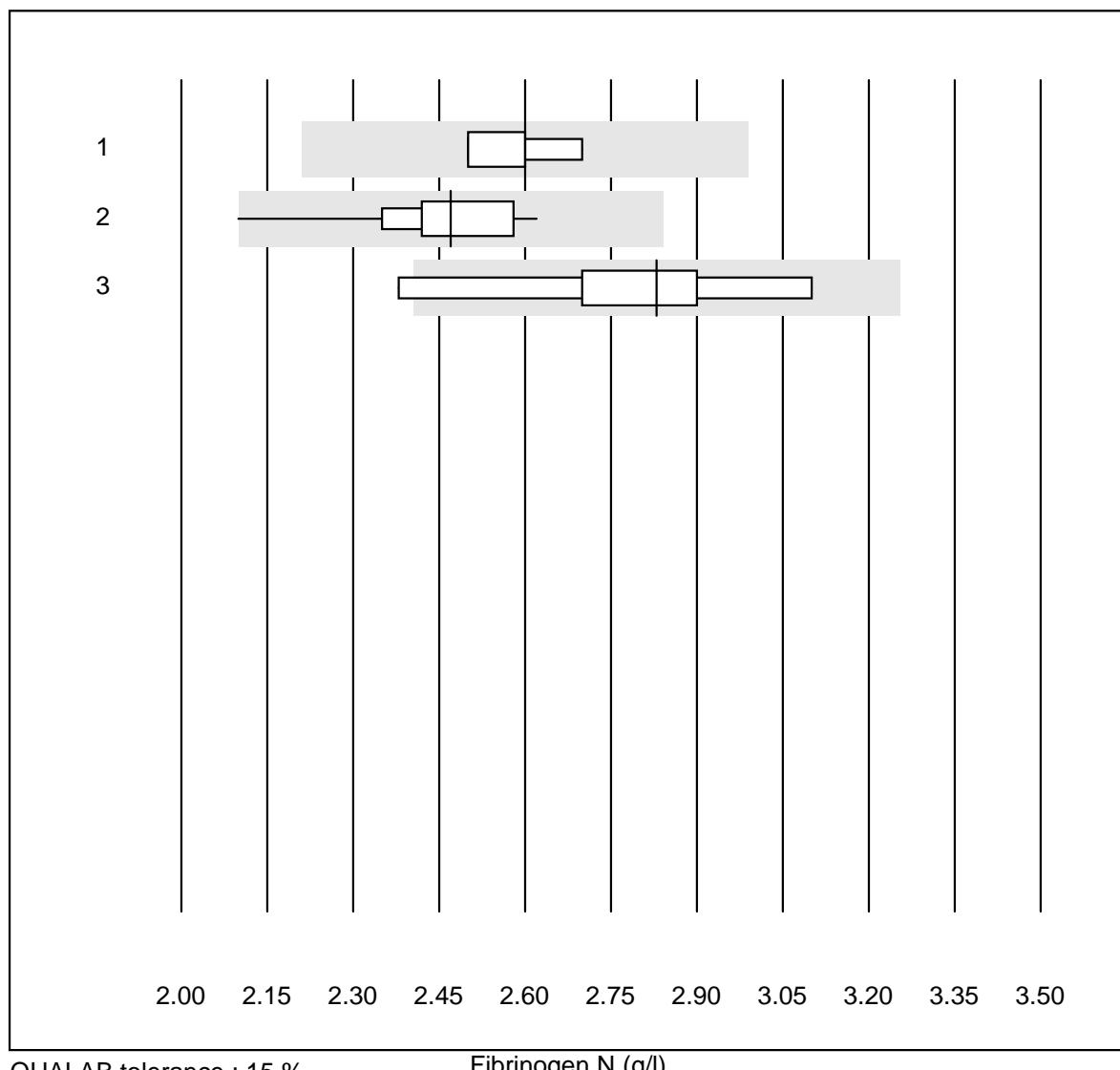


No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Other methods	14	85.8	7.1	7.1	41.5	13.4	e*
2 Actin FS	8	100.0	0.0	0.0	47.6	12.1	e*
3 Stago/STA	6	100.0	0.0	0.0	50.0	2.1	e

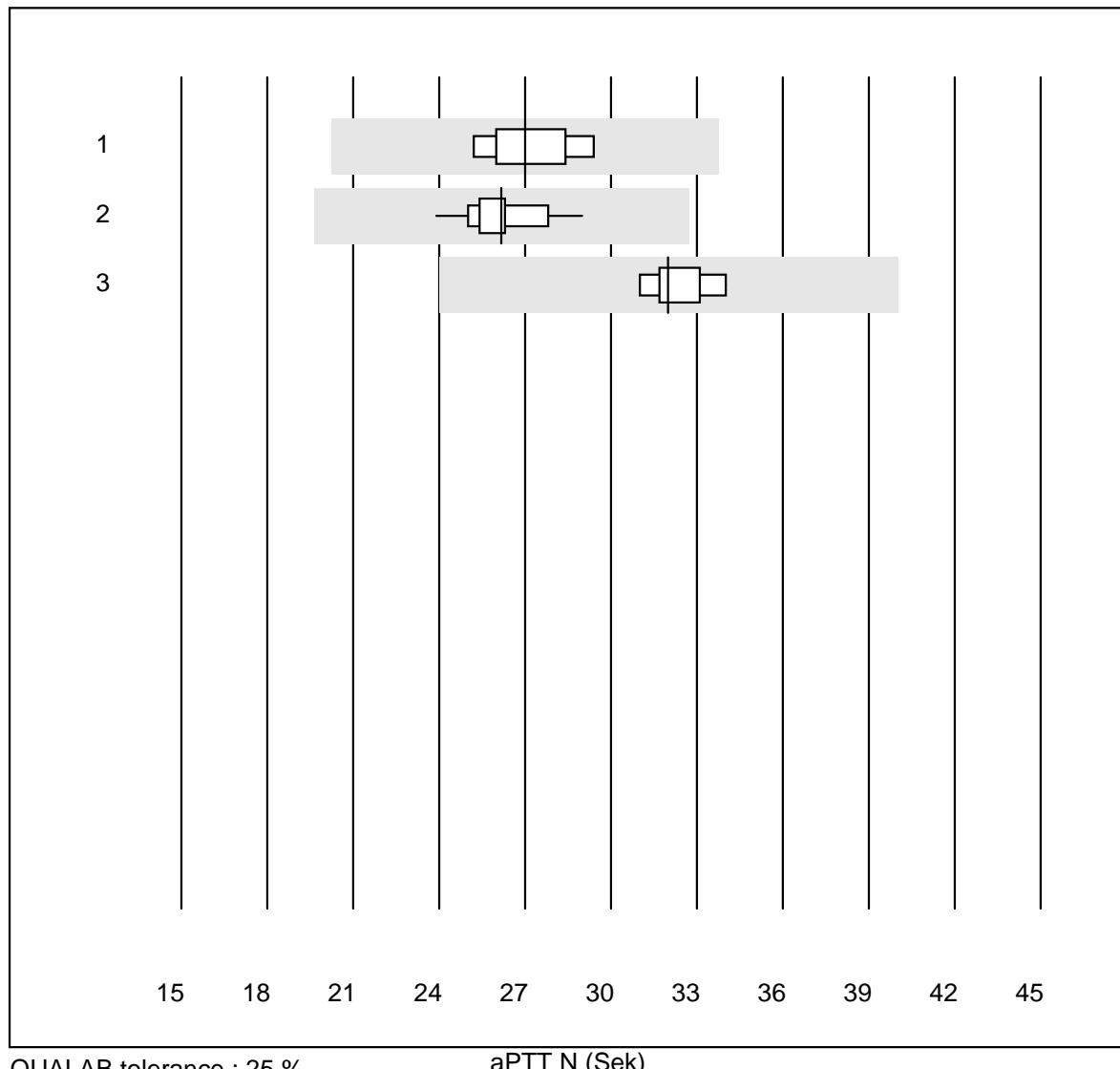
Prothrombin time NT



No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Neoplastin R	5	100.0	0.0	0.0	93	4.6	e*
2 Innovin	10	100.0	0.0	0.0	95	6.8	e*
3 all Participants	5	100.0	0.0	0.0	92	7.2	e*
4 Recombiplastin IL	6	100.0	0.0	0.0	100	0.0	e

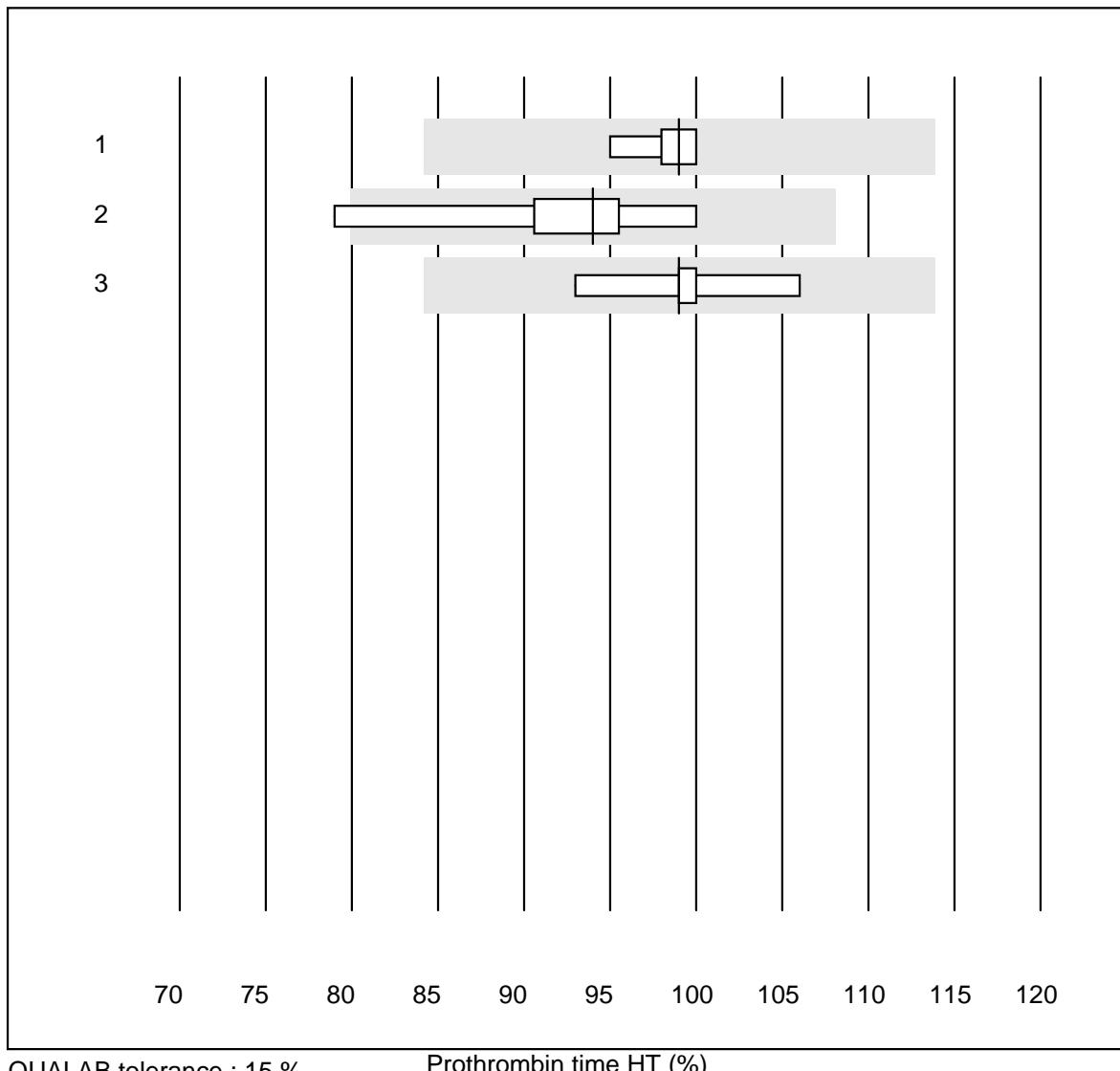
Fibrinogen N

No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Siemens Thrombin	4	100.0	0.0	0.0	2.60	3.1	e
2	Other methods	12	100.0	0.0	0.0	2.47	5.7	e
3	Stago/STA	8	87.5	12.5	0.0	2.83	7.4	e*

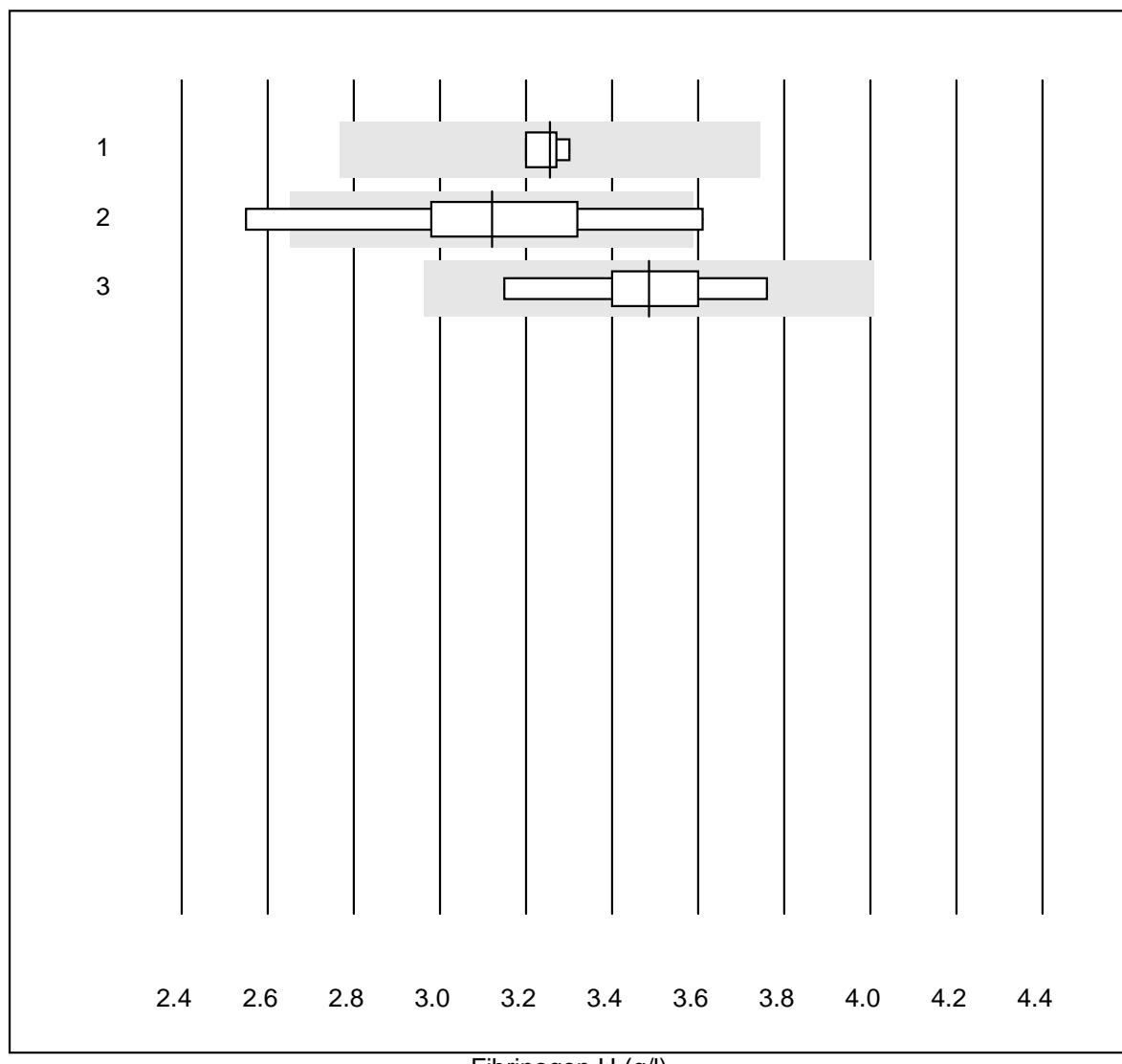
aPTT N

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Actin FS	9	88.9	0.0	11.1	27.0	5.3	e
2 Other methods	12	100.0	0.0	0.0	26.2	5.0	e
3 Stago/STA	7	100.0	0.0	0.0	32.0	3.2	e

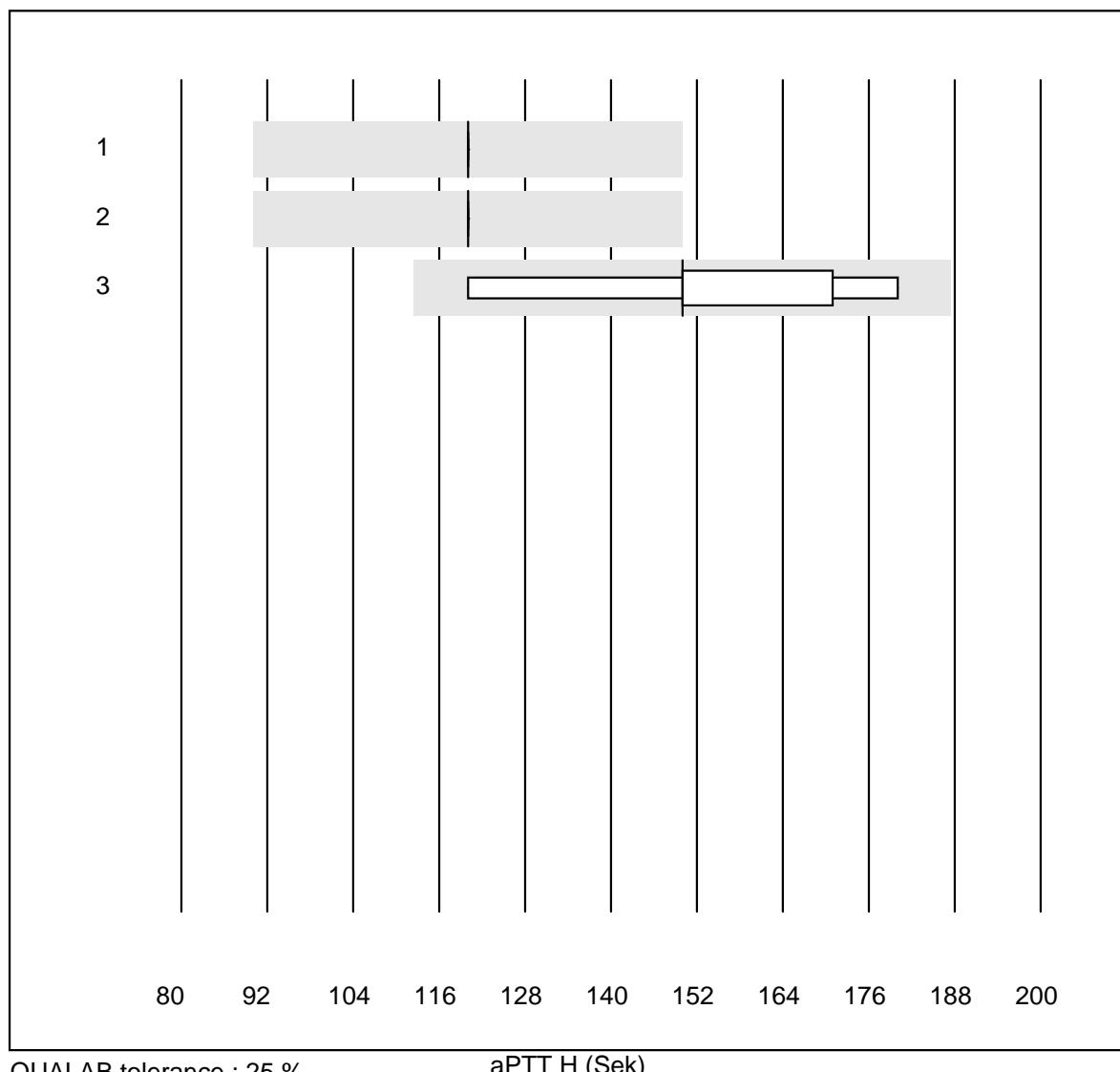
Prothrombin time HT



Fibrinogen H



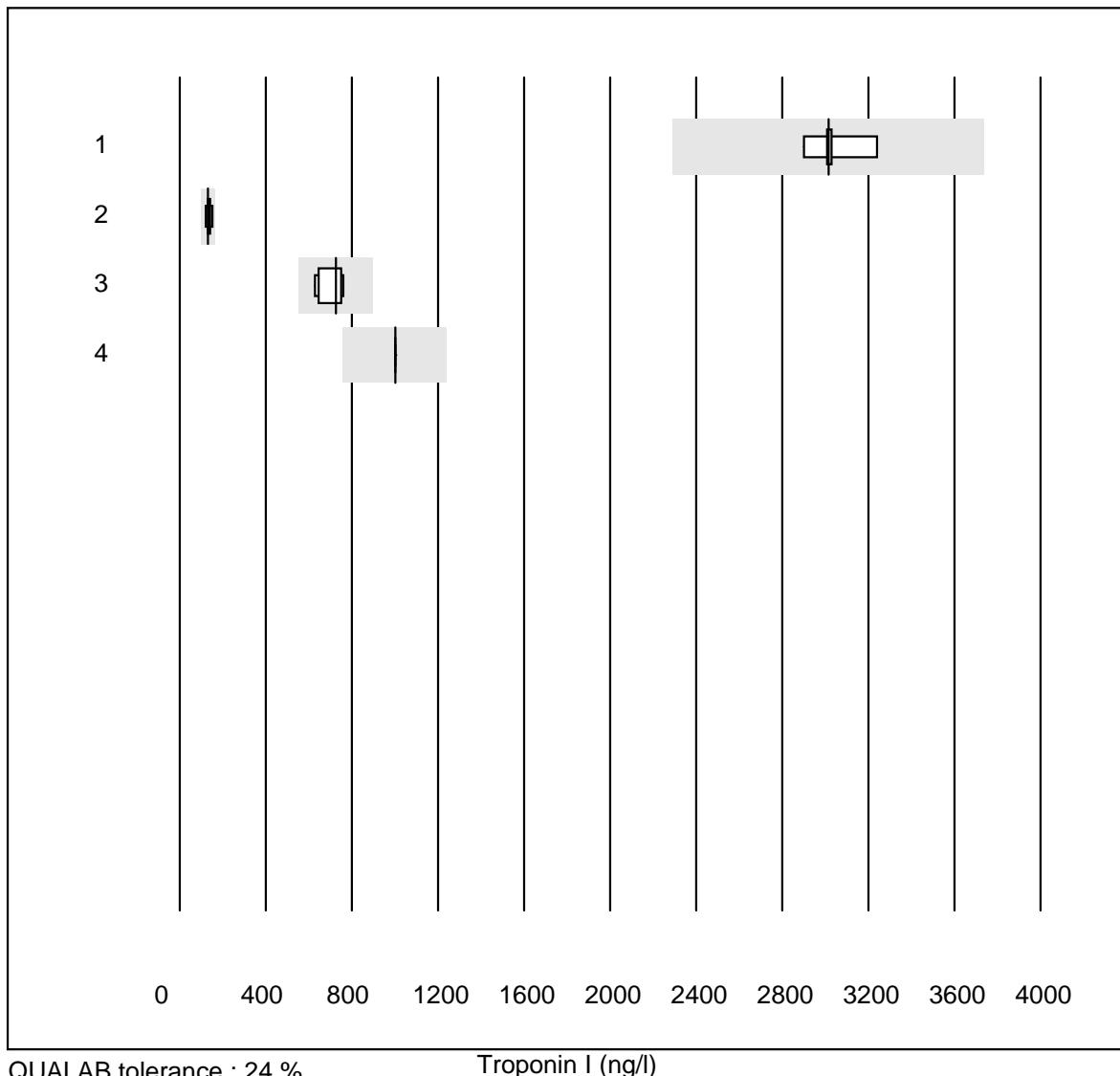
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Siemens Thrombin	4	100.0	0.0	0.0	3.26	1.3	e
2	Other methods	10	70.0	30.0	0.0	3.12	11.3	e*
3	Stago/STA	6	100.0	0.0	0.0	3.49	6.1	e*

aPTT H

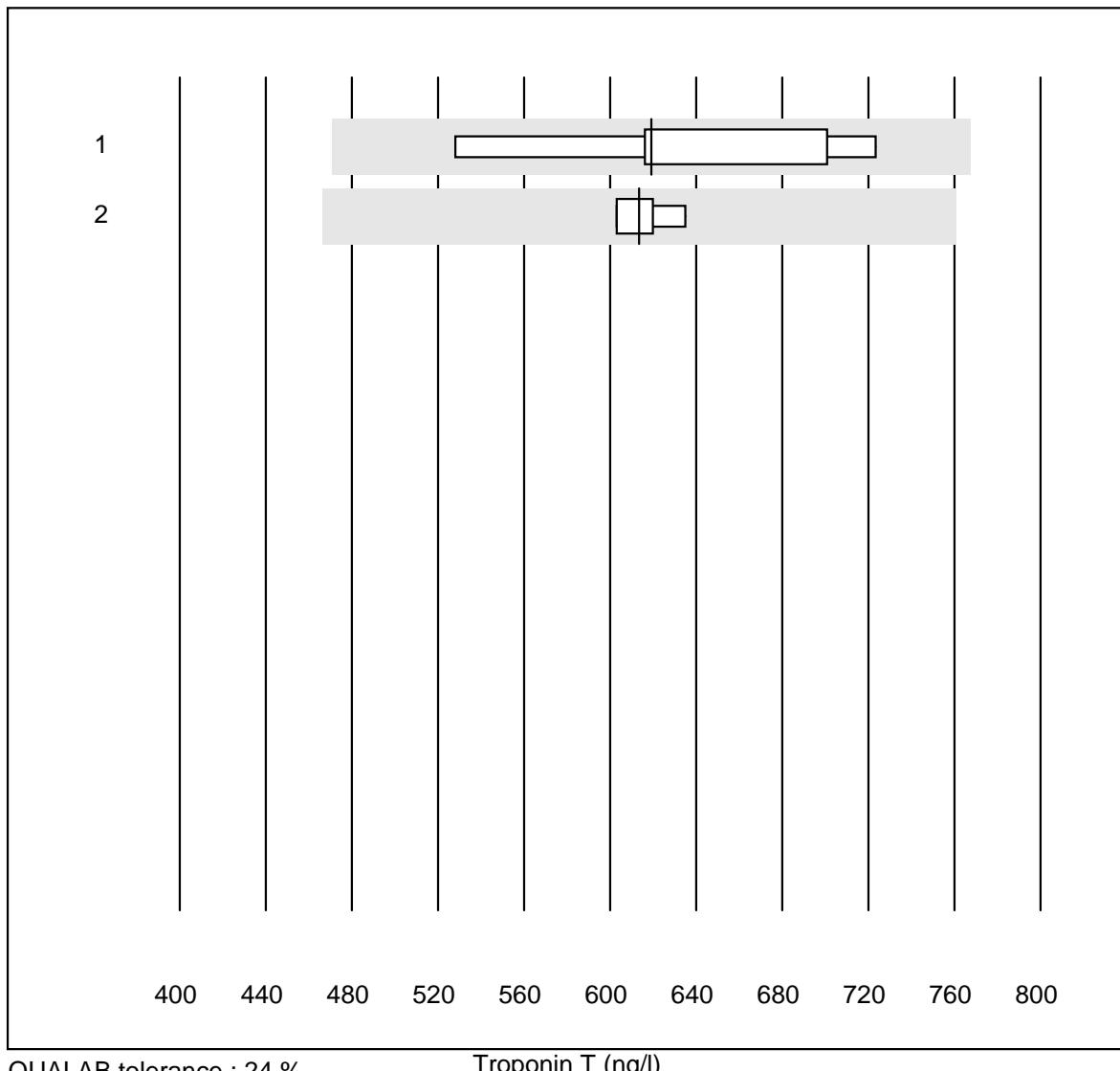
QUALAB tolerance : 25 %

aPTT H (Sek)

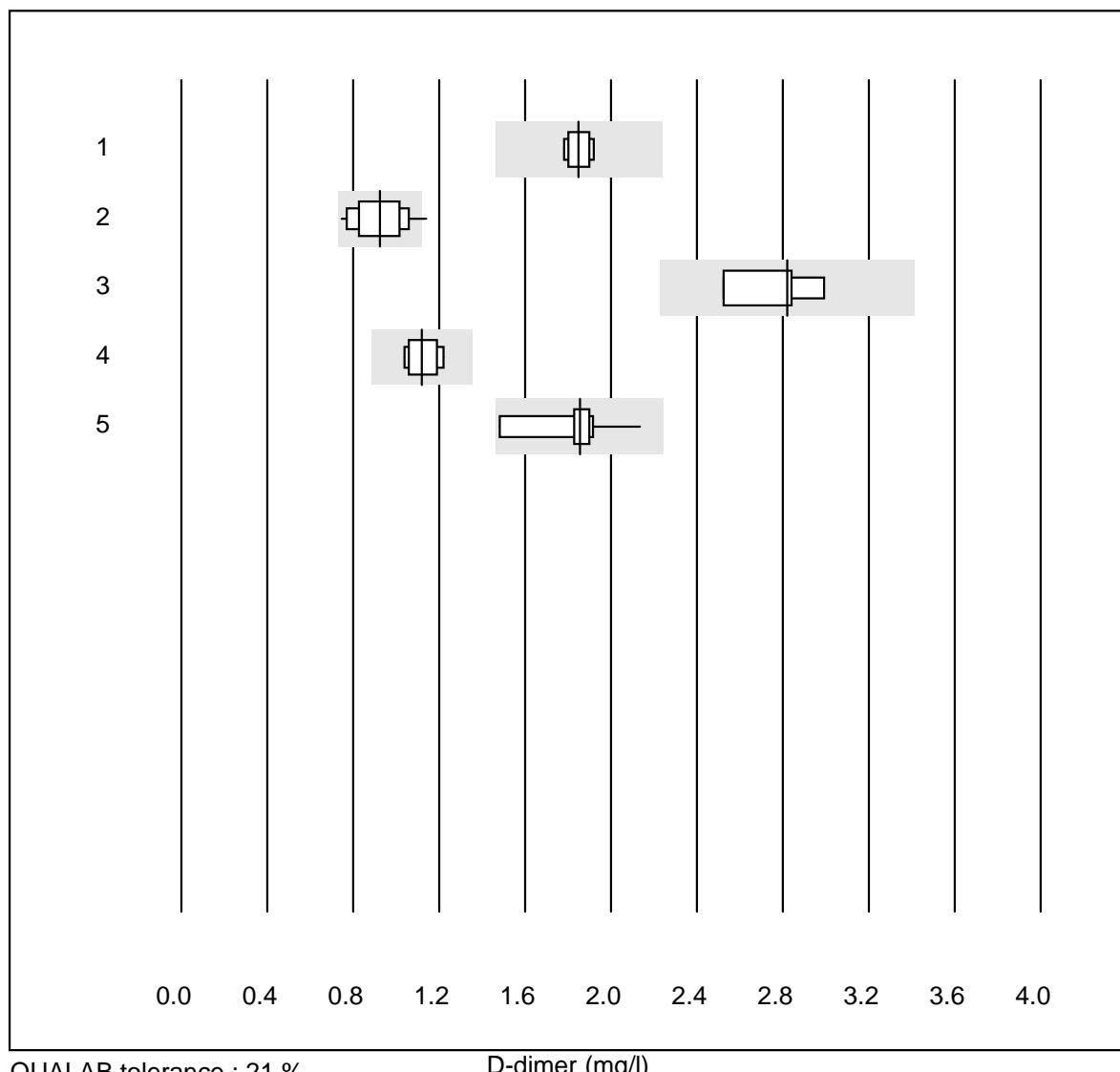
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Actin FS	7	100.0	0.0	0.0	120.0	0.0	e
2 Other methods	9	100.0	0.0	0.0	120.0	0.0	e
3 Stago/STA	5	100.0	0.0	0.0	150.0	15.0	e*

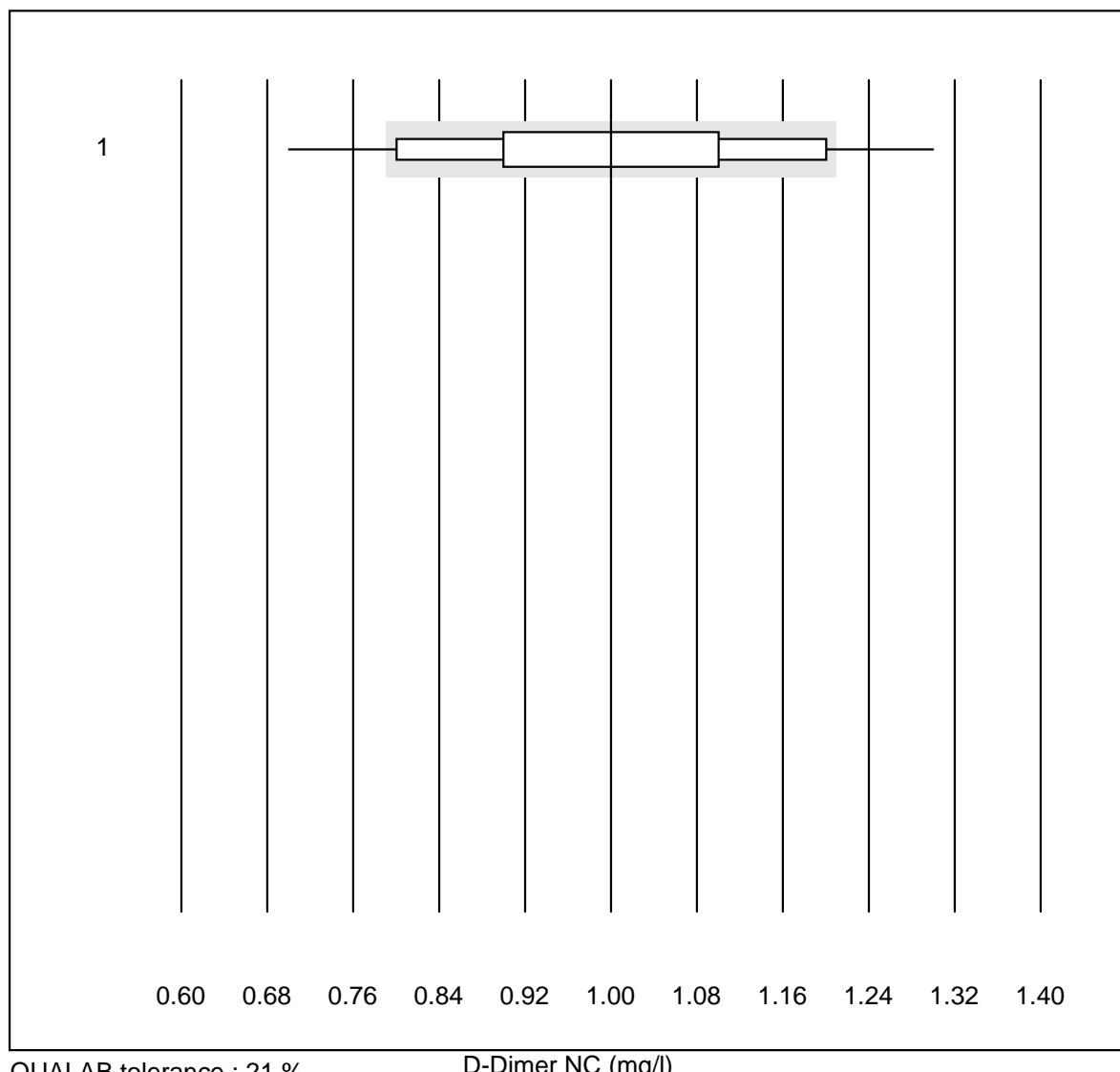
Troponin I

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Vidas	6	83.3	0.0	16.7	3015.0	4.1	e
2 AQT 90 FLEX	6	100.0	0.0	0.0	130.0	7.7	e*
3 ADVIA Centaur XP/CP	6	100.0	0.0	0.0	725.0	7.9	e*
4 Eurolyser	12	100.0	0.0	0.0	1000.0	0.0	e

Troponin T

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Cobas hs	5	100.0	0.0	0.0	619.00	12.2	e*
2 Cobas hs STAT	4	100.0	0.0	0.0	613.40	2.3	e

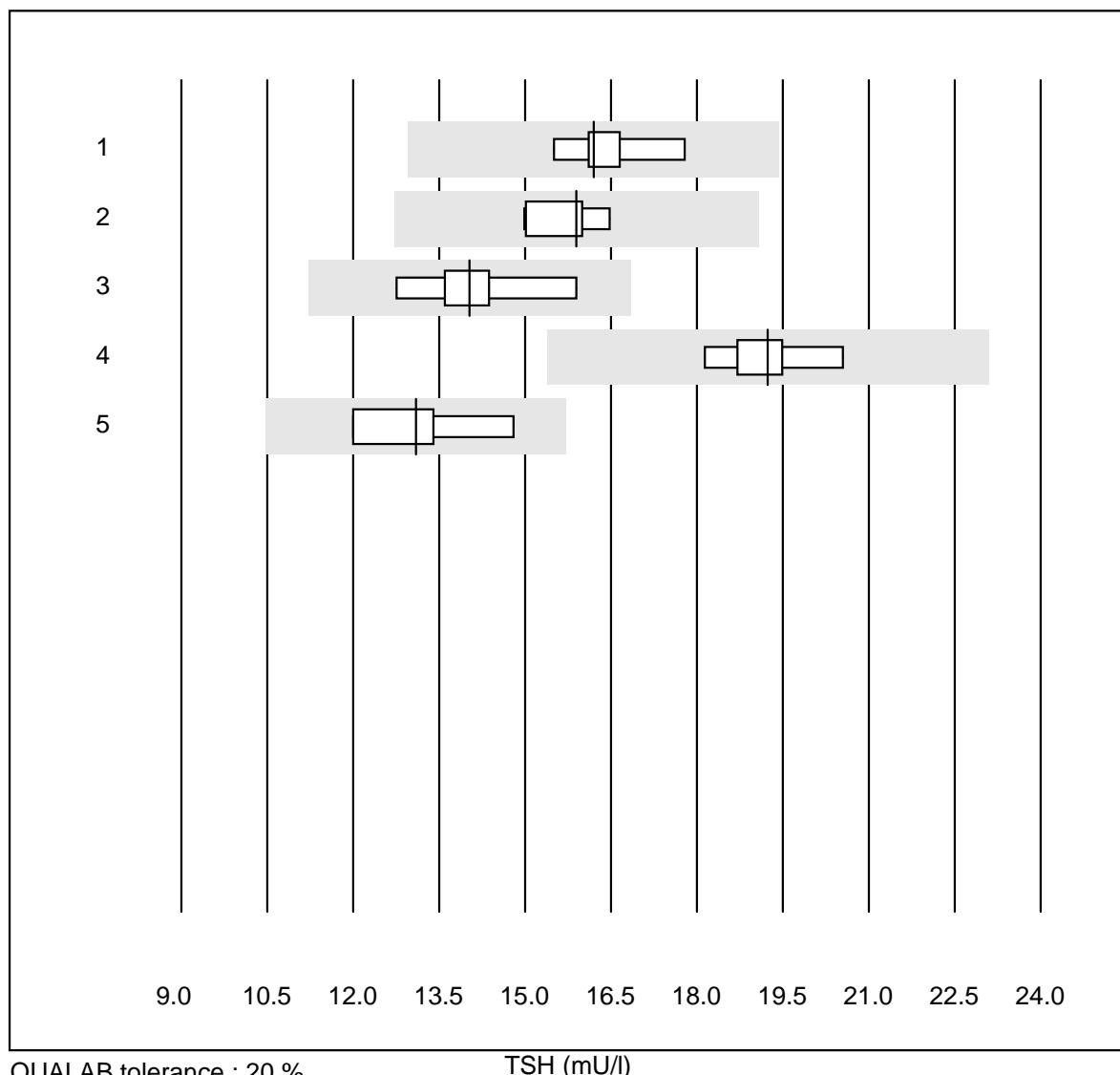
D-dimer

D-Dimer NC

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 NycoCard	37	78.4	10.8	10.8	1.00	16.5	e*

K6 Hormones

TSH



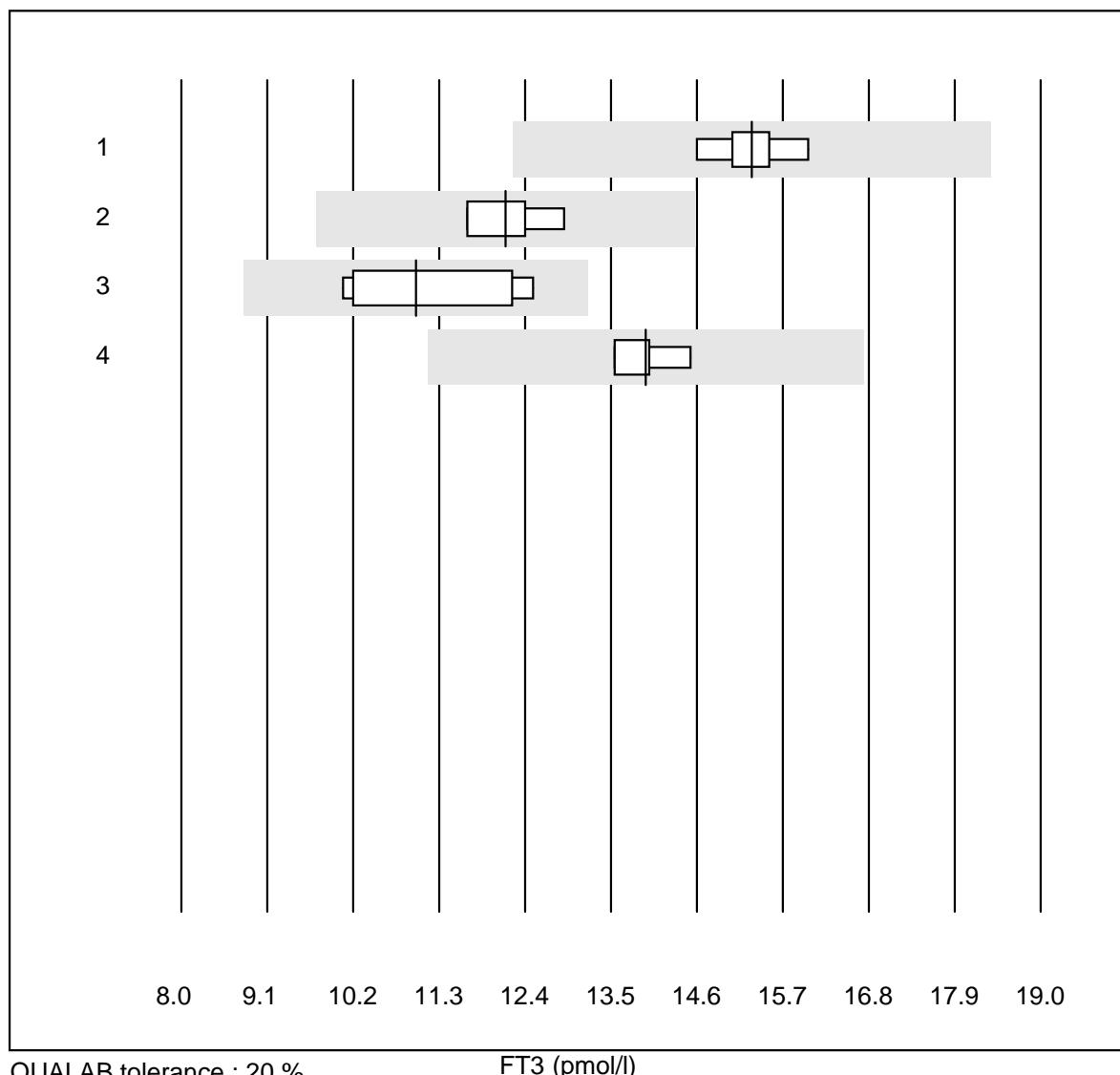
QUALAB tolerance : 20 %

TSH (mU/l)

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Cobas E / Elecsys	9	100.0	0.0	0.0	16.2	4.1	e
2 ADVIA Centaur XP/CP	5	100.0	0.0	0.0	15.9	4.2	e
3 Architect	8	100.0	0.0	0.0	14.0	6.6	e
4 Vidas	9	100.0	0.0	0.0	19.2	4.2	e
5 Qualigen	4	100.0	0.0	0.0	13.1	8.9	e*

