

Verein für
Association pour le
Associazione per il



medizinische Qualitätskontrolle
contrôle de qualité médical
controllo di qualità medico

Survey Report

2016 - 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 calculate the target values, we use the mean value of the method group. Values that differ more than 1.5 times the QUALAB-tolerance are outliers and are not used to calculate the target value. Starting point for the elimination of outliers are the values of our suitability tests. 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):

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

- 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.

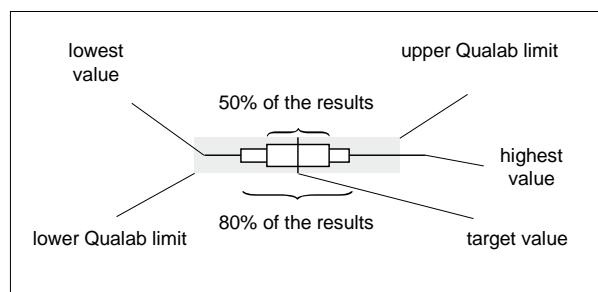
QUALAB and MQ tolerances

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:

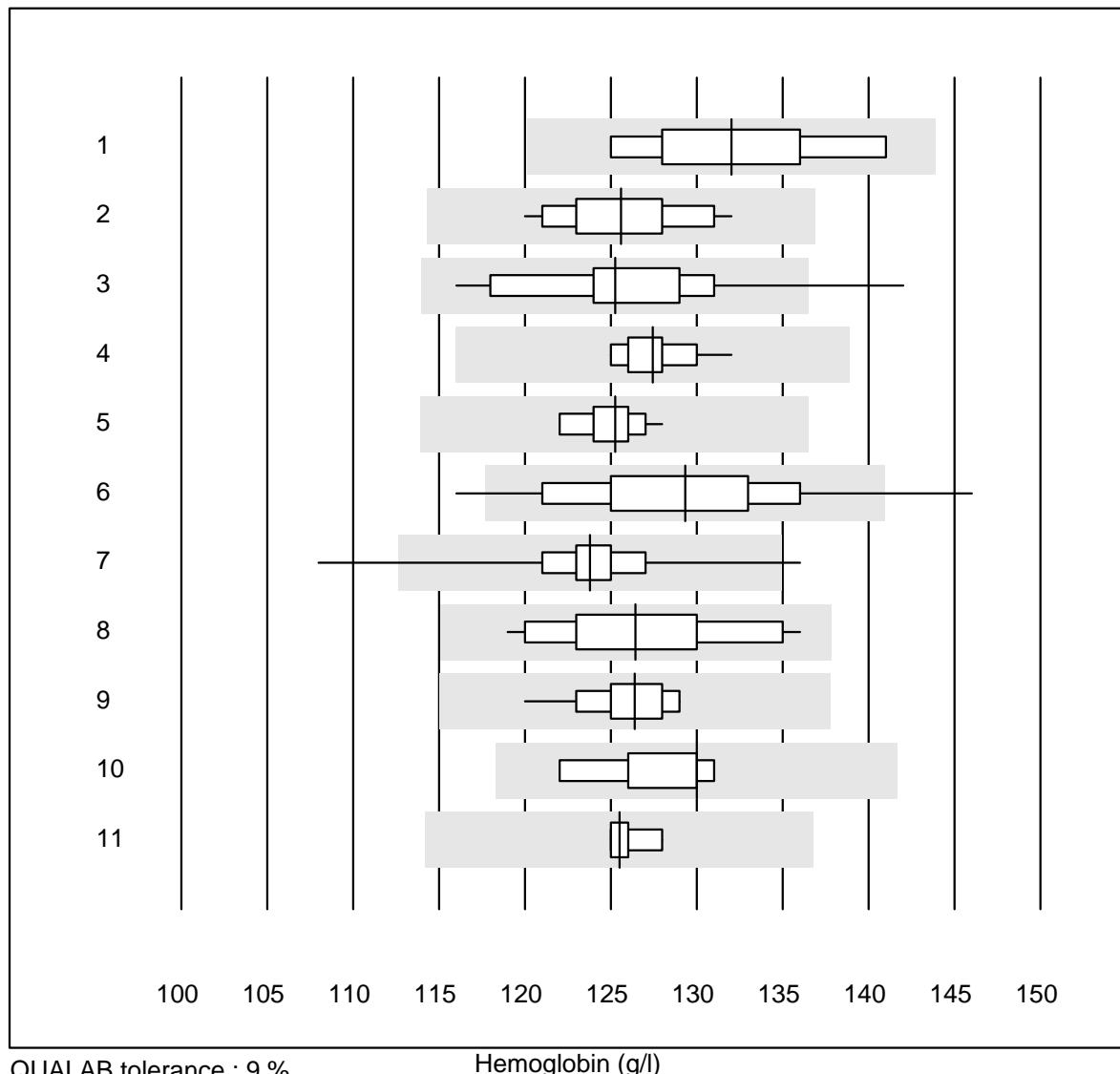
- The chemical control K1 is a ready-to-use commercial control serum. Even if the specimen is of human origin, it is possible that matrix effects occur. These are device-specific and result in different target values.
- Only one specimen was measured. Since the scatter of the results is dependent on 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, 4.7.2016

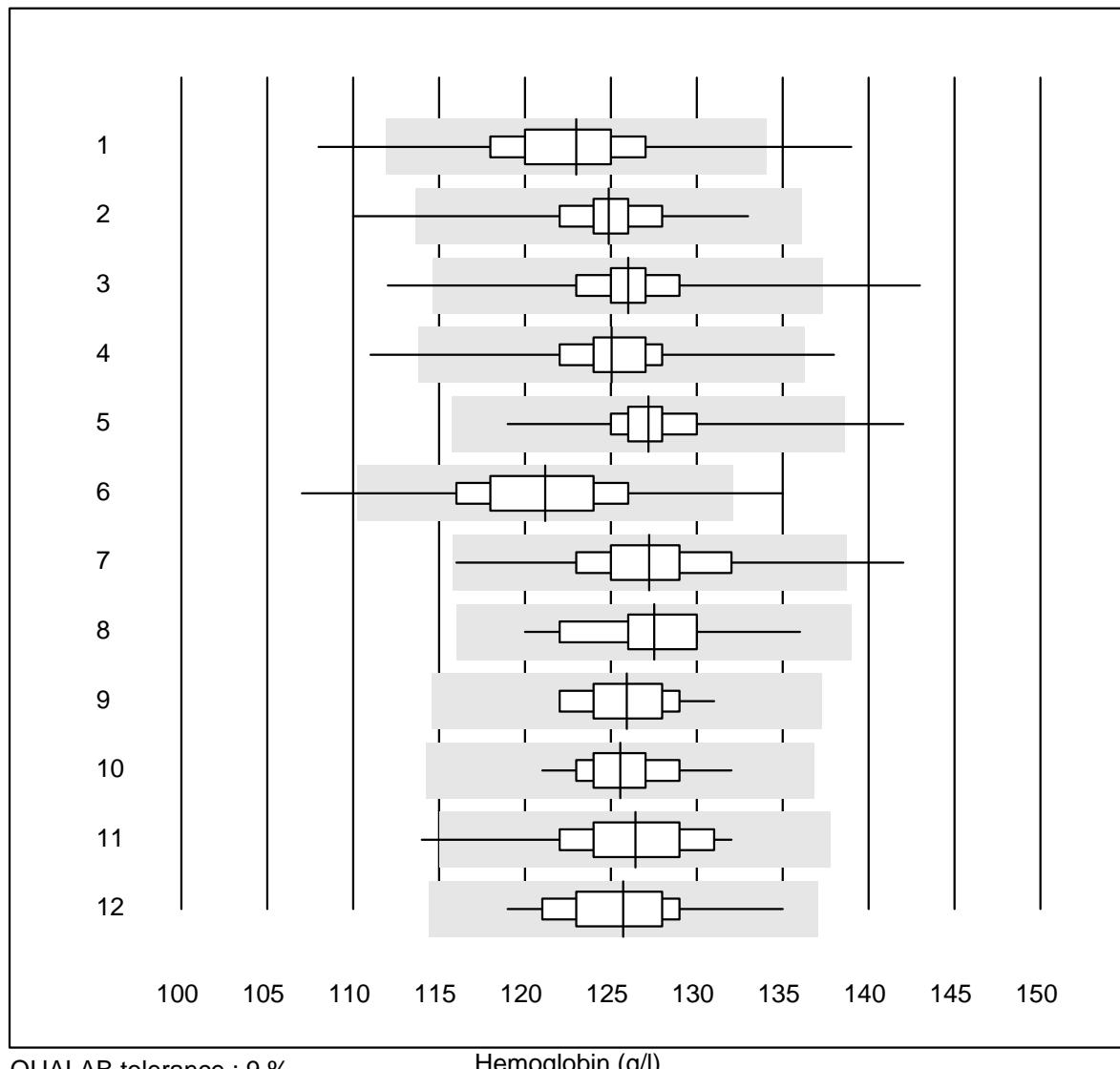
Dr. R. Fried
Survey Director

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

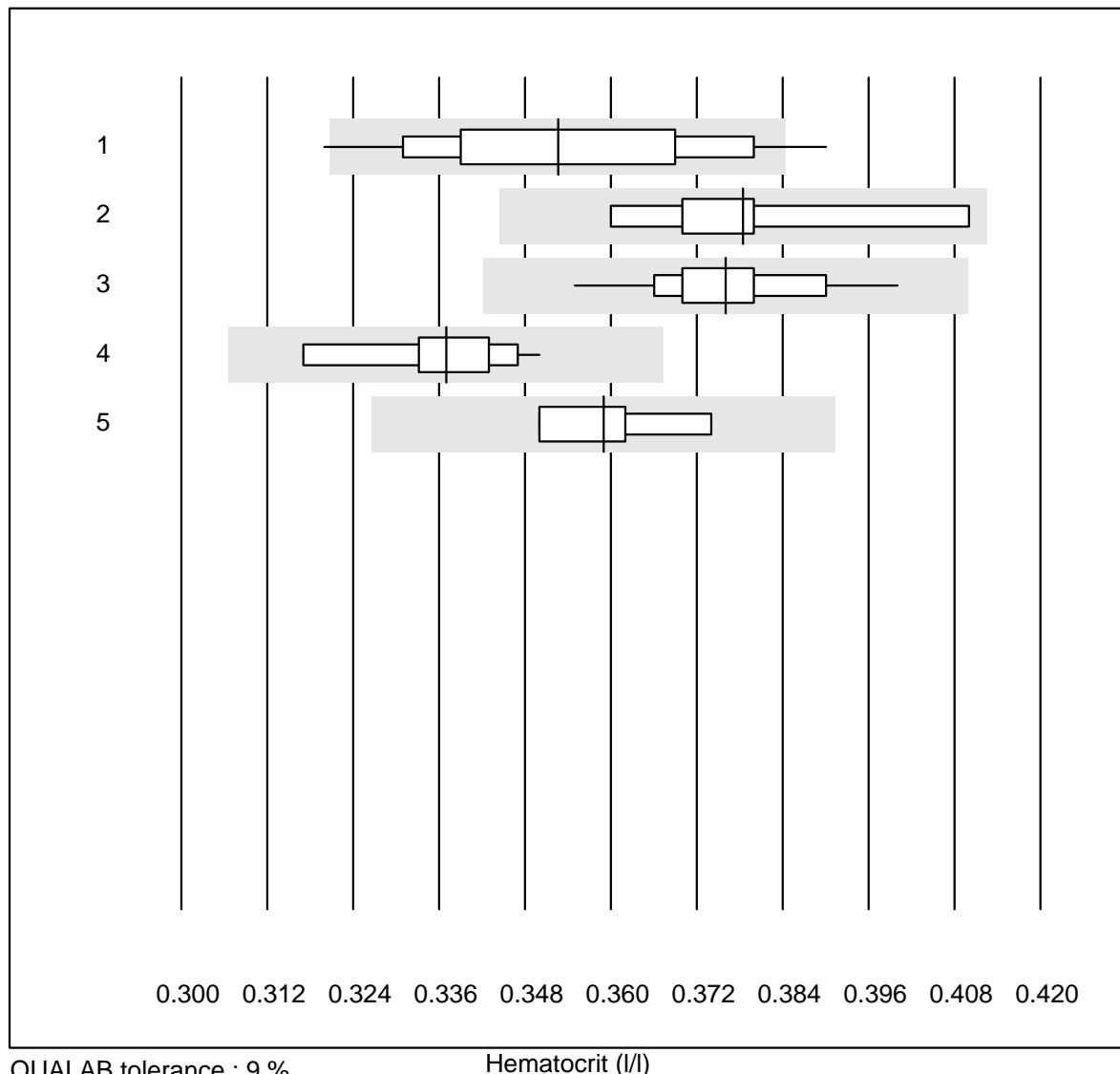


Hemoglobin

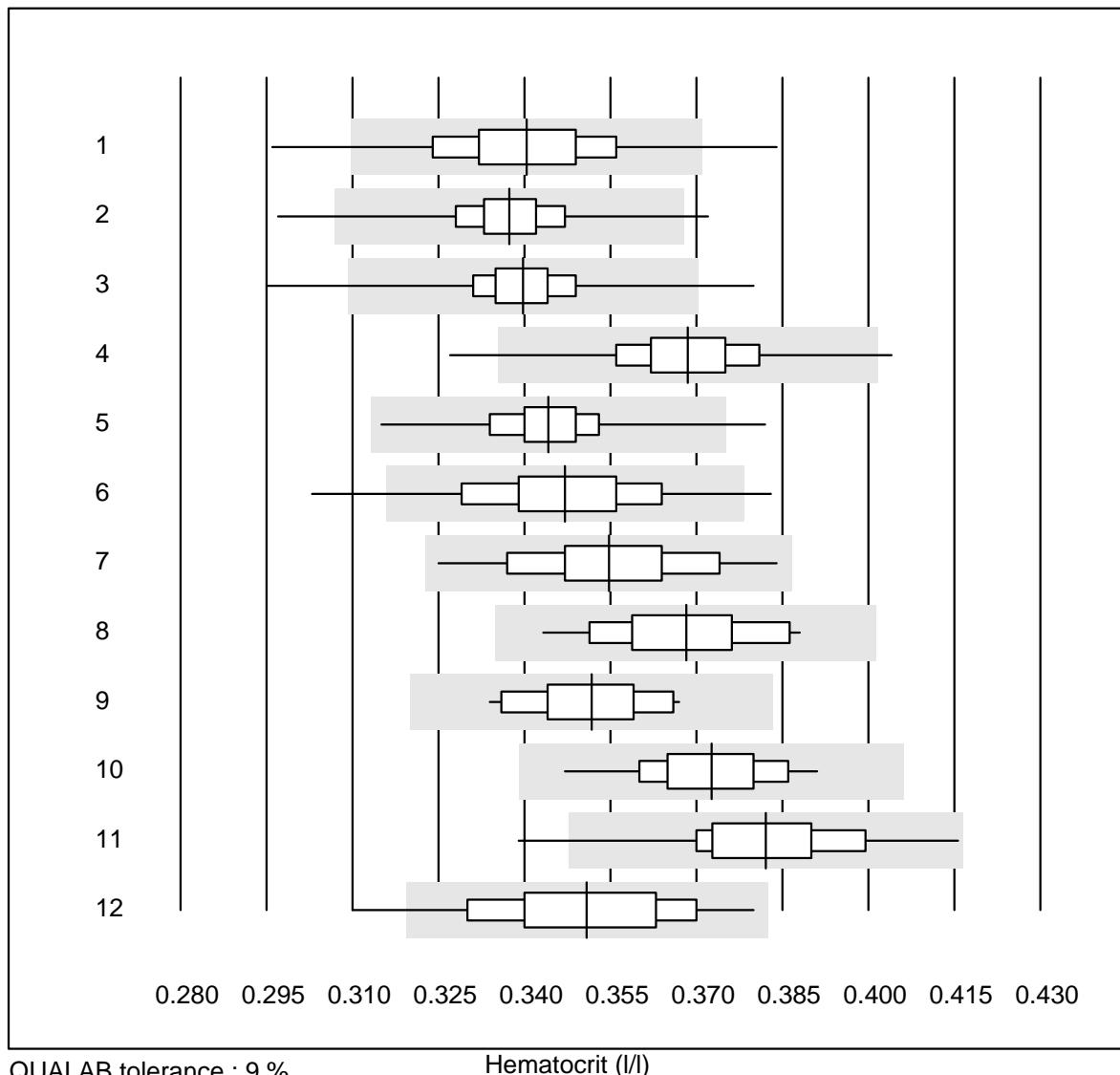


| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-------------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Abx Micros | 739 | 96.2 | 1.1 | 2.7 | 123.0 | 3.1 | e |
| 2 Microsemi | 366 | 99.2 | 0.3 | 0.5 | 124.9 | 2.1 | e |
| 3 Sysmex KX21 | 397 | 95.7 | 0.8 | 3.5 | 126.0 | 2.2 | e |
| 4 Sysmex Poch - 100i | 206 | 95.2 | 2.4 | 2.4 | 125.0 | 2.7 | e |
| 5 Sysmex XP 300 | 261 | 96.5 | 0.8 | 2.7 | 127.2 | 2.0 | e |
| 6 Mythic | 242 | 94.6 | 1.7 | 3.7 | 121.2 | 3.6 | e |
| 7 Swelab | 68 | 97.0 | 1.5 | 1.5 | 127.2 | 3.1 | e |
| 8 Abacus Junior | 12 | 100.0 | 0.0 | 0.0 | 127.5 | 3.2 | e |
| 9 Medonic | 15 | 93.3 | 0.0 | 6.7 | 125.9 | 2.1 | e |
| 10 Nihon Kohden Celltac | 35 | 97.1 | 0.0 | 2.9 | 125.6 | 2.1 | e |
| 11 Samsung HC10 | 45 | 93.4 | 4.4 | 2.2 | 126.4 | 3.2 | e |
| 12 Norma Icon 3 | 26 | 100.0 | 0.0 | 0.0 | 125.7 | 2.9 | e |