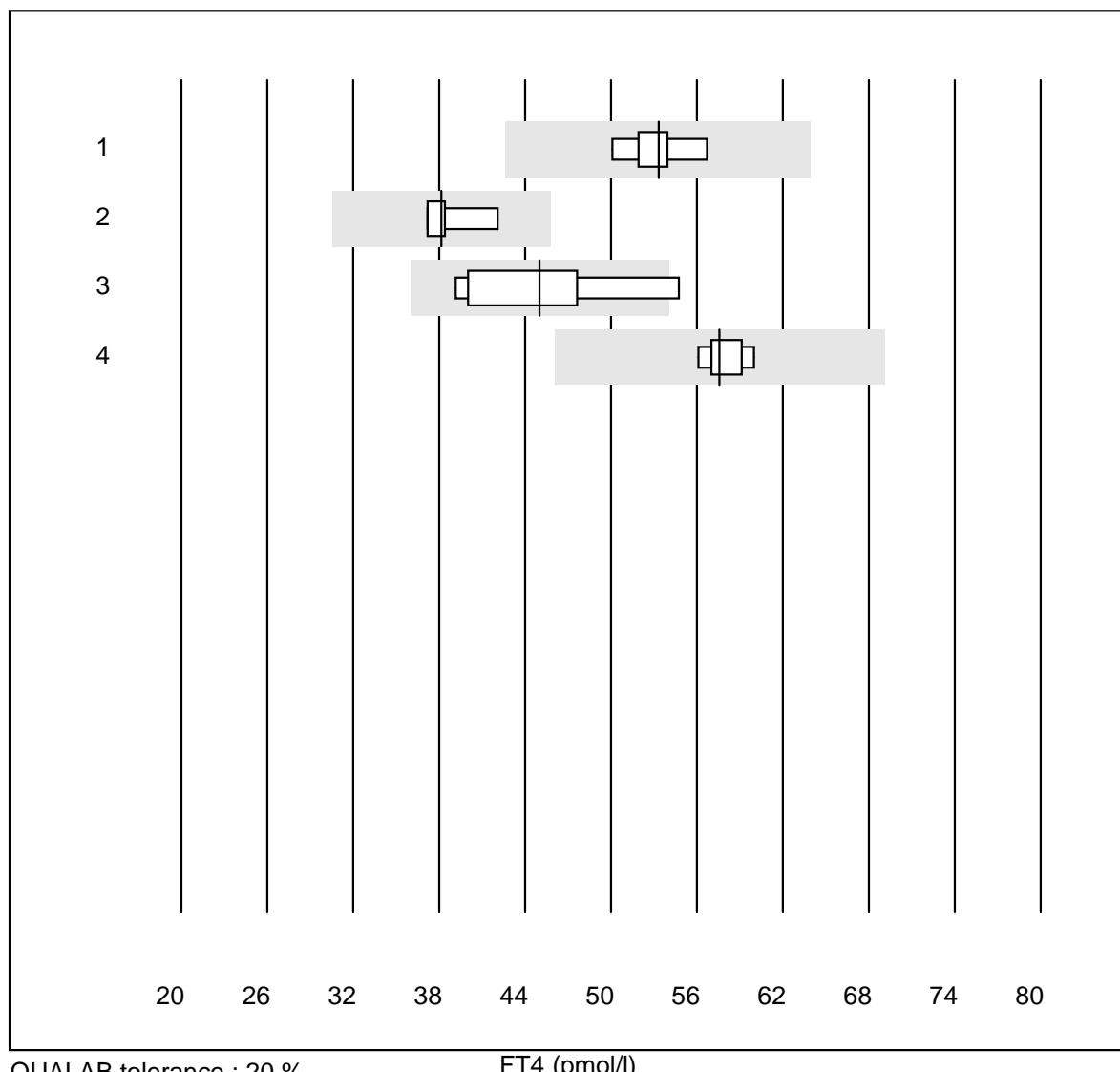
K6 Hormones

FT3

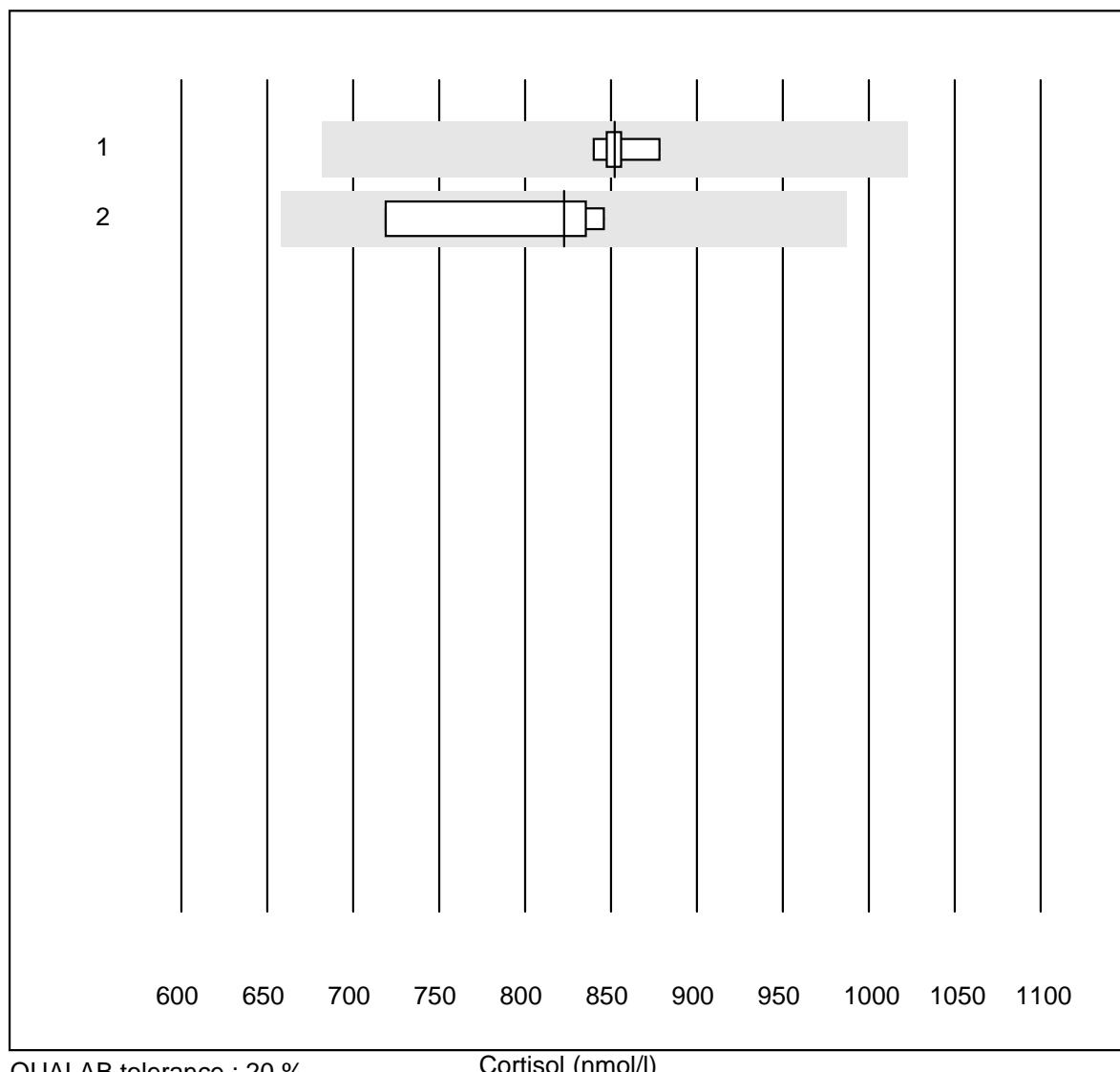


K6 Hormones

FT4



No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Cobas E / Elecsys	10	100.0	0.0	0.0	53.3	4.0	e
2	ADVIA Centaur XP	4	100.0	0.0	0.0	38.2	5.6	e*
3	Architect	8	75.0	12.5	12.5	45.0	11.8	e*
4	Vidas	6	100.0	0.0	0.0	57.6	2.5	e

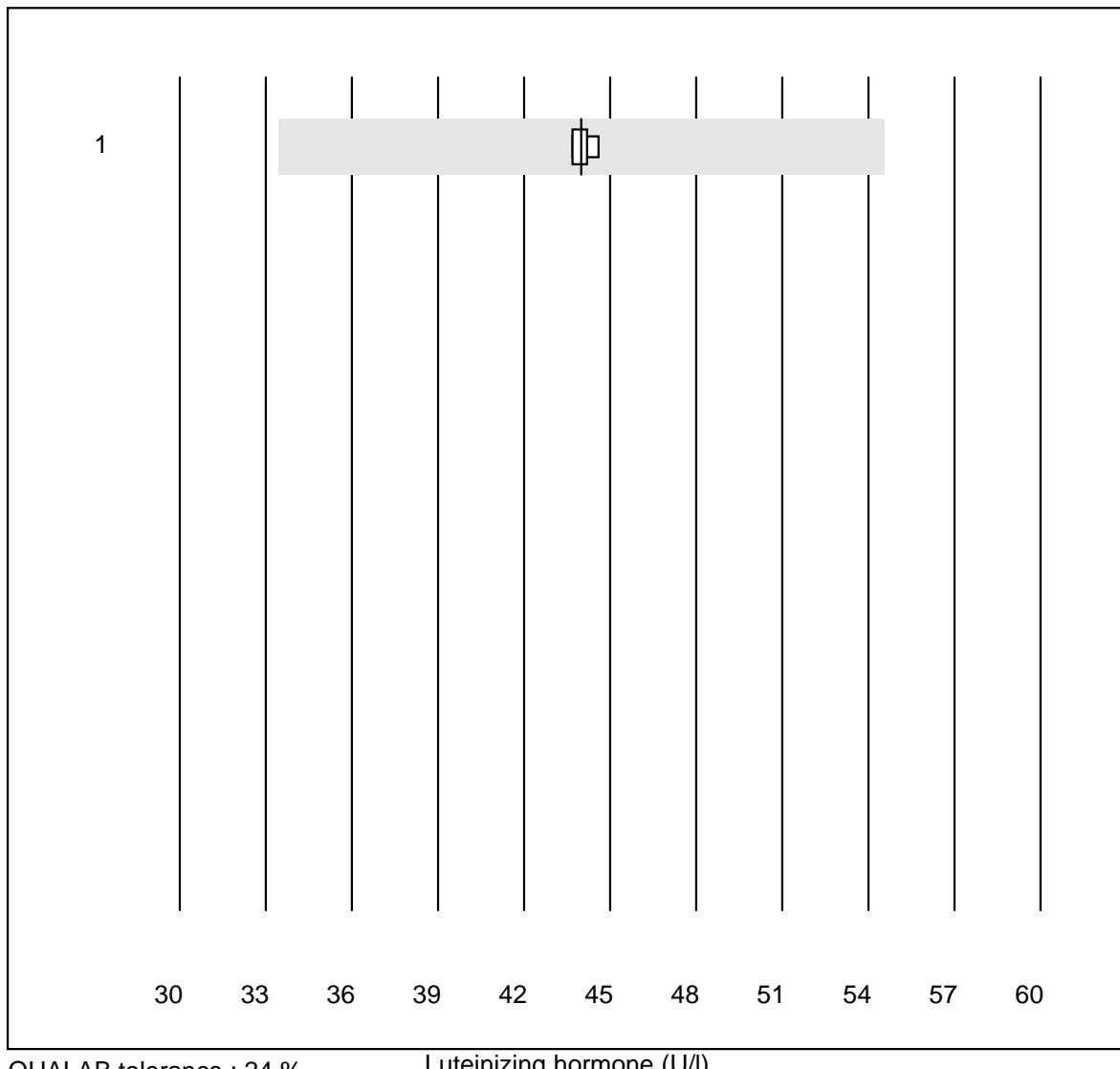
Cortisol

QUALAB tolerance : 20 %

Cortisol (nmol/l)

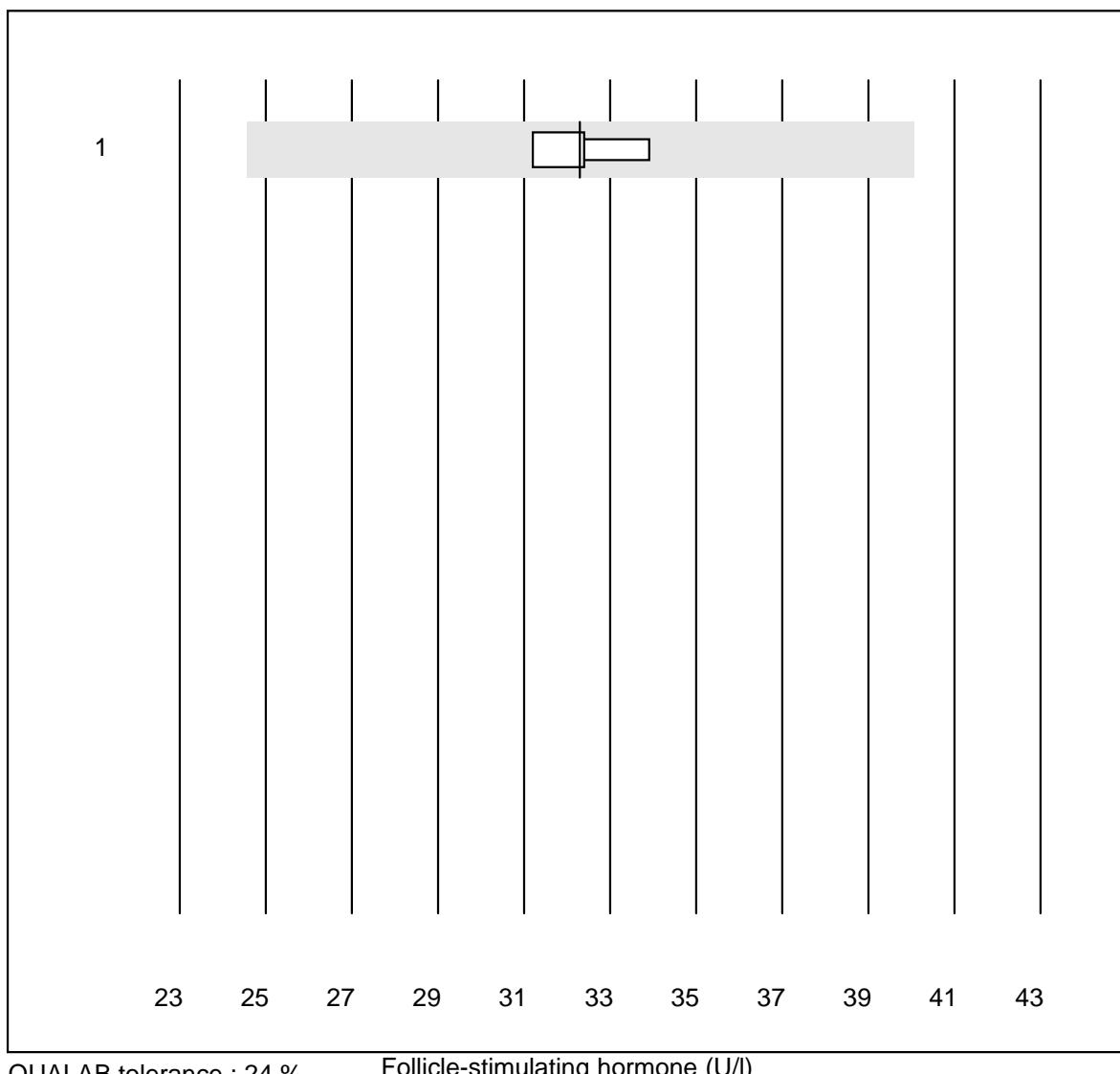
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Cobas E / Elecsys	6	100.0	0.0	0.0	852	1.5	e
2 ADVIA Centaur XP/CP	4	100.0	0.0	0.0	823	7.2	e*

Luteinizing hormone



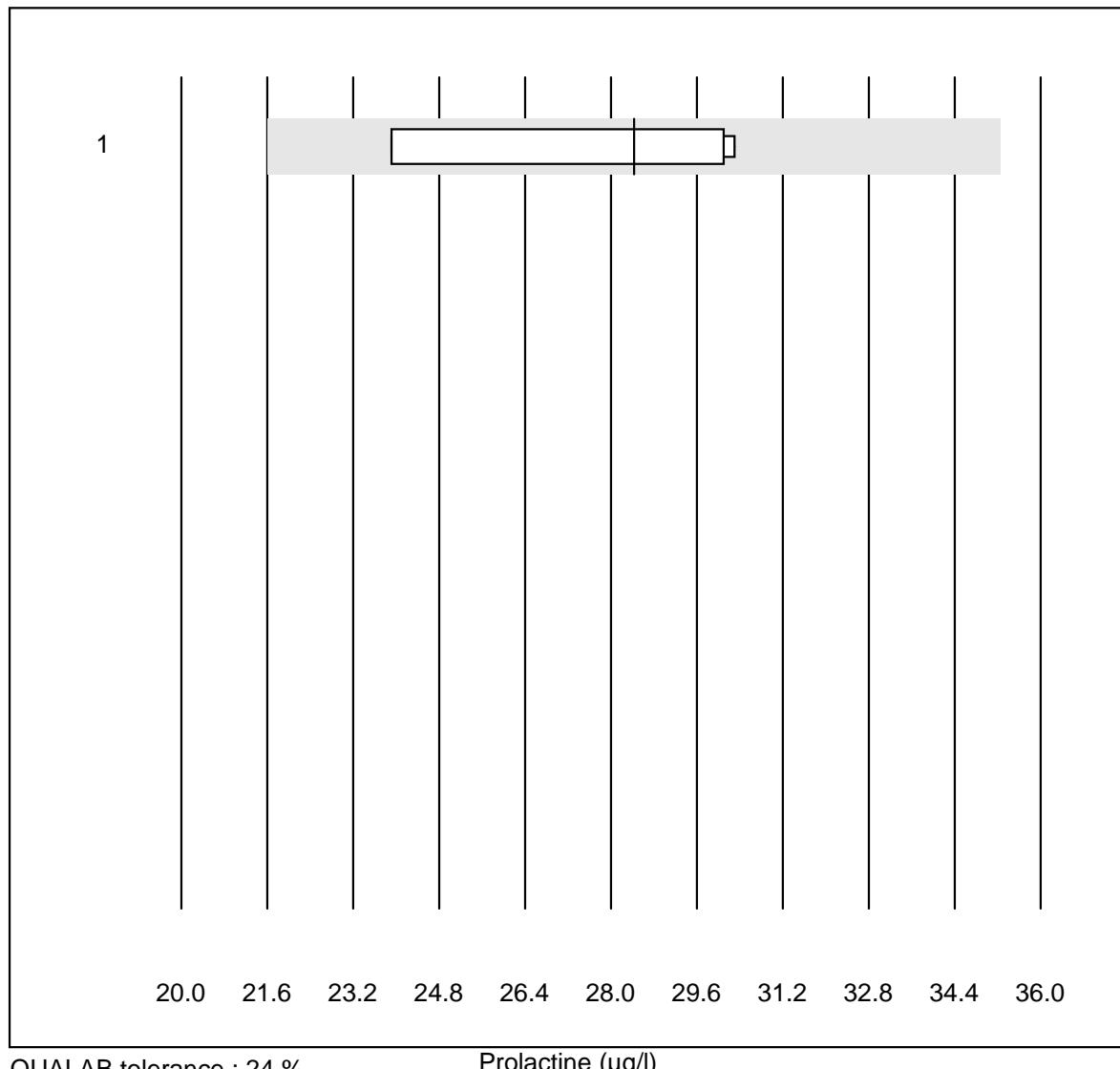
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	ADVIA Centaur XP/CP	4	100.0	0.0	0.0	44.0	0.9	e

Follicle-stimulating hormone



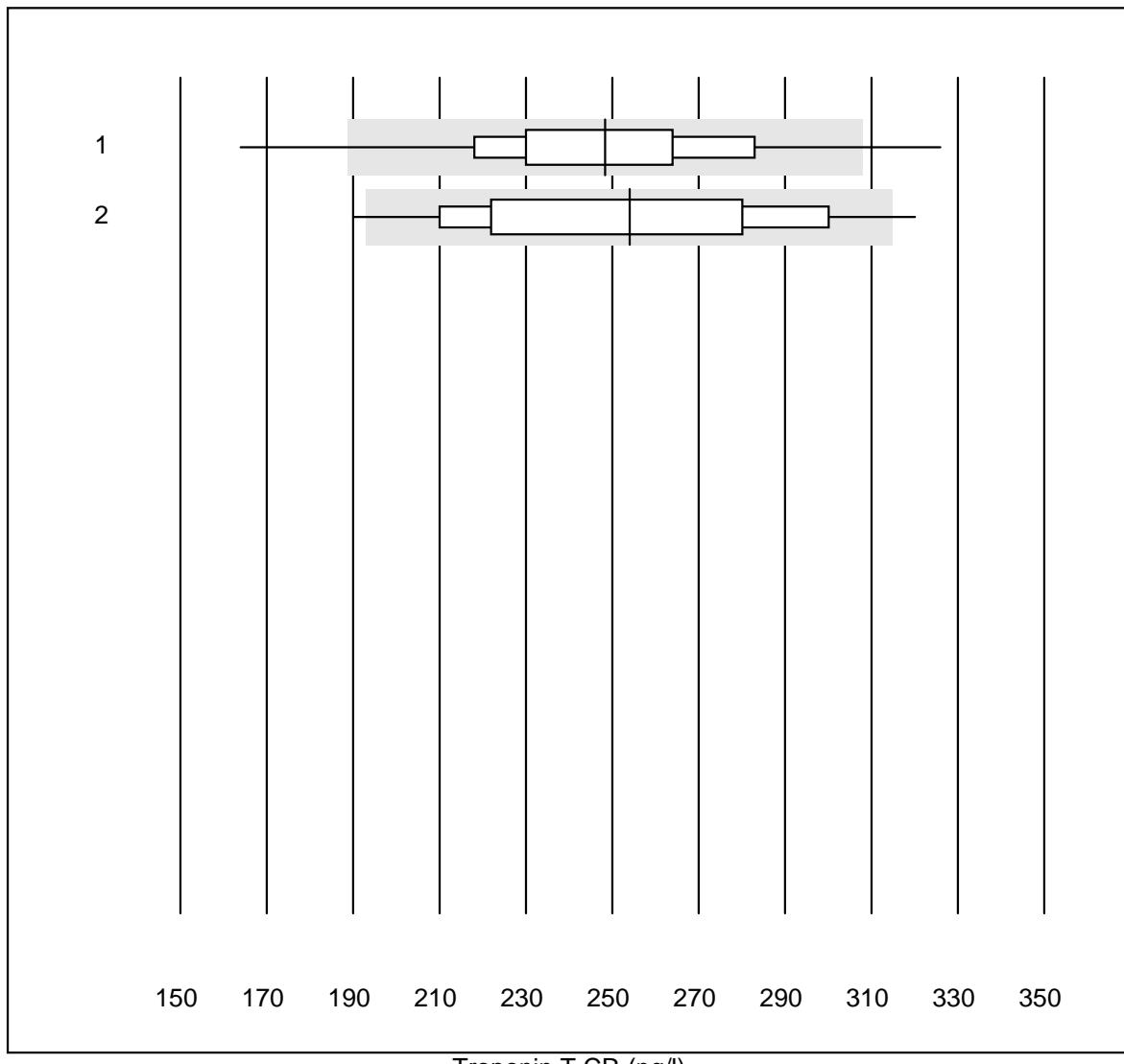
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	ADVIA Centaur XP/CP	4	100.0	0.0	0.0	32.3	3.4	e

Prolactine



No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	ADVIA Centaur XP/CP	4	100.0	0.0	0.0	28.4	11.0	e*

Troponin T CR

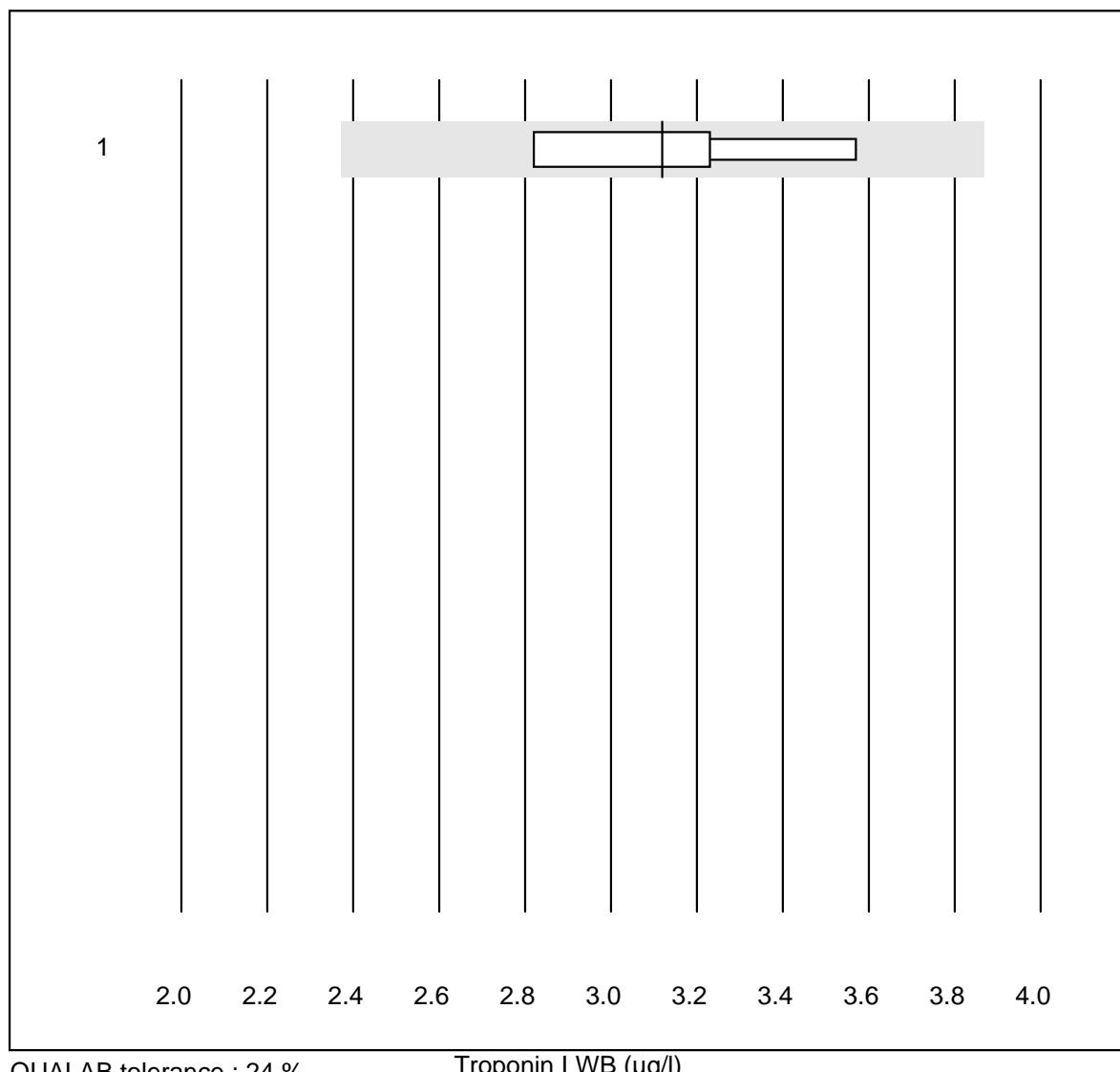


QUALAB tolerance : 24 %

Troponin T CR (ng/l)

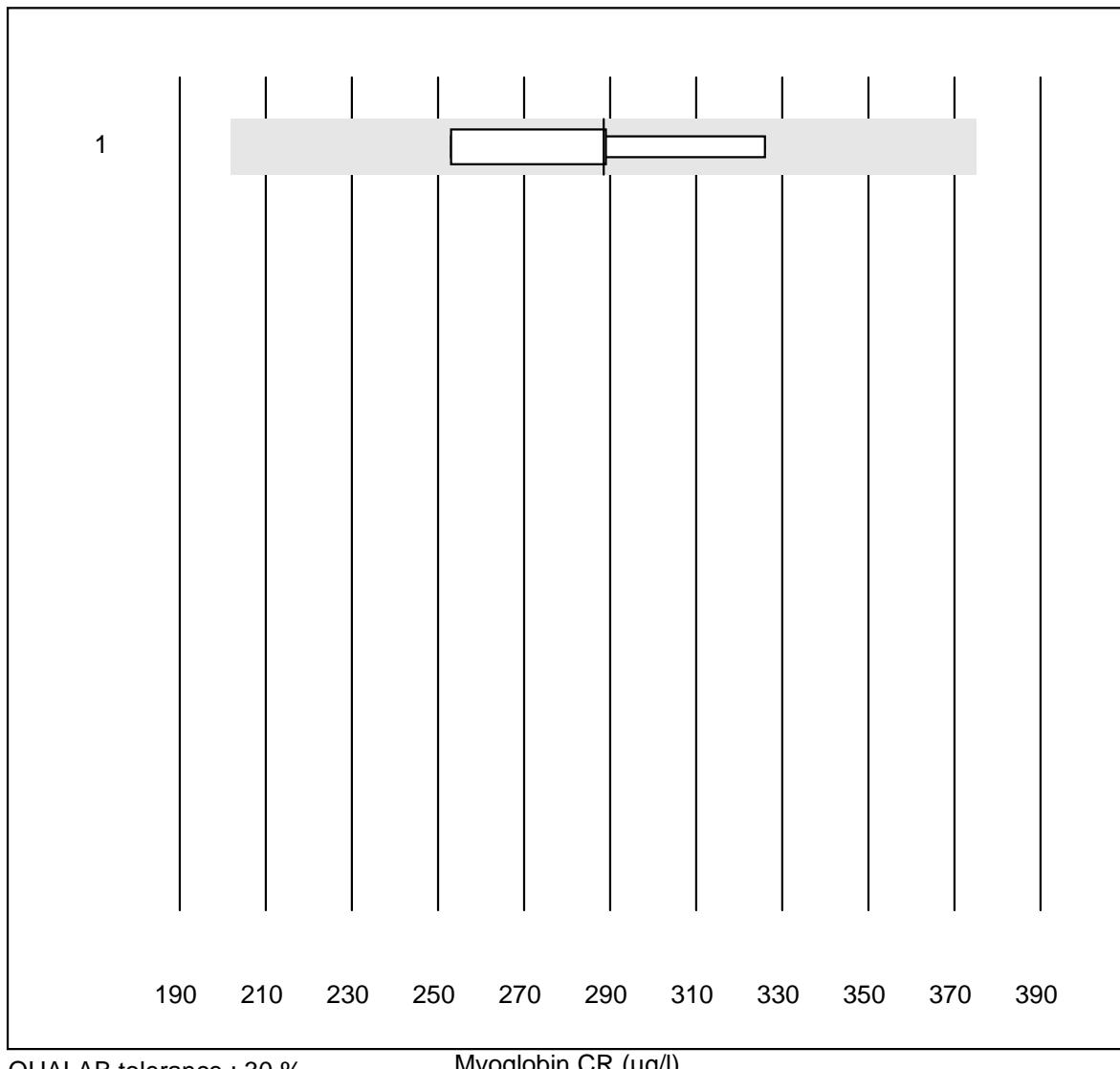
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Cobas h 232	765	96.2	3.0	0.8	248.32	10.4	e
2 Cardiac Reader	62	95.2	3.2	1.6	253.98	13.6	e

Troponin I WB

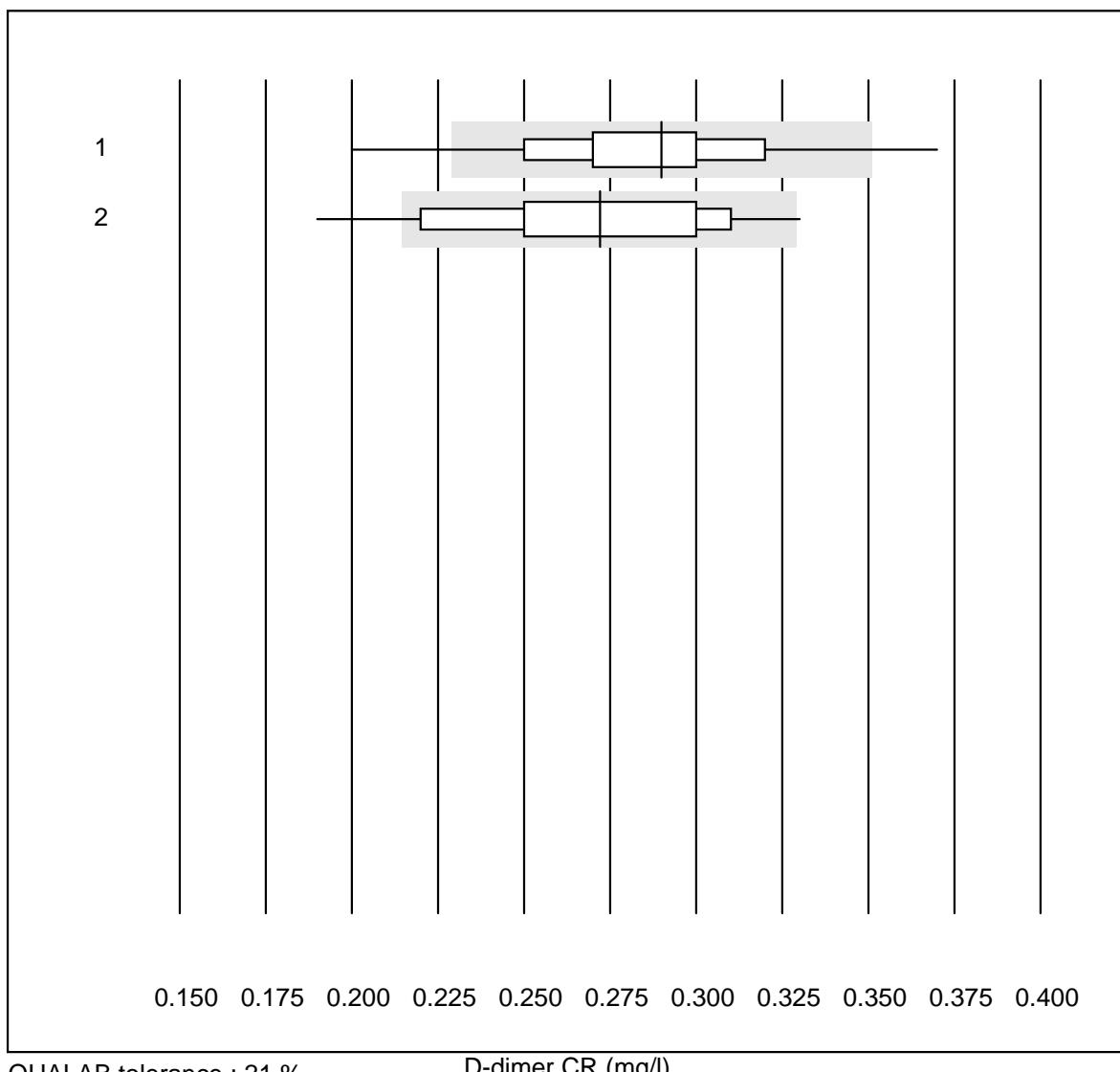


No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	iStat	4	100.0	0.0	0.0	3.12	10.2	e*

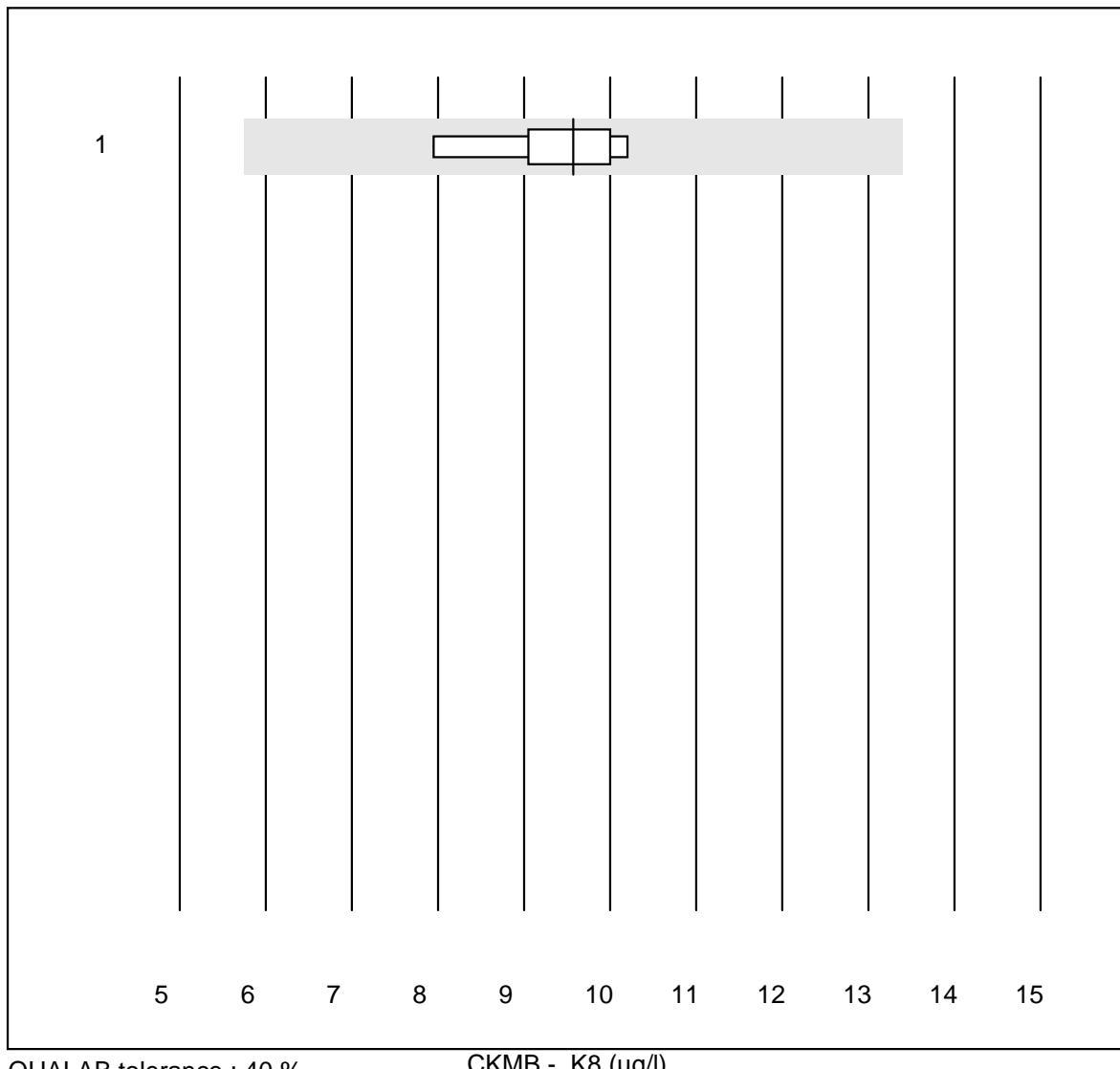
Myoglobin CR



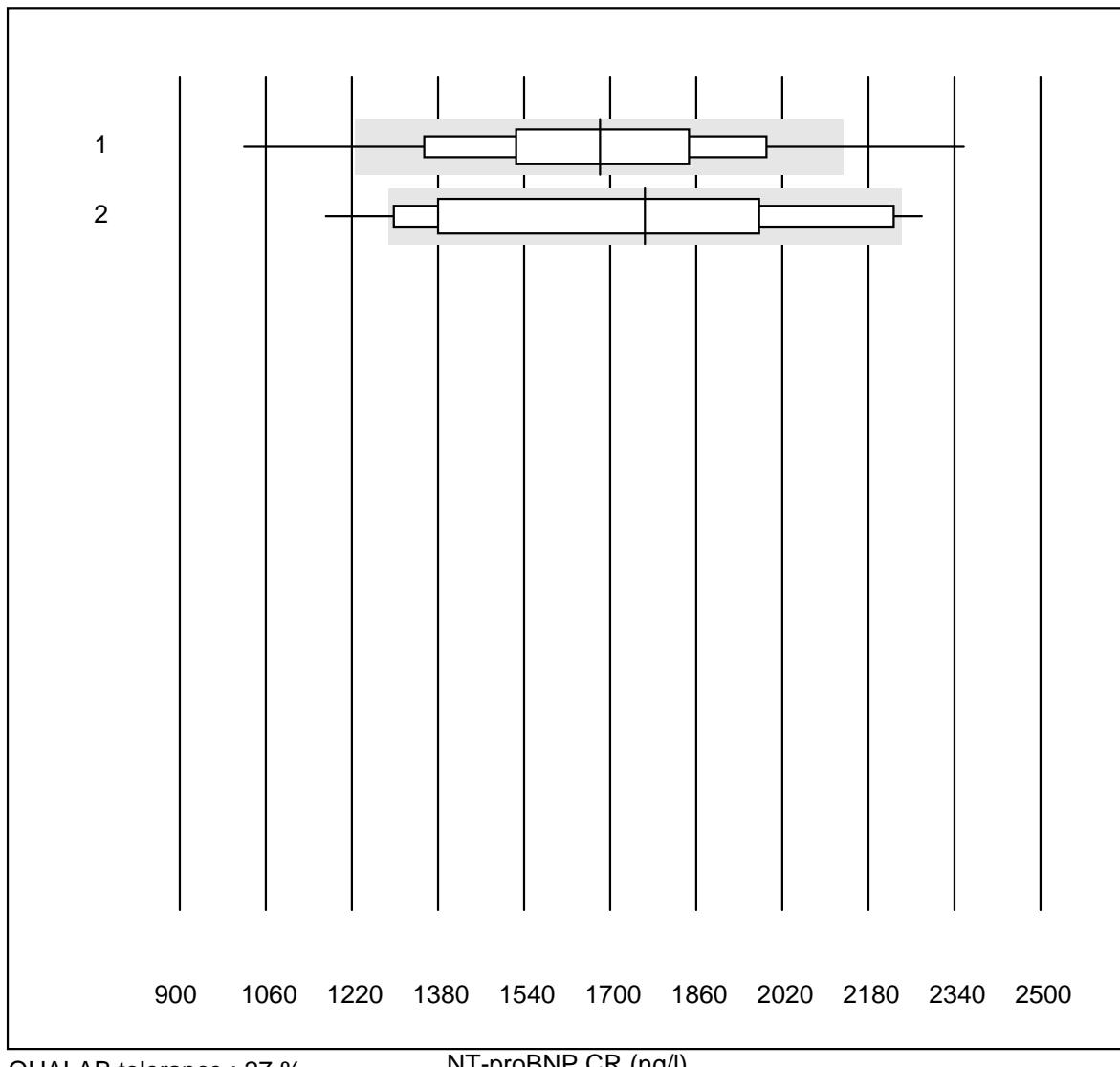
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Cobas h 232	4	100.0	0.0	0.0	288.5	10.3	e*

D-dimer CR

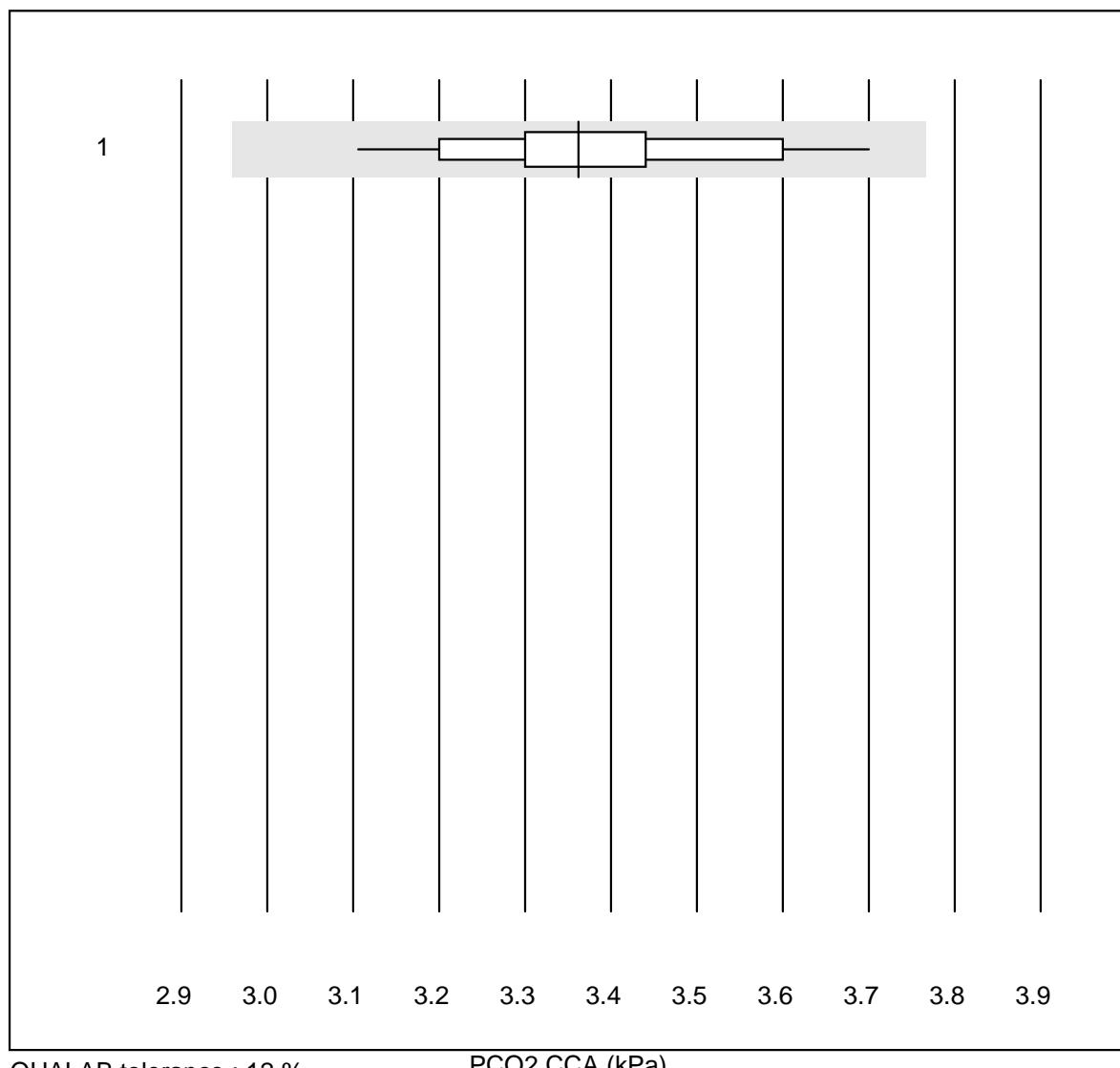
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Cobas h 232	784	96.3	2.9	0.8	0.29	9.6	e
2 Cardiac Reader	58	84.5	13.8	1.7	0.27	13.3	e

CKMB - K8

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Cobas h 232	7	100.0	0.0	0.0	9.6	8.4	e

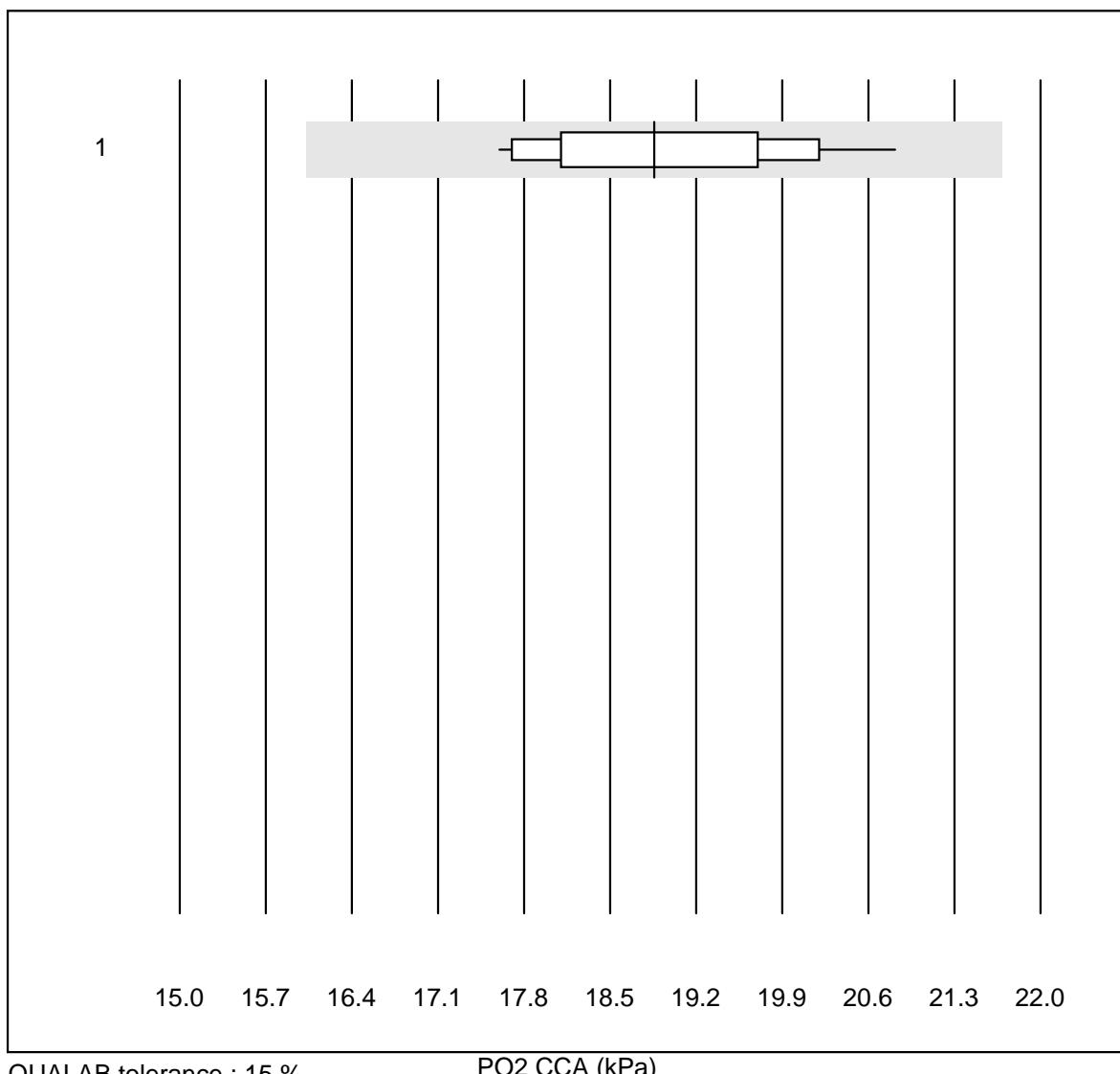
NT-proBNP CR

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Cobas h 232	488	89.7	7.0	3.3	1681	14.7	e
2 Cardiac Reader	20	85.0	10.0	5.0	1765	20.7	e*

PCO₂ CCA

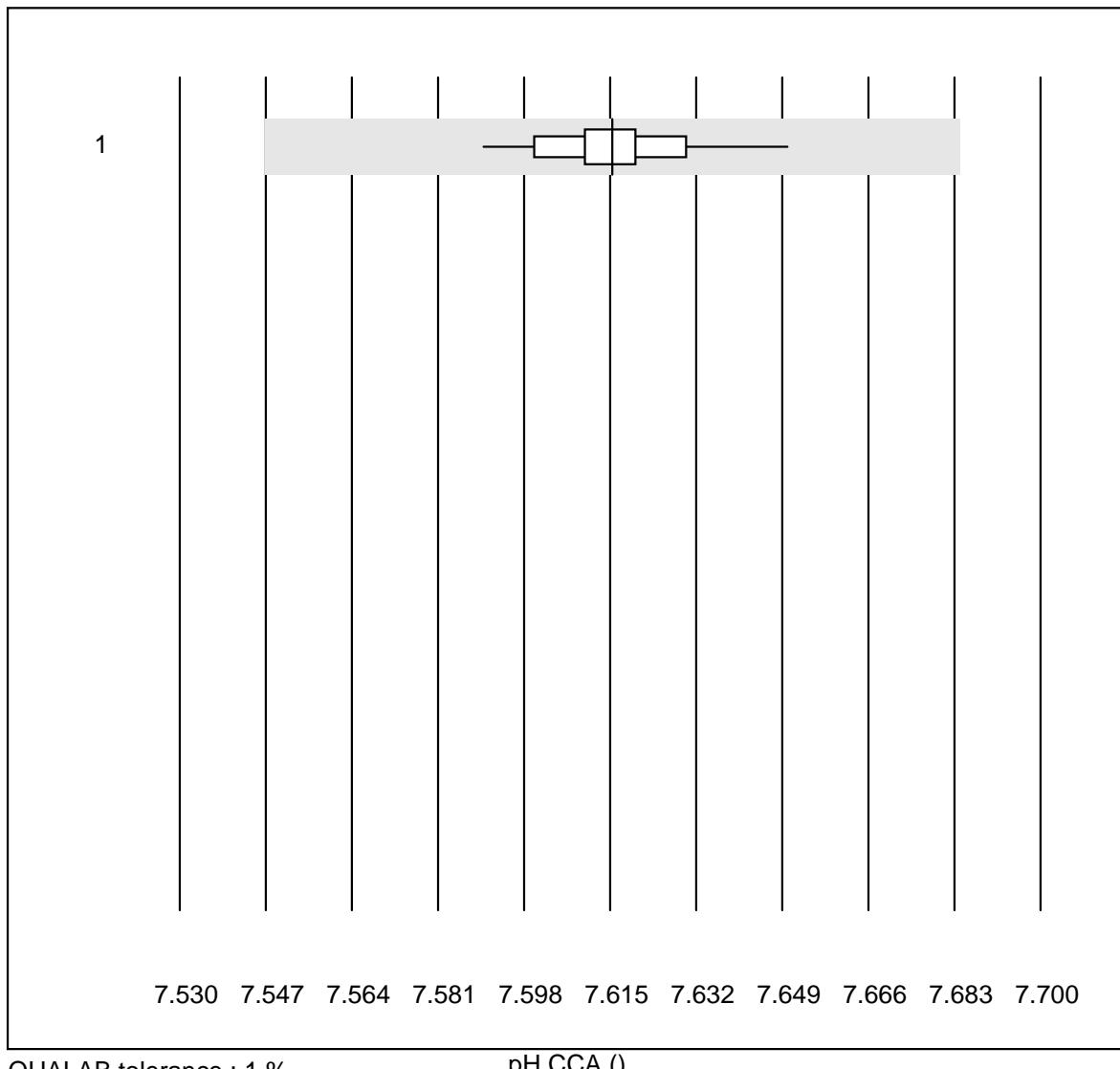
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	OPTI CCA	14	100.0	0.0	0.0	3.36	4.7	e

PO2 CCA



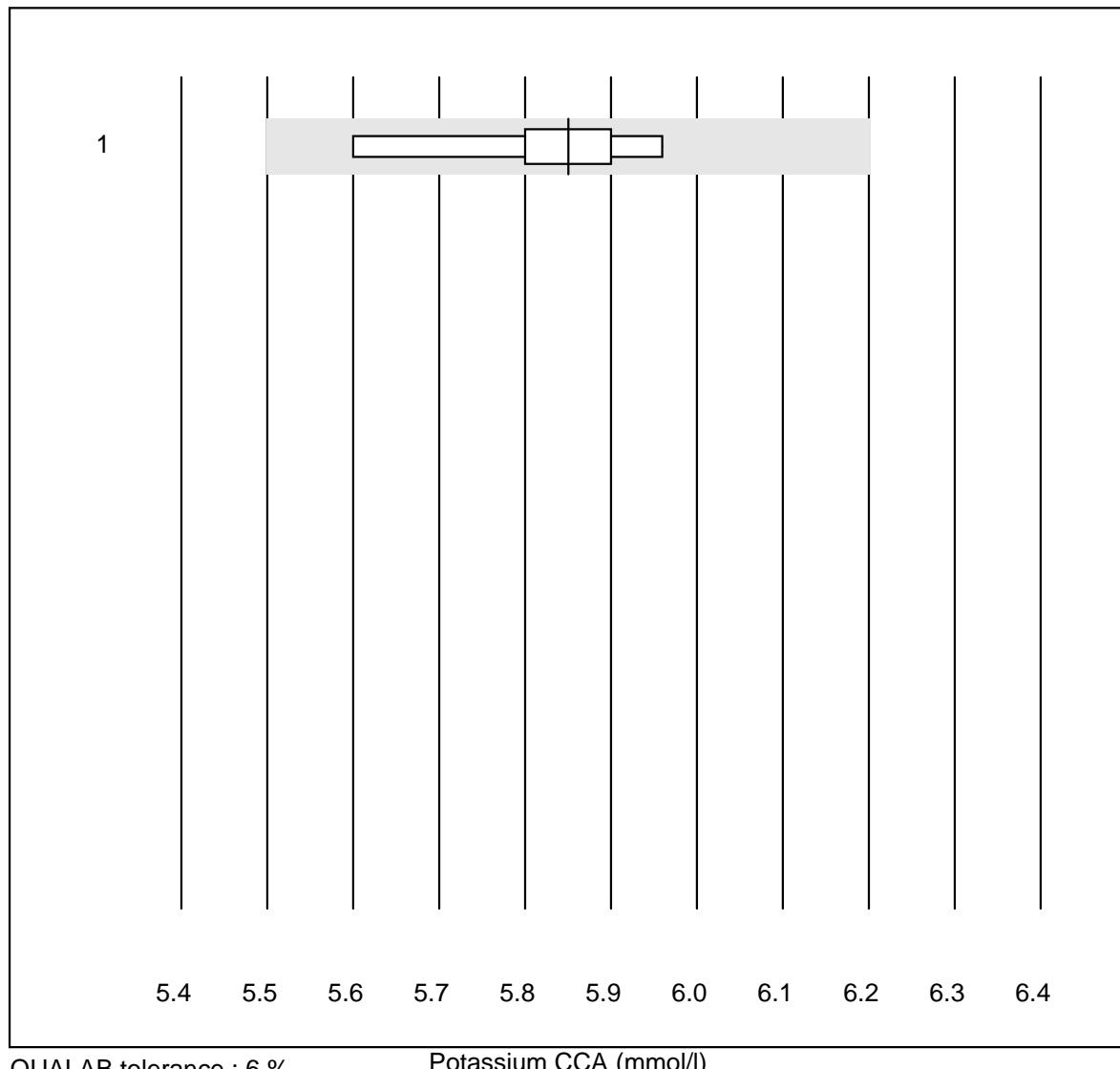
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	OPTI CCA	14	100.0	0.0	0.0	18.86	5.3	e

pH CCA



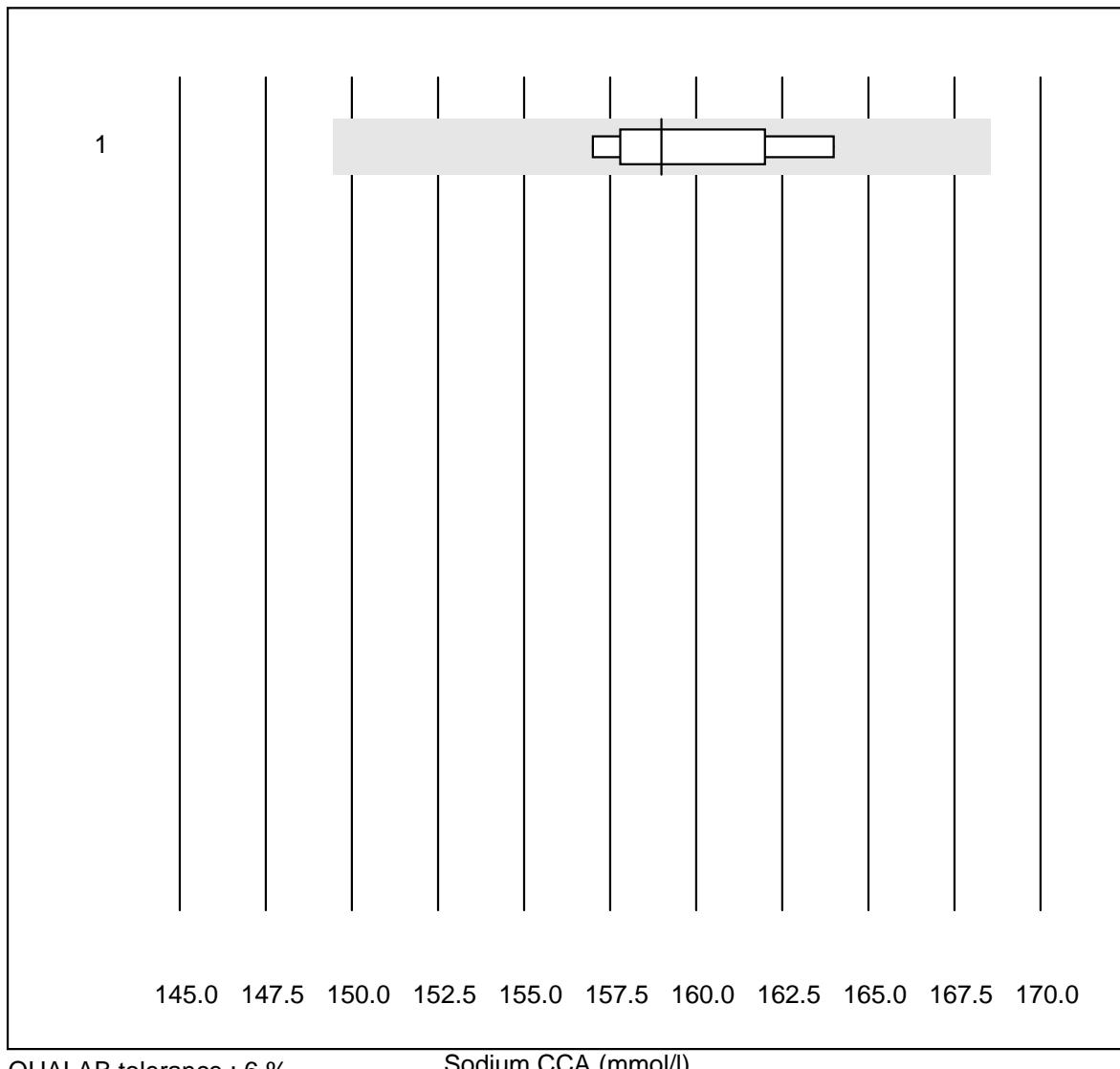
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 OPTI CCA	14	100.0	0.0	0.0	7.62	0.2	e

Potassium CCA

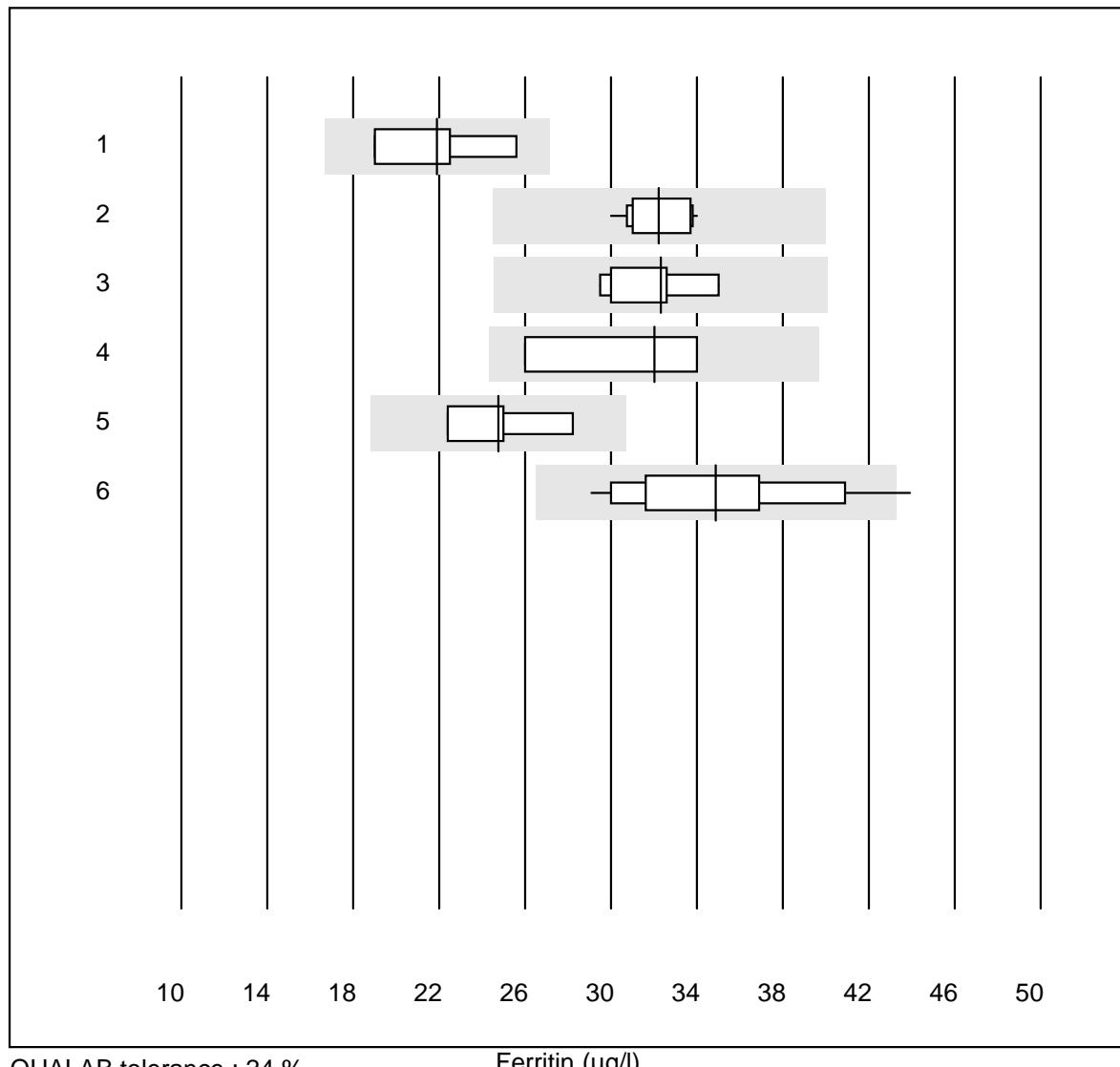


No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	OPTI CCA	8	100.0	0.0	0.0	5.9	1.9	e

Sodium CCA



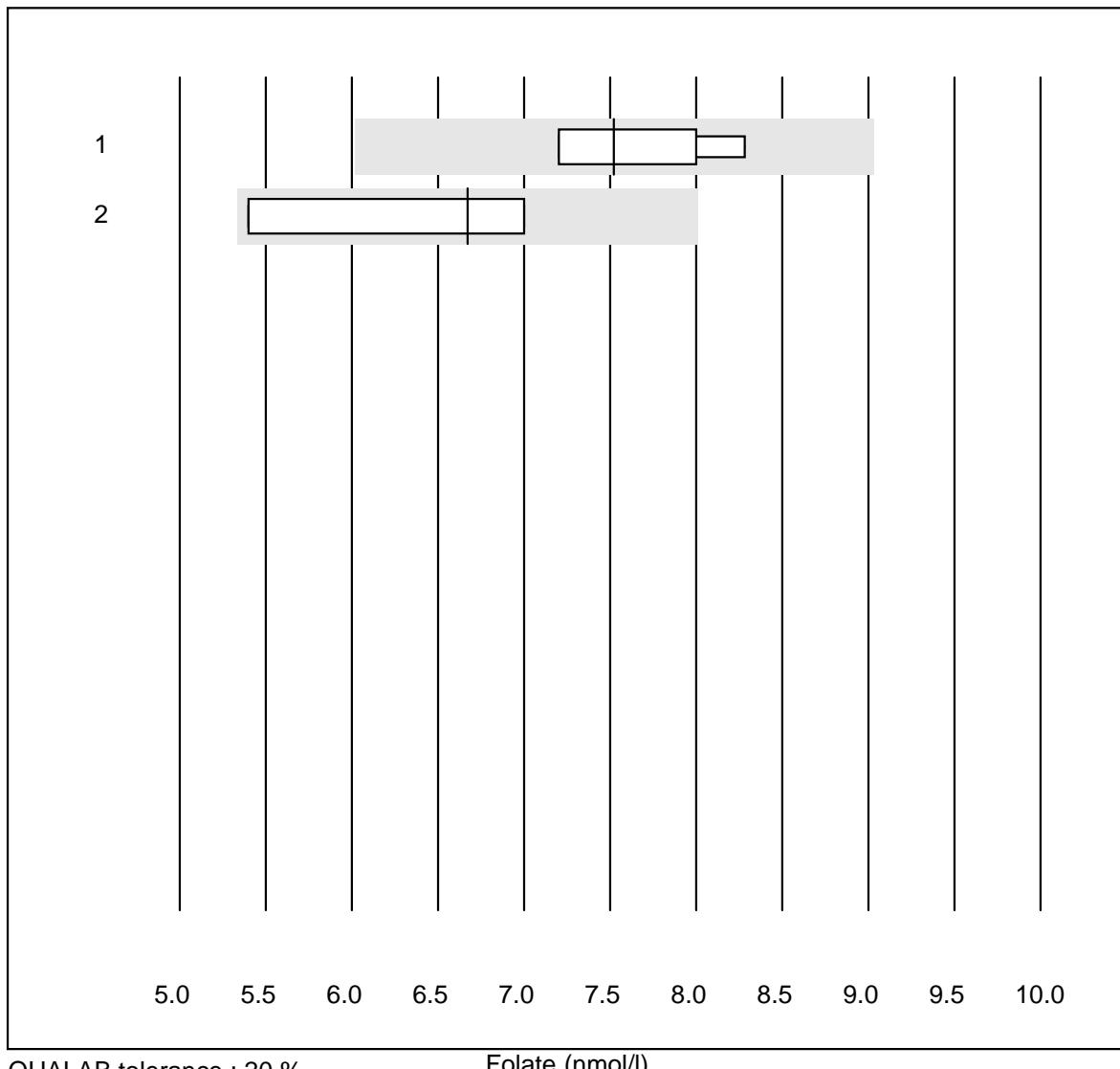
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	OPTI CCA	7	100.0	0.0	0.0	159.0	1.5	e

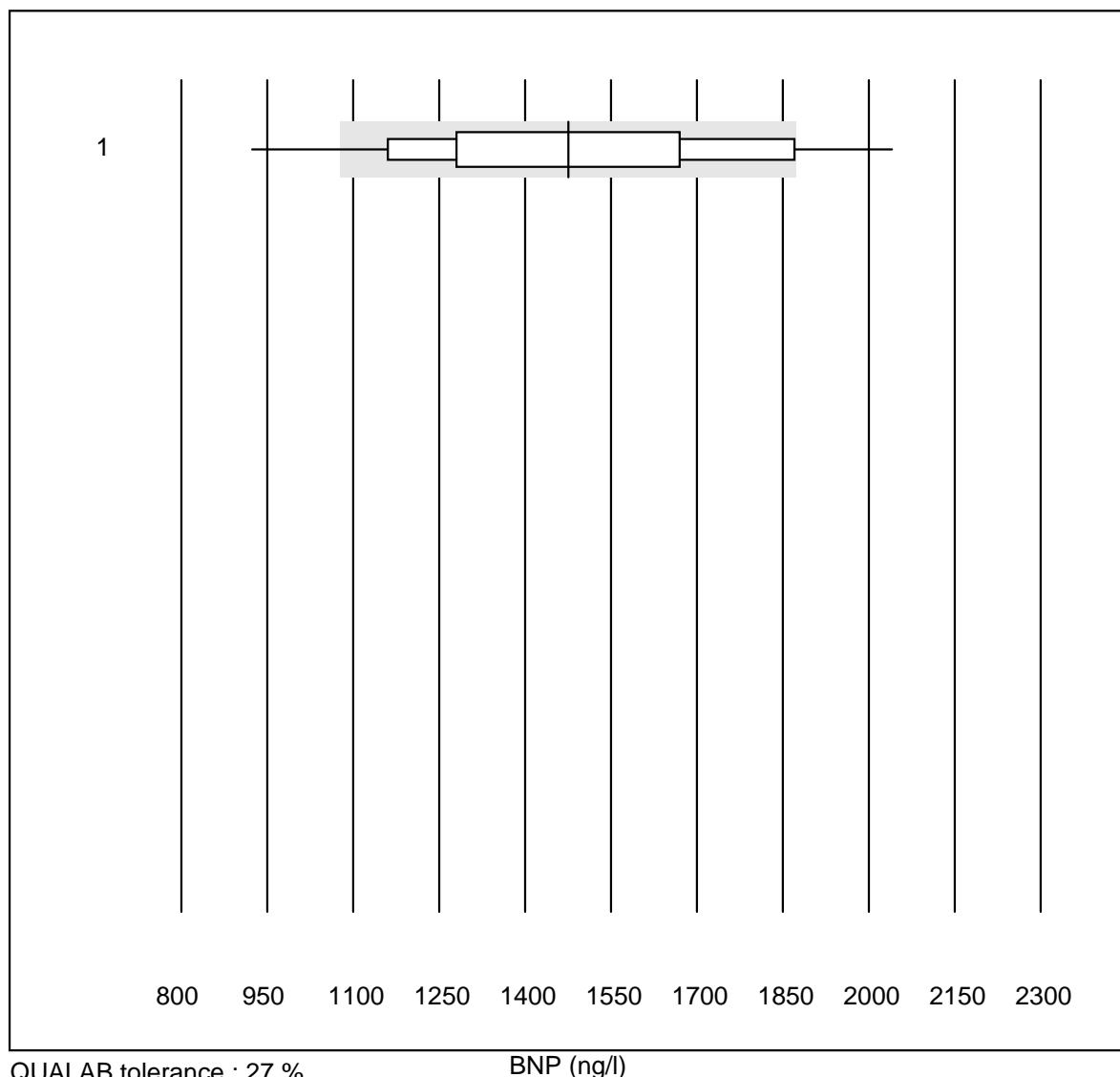
Ferritin

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Beckman	4	100.0	0.0	0.0	21.90	12.4	e*
2 Cobas E / Elecsys	11	100.0	0.0	0.0	32.23	4.3	e
3 Architect	5	100.0	0.0	0.0	32.33	7.0	e*
4 Mira/DiaSys	4	100.0	0.0	0.0	32.00	12.4	e*
5 Mini Vidas	4	100.0	0.0	0.0	24.74	9.6	e*
6 Eurolyser	18	88.8	5.6	5.6	34.89	11.9	e

Vitamin B12

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 ADVIA Centaur XP/CP	5	80.0	20.0	0.0	130.00	13.1	e*
2 Cobas E / Elecsys	7	100.0	0.0	0.0	211.39	6.5	e*
3 Architect	4	100.0	0.0	0.0	200.50	7.2	e*

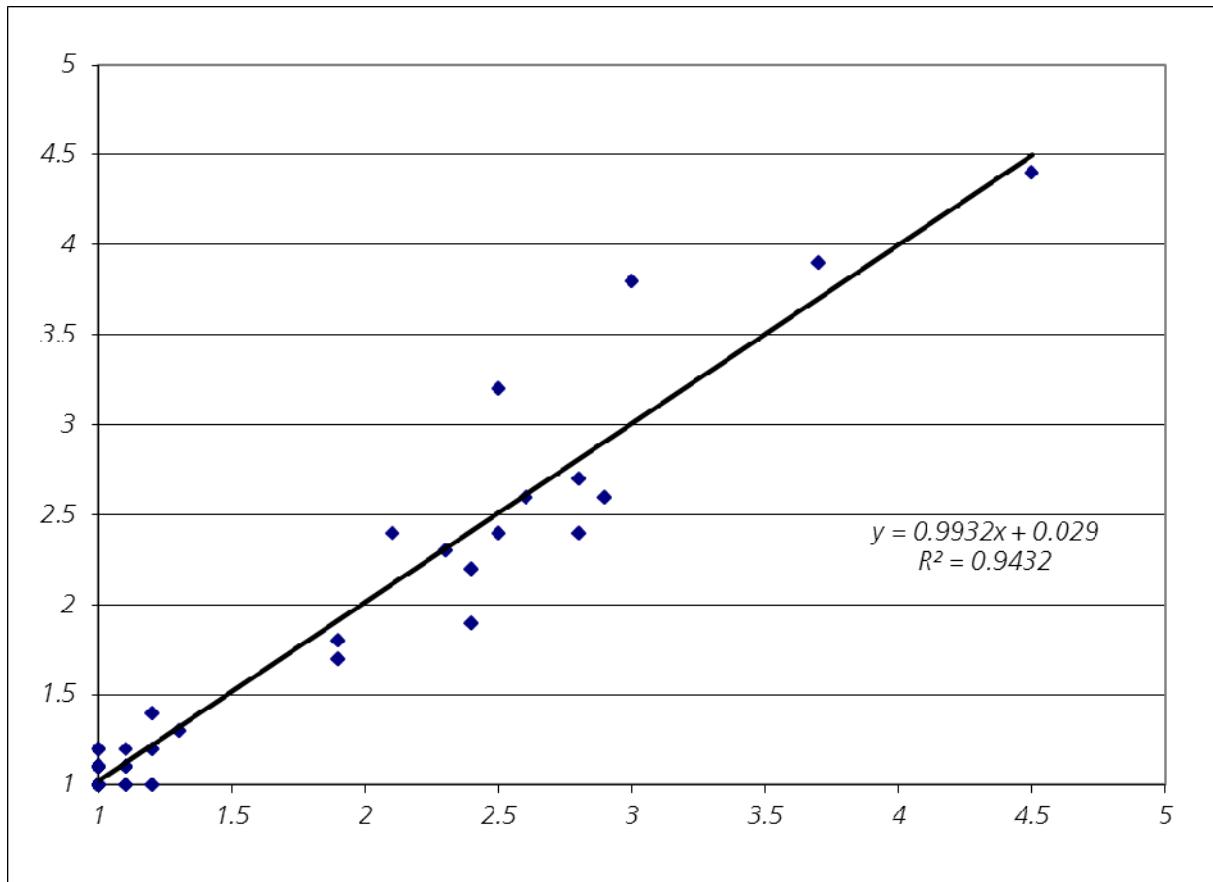
Folate

BNP

G10 INR INRatio

INR INRatio

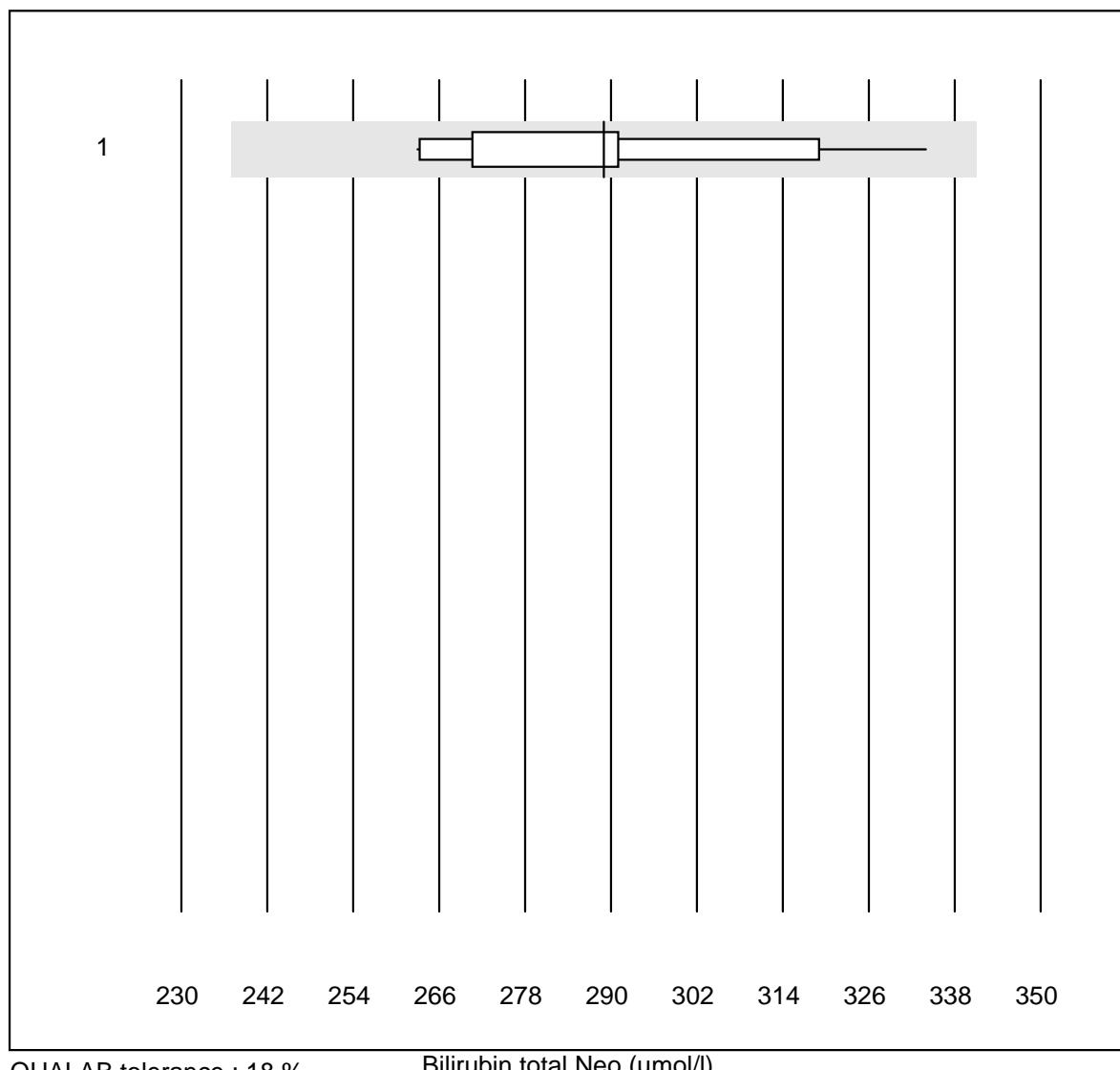
University Hospital Zuerich



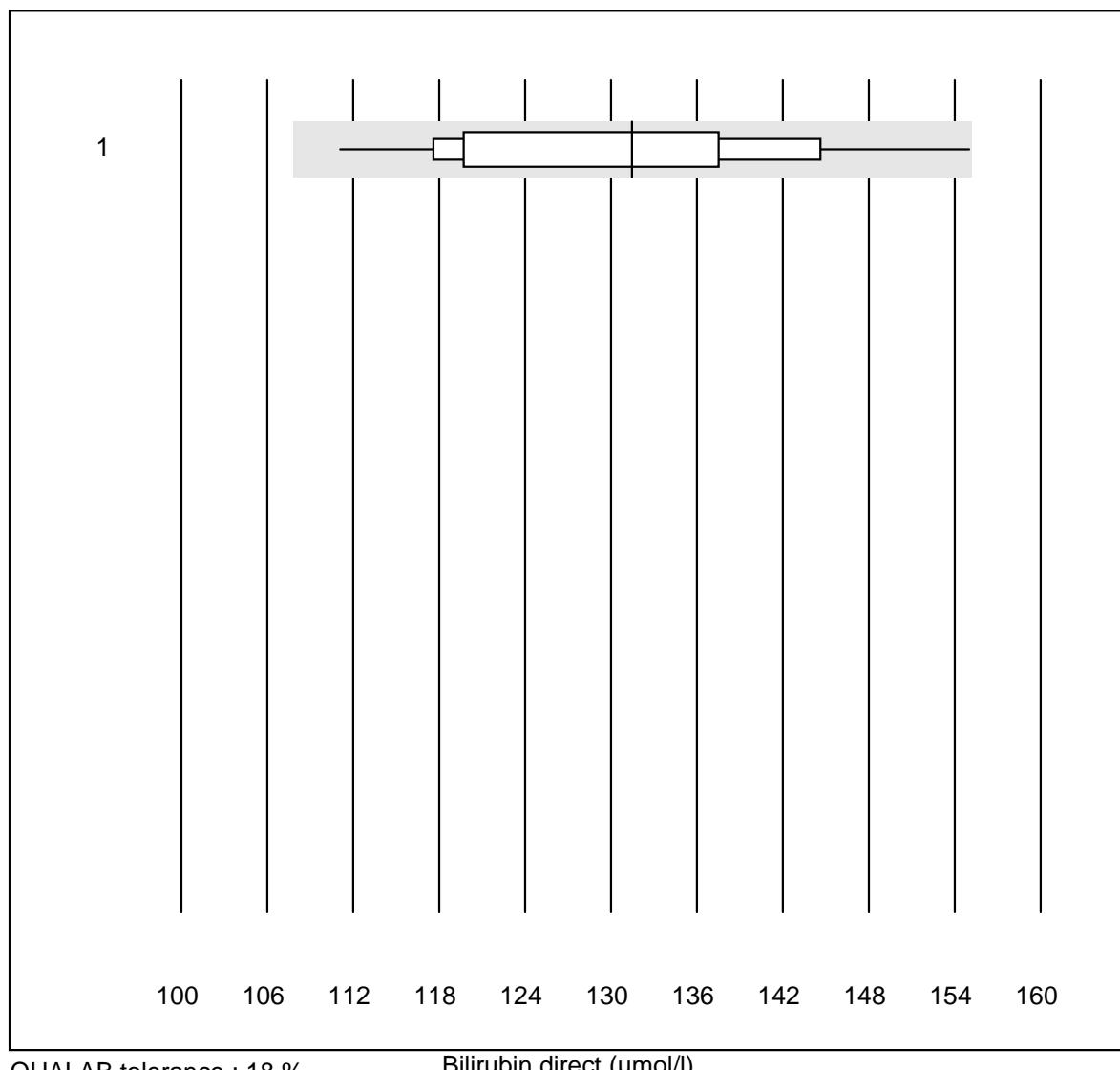
INRatio Participants

G10 is a split-sample survey. We compare INR-values from our participants with the corresponding plasma INR from University Hospital Zuerich.