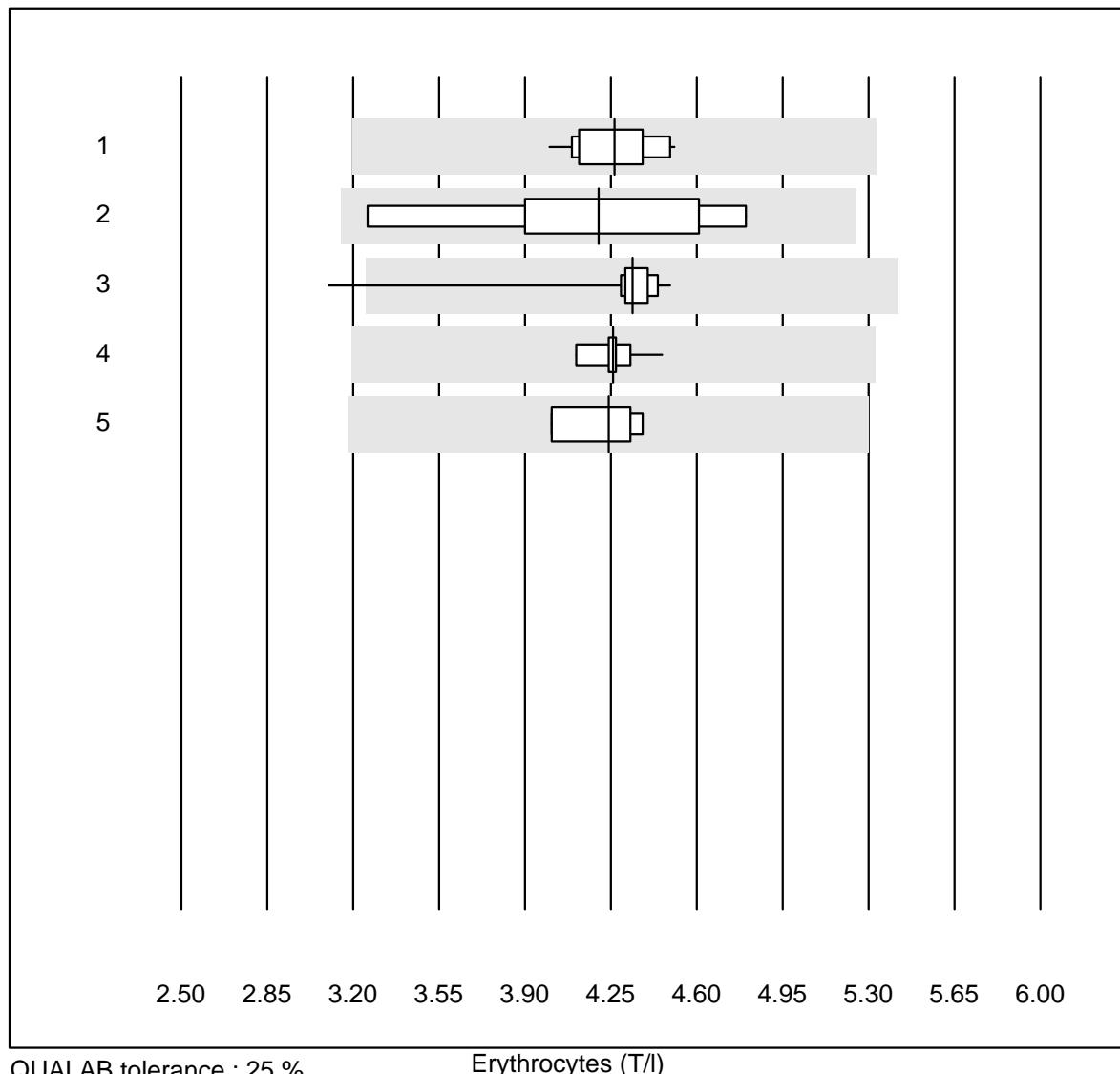
Hematocrit



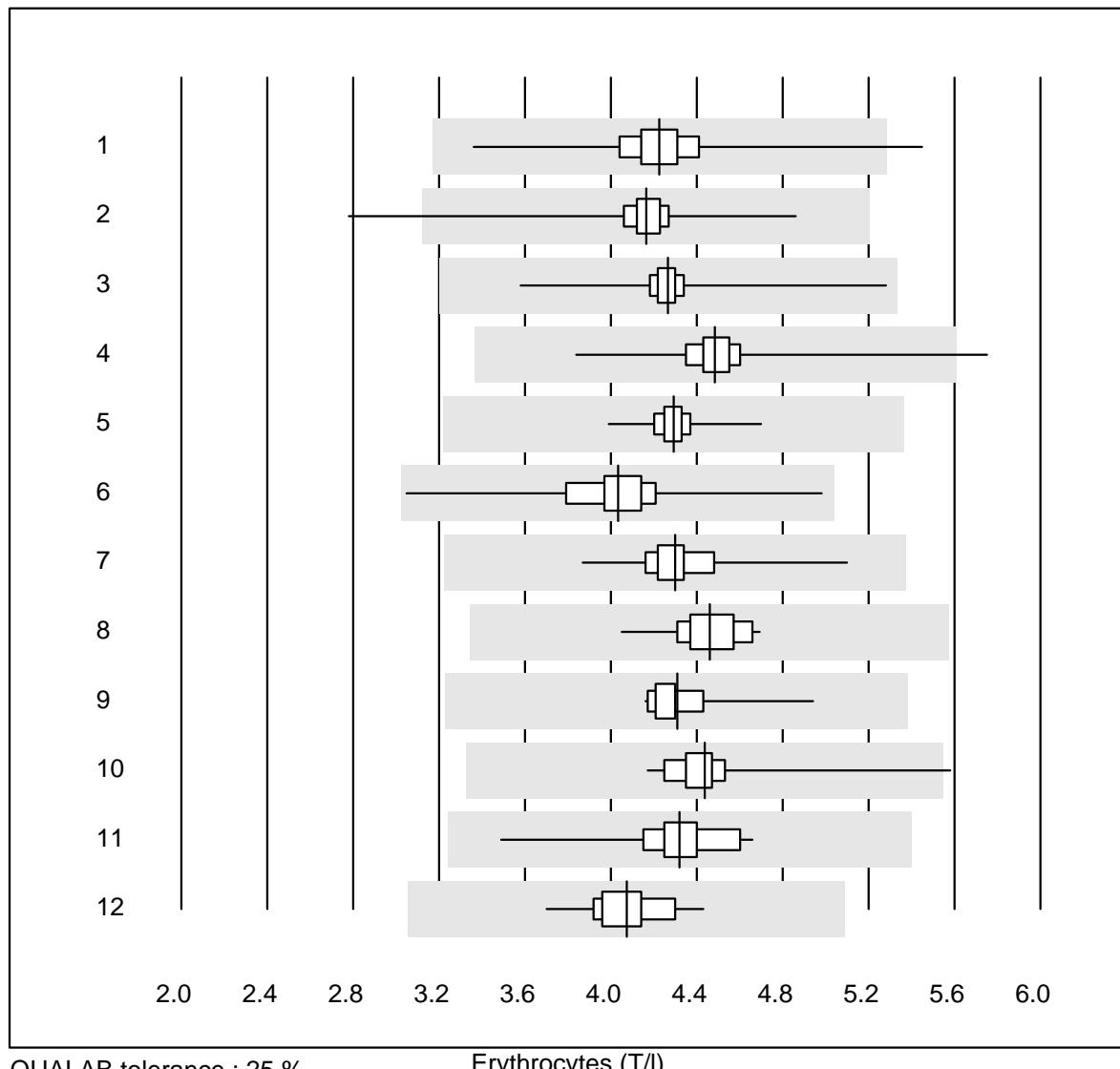
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Automat | 28 | 85.7 | 10.7 | 3.6 | 0.35 | 5.4 | e |
| 2 Centrifuge | 13 | 100.0 | 0.0 | 0.0 | 0.38 | 4.7 | e* |
| 3 Sysmex X | 38 | 97.4 | 0.0 | 2.6 | 0.38 | 2.6 | e |
| 4 ABX Pentra | 10 | 100.0 | 0.0 | 0.0 | 0.34 | 2.8 | e |
| 5 Celldyn | 4 | 100.0 | 0.0 | 0.0 | 0.36 | 2.8 | e* |

Hematocrit

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-------------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Abx Micros | 739 | 94.4 | 2.4 | 3.2 | 0.34 | 3.8 | e |
| 2 Microsemi | 363 | 98.6 | 0.8 | 0.6 | 0.34 | 2.4 | e |
| 3 Sysmex KX21 | 397 | 95.7 | 0.8 | 3.5 | 0.34 | 2.5 | e |
| 4 Sysmex Poch - 100i | 206 | 98.0 | 1.5 | 0.5 | 0.37 | 3.1 | e |
| 5 Sysmex XP 300 | 257 | 98.0 | 1.2 | 0.8 | 0.34 | 2.4 | e |
| 6 Mythic | 242 | 88.4 | 3.3 | 8.3 | 0.35 | 4.0 | e |
| 7 Swelab | 68 | 98.5 | 0.0 | 1.5 | 0.35 | 3.8 | e |
| 8 Abacus Junior | 12 | 100.0 | 0.0 | 0.0 | 0.37 | 3.6 | e |
| 9 Medonic | 15 | 93.3 | 0.0 | 6.7 | 0.35 | 3.0 | e |
| 10 Nihon Kohden Celltac | 36 | 94.4 | 0.0 | 5.6 | 0.37 | 3.0 | e |
| 11 Samsung HC10 | 45 | 93.4 | 4.4 | 2.2 | 0.38 | 4.0 | e |
| 12 Norma Icon 3 | 26 | 96.2 | 3.8 | 0.0 | 0.35 | 4.7 | e |