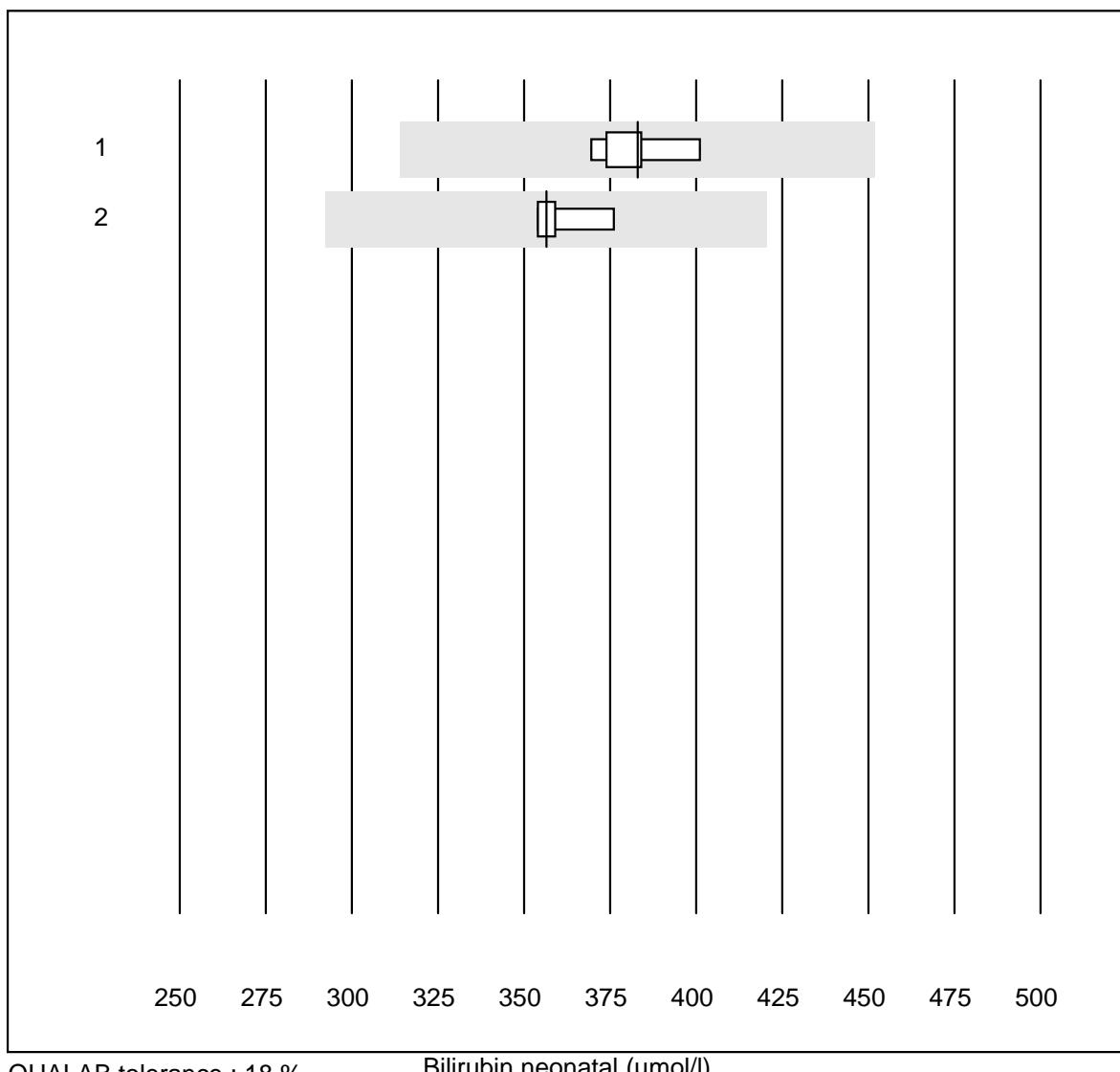
Nr.	Device	Total	% good	% insufficient	% outlier
1	58	93.1	5.17	1.72	58

Bilirubin total Neo

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 all Participants	12	100.0	0.0	0.0	289	7.5	e*

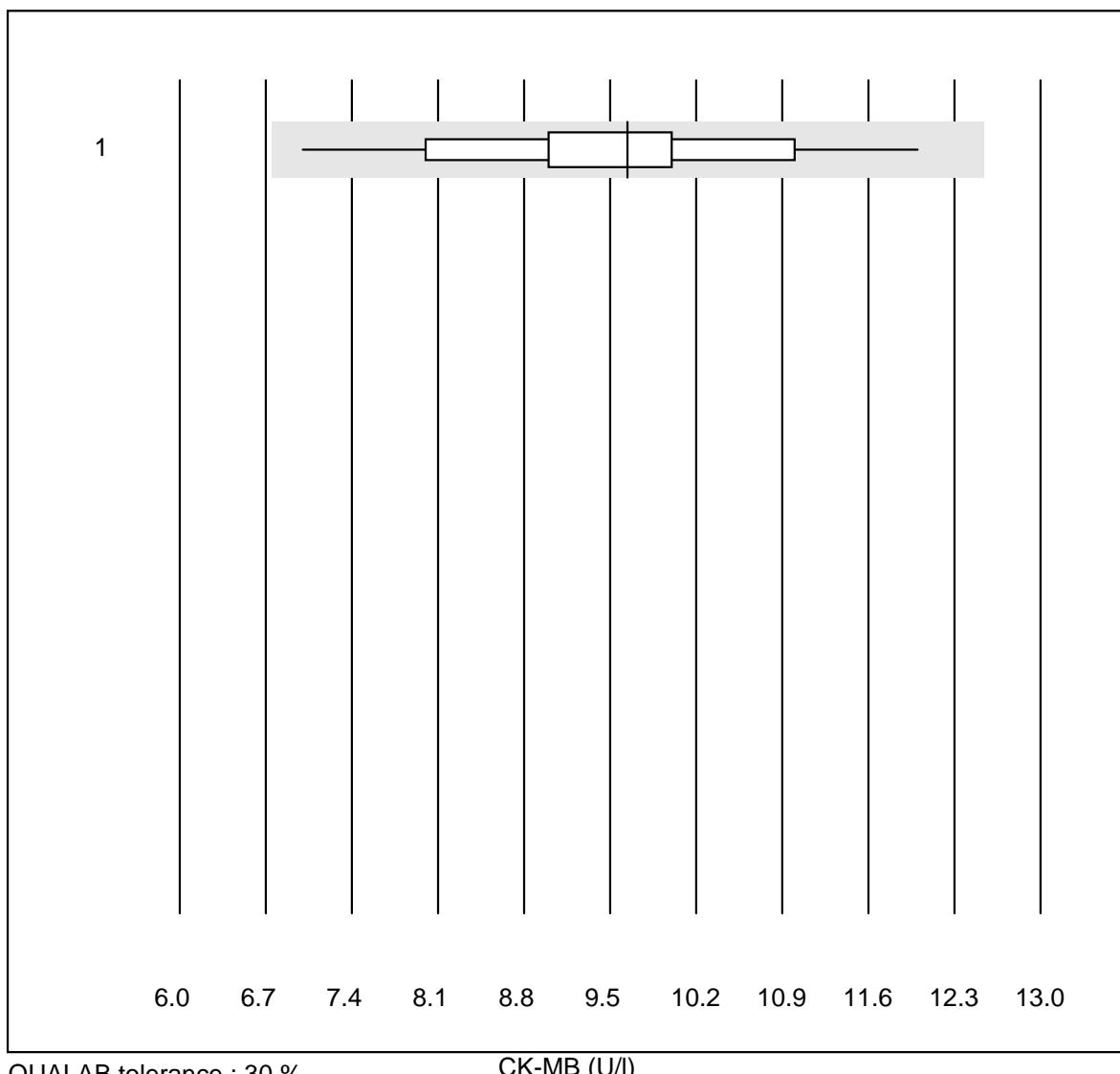
Bilirubin direct

Bilirubin neonatal



K15 Creatinkinase Activity

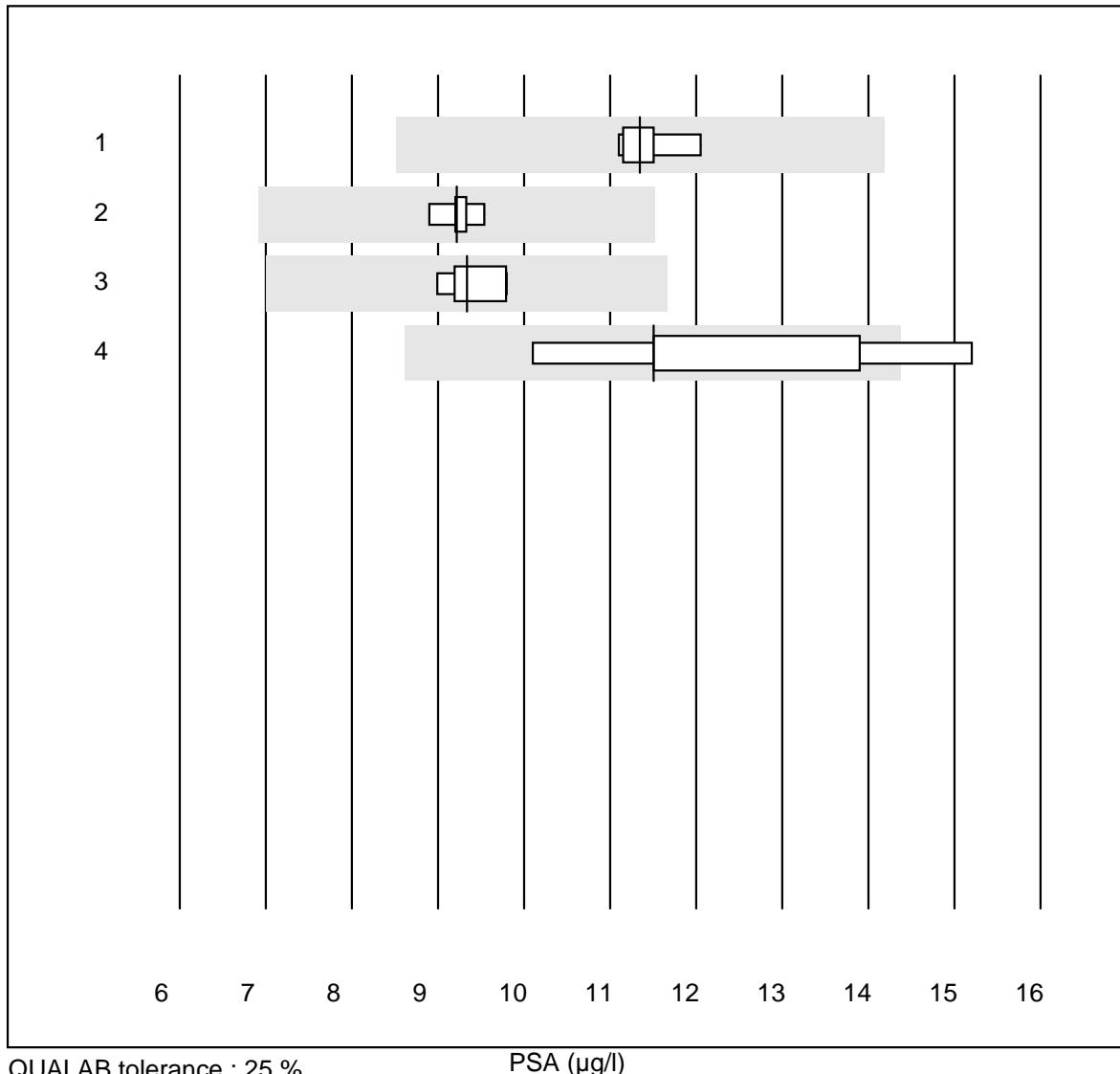
CK-MB



No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Fuji Dri-Chem	42	92.9	0.0	7.1	9.6	11.0	e

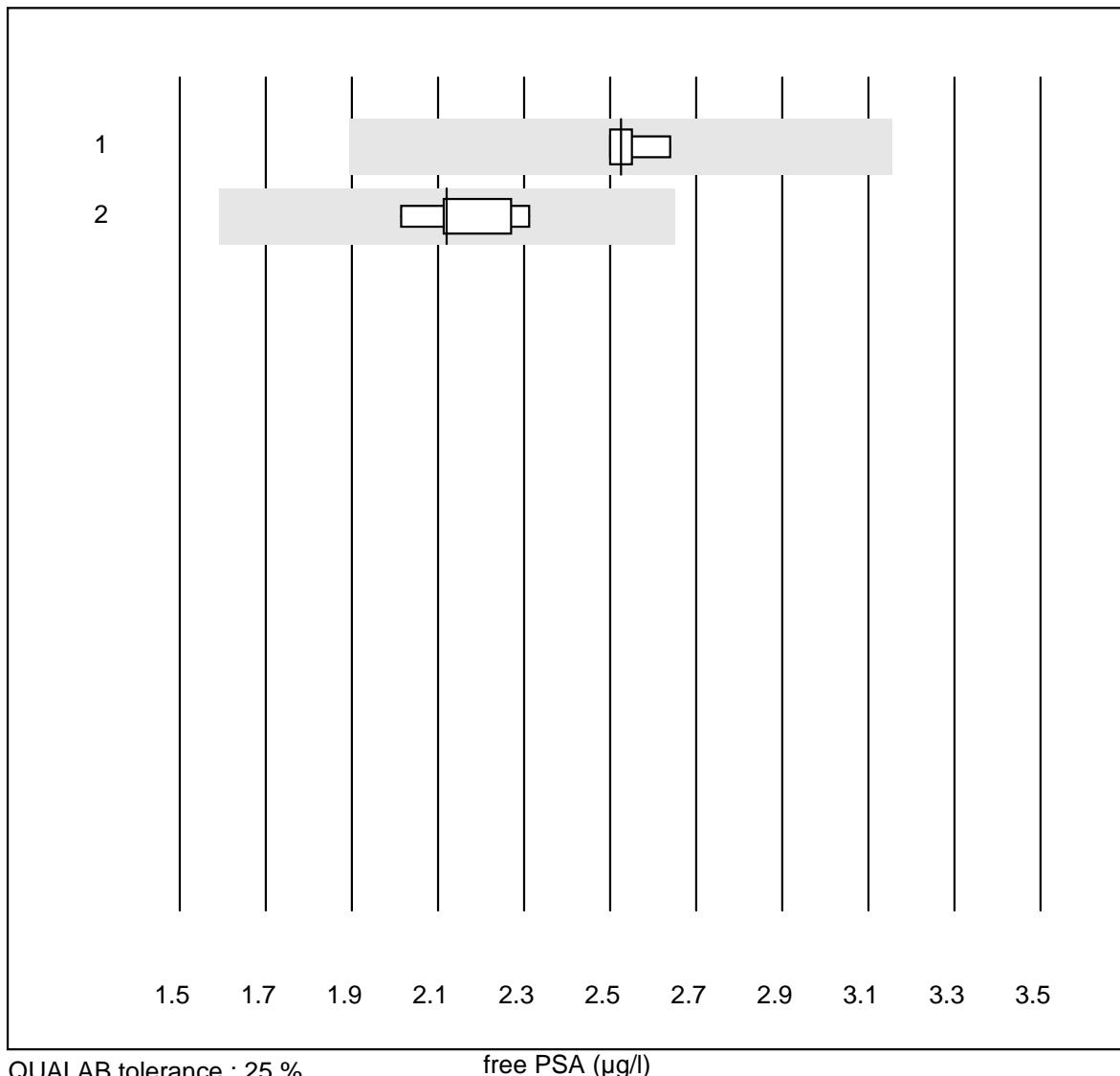
K14 Tumor Markers

PSA



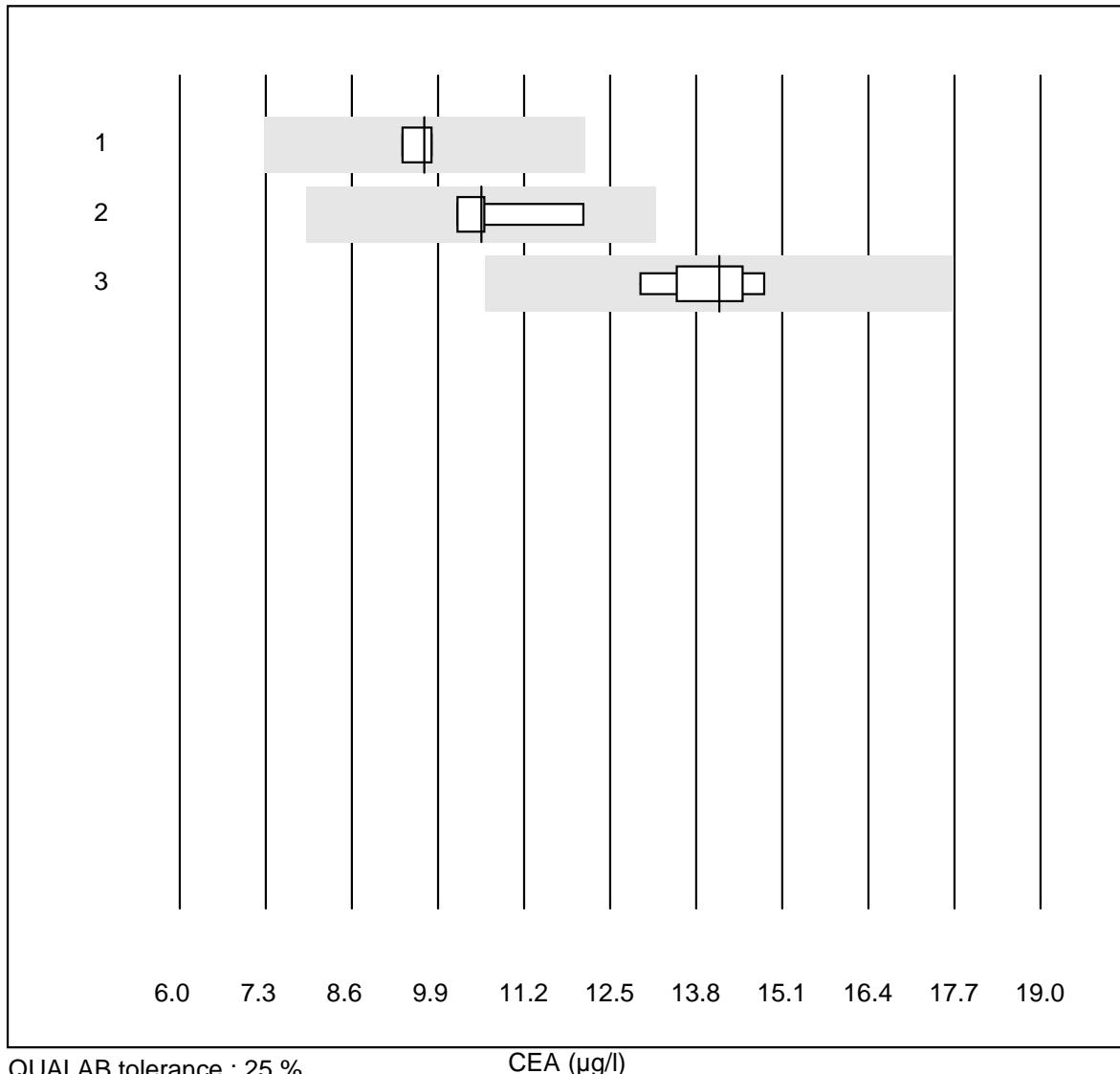
K14 Tumor Markers

free PSA



K14 Tumor Markers

CEA



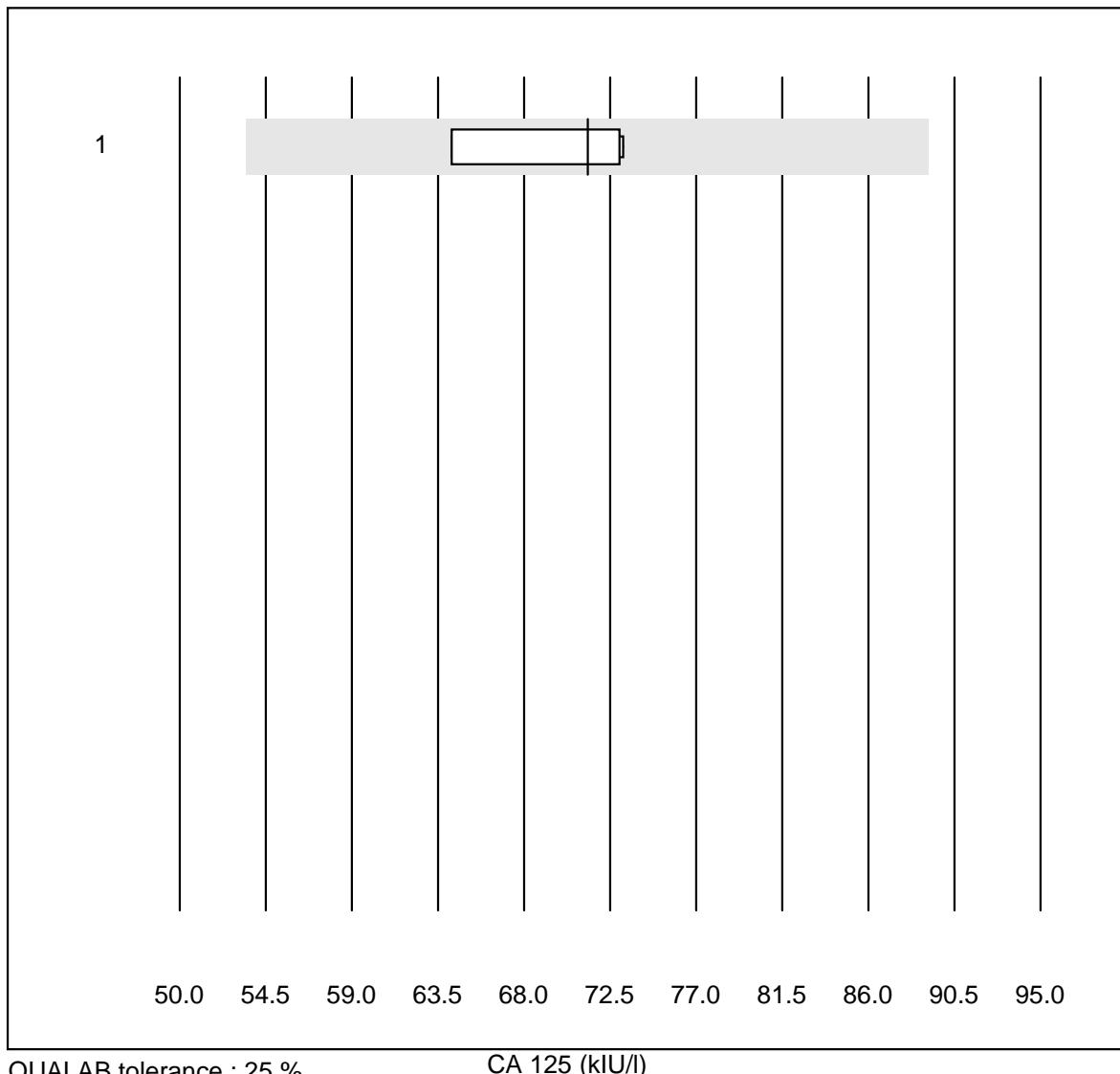
QUALAB tolerance : 25 %

CEA ($\mu\text{g/l}$)

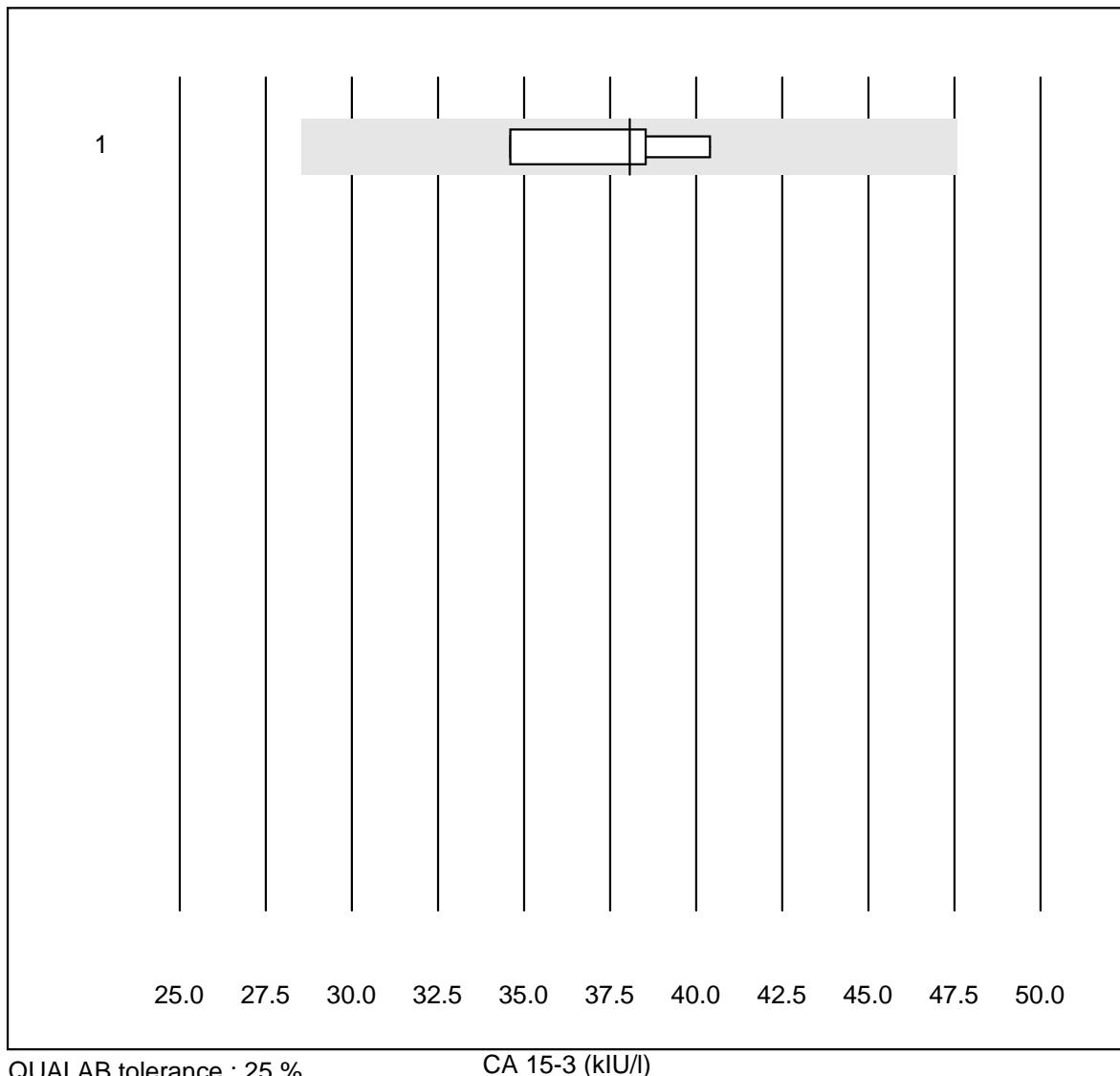
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Cobas E / Elecsys	4	100.0	0.0	0.0	9.7	2.2	e
2	ADVIA Centaur XP/CP	4	100.0	0.0	0.0	10.6	7.8	e*
3	Architect	5	100.0	0.0	0.0	14.2	5.4	e

K14 Tumor Markers

CA 125



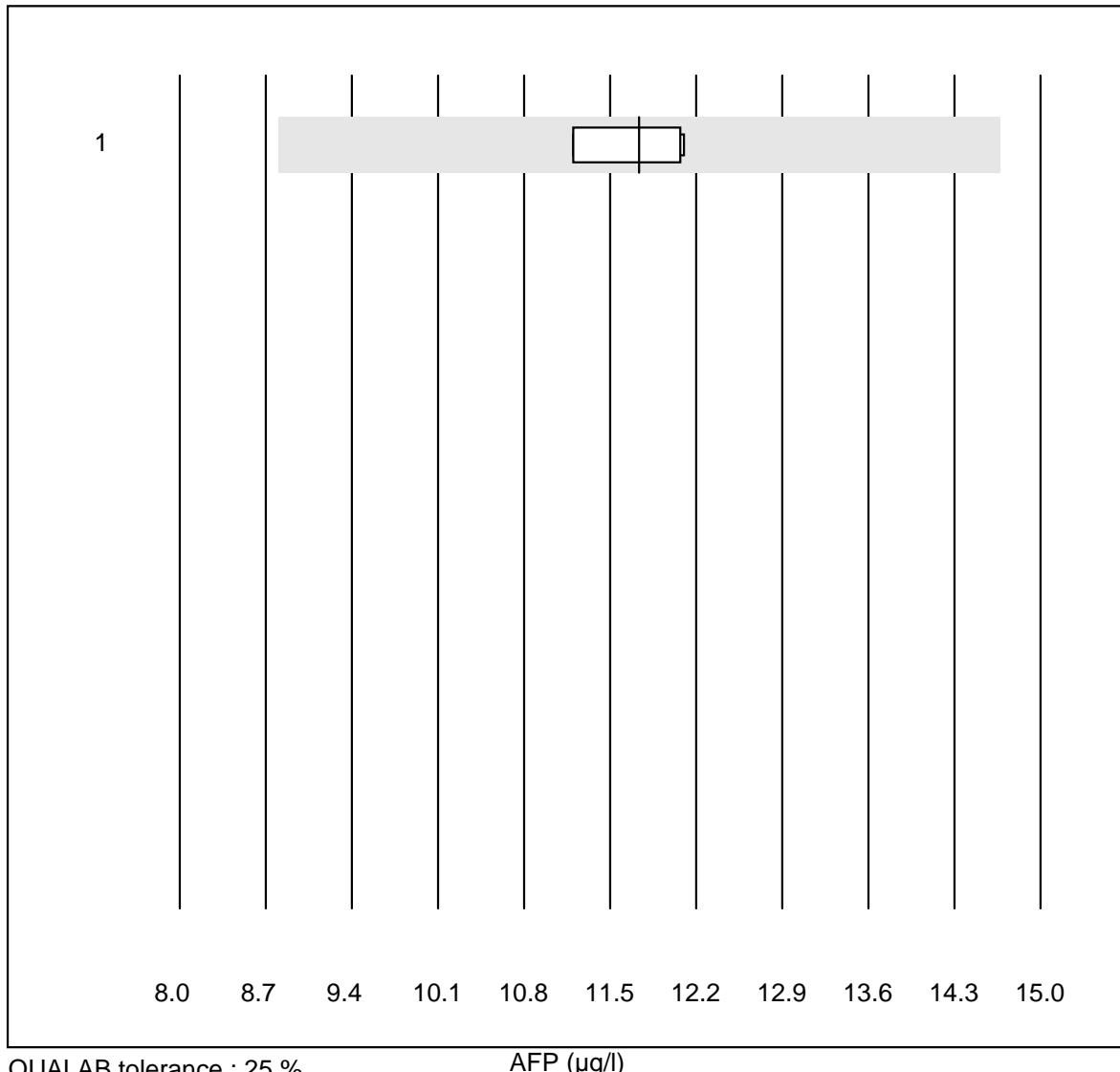
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Architect	4	100.0	0.0	0.0	71.3	6.0	e

CA 15-3

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Architect	4	100.0	0.0	0.0	38.1	6.4	e*

K14 Tumor Markers

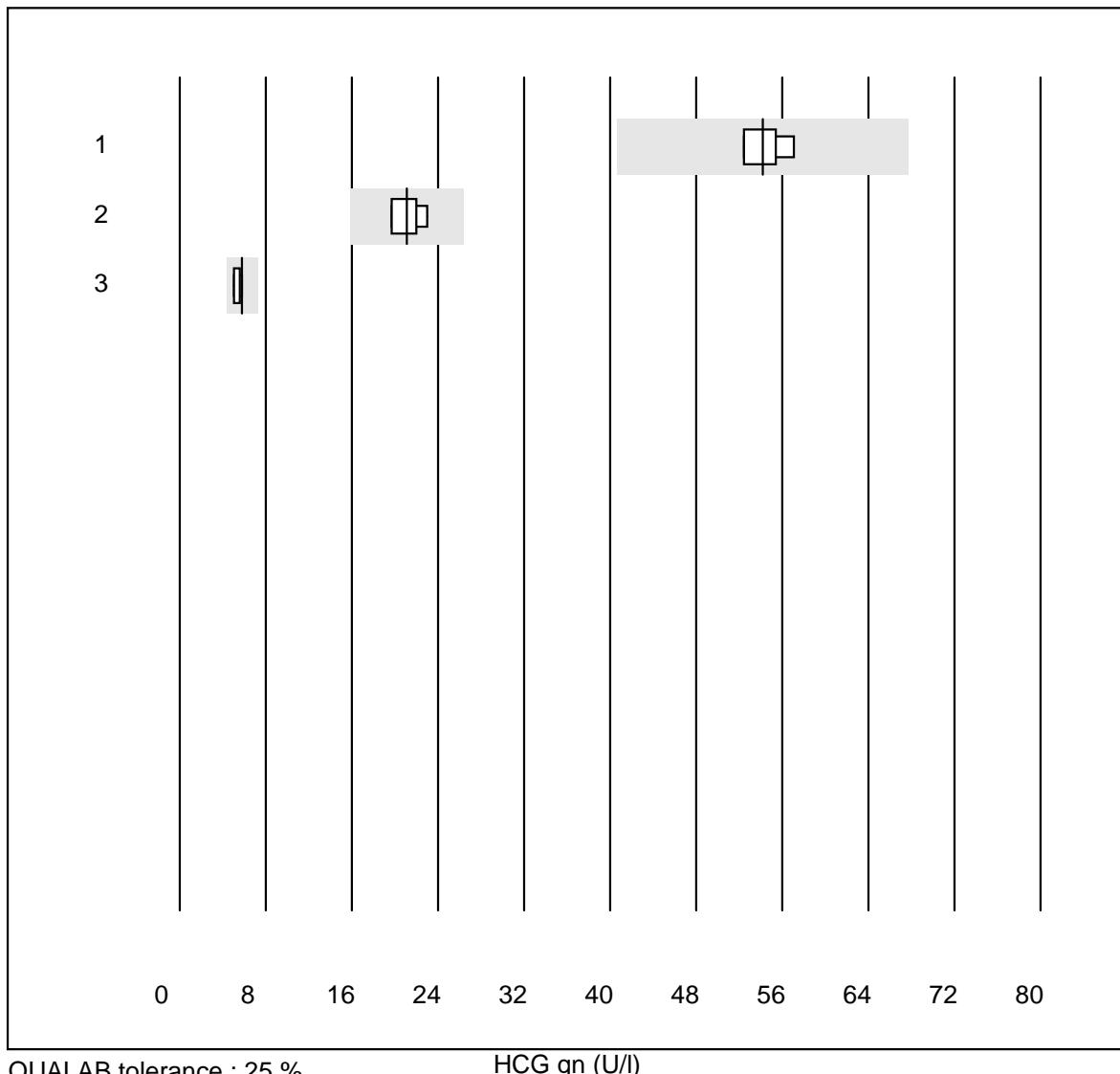
AFP



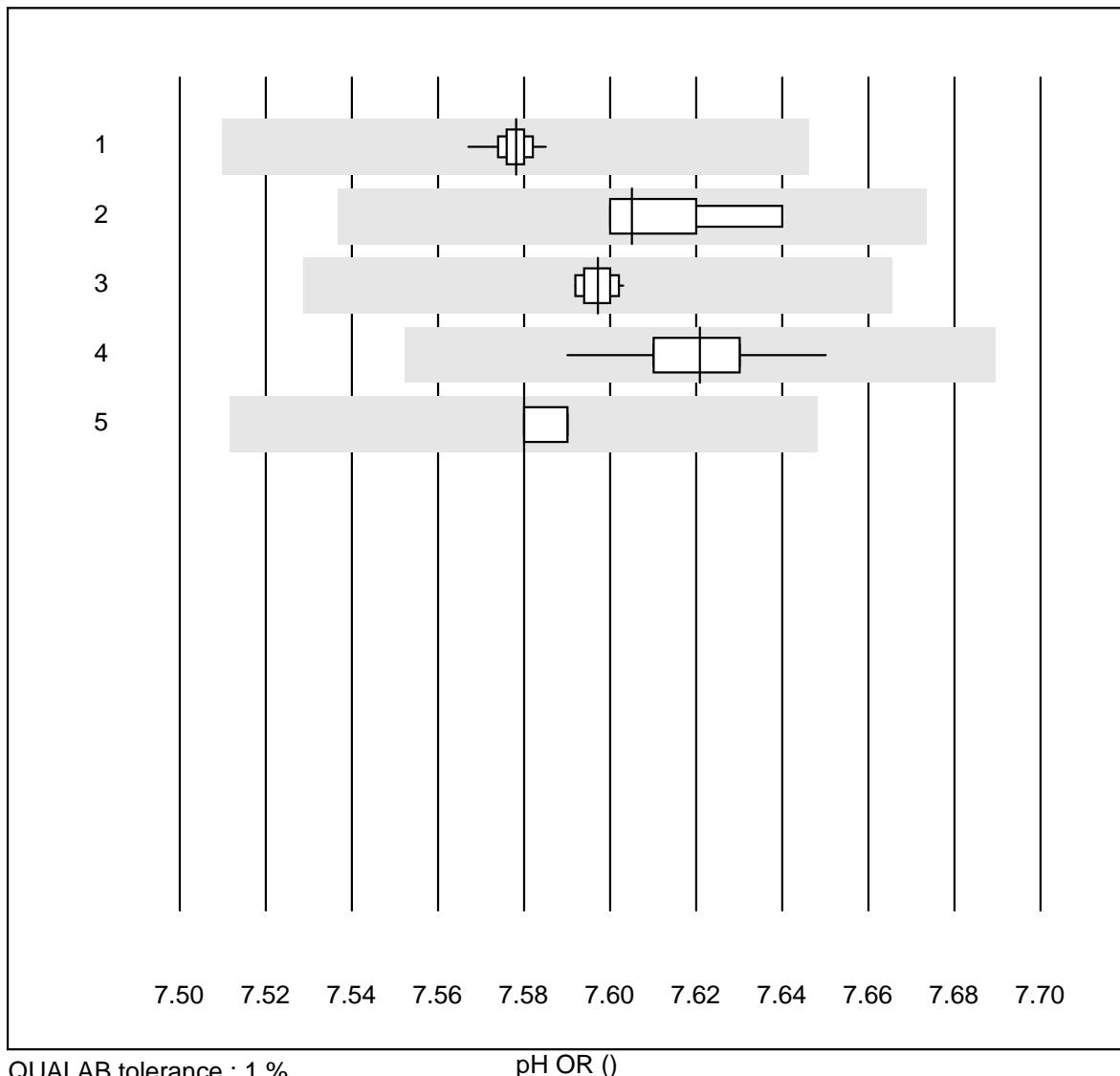
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Architect	4	100.0	0.0	0.0	12	3.9	e

K14 Tumor Markers

HCG qn



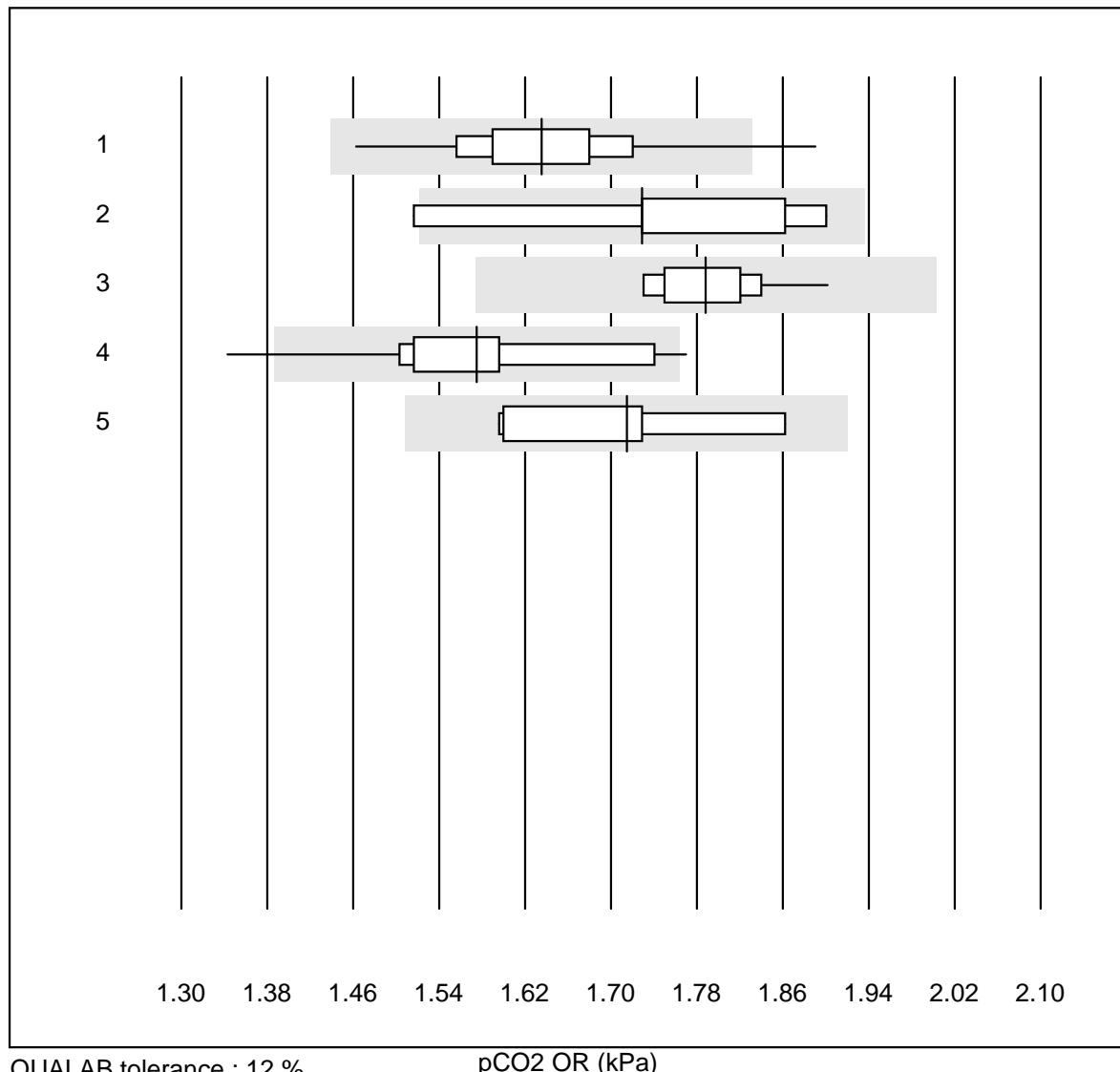
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Cobas E / Elecsys	4	100.0	0.0	0.0	54	4.0	e
2	ADVIA Centaur XP/CP	4	100.0	0.0	0.0	21	7.3	e*
3	Vidas	4	100.0	0.0	0.0	6	6.5	a

pH OR

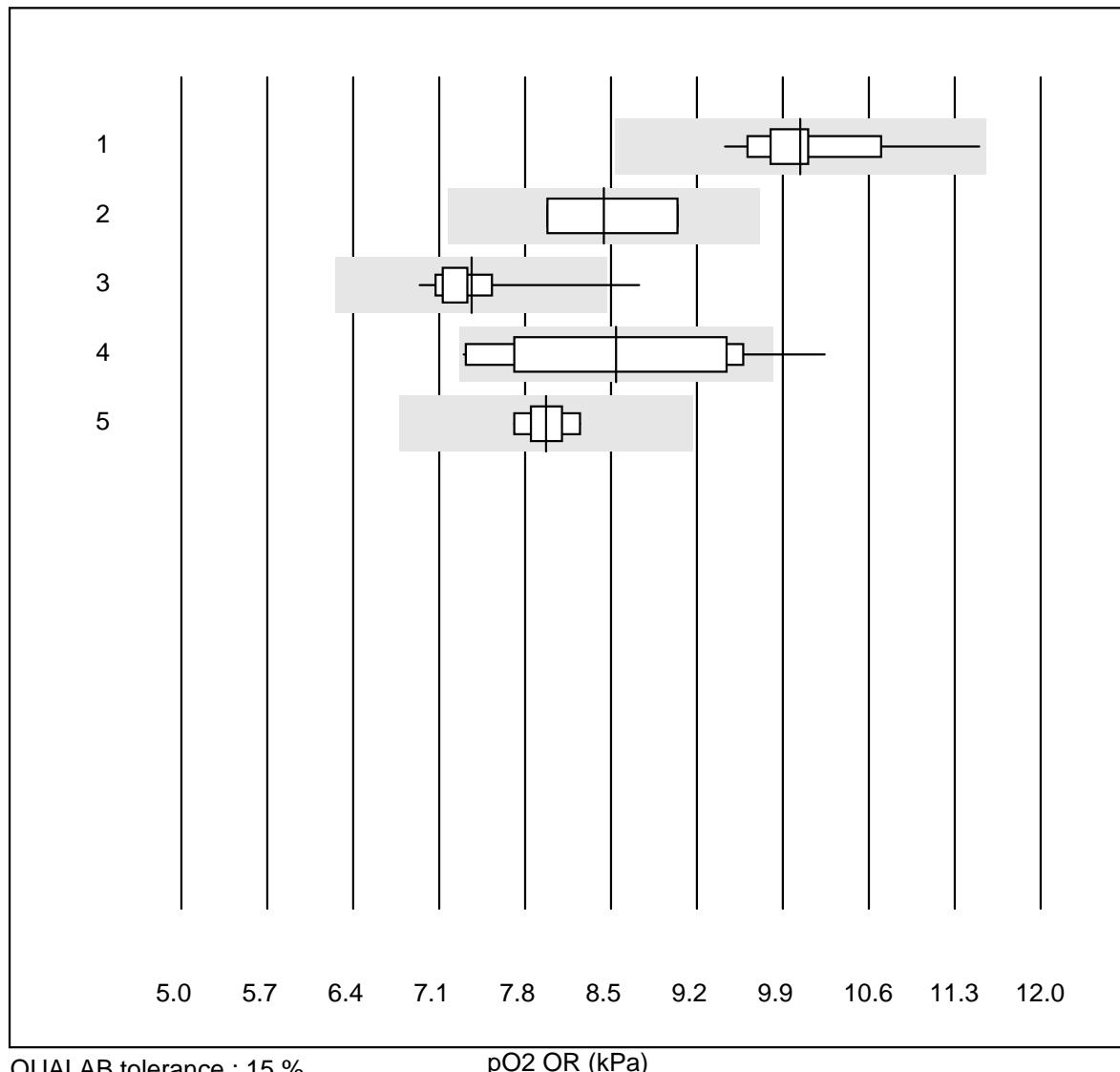
QUALAB tolerance : 1 %

pH OR ()

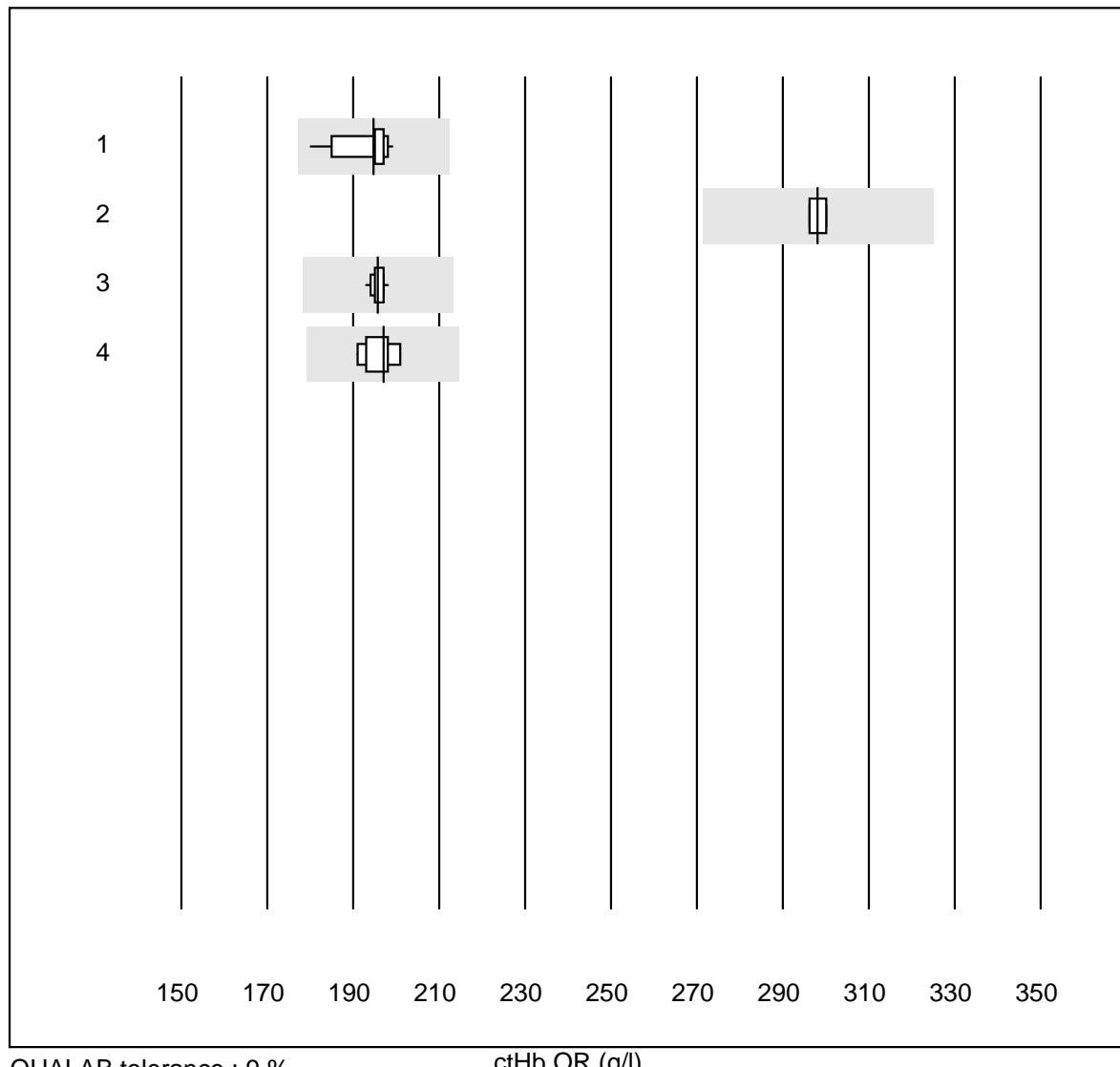
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	ABL700/800 Radiomete	78	98.7	0.0	1.3	7.58	0.0	e
2	Radiometer NPT-7	6	100.0	0.0	0.0	7.61	0.2	e
3	ABL 90	25	100.0	0.0	0.0	7.60	0.0	e
4	ABL 80 / Coox	13	100.0	0.0	0.0	7.62	0.2	e
5	ABL 5	6	100.0	0.0	0.0	7.58	0.1	e

pCO₂ OR

No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	ABL700/800 Radiomete	77	97.4	2.6	0.0	1.64	4.5	e
2	Radiometer NPT-7	6	66.6	16.7	16.7	1.73	8.6	e*
3	ABL 90	25	100.0	0.0	0.0	1.79	2.5	e
4	ABL 80 / Coox	13	84.6	15.4	0.0	1.58	7.3	e*
5	ABL 5	6	100.0	0.0	0.0	1.71	5.8	e*

pO₂ OR

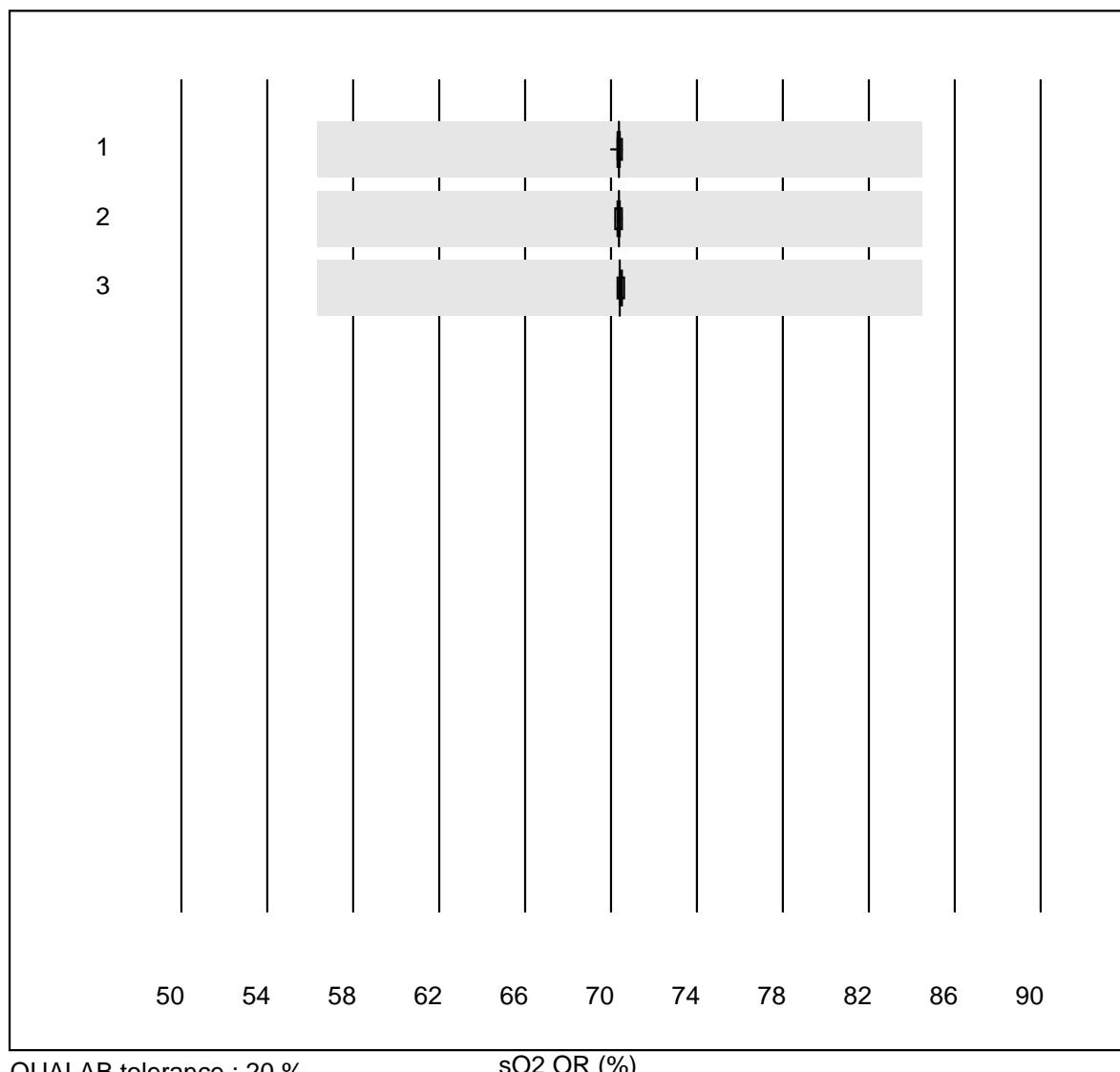
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	ABL700/800 Radiomete	77	92.2	0.0	7.8	10.04	4.5	e
2	Radiometer NPT-7	6	100.0	0.0	0.0	8.44	5.6	e*
3	ABL 90	25	72.0	8.0	20.0	7.36	6.2	e
4	ABL 80 / Coox	13	84.6	7.7	7.7	8.54	11.9	e*
5	ABL 5	6	100.0	0.0	0.0	7.97	2.5	e

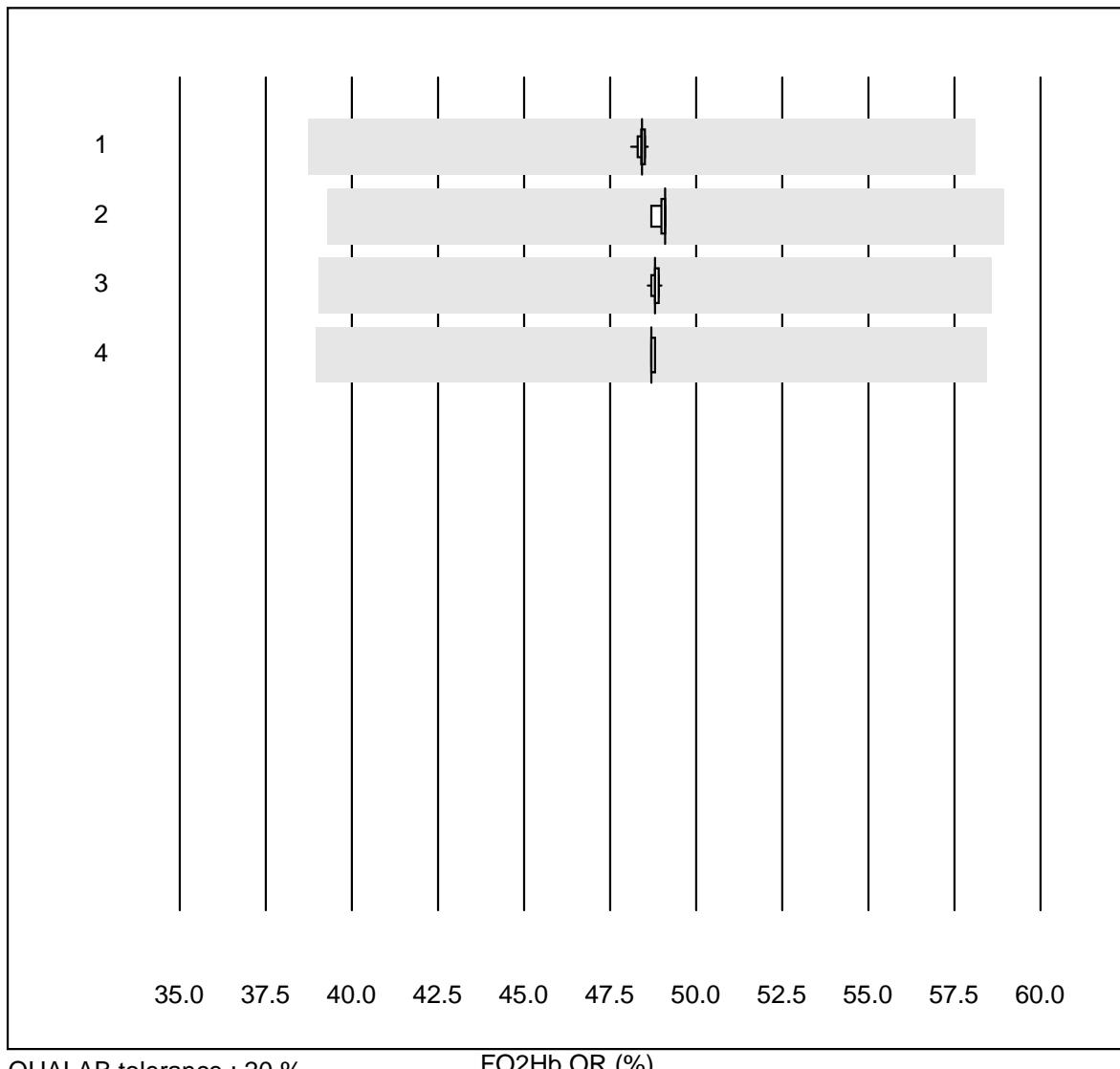
ctHb OR

QUALAB tolerance : 9 %

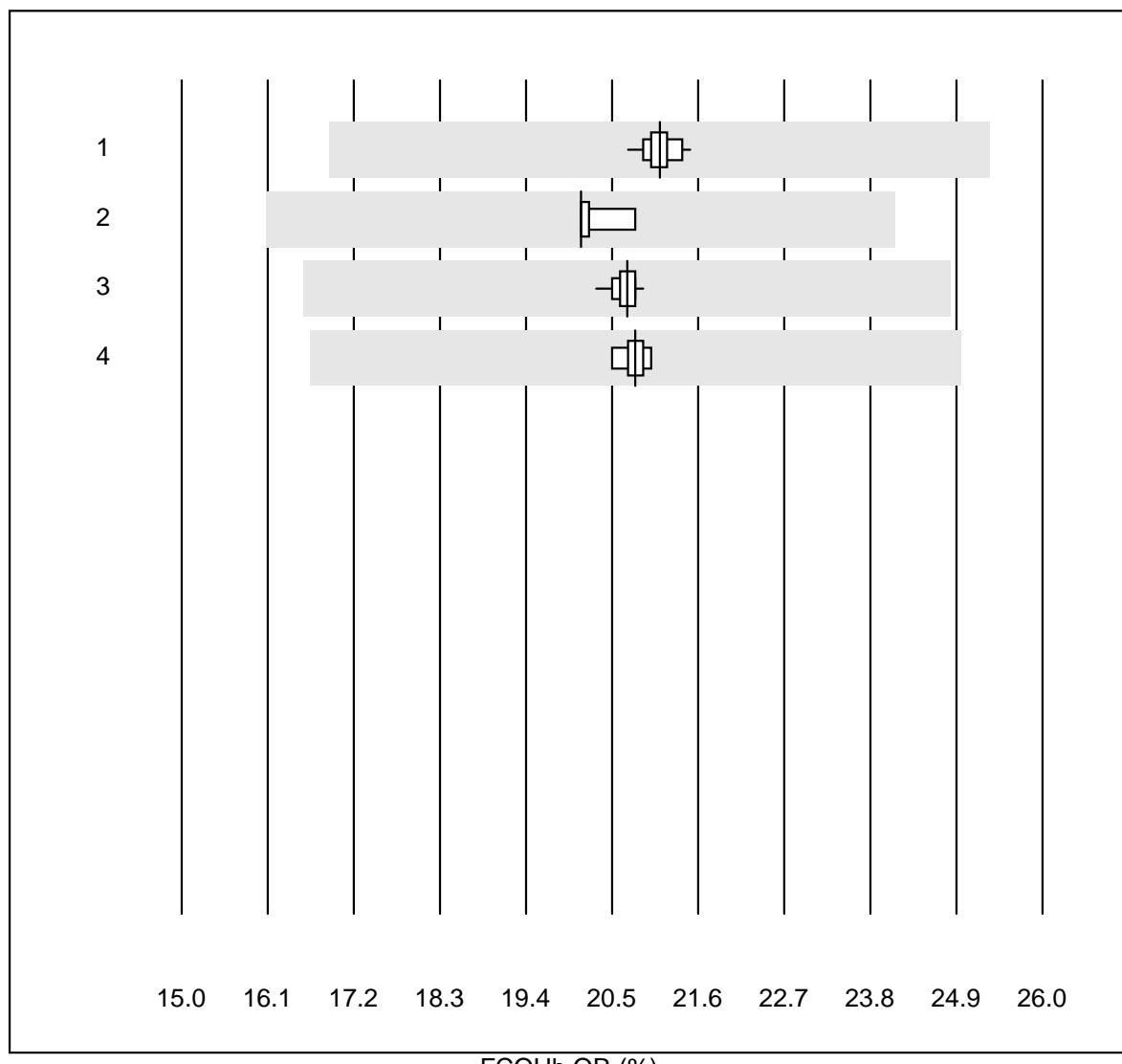
ctHb OR (g/l)

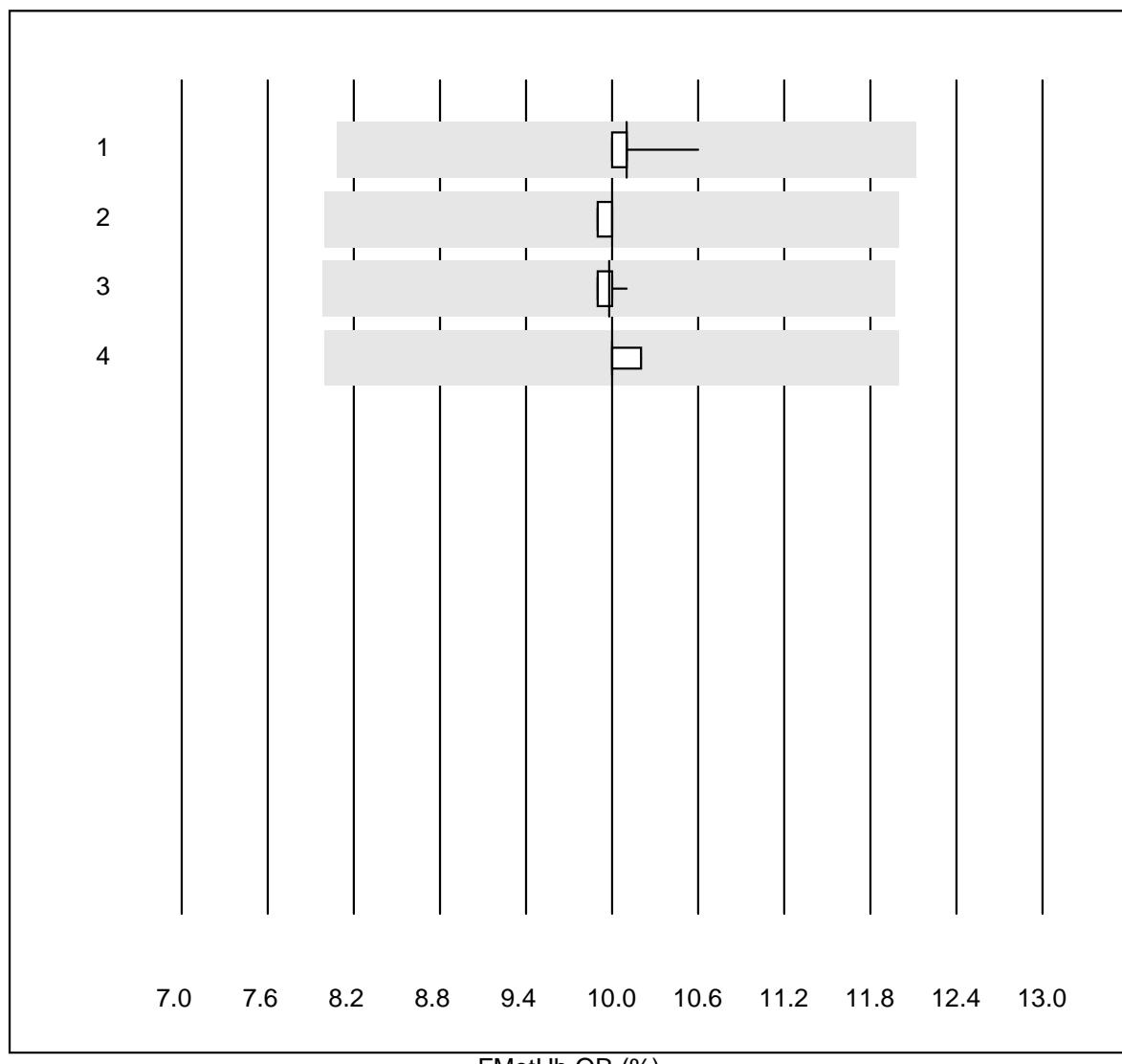
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	ABL700/800 Radiomete	70	95.7	0.0	4.3	194.8	2.3	e
2	Radiometer NPT-7	4	75.0	0.0	25.0	298.1	0.7	e
3	ABL 90	25	92.0	0.0	8.0	195.8	0.7	e
4	ABL 80 / Coox	9	100.0	0.0	0.0	197.0	1.7	e

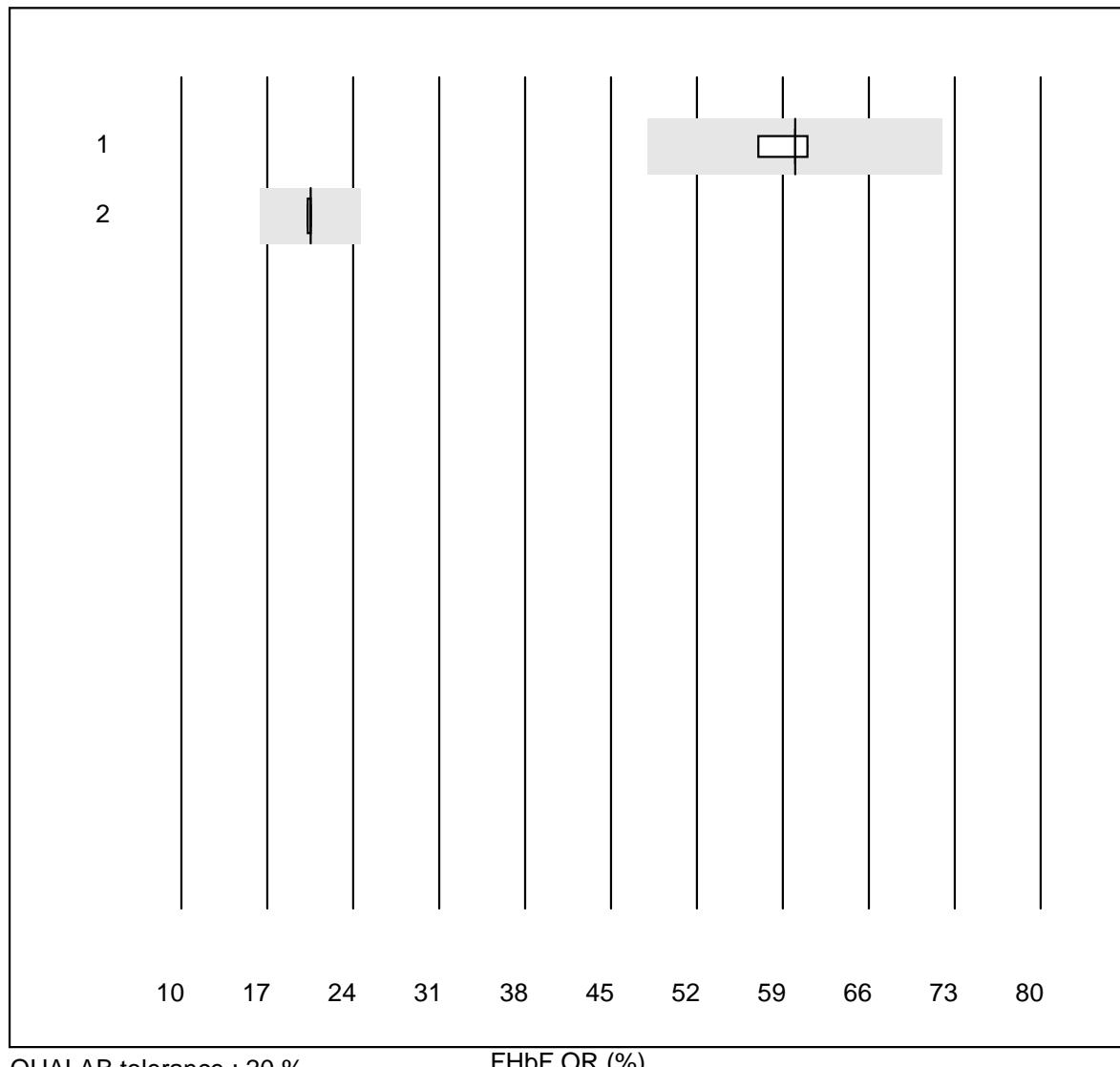
sO₂ OR

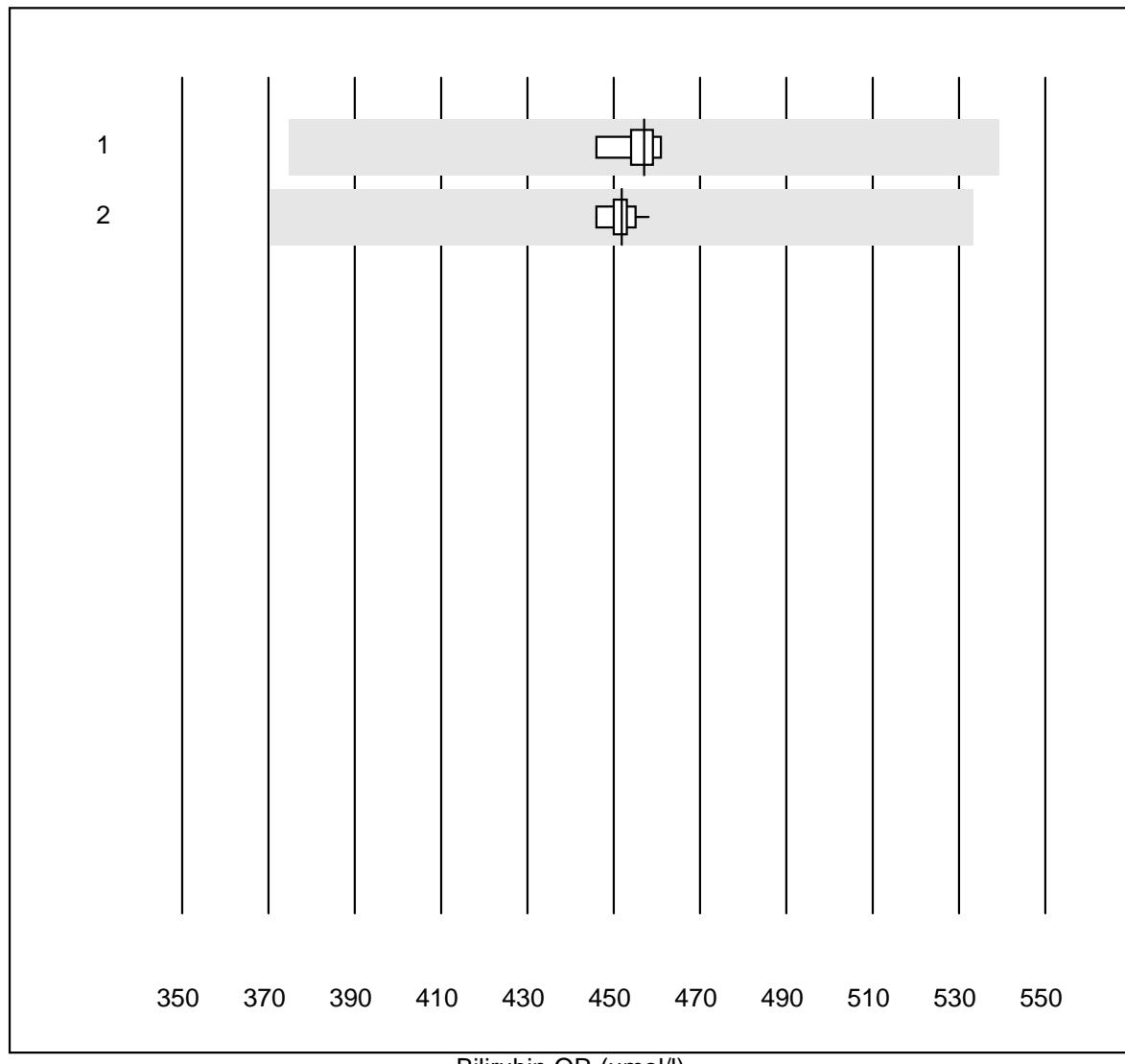
FO2Hb OR

No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	ABL700/800 Radiomete	49	100.0	0.0	0.0	48.422	0.2	e
2	Radiometer NPT-7	5	100.0	0.0	0.0	49.100	0.4	e
3	ABL 90	23	100.0	0.0	0.0	48.804	0.2	e
4	ABL 80 / Coox	9	100.0	0.0	0.0	48.700	0.1	e

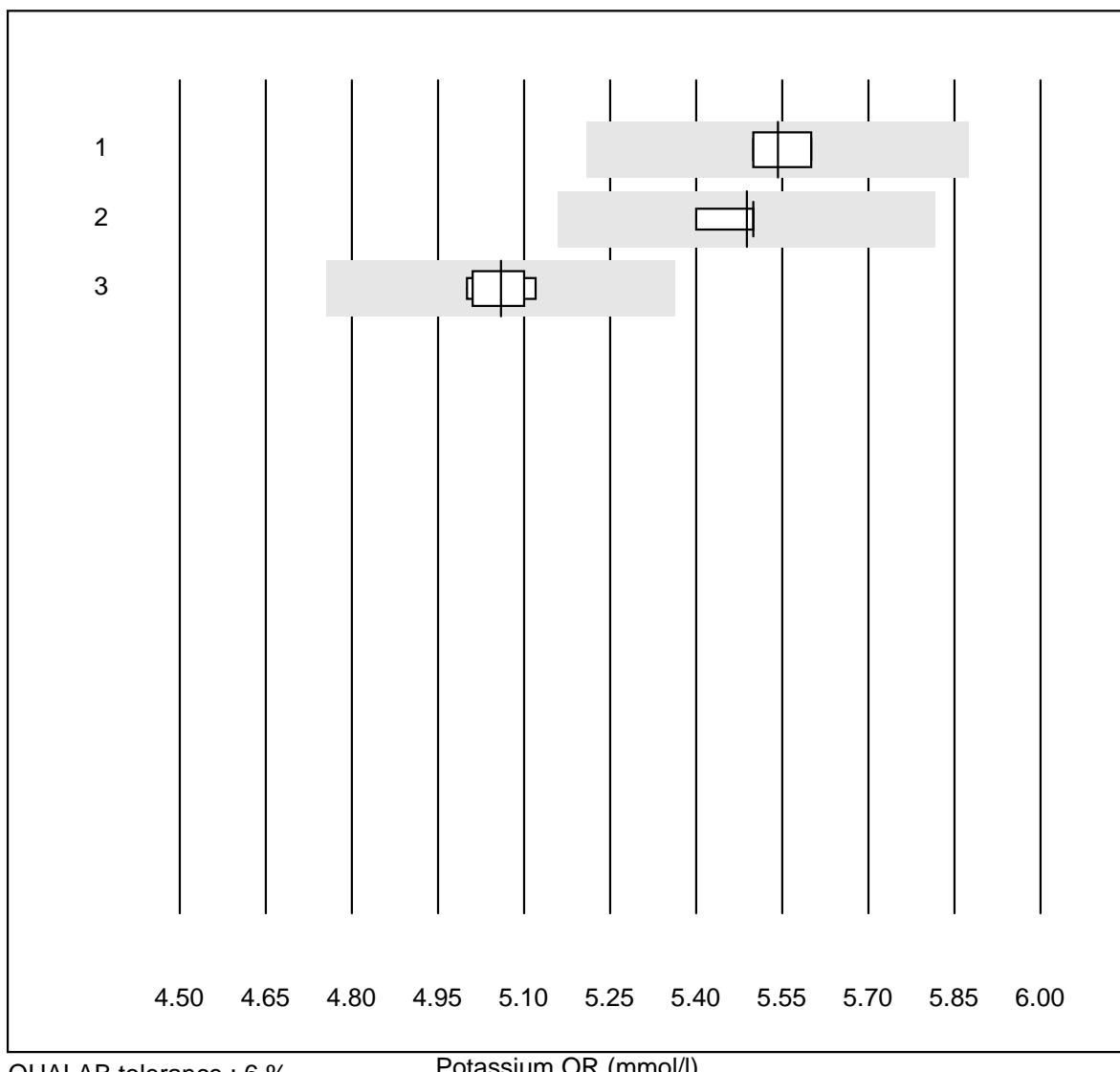
FCOHb OR

FMetHb OR

FHbF OR

Bilirubin OR

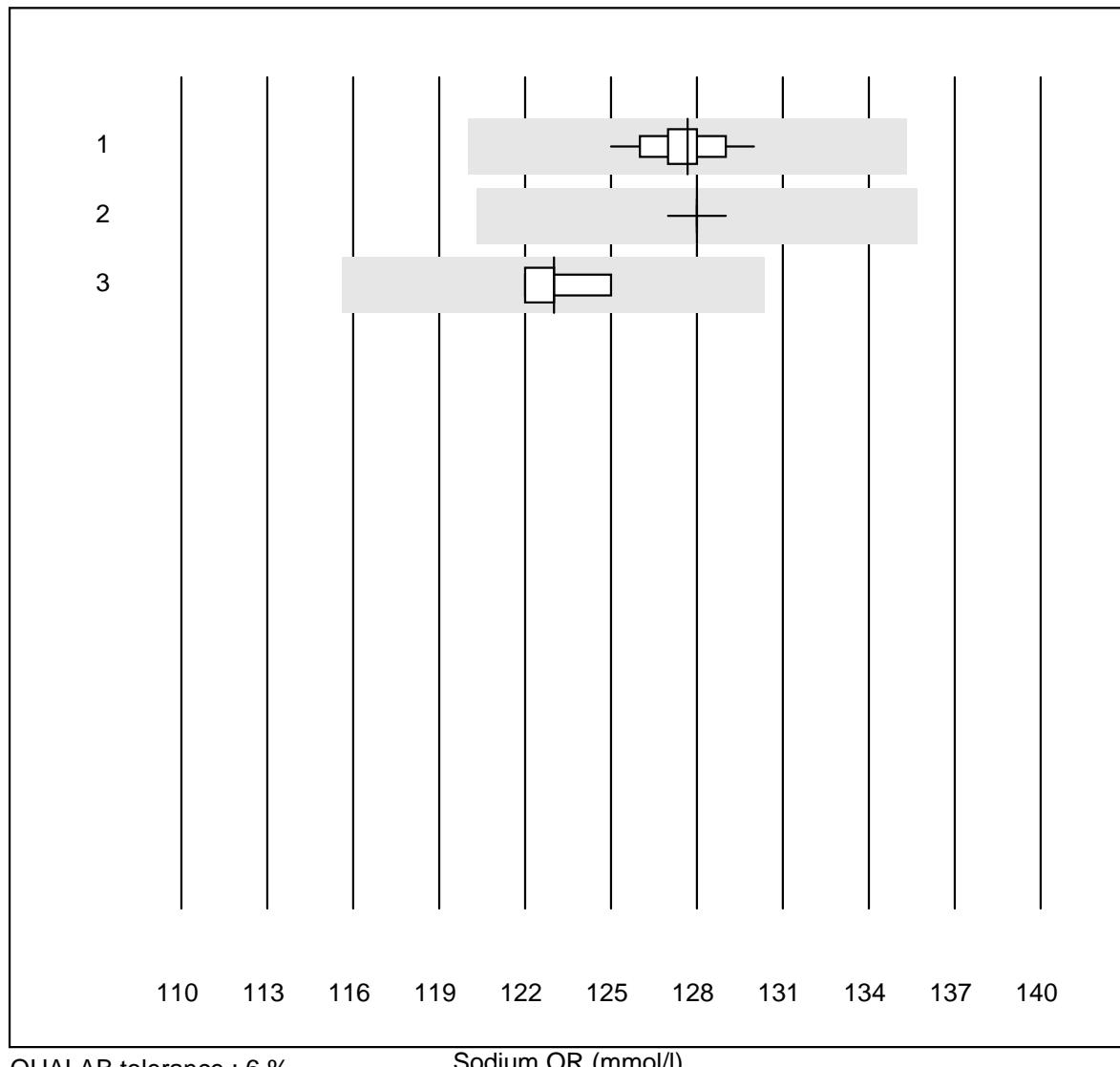
Potassium OR



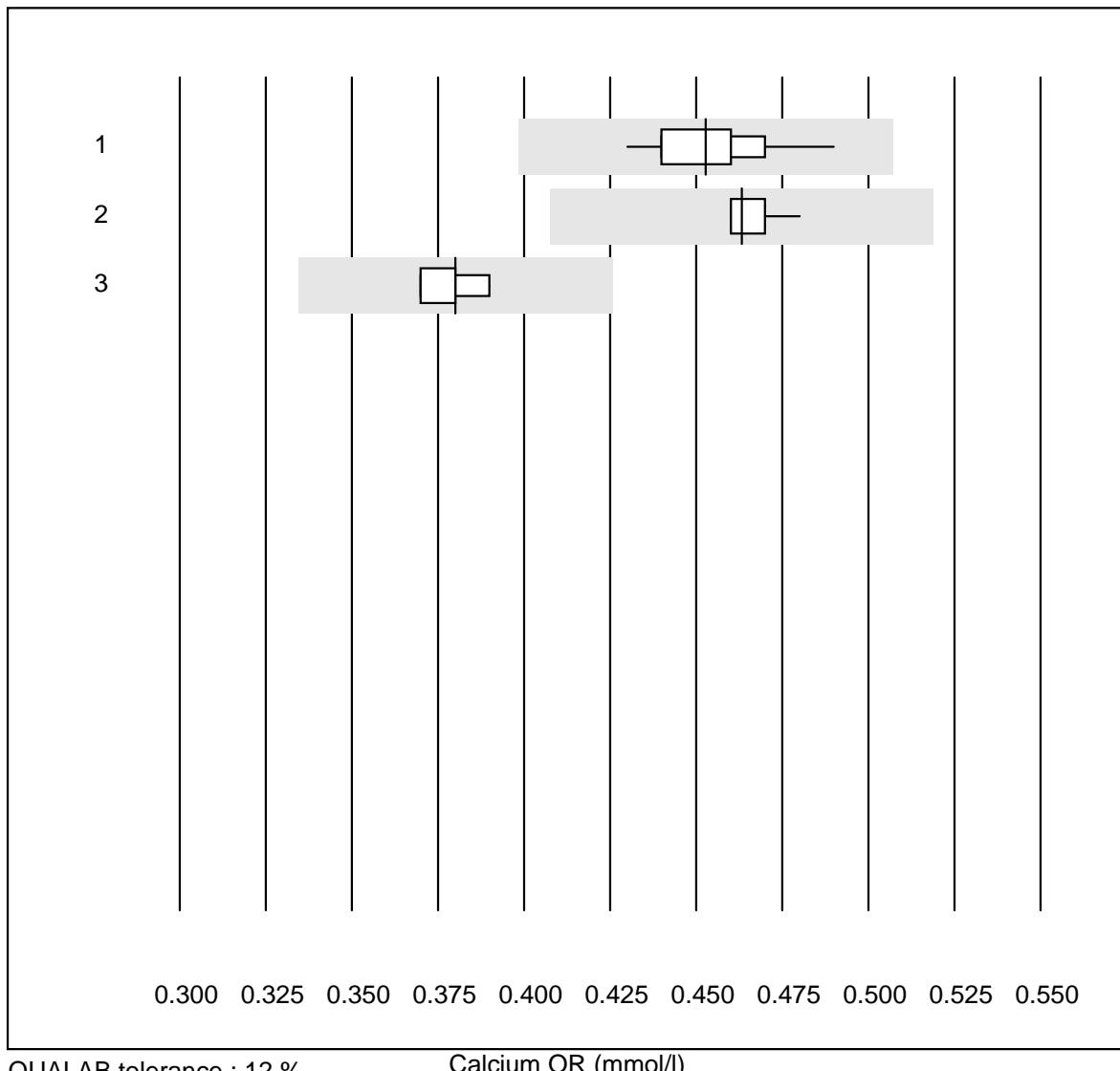
QUALAB tolerance : 6 %

Potassium OR (mmol/l)

No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	ABL700/800 Radiomete	65	98.5	0.0	1.5	5.5	0.9	e
2	ABL 90	25	100.0	0.0	0.0	5.5	0.6	e
3	ABL 80 / Coox	5	100.0	0.0	0.0	5.1	1.0	e

Sodium OR

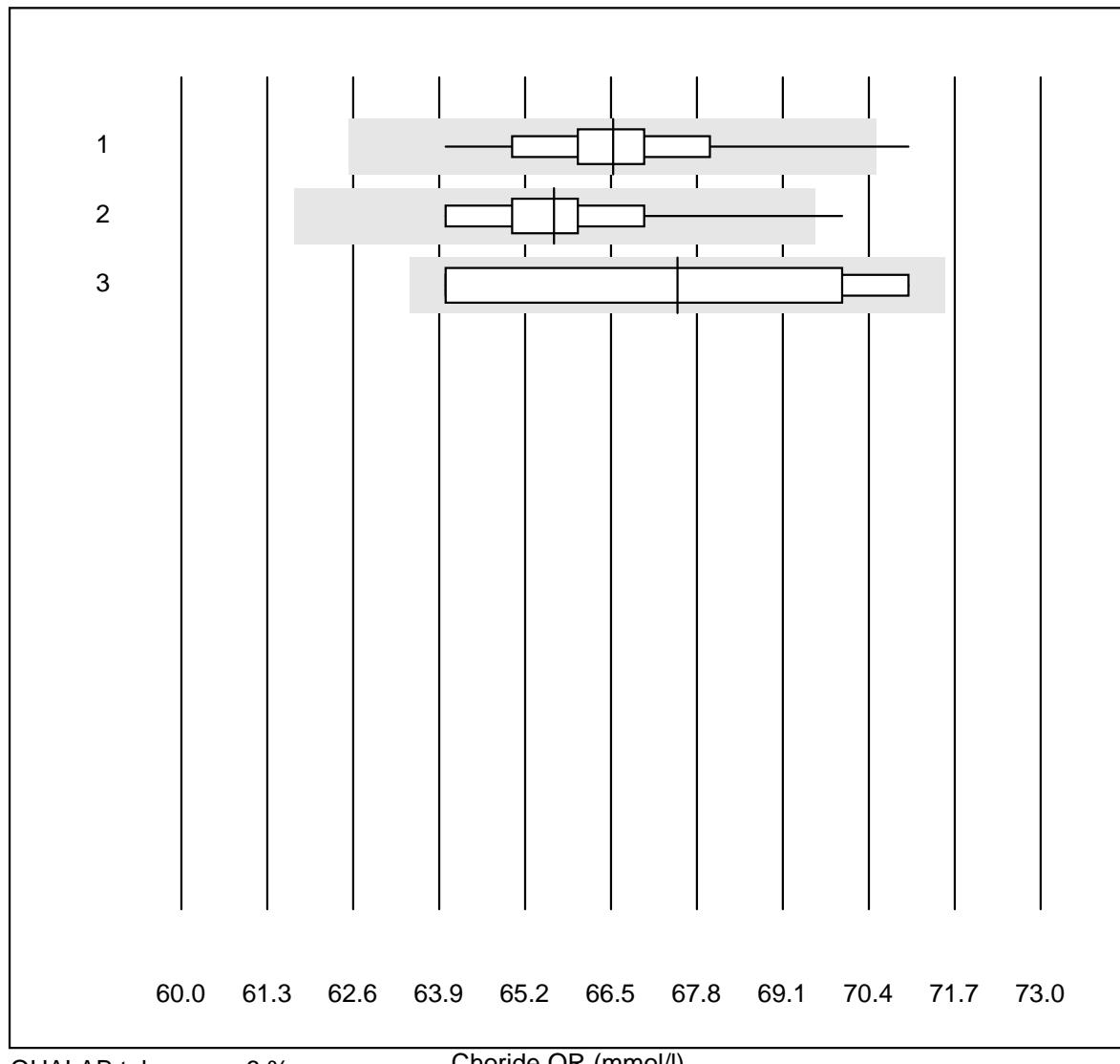
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	ABL700/800 Radiomete	63	100.0	0.0	0.0	127.7	0.7	e
2	ABL 90	25	100.0	0.0	0.0	128.0	0.3	e
3	ABL 80 / Coox	4	100.0	0.0	0.0	123.0	1.0	e

Calcium OR

QUALAB tolerance : 12 %

Calcium OR (mmol/l)

No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	ABL700/800 Radiomete	64	98.4	0.0	1.6	0.45	2.8	e
2	ABL 90	25	100.0	0.0	0.0	0.46	1.2	e
3	ABL 80 / Coox	5	100.0	0.0	0.0	0.38	2.2	e

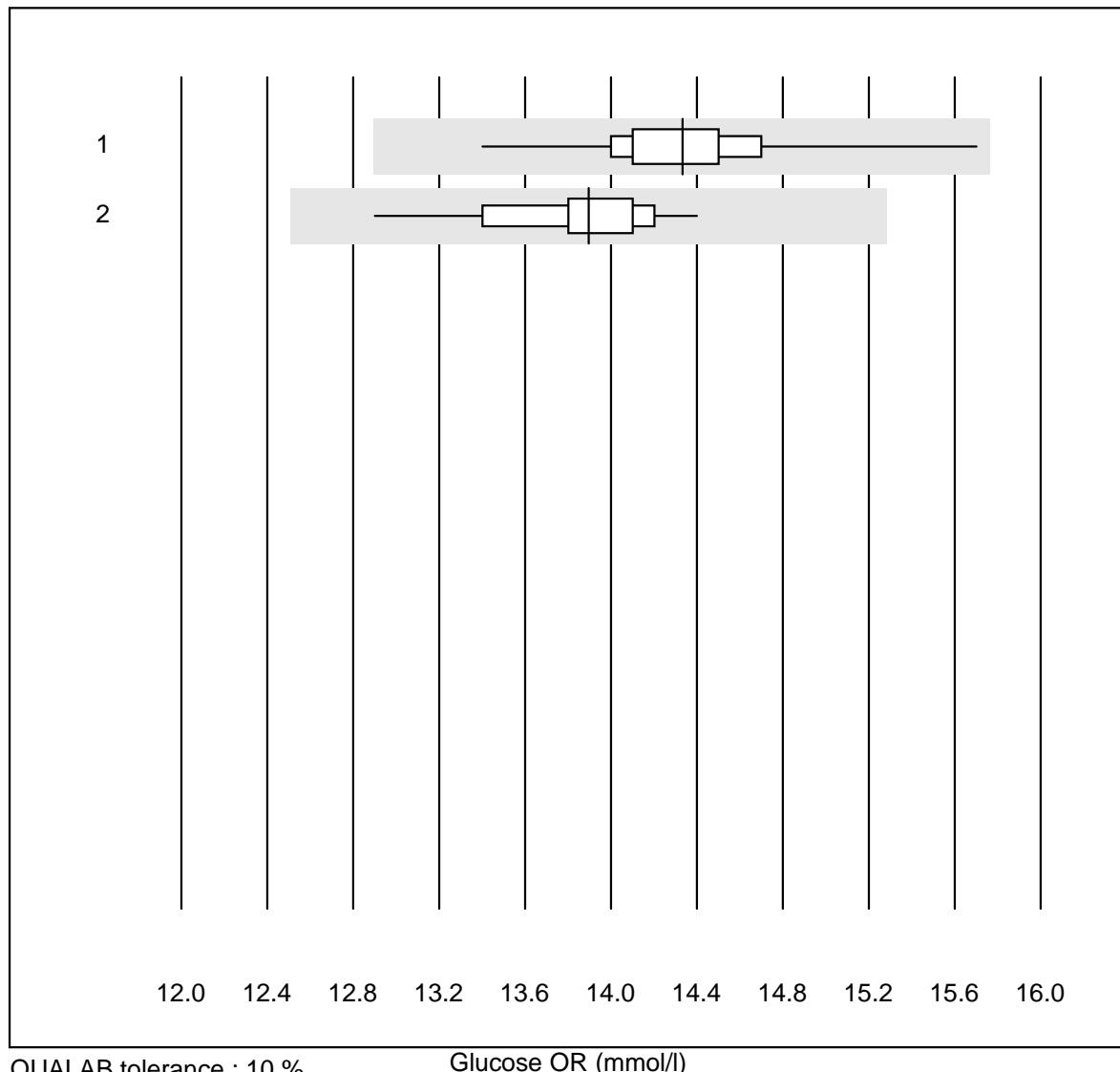
Choride OR

QUALAB tolerance : 6 %

Choride OR (mmol/l)