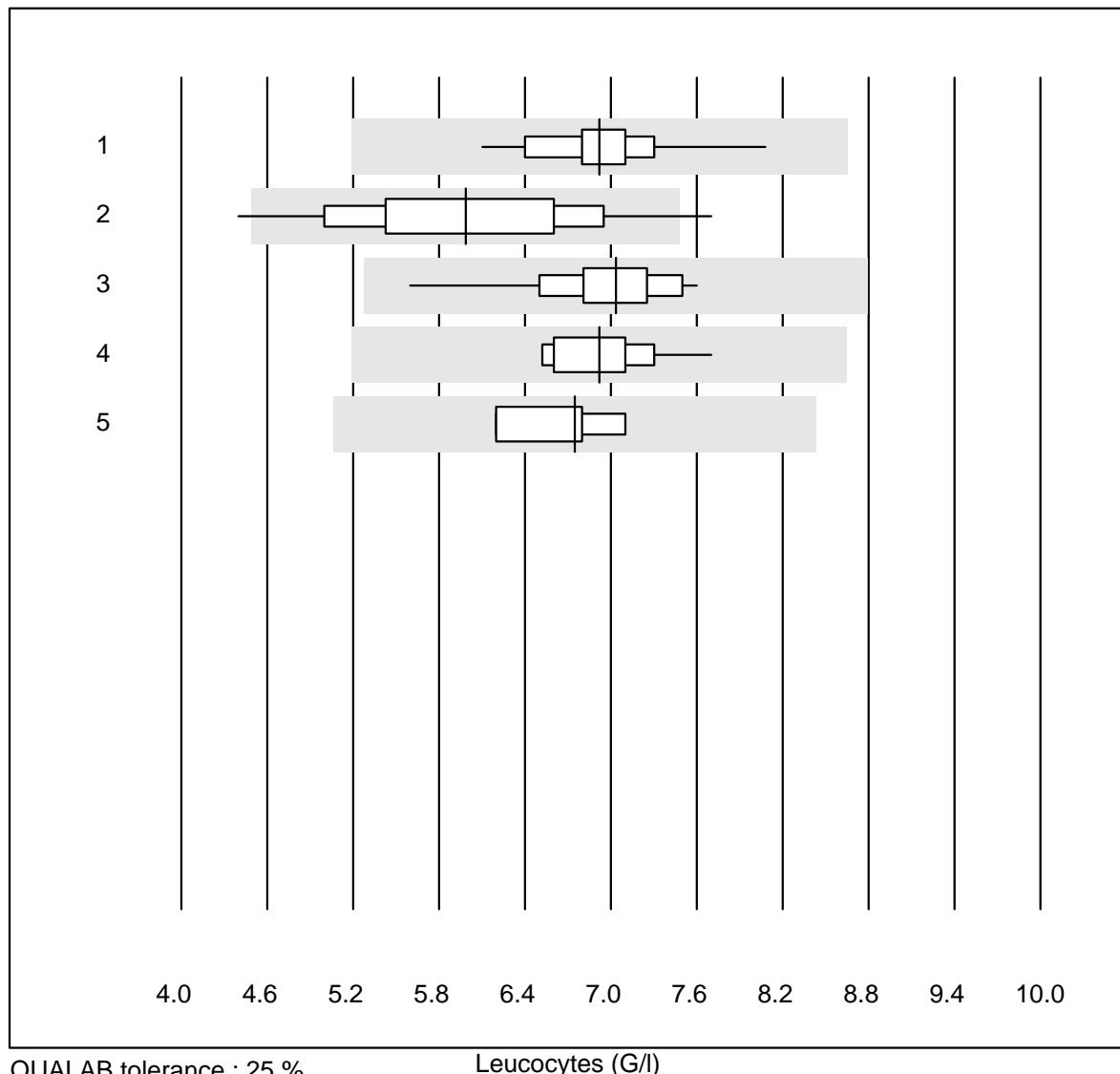
Erythrocytes

Erythrocytes

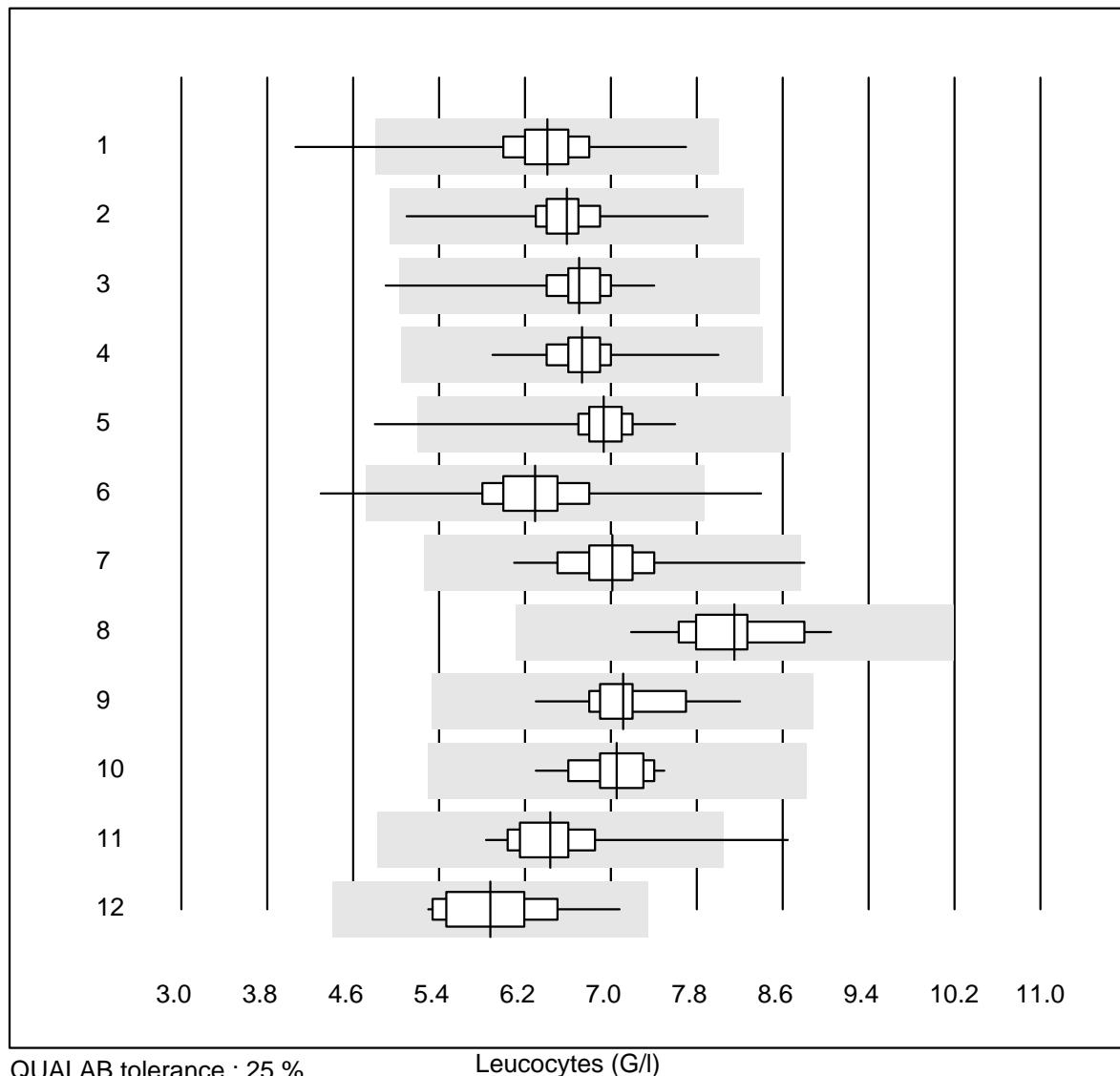


| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-------------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Abx Micros | 739 | 98.3 | 0.3 | 1.4 | 4.23 | 4.3 | e |
| 2 Microsemi | 365 | 99.2 | 0.5 | 0.3 | 4.16 | 3.6 | e |
| 3 Sysmex KX21 | 397 | 97.7 | 0.0 | 2.3 | 4.27 | 2.7 | e |
| 4 Sysmex Poch - 100i | 206 | 98.5 | 0.5 | 1.0 | 4.48 | 3.3 | e |
| 5 Sysmex XP 300 | 259 | 98.8 | 0.0 | 1.2 | 4.29 | 2.0 | e |
| 6 Mythic | 242 | 97.5 | 0.0 | 2.5 | 4.03 | 5.2 | e |
| 7 Swelab | 68 | 100.0 | 0.0 | 0.0 | 4.30 | 3.9 | e |
| 8 Abacus Junior | 12 | 100.0 | 0.0 | 0.0 | 4.46 | 3.8 | e |
| 9 Medonic | 15 | 100.0 | 0.0 | 0.0 | 4.31 | 4.4 | e |
| 10 Nihon Kohden Celltac | 35 | 97.1 | 2.9 | 0.0 | 4.44 | 5.1 | e |
| 11 Samsung HC10 | 45 | 100.0 | 0.0 | 0.0 | 4.32 | 4.7 | e |
| 12 Norma Icon 3 | 26 | 100.0 | 0.0 | 0.0 | 4.07 | 3.8 | e |