No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	ABL700/800 Radiomete	53	98.1	1.9	0.0	66.53	1.9	e
2	ABL 90	25	96.0	4.0	0.0	65.64	2.1	e
3	ABL 80 / Coox	4	100.0	0.0	0.0	67.50	5.2	e*

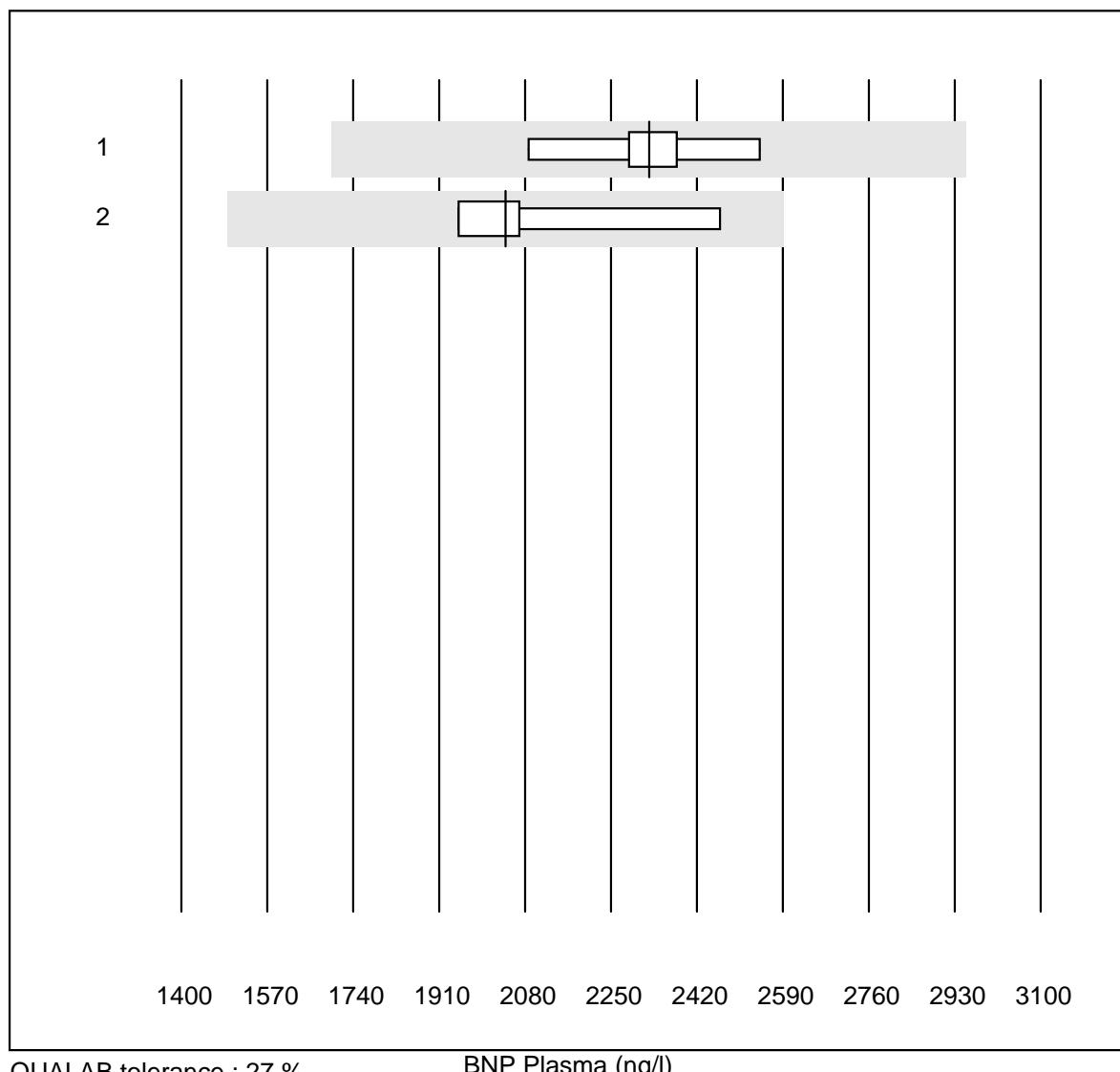
Glucose OR

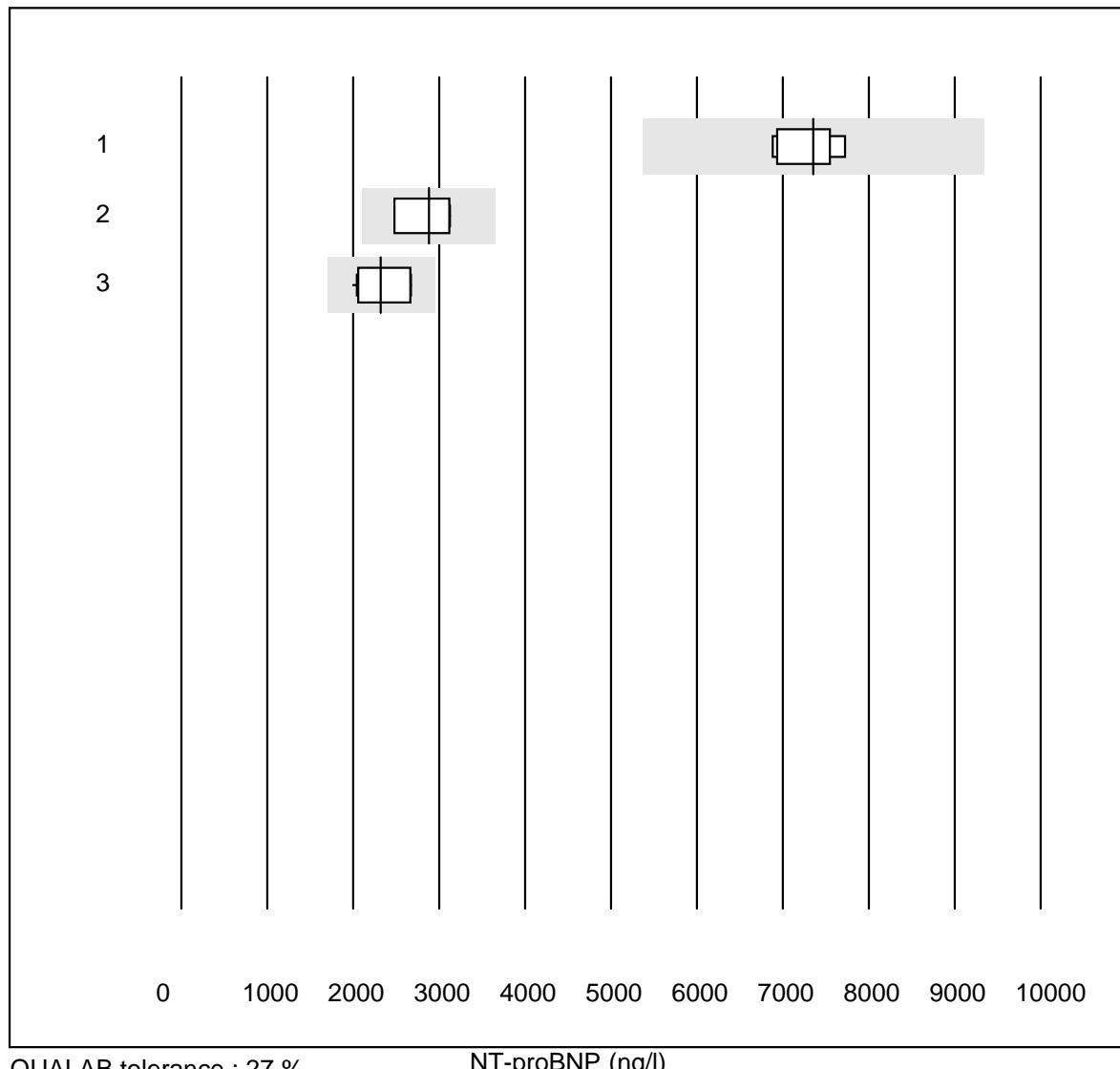


Lactate OR

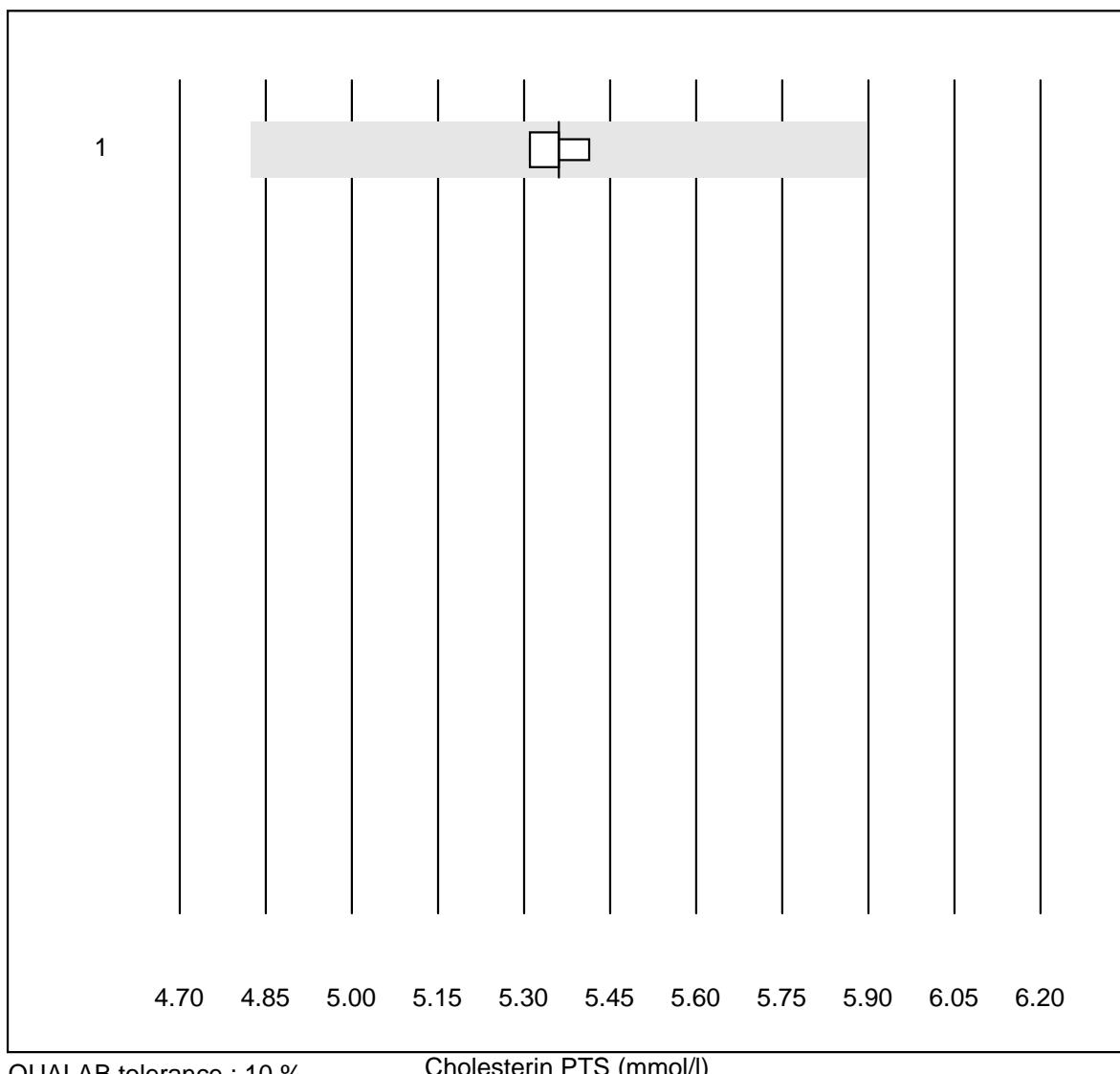


No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	ABL700/800 Radiomete	67	95.5	1.5	3.0	10.38	4.1	e
2	ABL 90	25	96.0	4.0	0.0	10.35	3.7	e

BNP Plasma

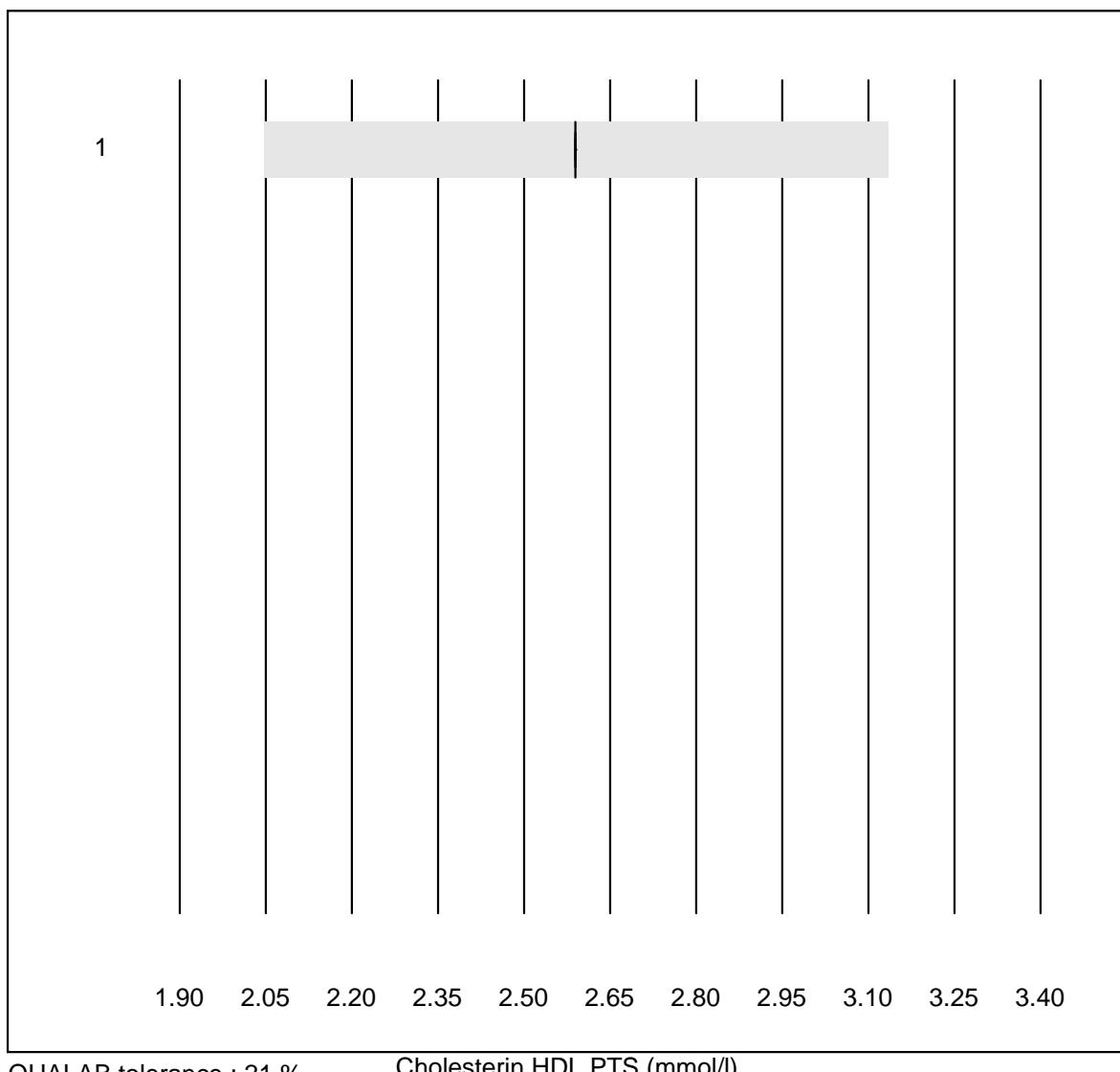
NT-proBNP

Cholesterin PTS



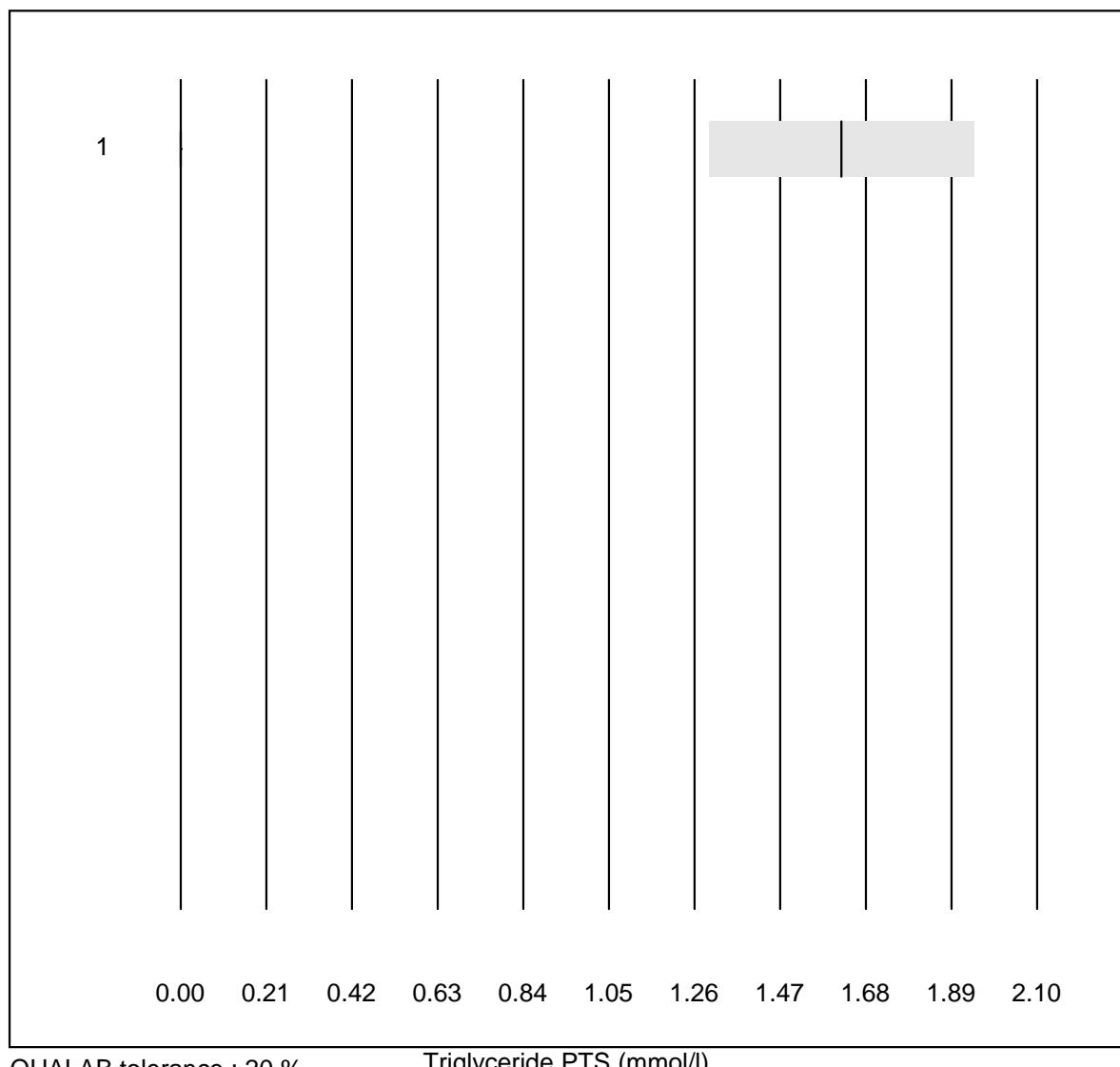
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 CardioChek	4	100.0	0.0	0.0	5.4	0.8	e

Cholesterin HDL PTS

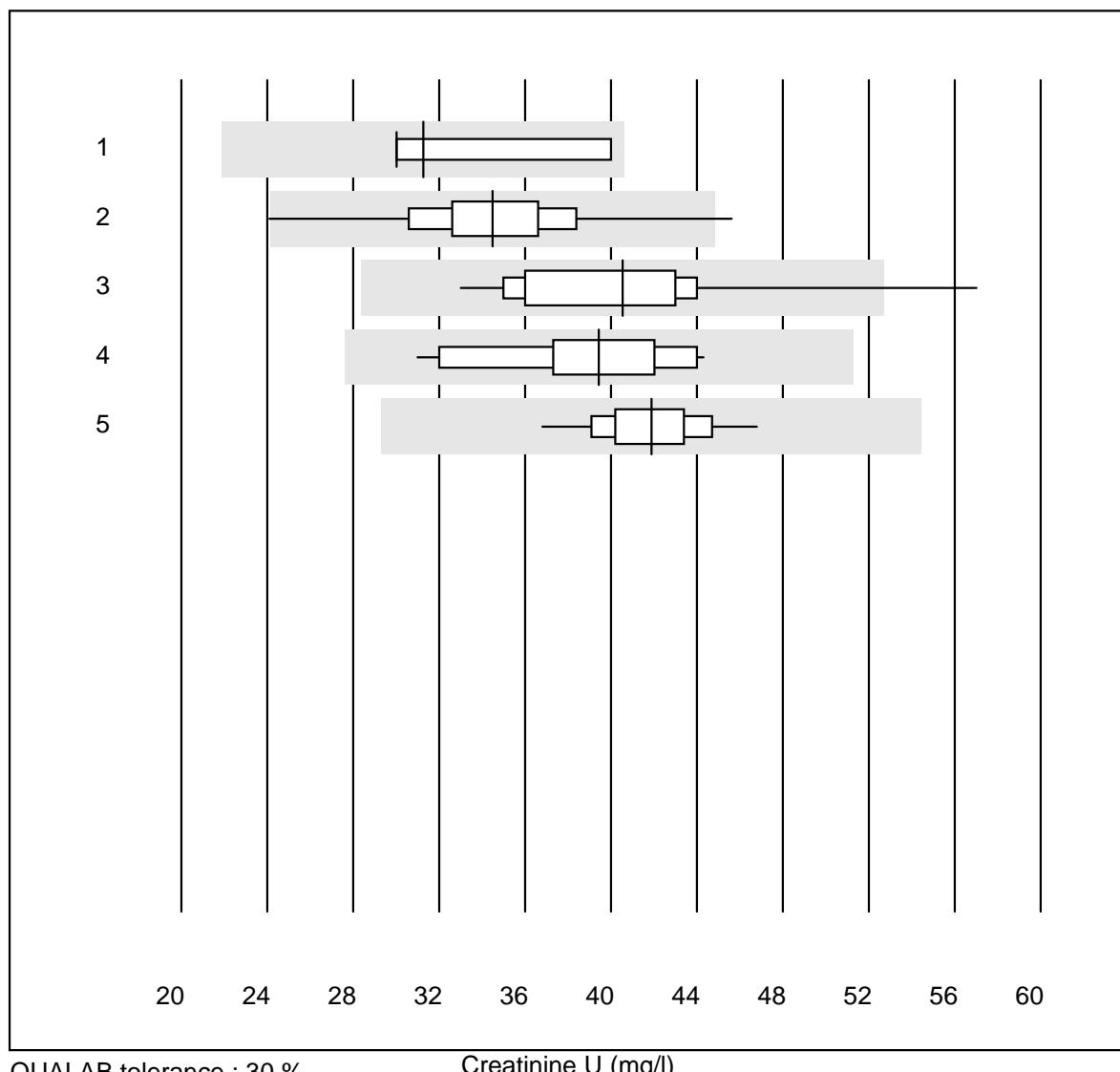


No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 CardioChek	4	100.0	0.0	0.0	2.6	0.0	e

Triglyceride PTS

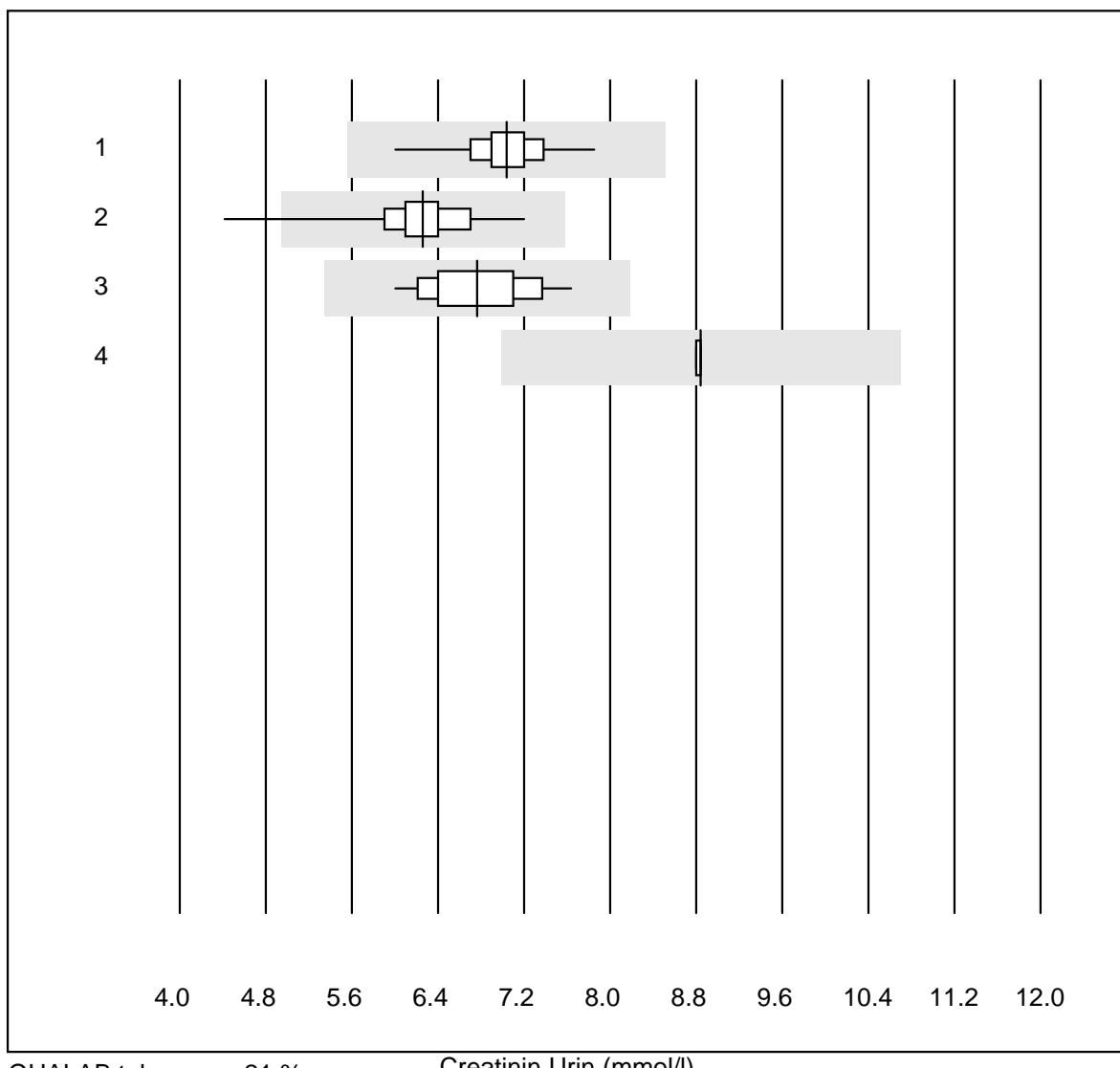


No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 CardioChek	4	0.0	0.0	100.0	1.62	0.0	e

Creatinine U

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Siemens Clinitek	10	90.0	0.0	10.0	31.3	10.7	e
2 Afinion	323	97.5	0.6	1.9	34.5	9.2	e
3 NycoCard	16	74.9	6.3	18.8	40.5	15.2	e*
4 Turbidimetry	16	100.0	0.0	0.0	39.4	10.0	e
5 DCA2000/Vantage	112	97.3	0.0	2.7	41.9	5.3	e

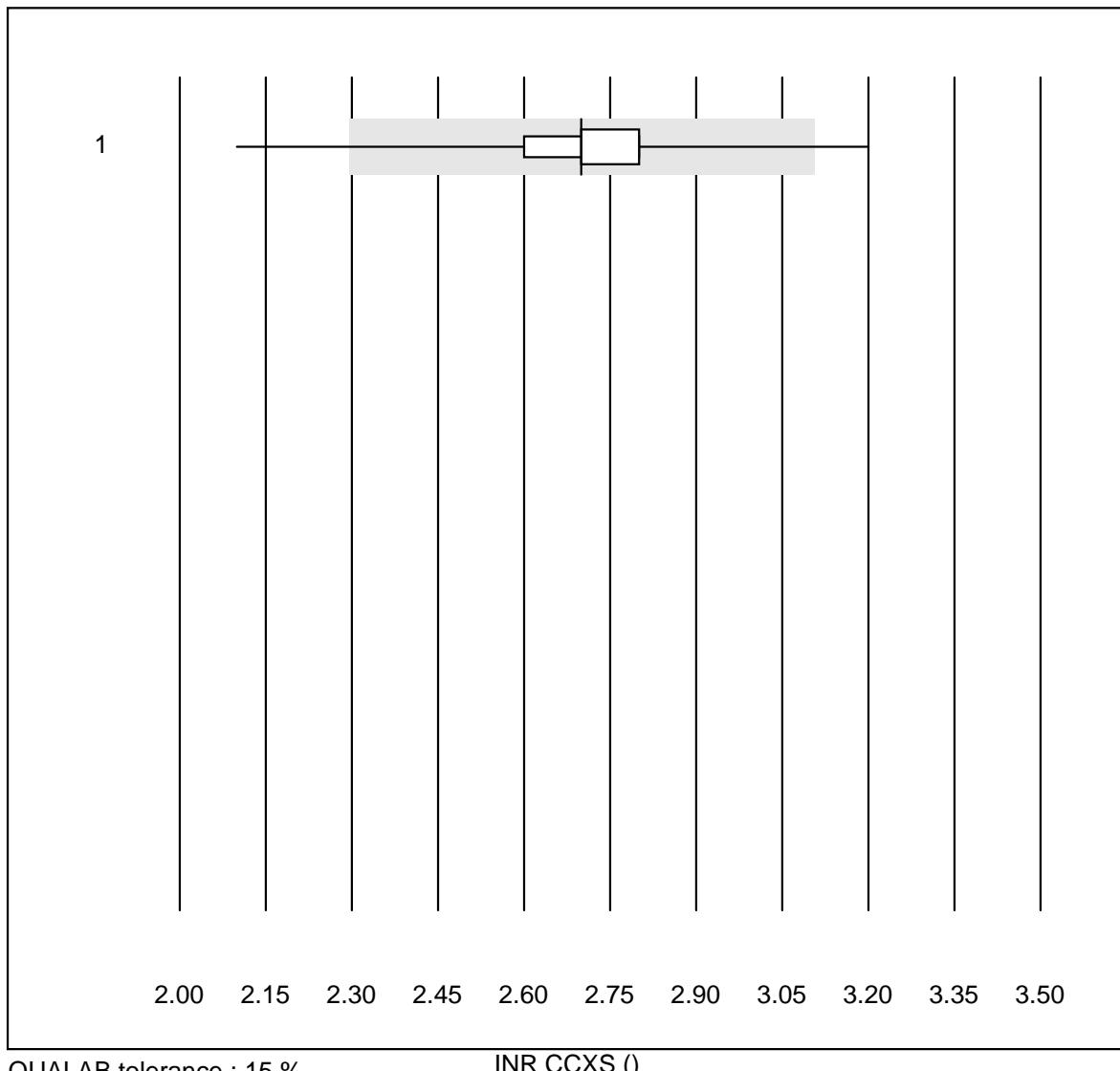
Creatinin Urin



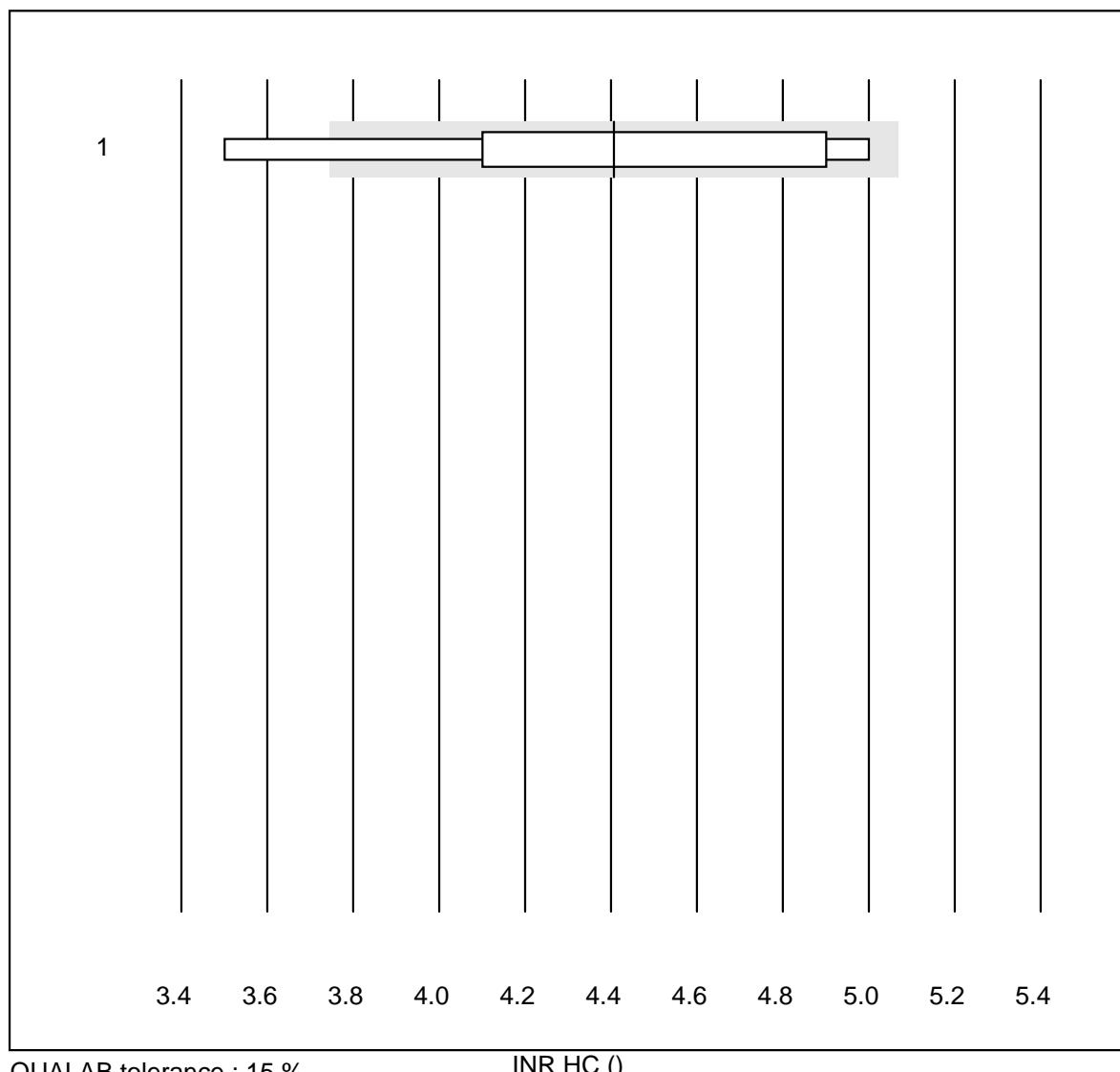
QUALAB tolerance : 21 %

Creatinin Urin (mmol/l)

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 DCA2000/Vantage	112	97.3	0.0	2.7	7.0	4.0	e
2 Afinion	323	99.1	0.3	0.6	6.3	5.2	e
3 Standard chemistry	26	100.0	0.0	0.0	6.8	6.4	e
4 Siemens Clinitek	9	88.9	0.0	11.1	8.8	0.2	e

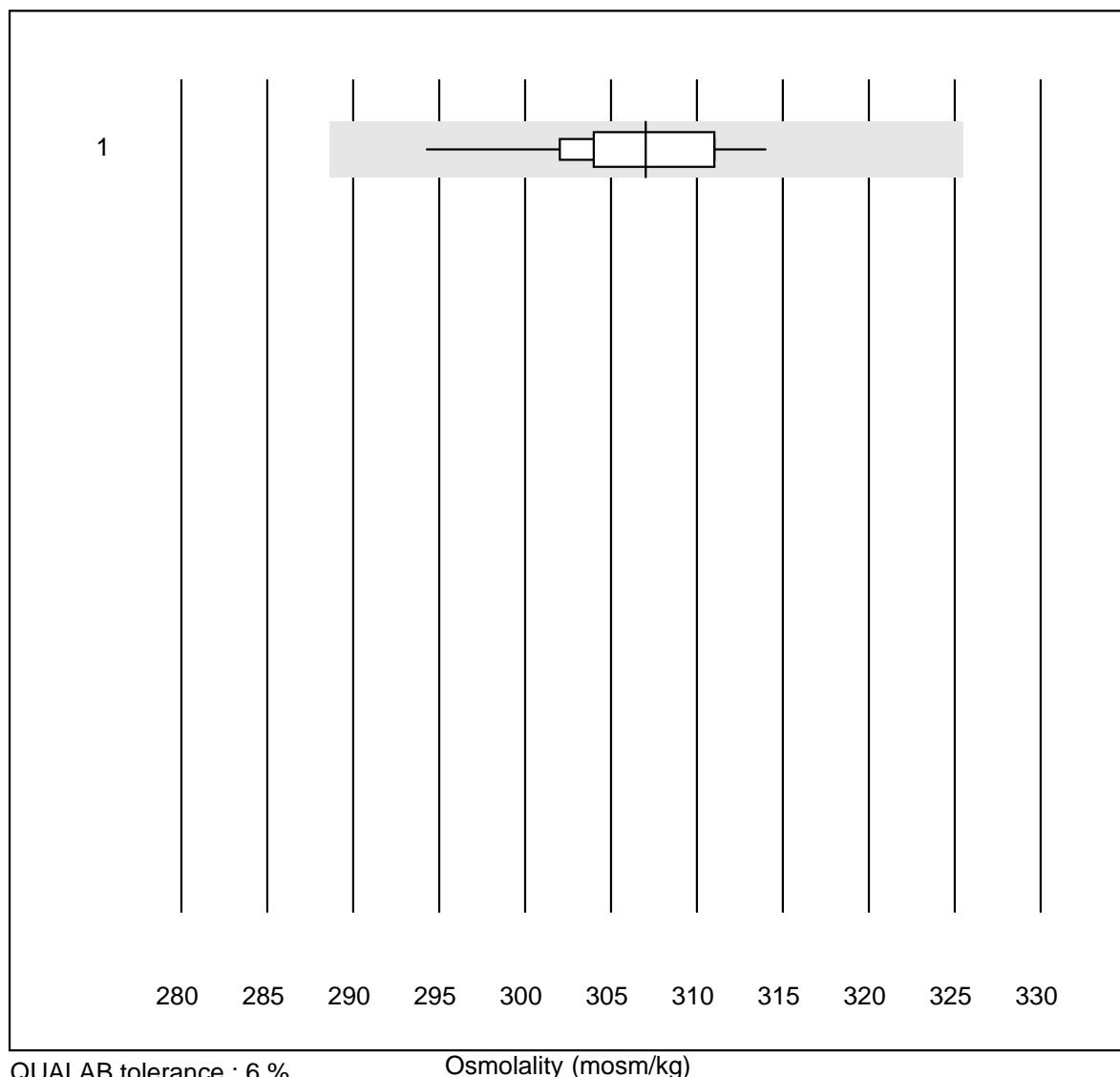
INR CCXS

No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	CoaguChek XS	2336	98.8	0.7	0.5	2.7	4.2	e

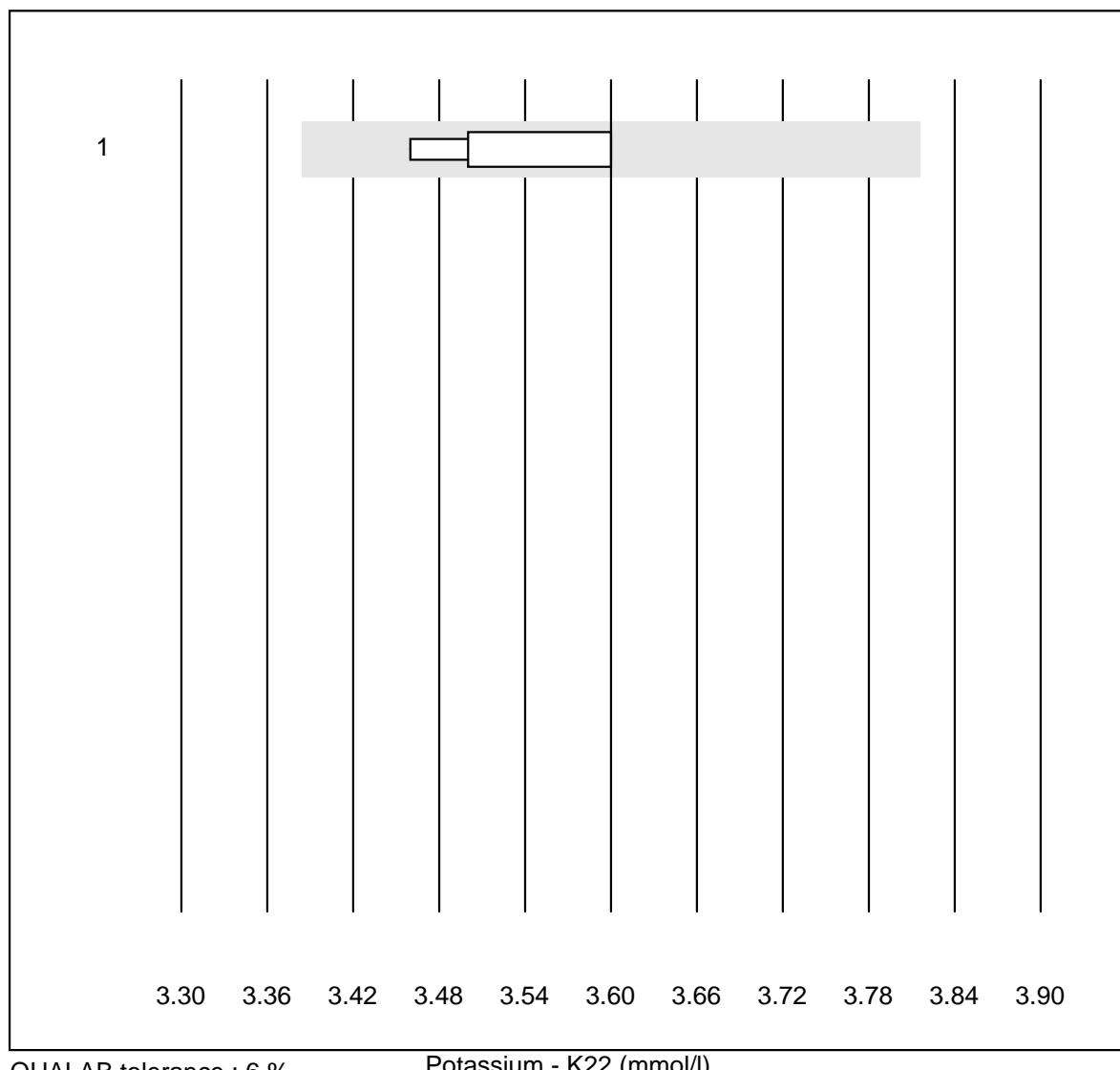
INR HC

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Hemochron j.	21	71.5	9.5	19.0	4.4	11.8	e*

Osmolality



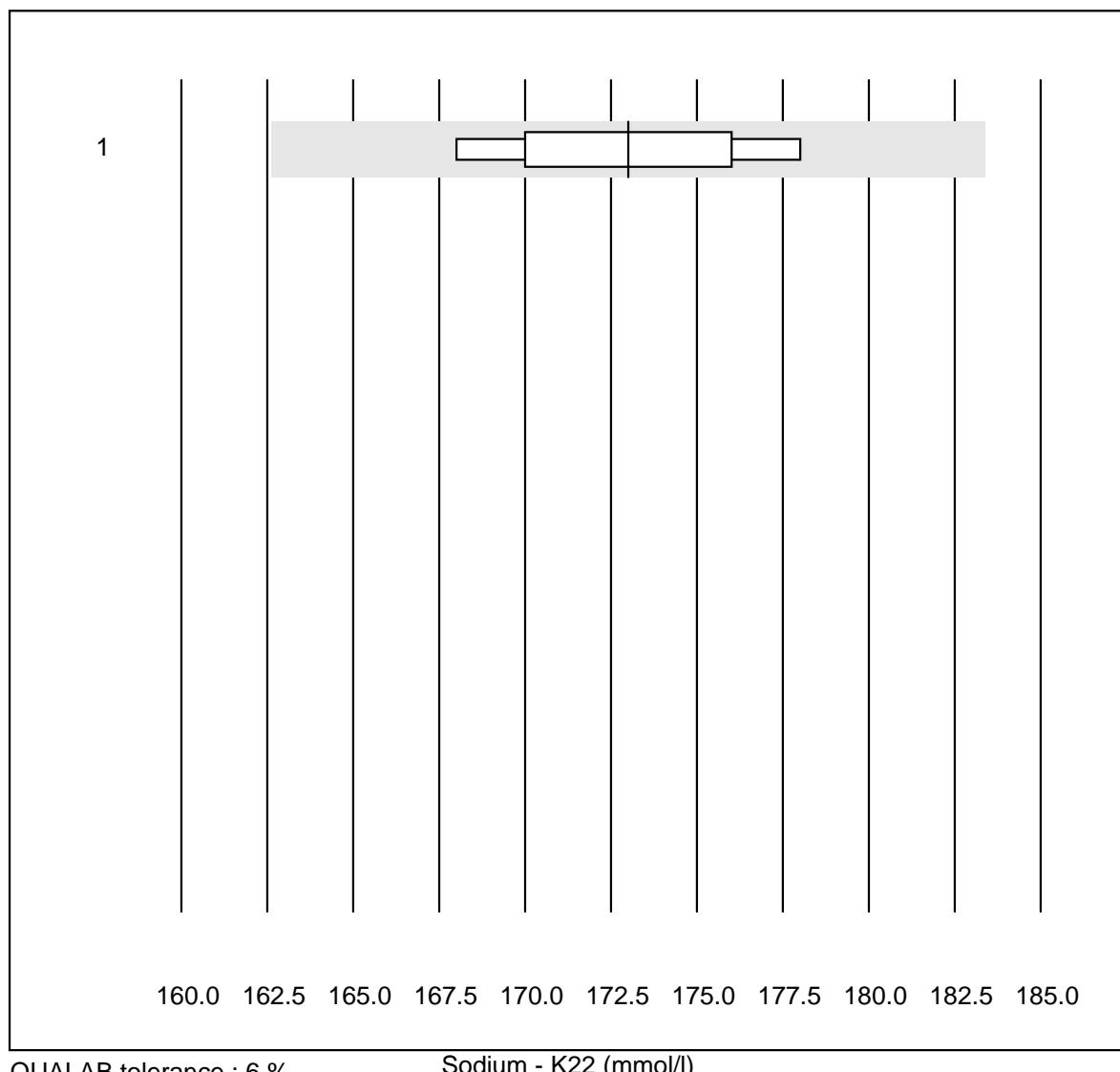
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Cryoskopy	11	100.0	0.0	0.0	307	1.8	e

Potassium - K22

No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	ISE	9	100.0	0.0	0.0	3.6	1.7	e

K22 Osmolality

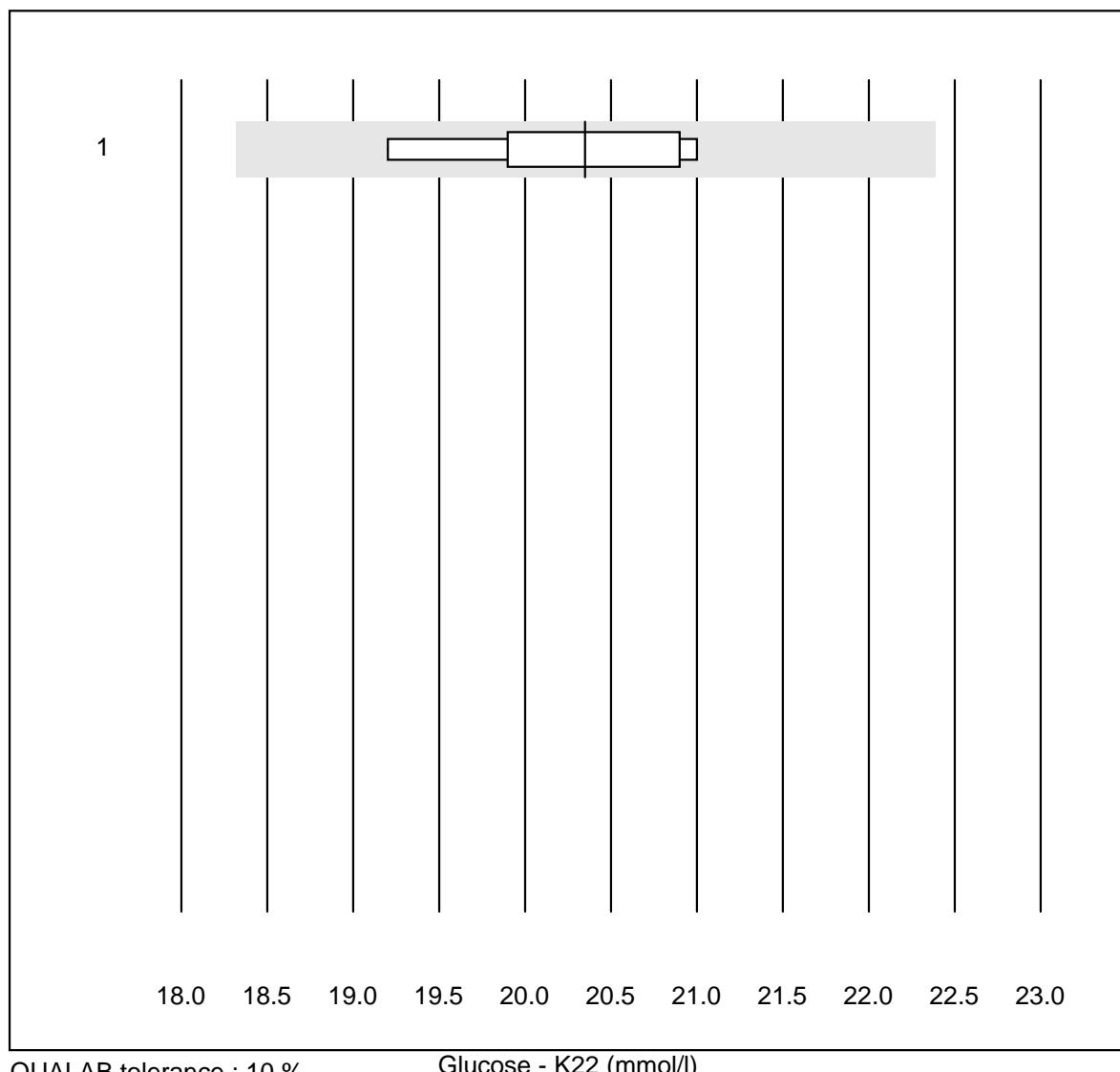
Sodium - K22



No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	ISE	8	100.0	0.0	0.0	173	2.1	e*

K22 Osmolality

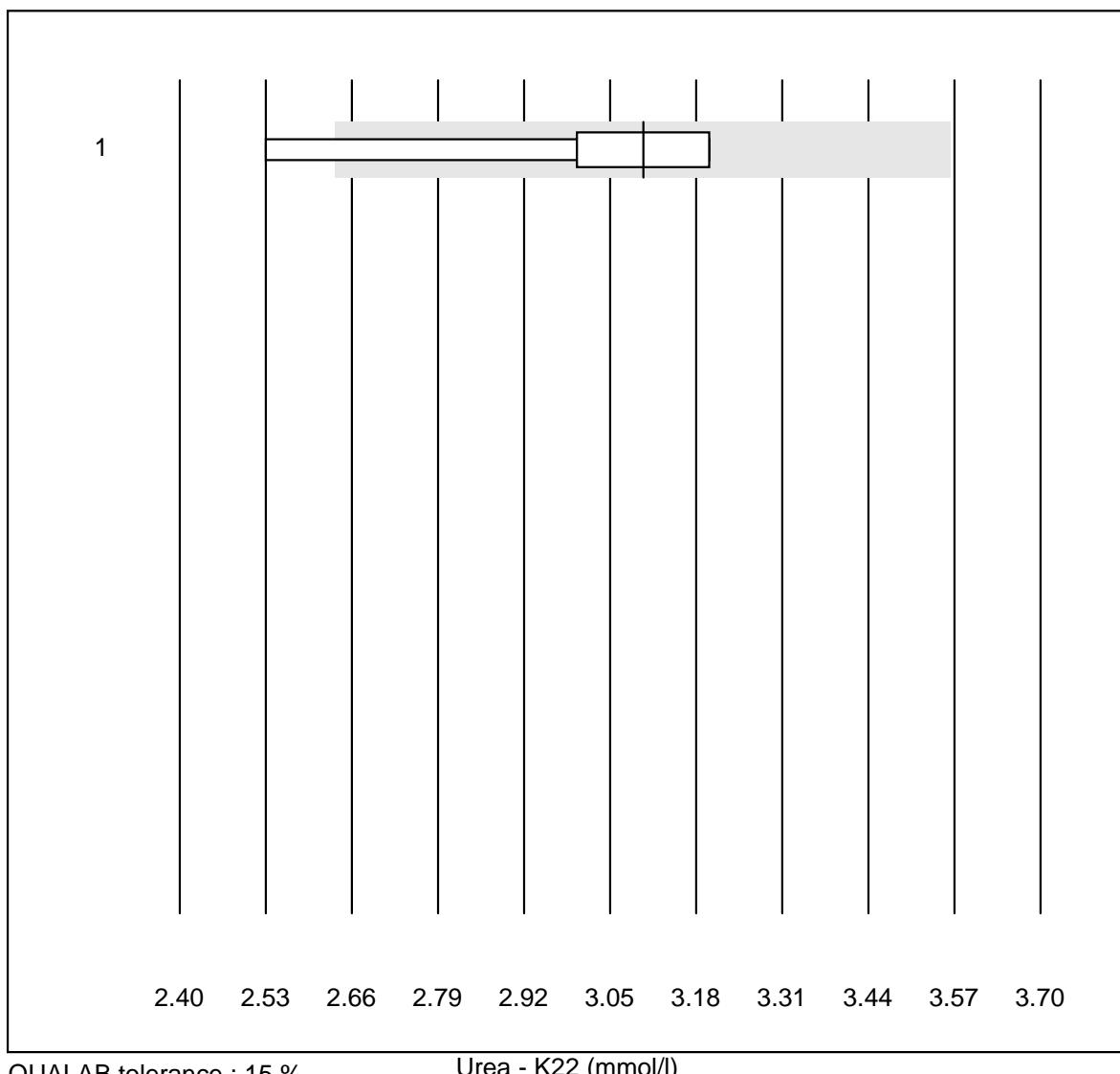
Glucose - K22



No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Standard chemistry	8	100.0	0.0	0.0	20.4	3.0	e

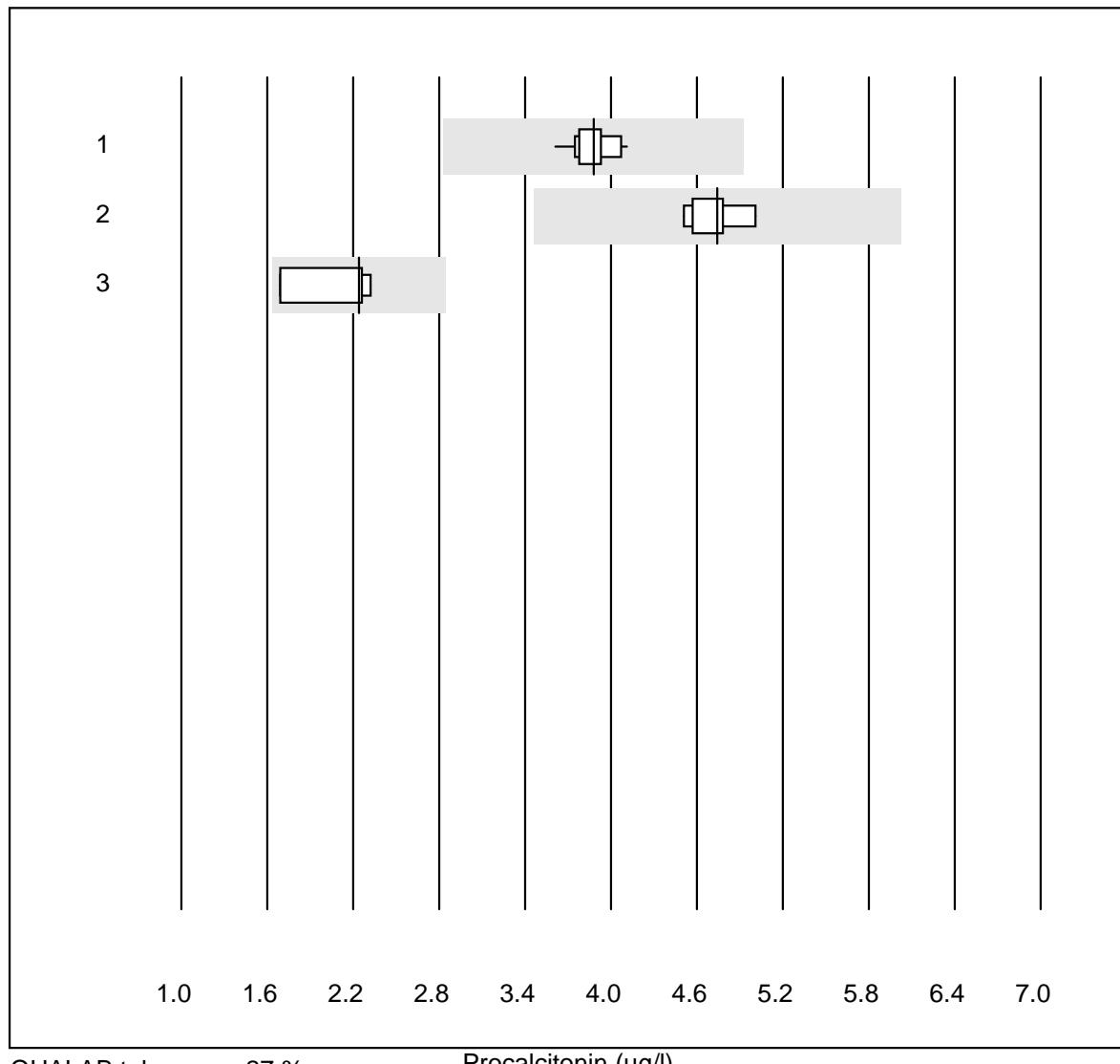
K22 Osmolality

Urea - K22

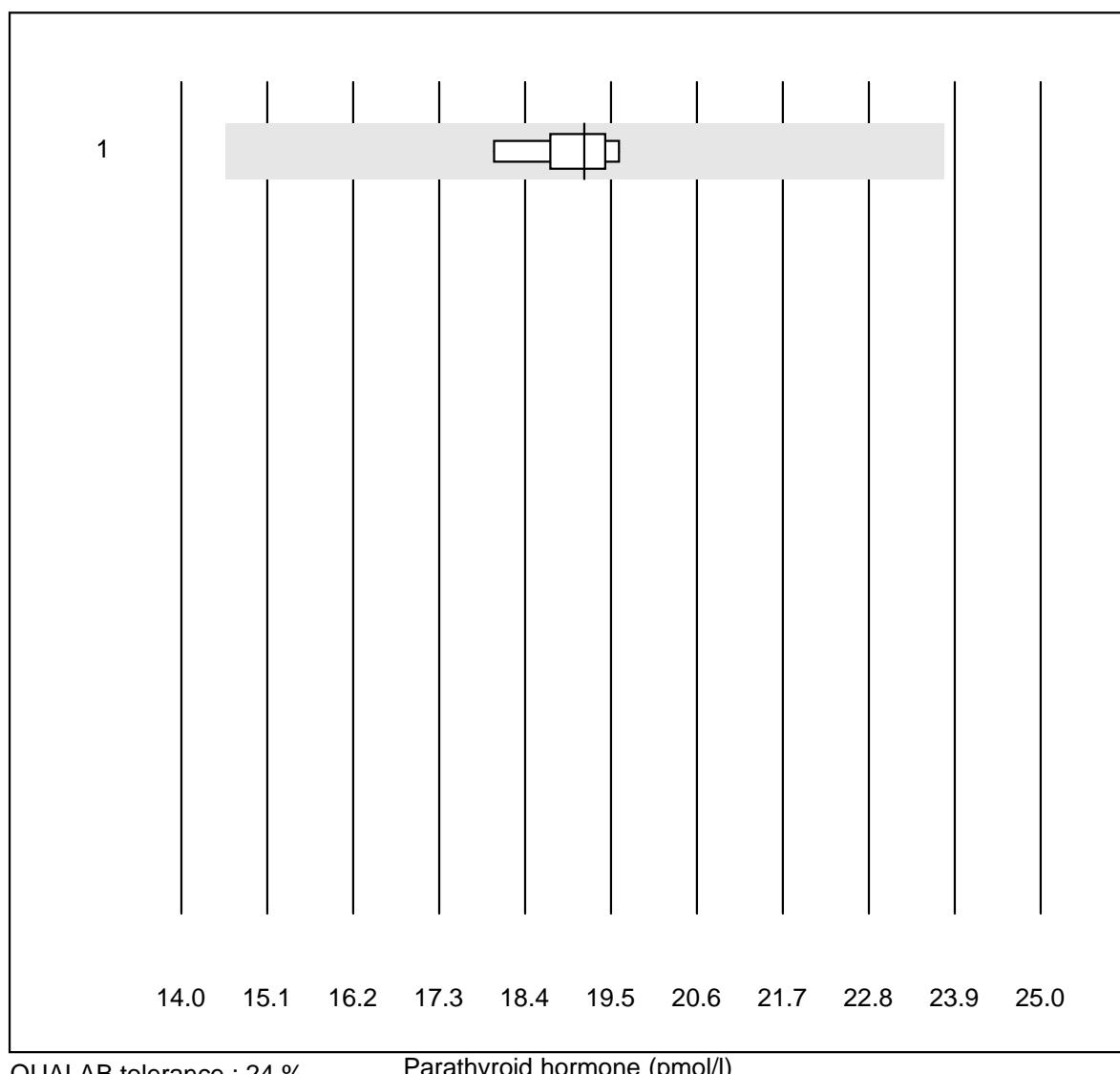


No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Standard chemistry	8	87.5	12.5	0.0	3.1	7.3	e*

Procalcitonin

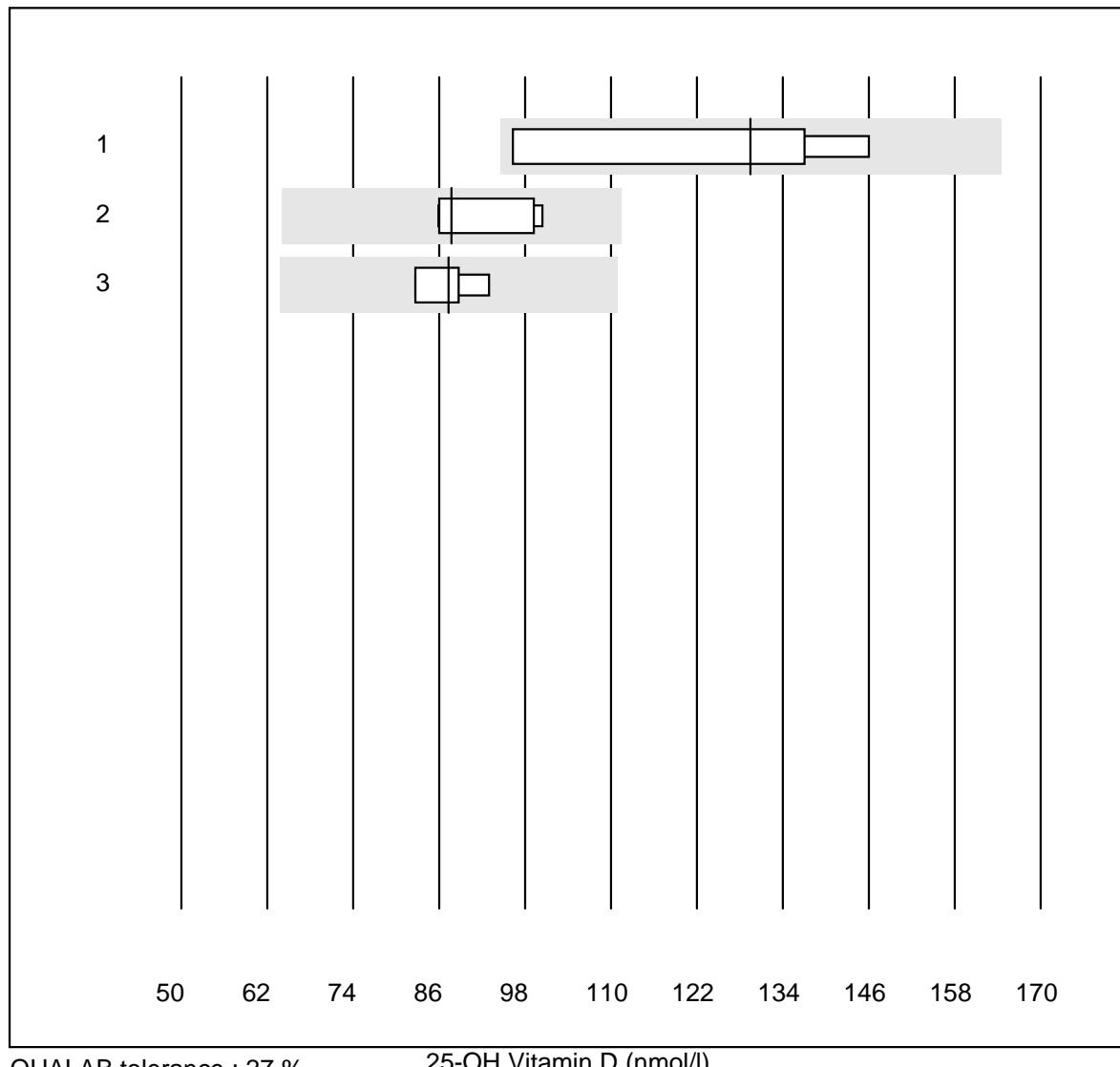


Parathyroid hormone



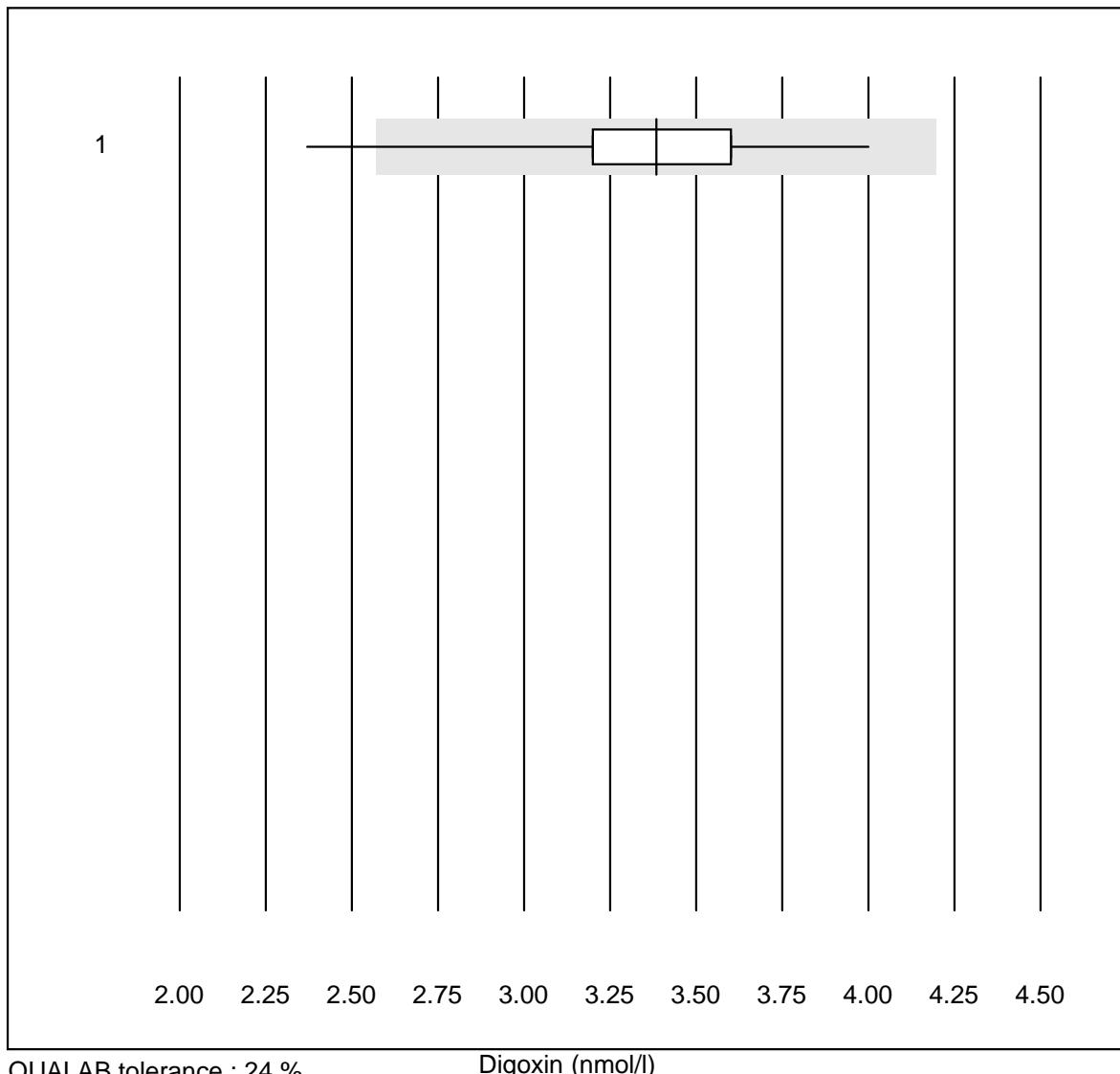
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Cobas PTH STAT	5	100.0	0.0	0.0	19.2	3.4	e

25-OH Vitamin D

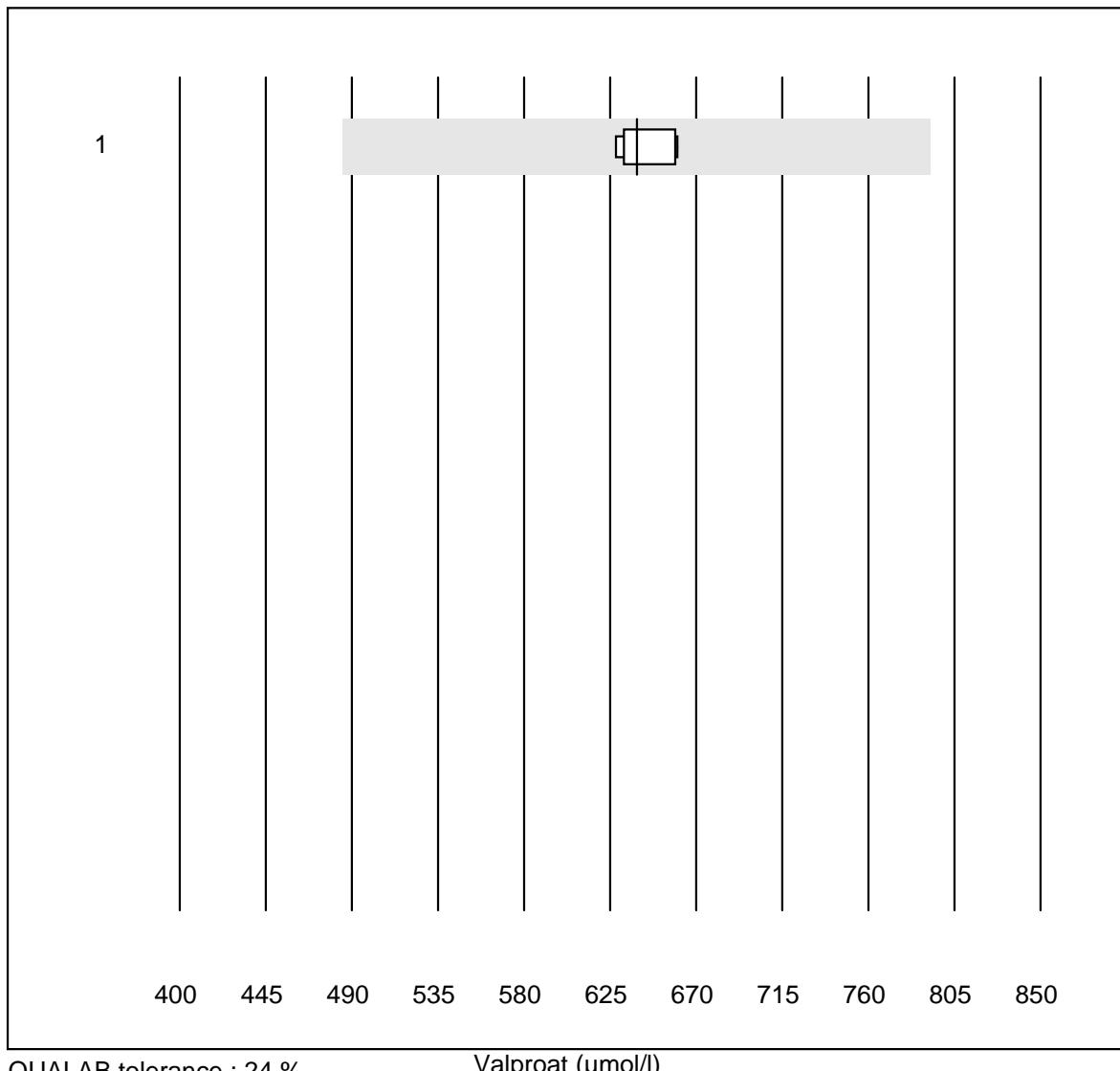


No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Qualigen	4	100.0	0.0	0.0	129.5	17.3	e*
2 Cobas	8	87.5	0.0	12.5	87.7	7.1	e
3 Architect	4	100.0	0.0	0.0	87.4	5.0	e

Digoxin

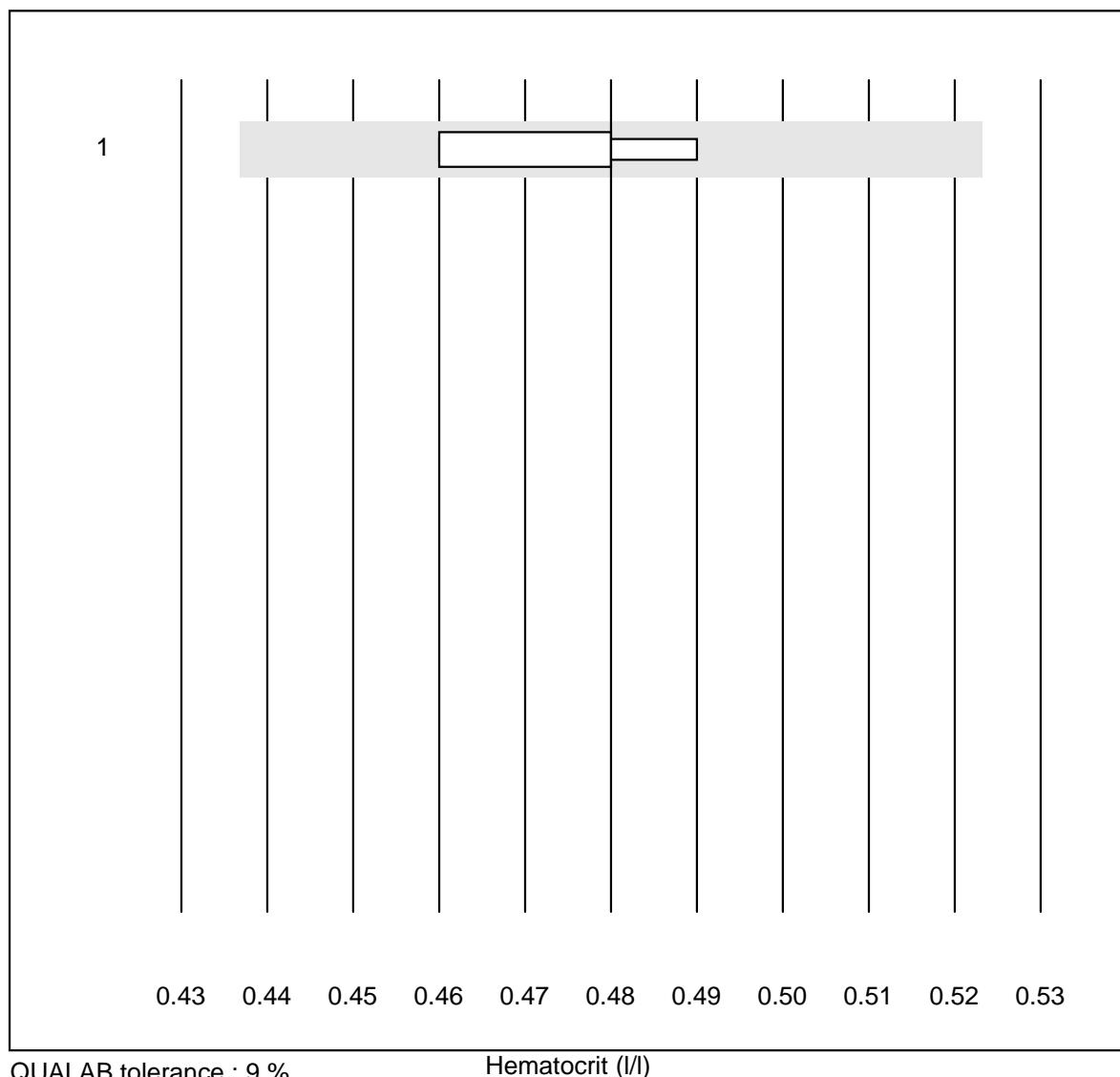


No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Other methods	11	90.9	9.1	0.0	3.38	12.0	e*

Valproat

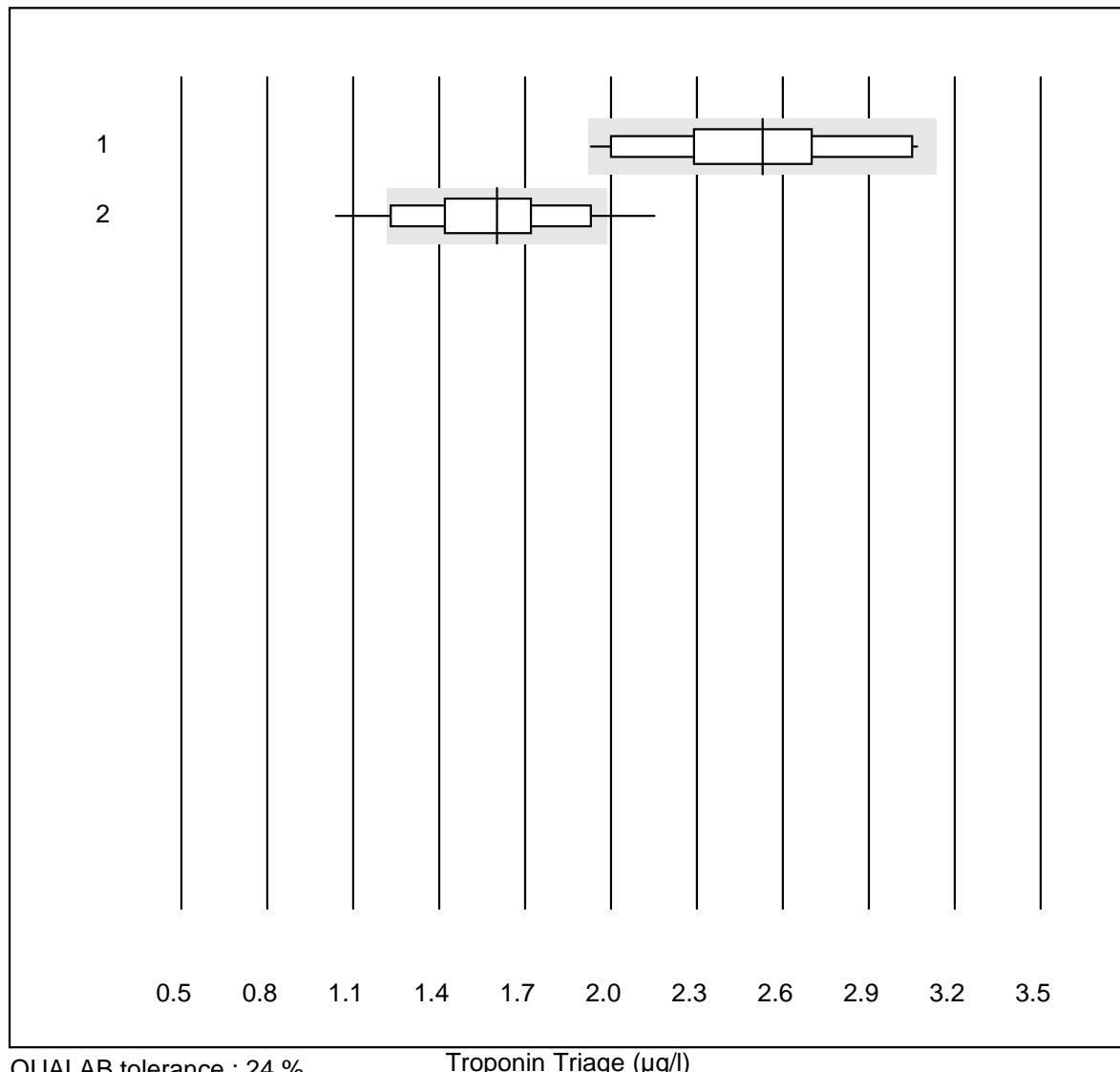
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 all Participants	5	100.0	0.0	0.0	639.0	2.3	e

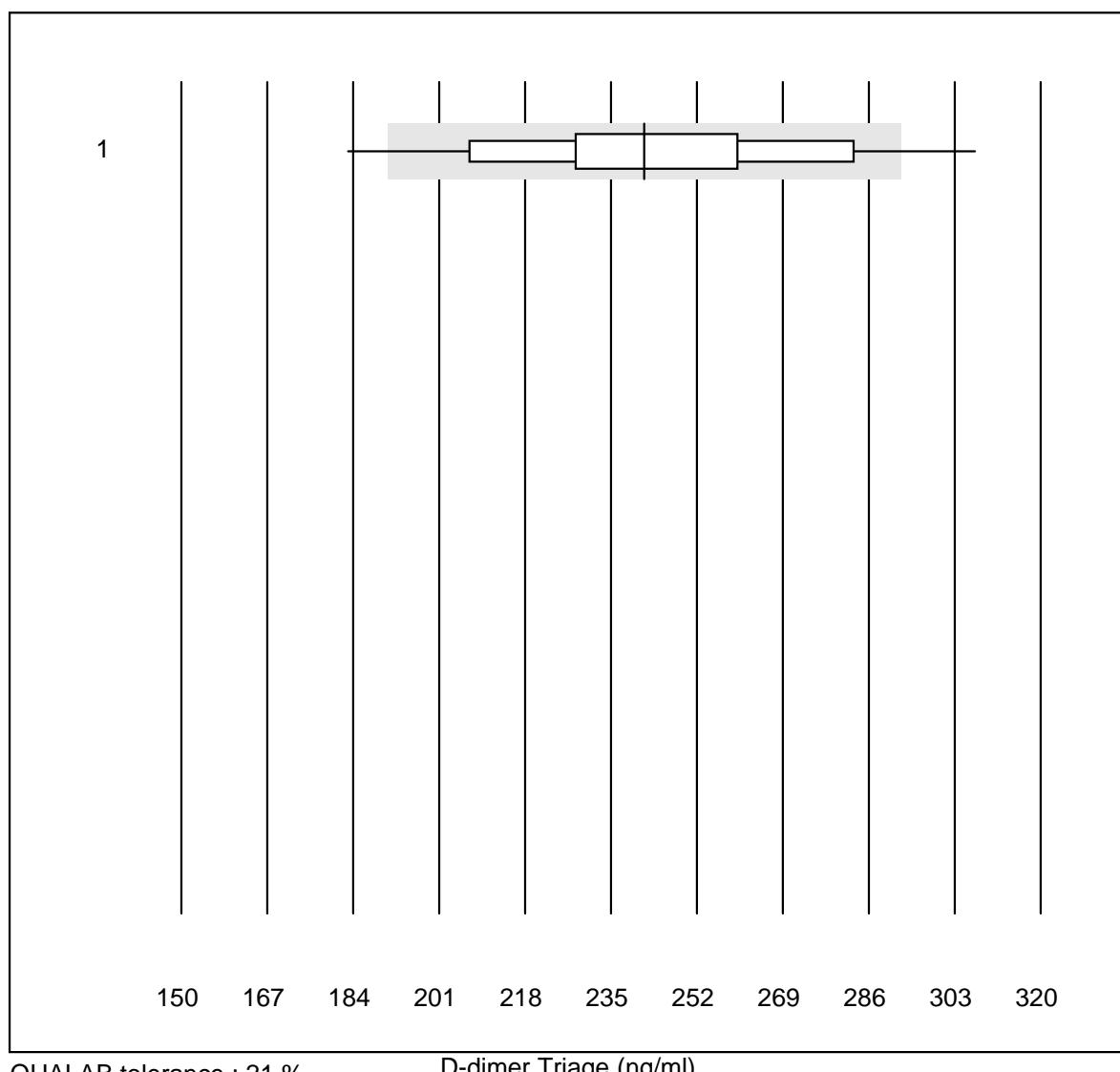
Hematocrit



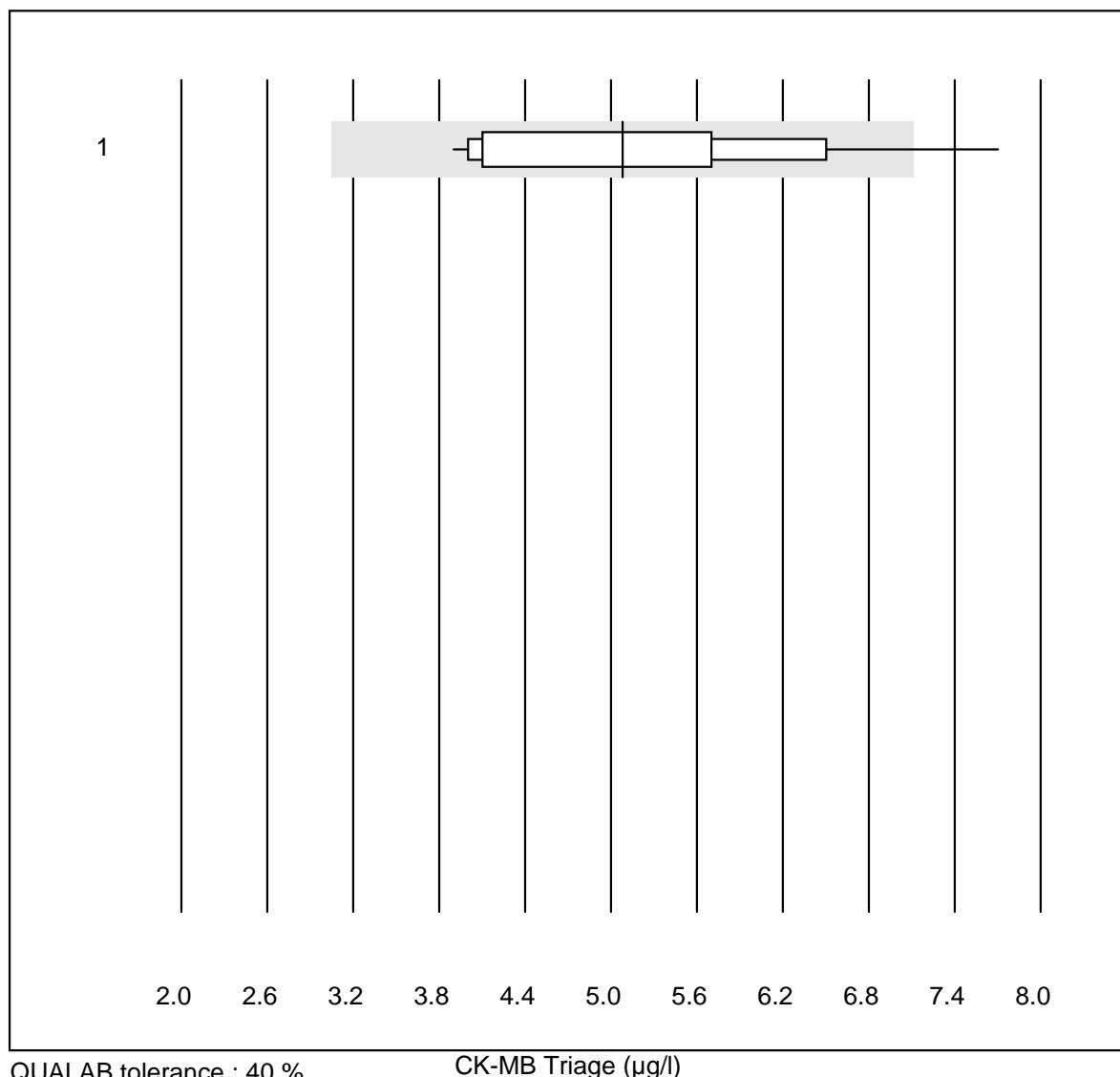
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	EPOC	4	100.0	0.0	0.0	0.48	2.7	e*