Leucocytes

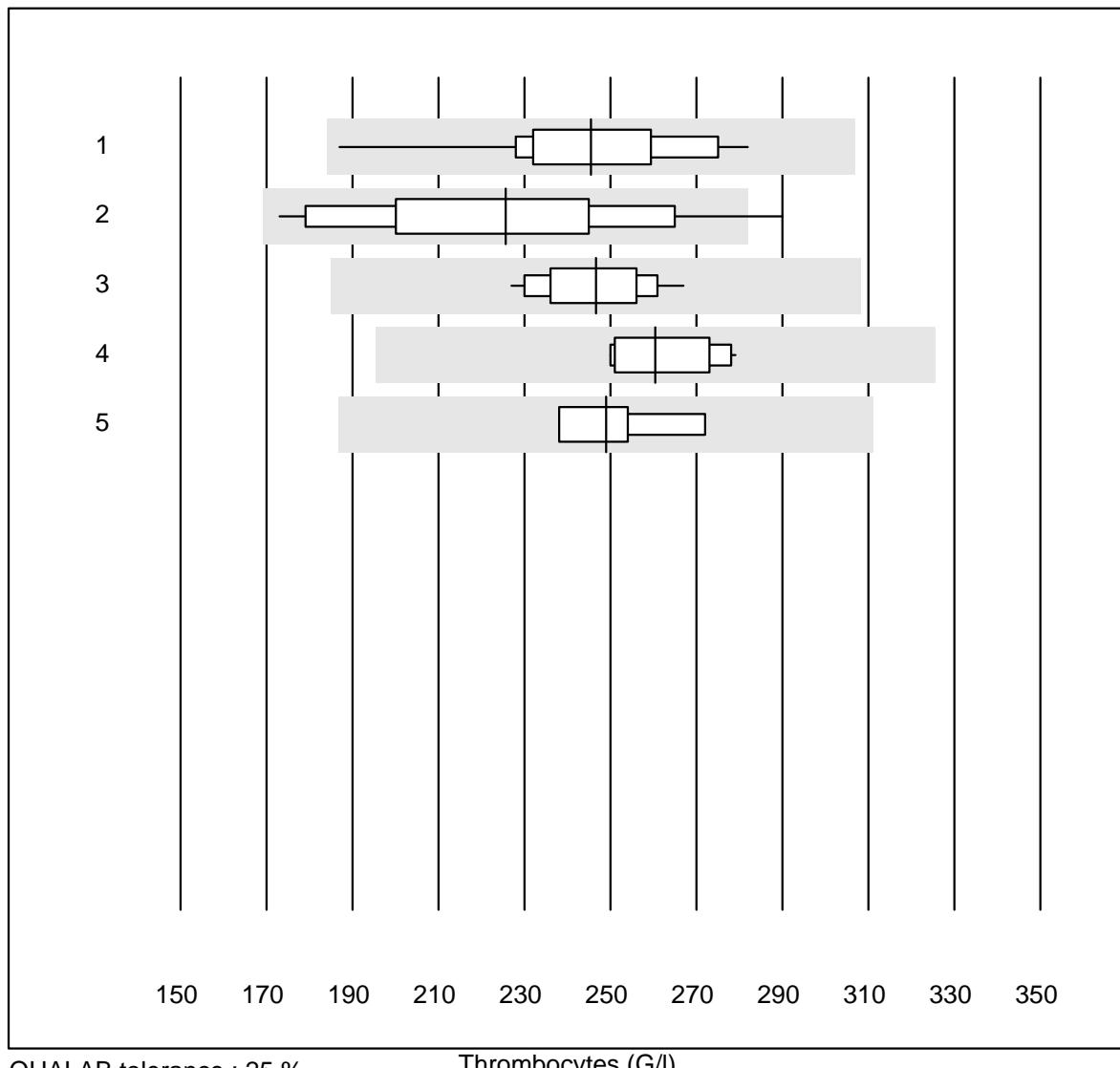


Leucocytes



| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-------------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Abx Micros | 739 | 99.2 | 0.3 | 0.5 | 6.41 | 5.5 | e |
| 2 Microsemi | 366 | 100.0 | 0.0 | 0.0 | 6.59 | 4.0 | e |
| 3 Sysmex KX21 | 397 | 99.2 | 0.3 | 0.5 | 6.71 | 4.2 | e |
| 4 Sysmex Poch - 100i | 206 | 100.0 | 0.0 | 0.0 | 6.73 | 4.3 | e |
| 5 Sysmex XP 300 | 261 | 99.2 | 0.4 | 0.4 | 6.93 | 4.1 | e |
| 6 Mythic | 239 | 97.0 | 1.7 | 1.3 | 6.30 | 6.9 | e |
| 7 Swelab | 68 | 98.5 | 1.5 | 0.0 | 7.01 | 5.9 | e |
| 8 Abacus Junior | 12 | 100.0 | 0.0 | 0.0 | 8.15 | 6.3 | e |
| 9 Medonic | 15 | 100.0 | 0.0 | 0.0 | 7.11 | 6.0 | e |
| 10 Nihon Kohden Celltac | 35 | 100.0 | 0.0 | 0.0 | 7.06 | 4.2 | e |
| 11 Samsung HC10 | 45 | 97.8 | 2.2 | 0.0 | 6.44 | 6.9 | e |
| 12 Norma Icon 3 | 26 | 100.0 | 0.0 | 0.0 | 5.88 | 8.1 | e |

Thrombocytes

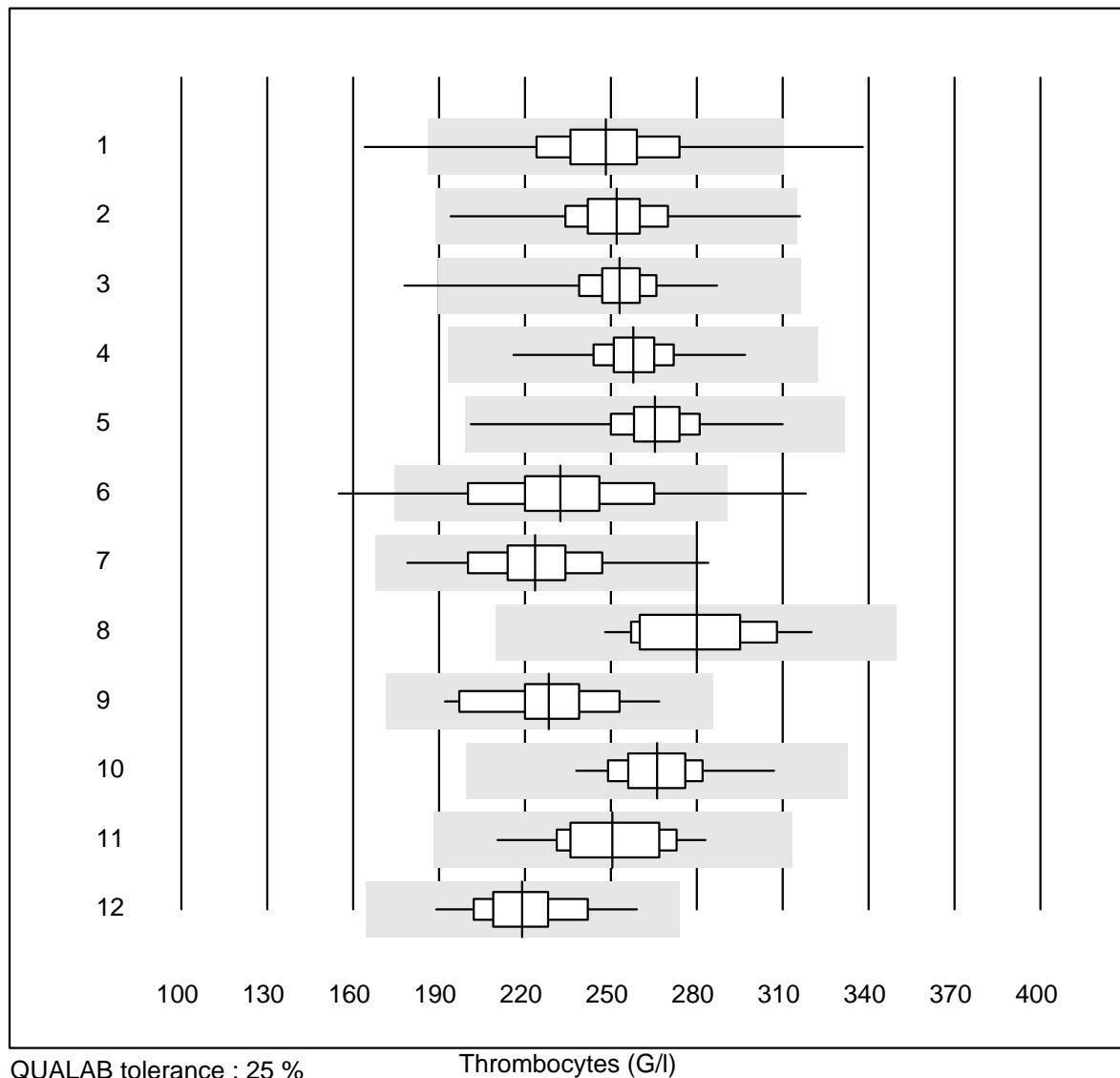


QUALAB tolerance : 25 %

Thrombocytes (G/l)

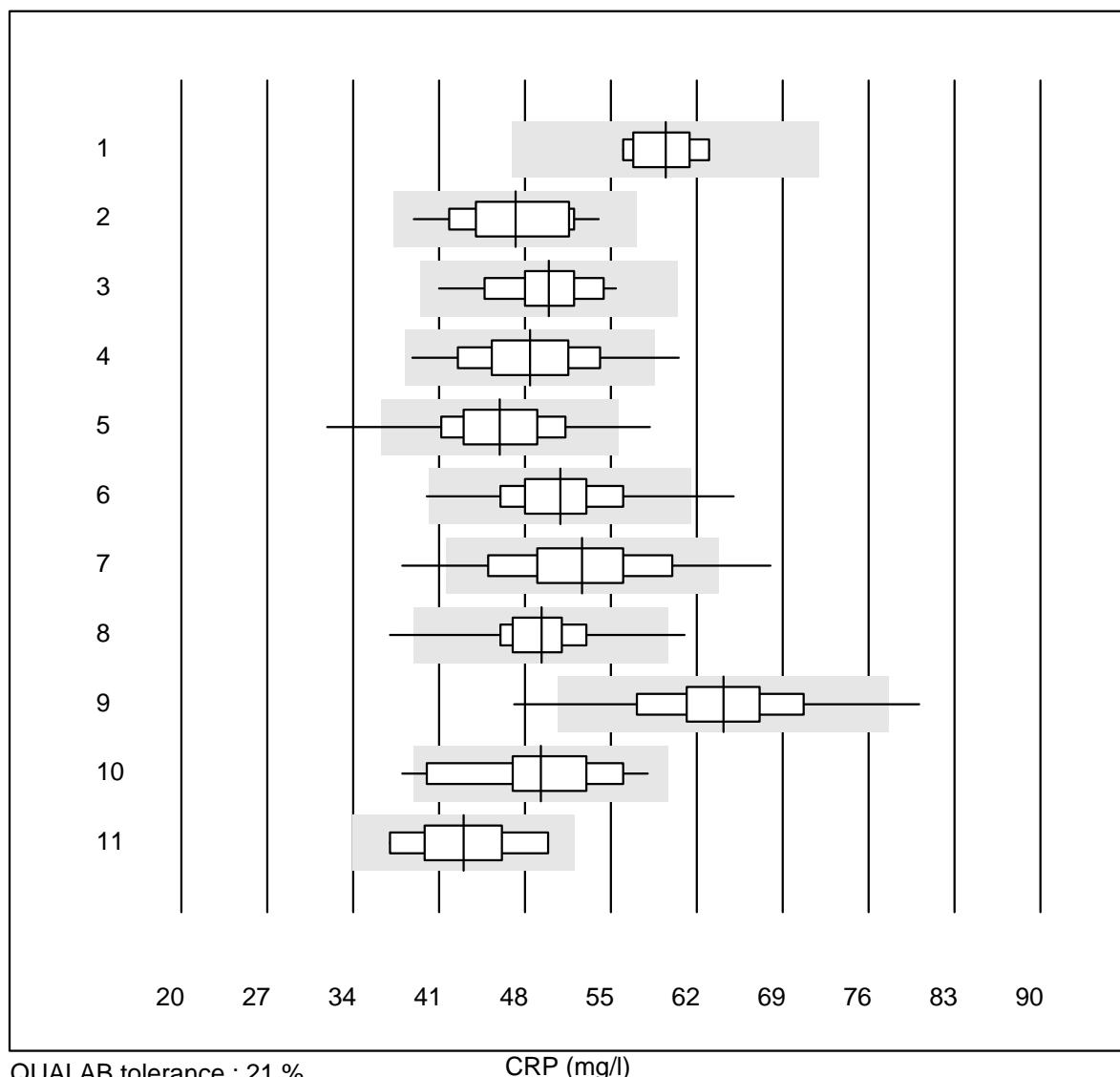
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|---------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 Automat | 23 | 100.0 | 0.0 | 0.0 | 245.5 | 9.0 | e |
| 2 Microscopic | 28 | 92.8 | 3.6 | 3.6 | 225.6 | 14.2 | e |
| 3 Sysmex X | 39 | 100.0 | 0.0 | 0.0 | 246.7 | 4.7 | e |
| 4 ABX Pentra | 10 | 100.0 | 0.0 | 0.0 | 260.4 | 4.5 | e |
| 5 Celldyn | 4 | 100.0 | 0.0 | 0.0 | 249.0 | 5.9 | e |

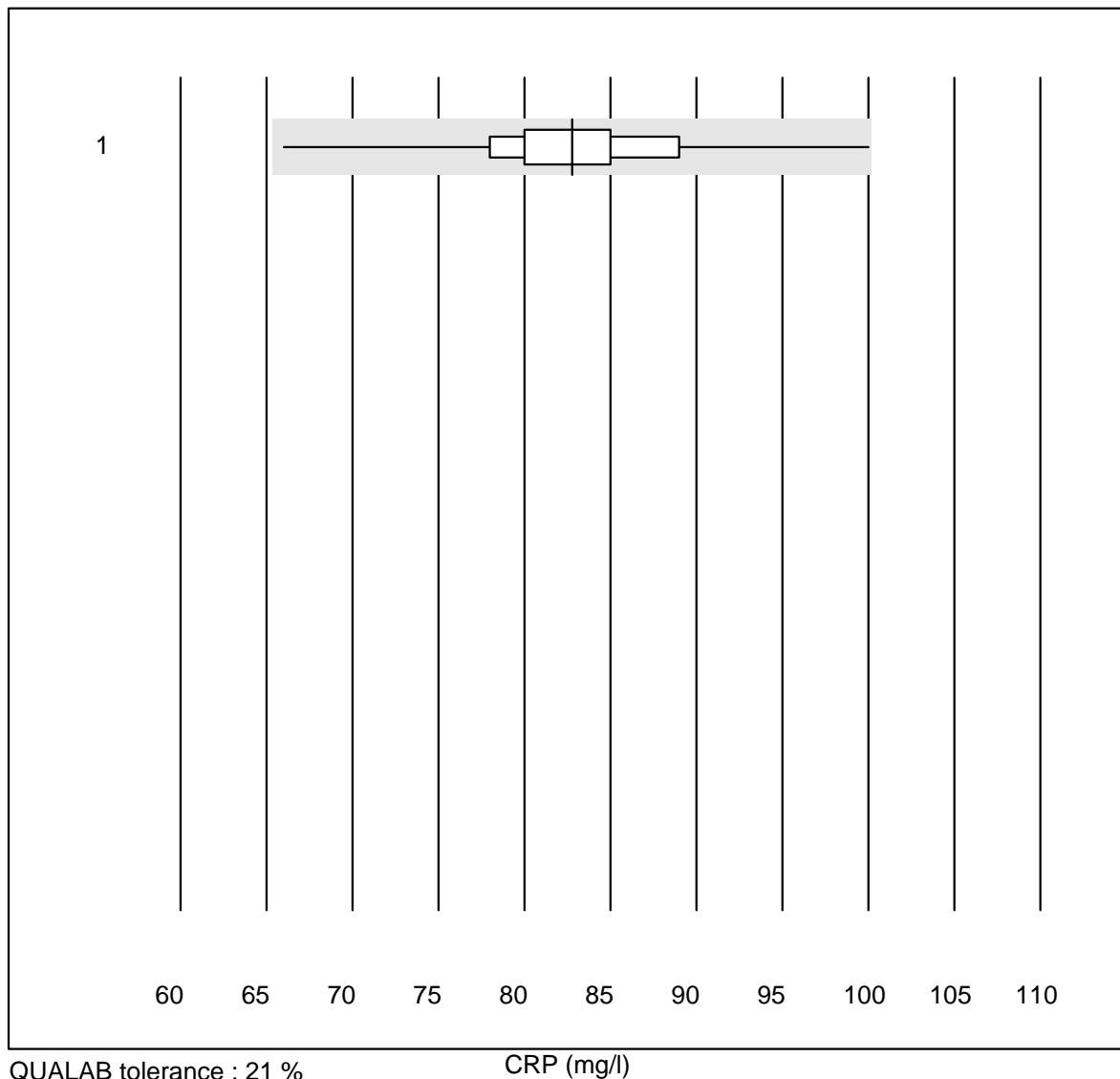
Thrombocytes



| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-------------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 Abx Micros | 739 | 97.6 | 1.6 | 0.8 | 248.1 | 8.5 | e |
| 2 Microsemi | 365 | 99.4 | 0.3 | 0.3 | 252.1 | 5.9 | e |
| 3 Sysmex KX21 | 397 | 99.2 | 0.5 | 0.3 | 253.0 | 4.9 | e |
| 4 Sysmex Poch - 100i | 206 | 100.0 | 0.0 | 0.0 | 257.8 | 4.5 | e |
| 5 Sysmex XP 300 | 260 | 100.0 | 0.0 | 0.0 | 265.4 | 5.1 | e |
| 6 Mythic | 242 | 95.5 | 3.3 | 1.2 | 232.5 | 10.7 | e |
| 7 Swelab | 68 | 97.0 | 1.5 | 1.5 | 223.6 | 8.8 | e |
| 8 Abacus Junior | 12 | 100.0 | 0.0 | 0.0 | 279.9 | 8.1 | e |
| 9 Medonic | 15 | 100.0 | 0.0 | 0.0 | 228.4 | 8.8 | e |
| 10 Nihon Kohden Celltac | 35 | 97.1 | 0.0 | 2.9 | 266.1 | 5.5 | e |
| 11 Samsung HC10 | 45 | 100.0 | 0.0 | 0.0 | 250.5 | 7.2 | e |
| 12 Norma Icon 3 | 26 | 100.0 | 0.0 | 0.0 | 219.1 | 7.0 | e |