Troponin Triage



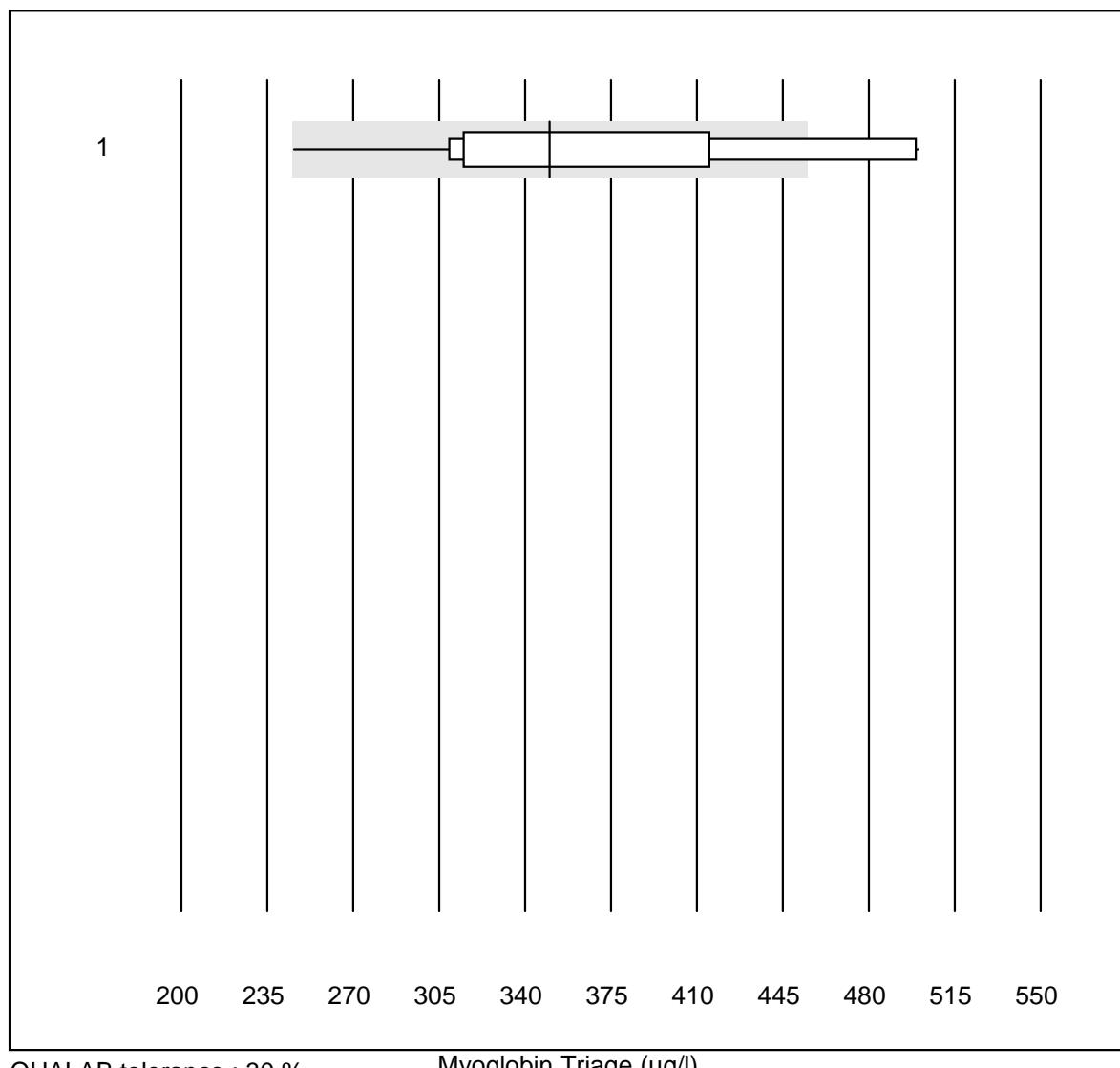
D-dimer Triage

No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Triage Meter	39	79.5	7.7	12.8	241.56	12.0	e

CK-MB Triage

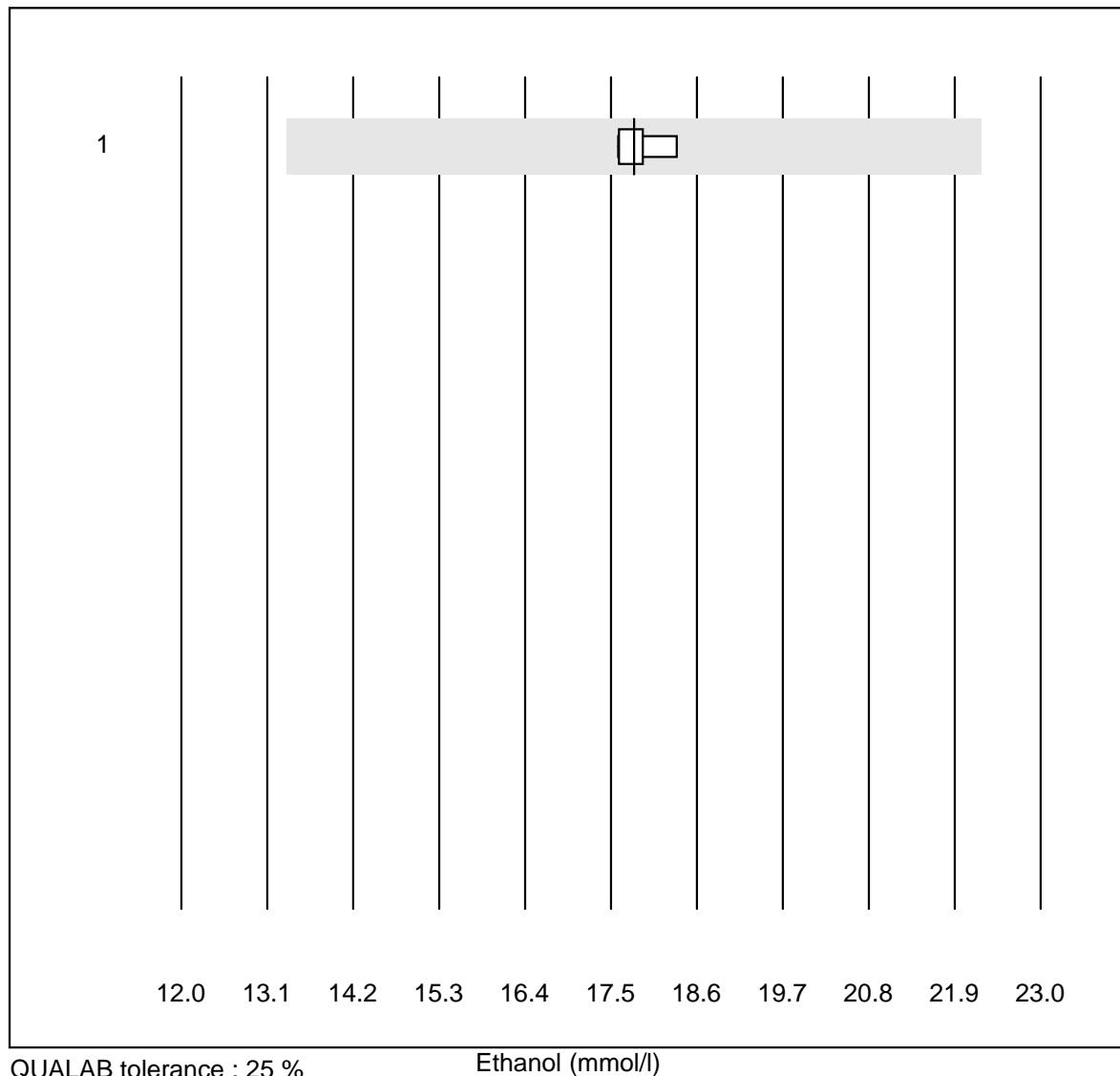
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Triage Meter	20	90.0	5.0	5.0	5.1	20.3	e

Myoglobin Triage



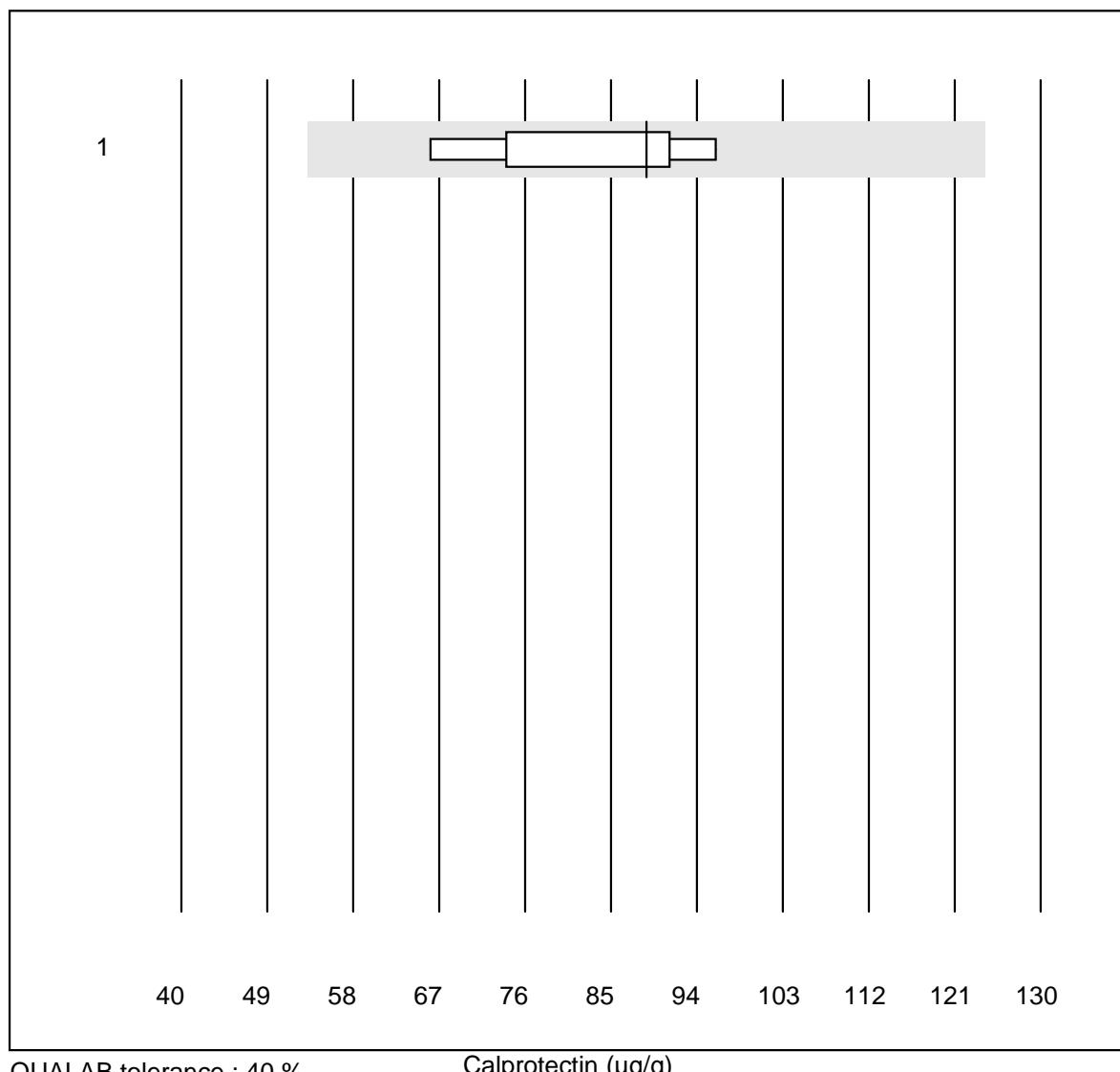
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Triage Meter	18	88.9	11.1	0.0	350.0	18.1	e*

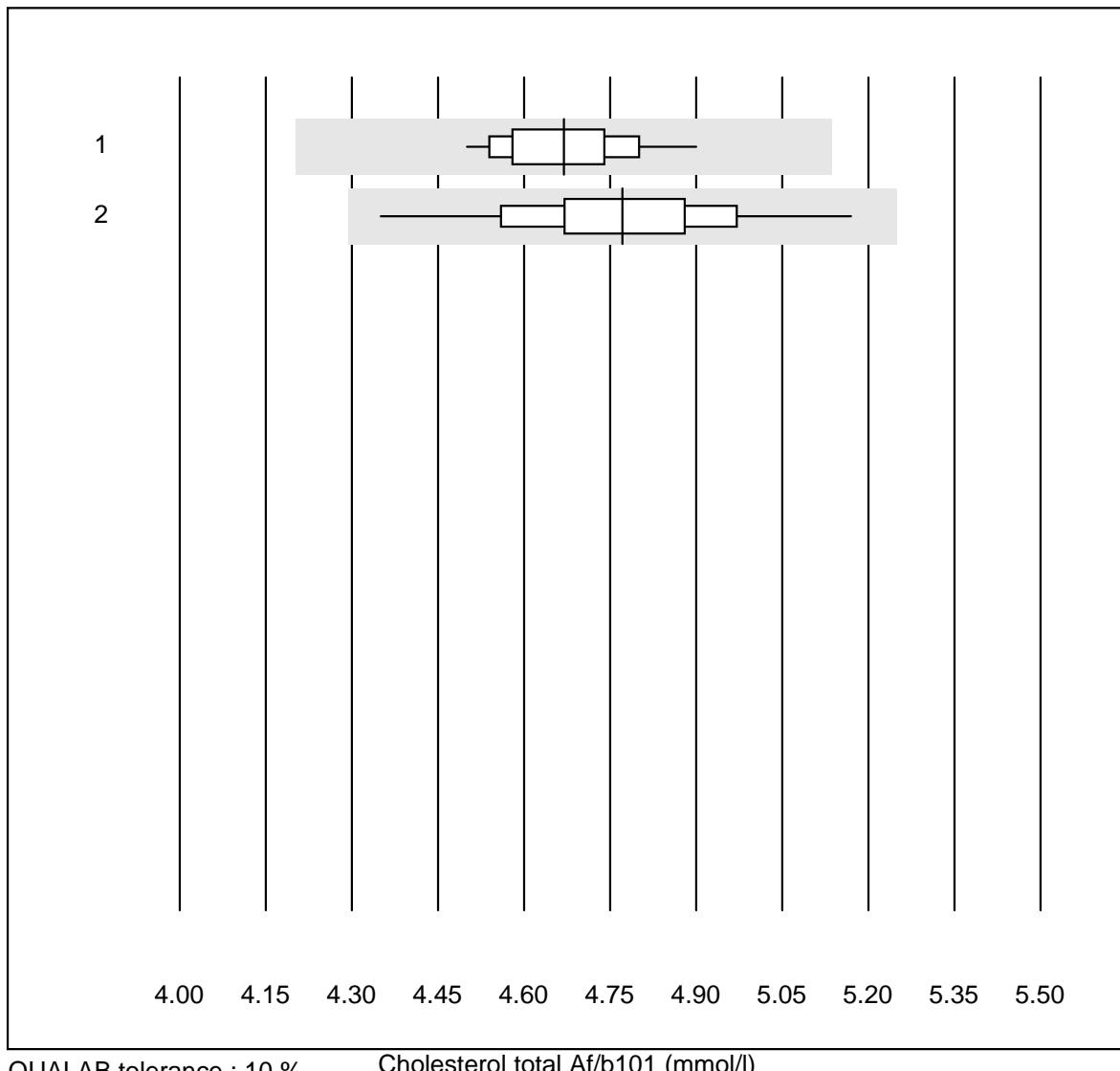
Ethanol



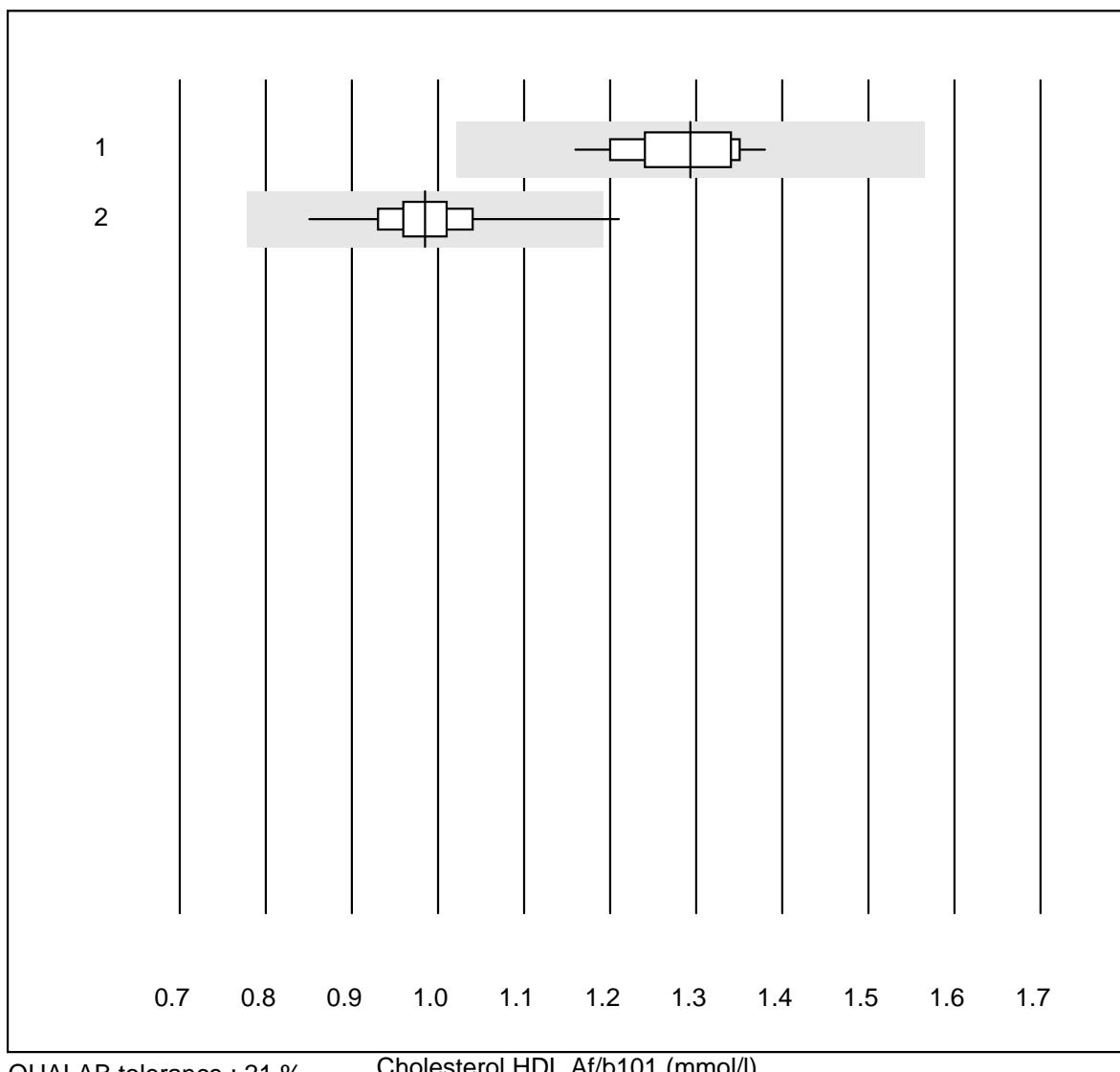
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 all Participants	5	100.0	0.0	0.0	17.8	1.7	e

Calprotectin



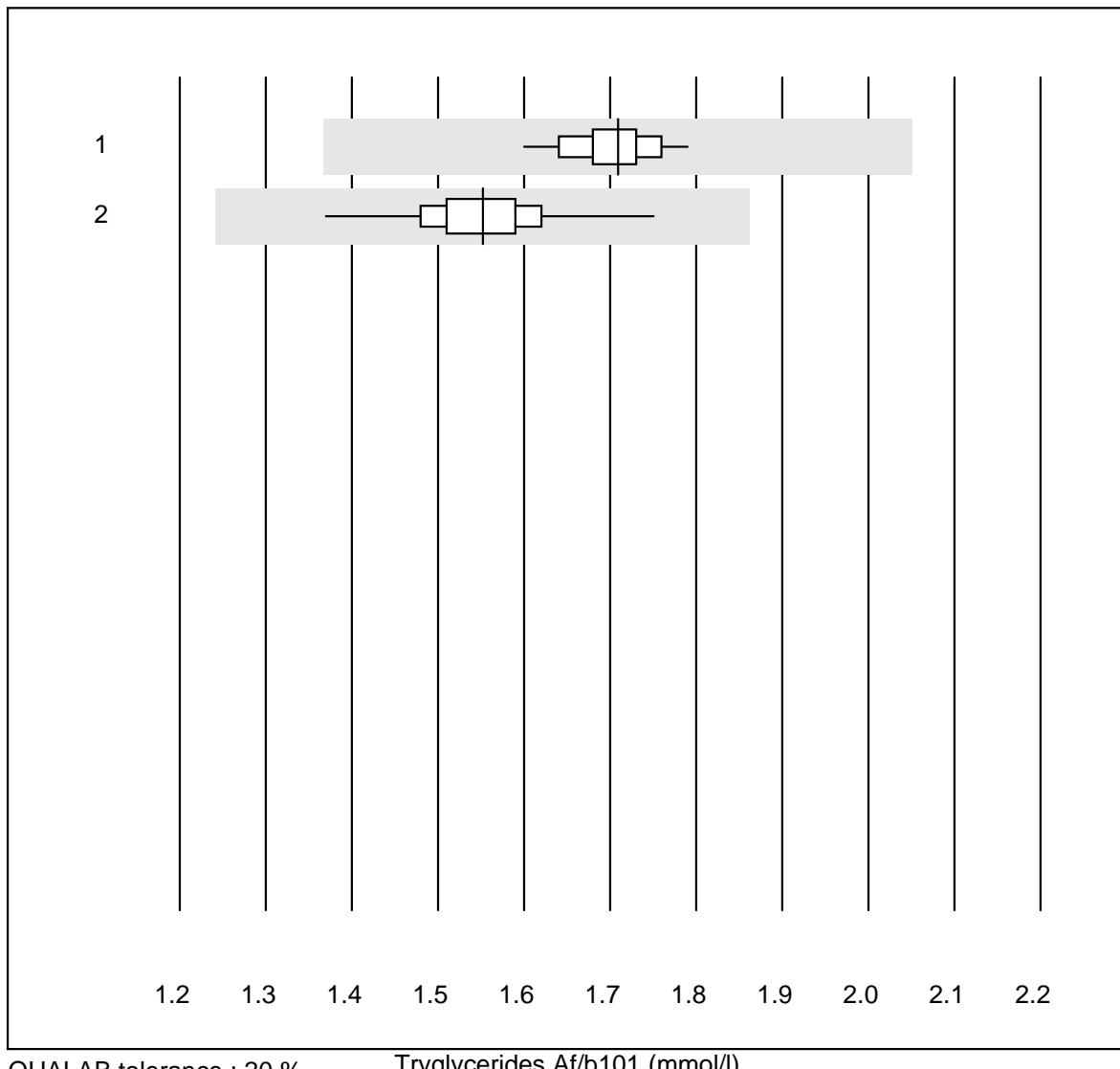
Cholesterol total Af/b101

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Cobas b101	28	100.0	0.0	0.0	4.7	2.2	e
2 Afinion	233	99.6	0.0	0.4	4.8	3.4	e

Cholesterol HDL Af/b101

No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Cobas b101	28	100.0	0.0	0.0	1.3	4.4	e
2	Afinion	233	97.0	0.4	2.6	1.0	4.8	e

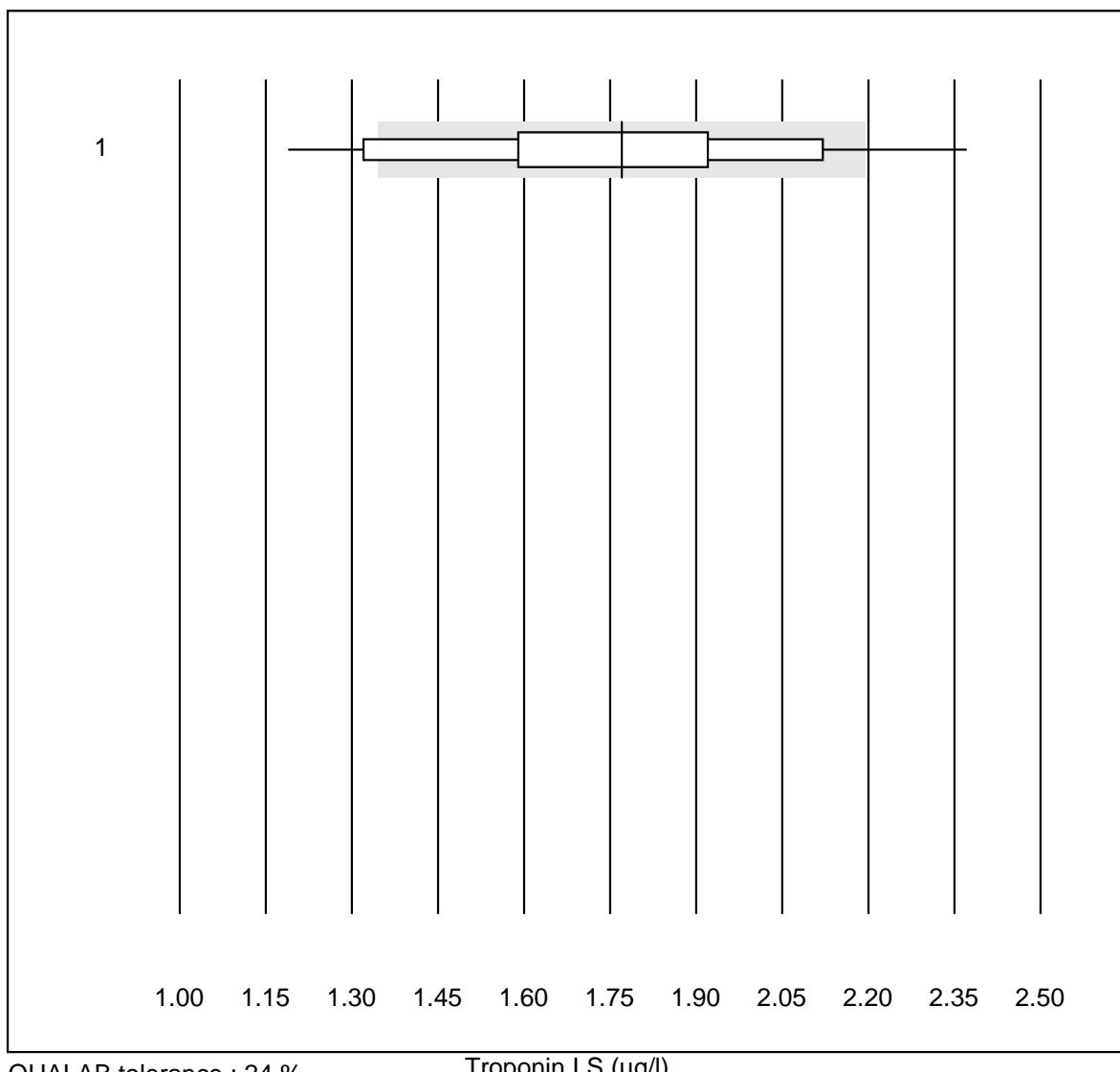
Tryglycerides Af/b101



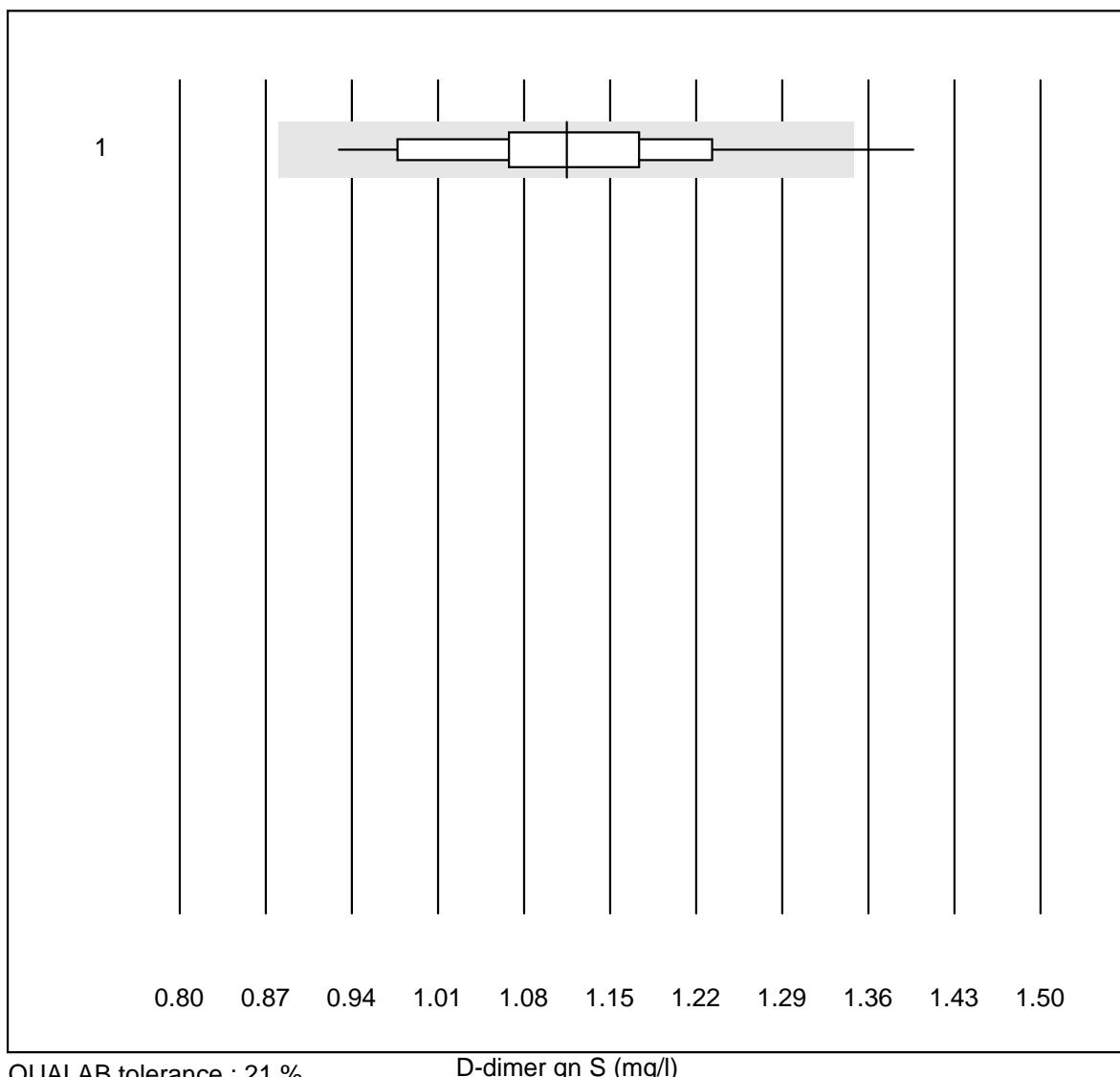
QUALAB tolerance : 20 %

Tryglycerides Af/b101 (mmol/l)

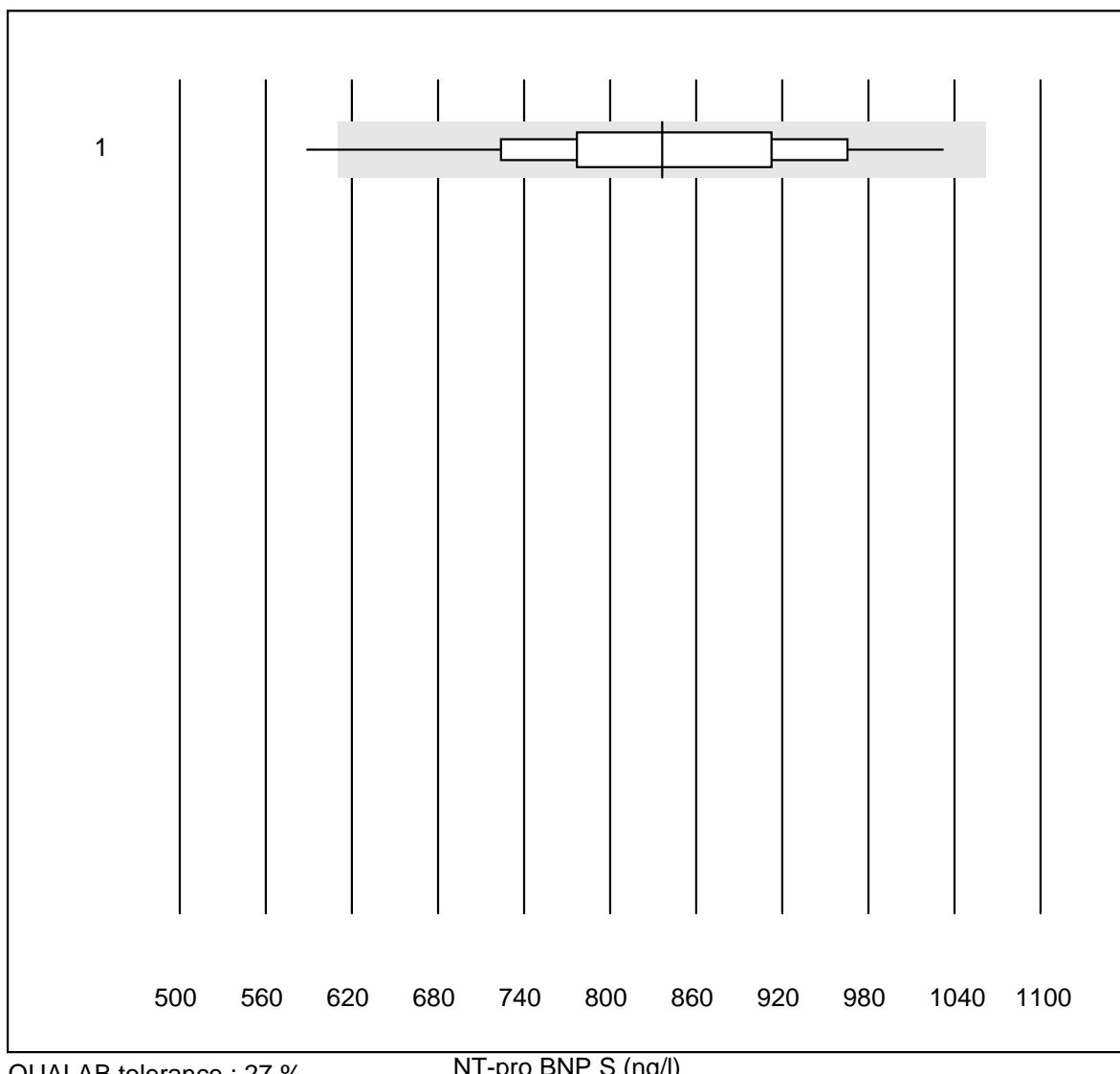
No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Cobas b101	28	100.0	0.0	0.0	1.71	2.5	e
2 Afinion	233	99.6	0.0	0.4	1.55	4.0	e

Troponin I S

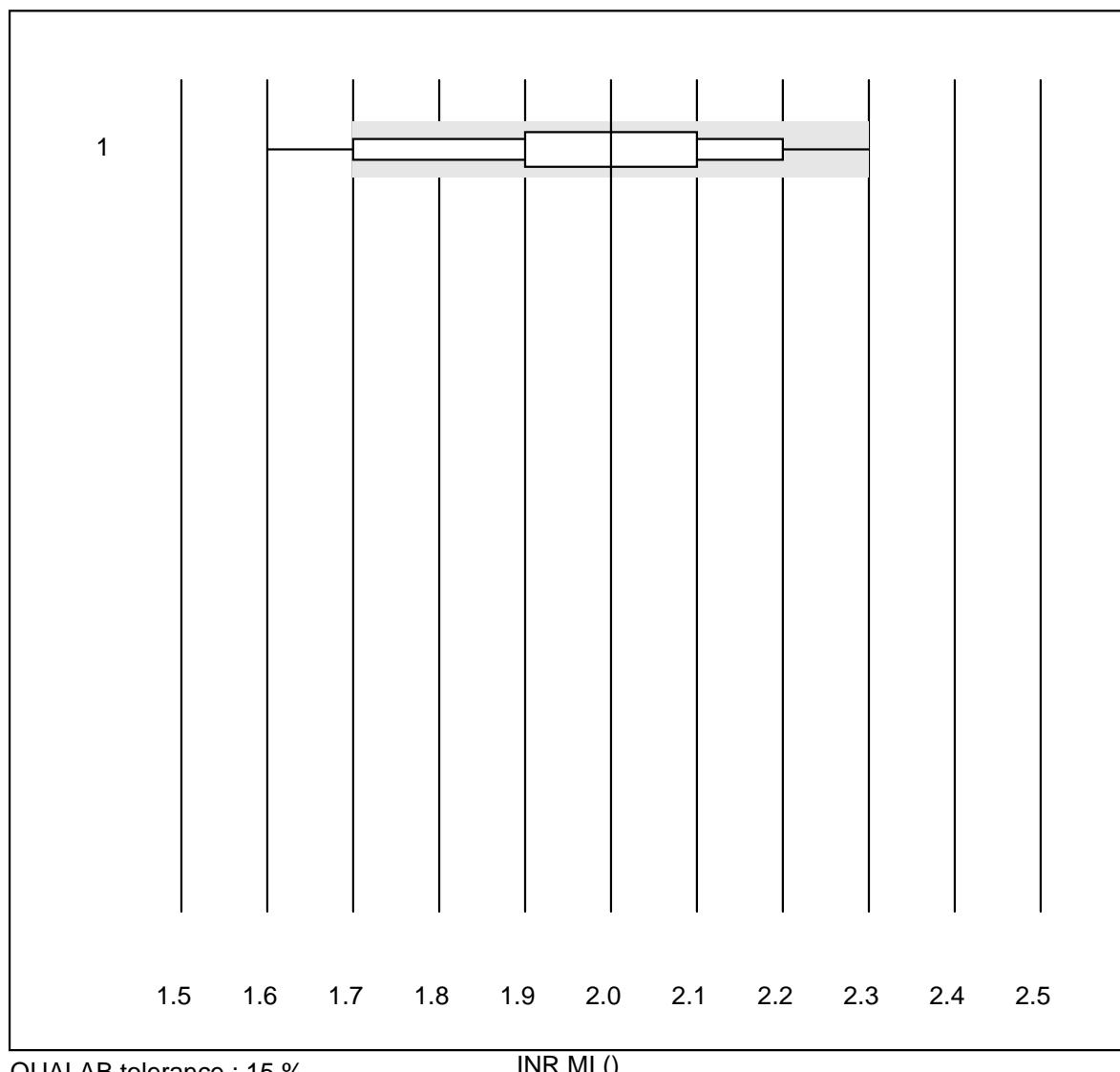
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Samsung LABGEO IB10	71	85.9	12.7	1.4	1.77	15.5	e

D-dimer qn S

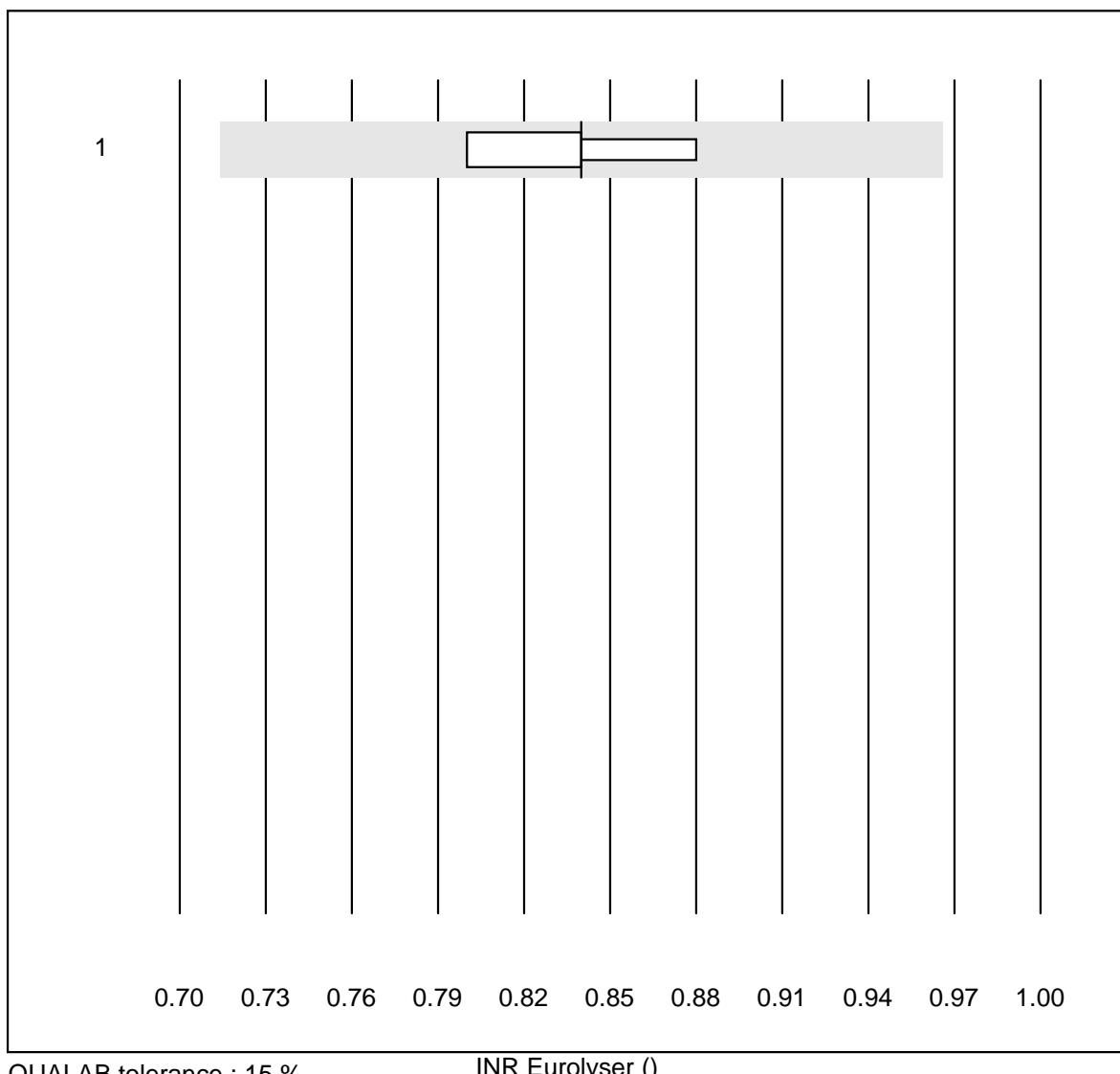
No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Samsung LABGEO IB10	85	97.6	1.2	1.2	1.11	8.2	e

NT-pro BNP S

No.	Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1	Samsung LABGEO IB10	54	96.2	1.9	1.9	836.5	11.6	e

INR MI

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 microINR	62	66.1	14.5	19.4	2.0	9.3	e

INR Eurolyser

No. Methode	Total	% good	% insuff.	% outlier	Target value	CV%	Typ
1 Eurolyser	7	71.4	0.0	28.6	0.8	4.2	e