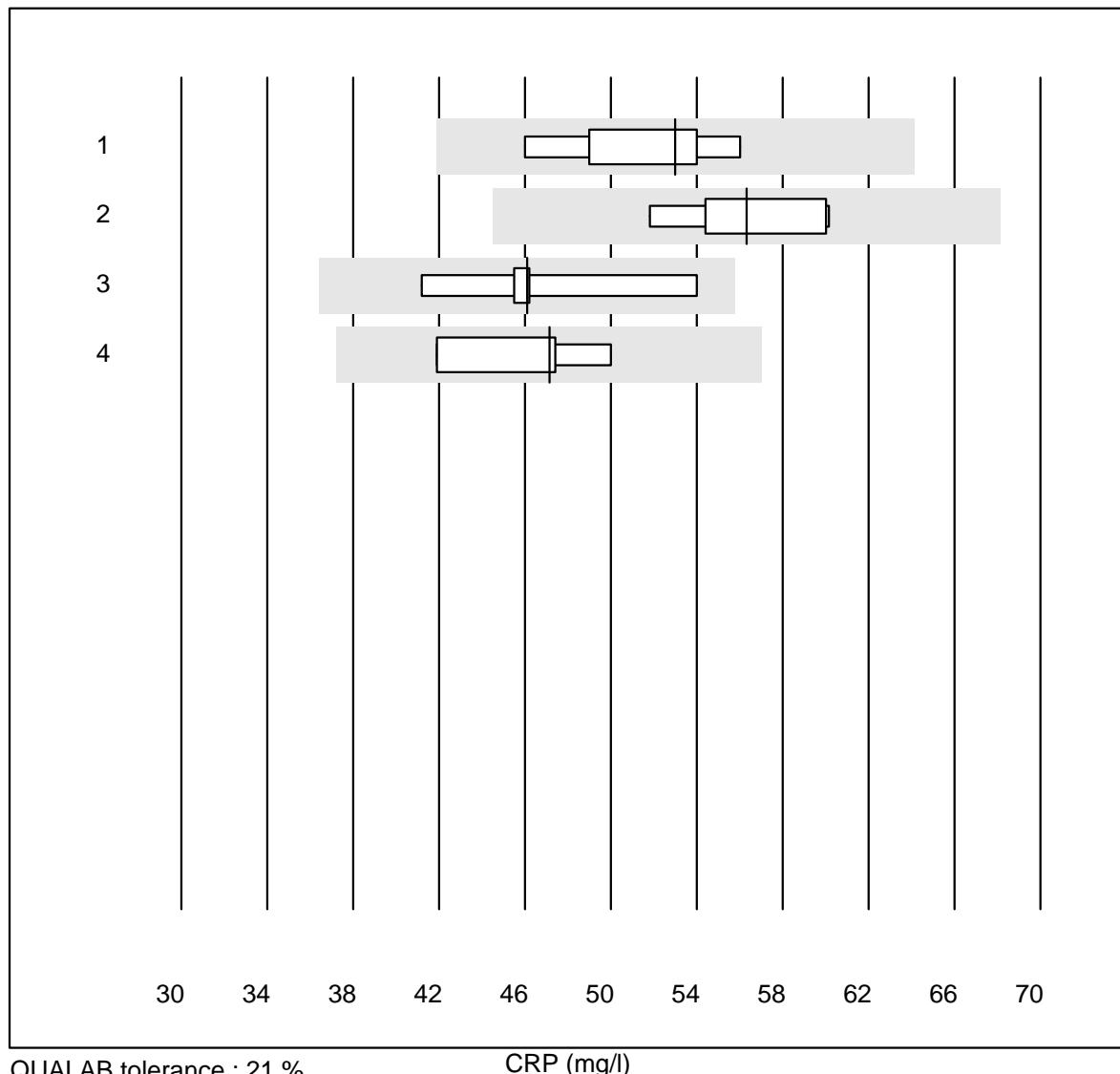
CRP

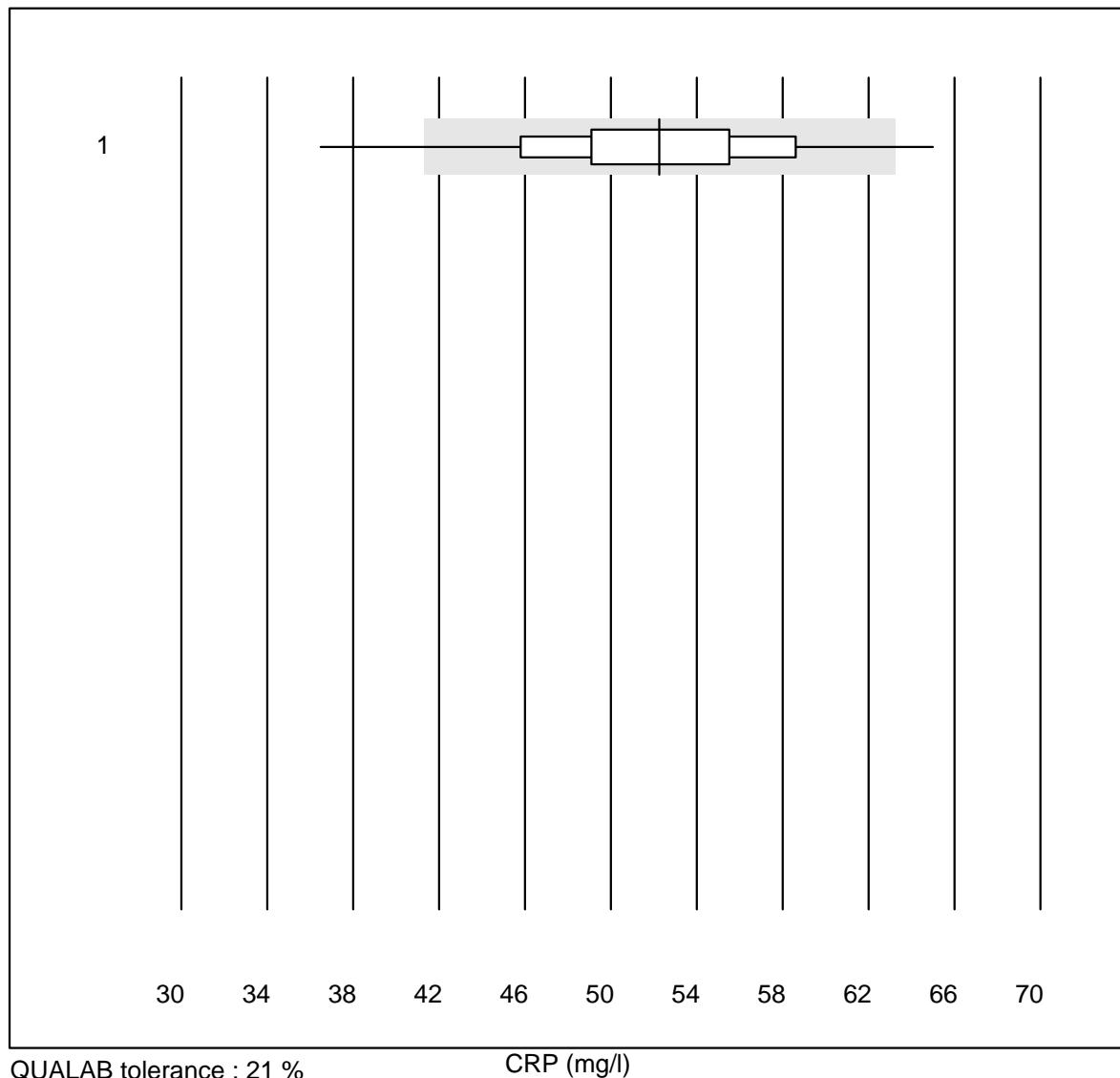


CRP

| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | QuikRead (Vollblut) | 143 | 99.3 | 0.0 | 0.7 | 82.8 | 6.0 | e |

CRP

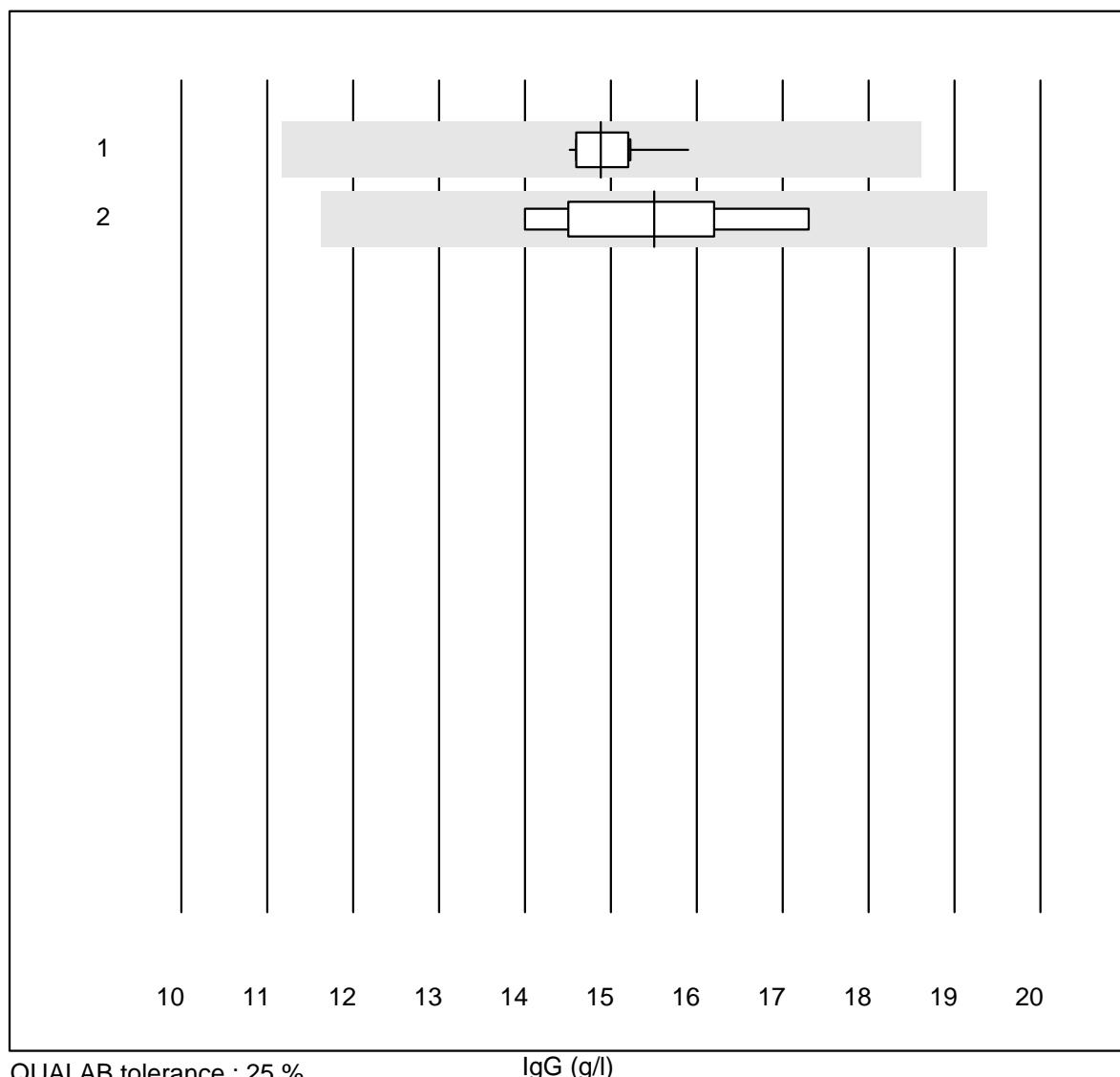


CRP

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Microsemi | 362 | 95.0 | 3.6 | 1.4 | 52.3 | 9.8 | e |

I2 Plasmaproteins

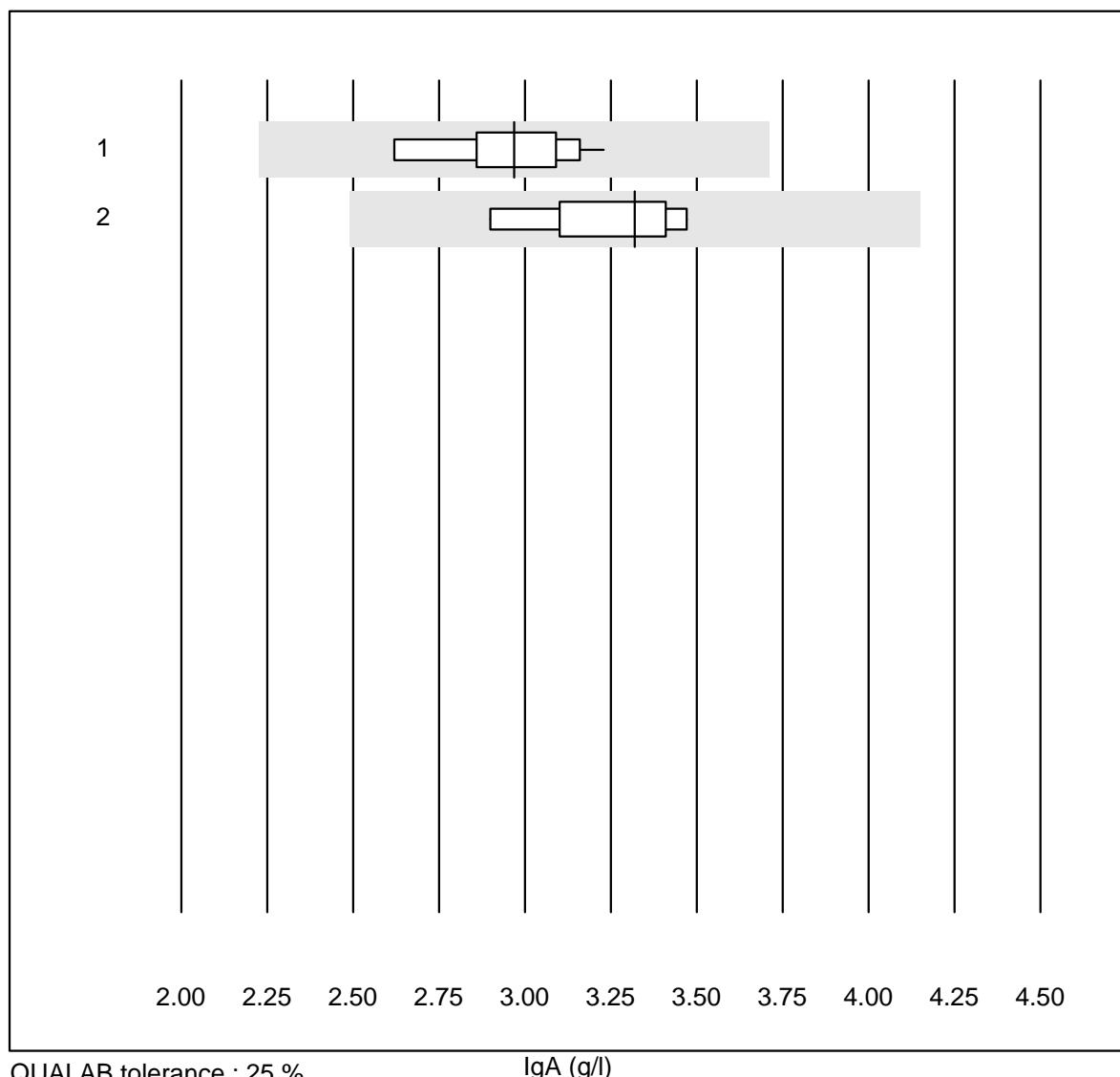
IgG



| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Turbidimetry | 11 | 100.0 | 0.0 | 0.0 | 14.9 | 2.8 | e |
| 2 Nephelometry | 7 | 100.0 | 0.0 | 0.0 | 15.5 | 7.4 | e |

I2 Plasmaproteins

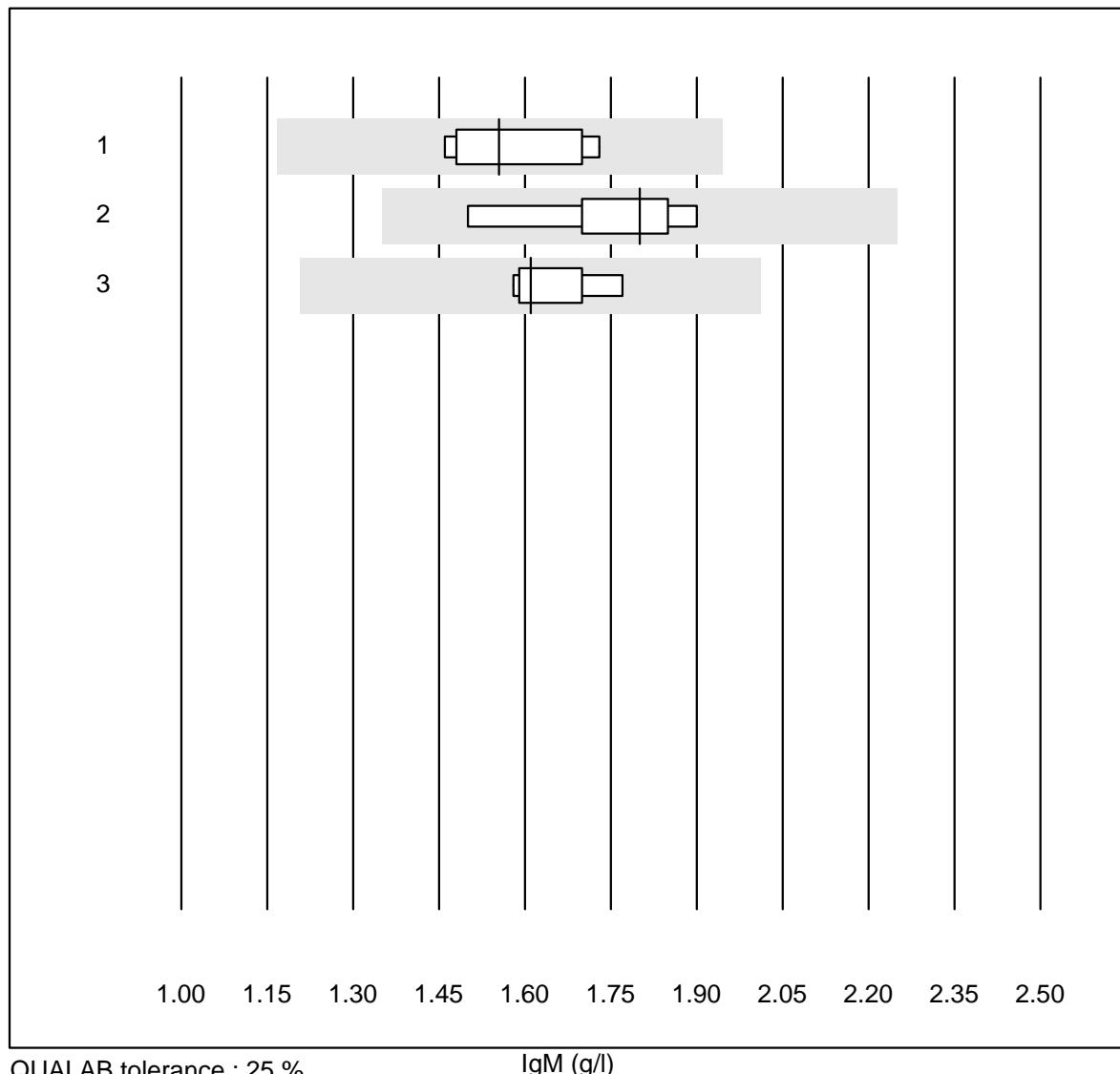
IgA



| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Turbidimetry | 10 | 100.0 | 0.0 | 0.0 | 3.0 | 5.9 | e |
| 2 Nephelometry | 7 | 100.0 | 0.0 | 0.0 | 3.3 | 6.0 | e |

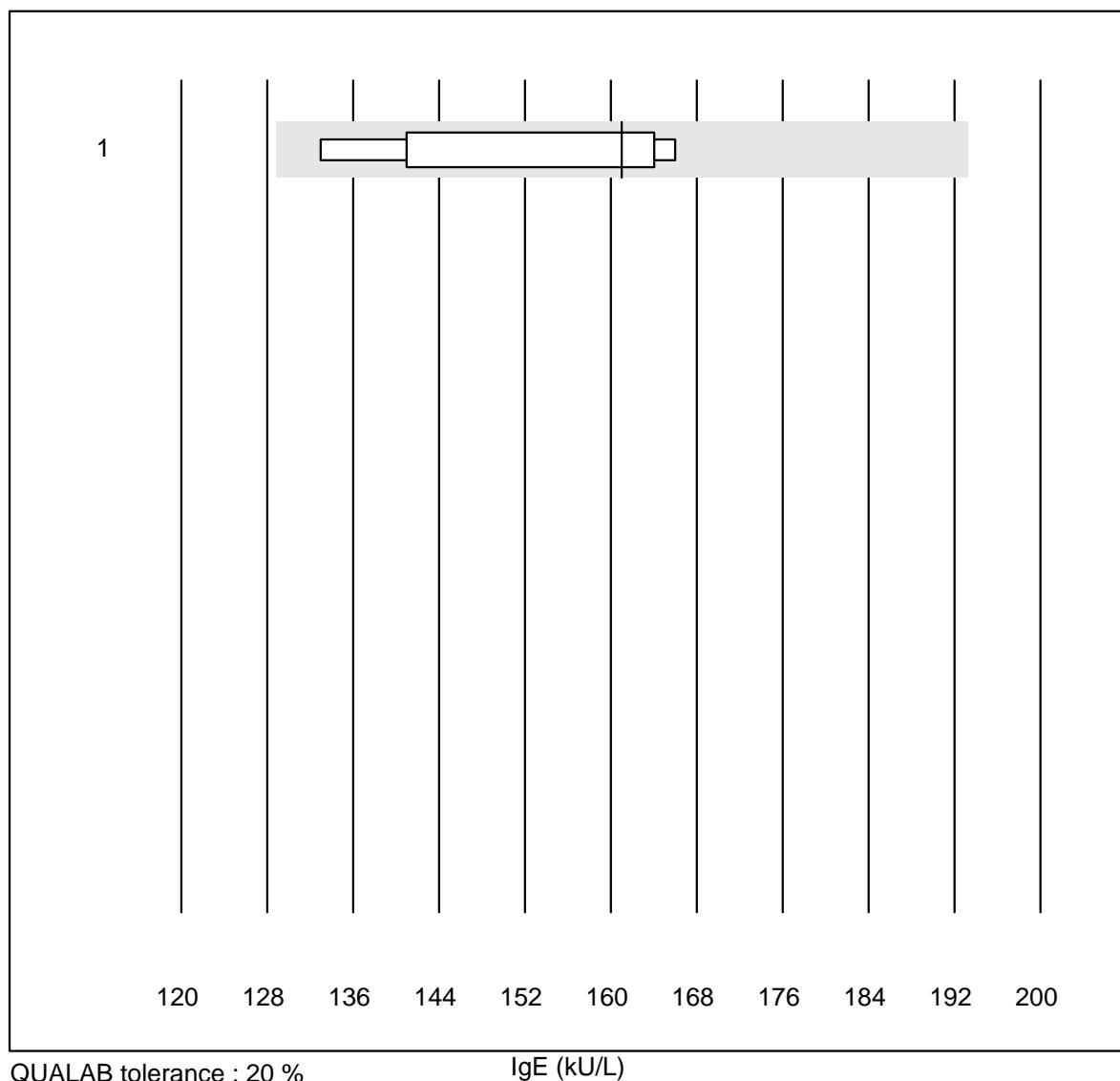
I2 Plasmaproteins

IgM



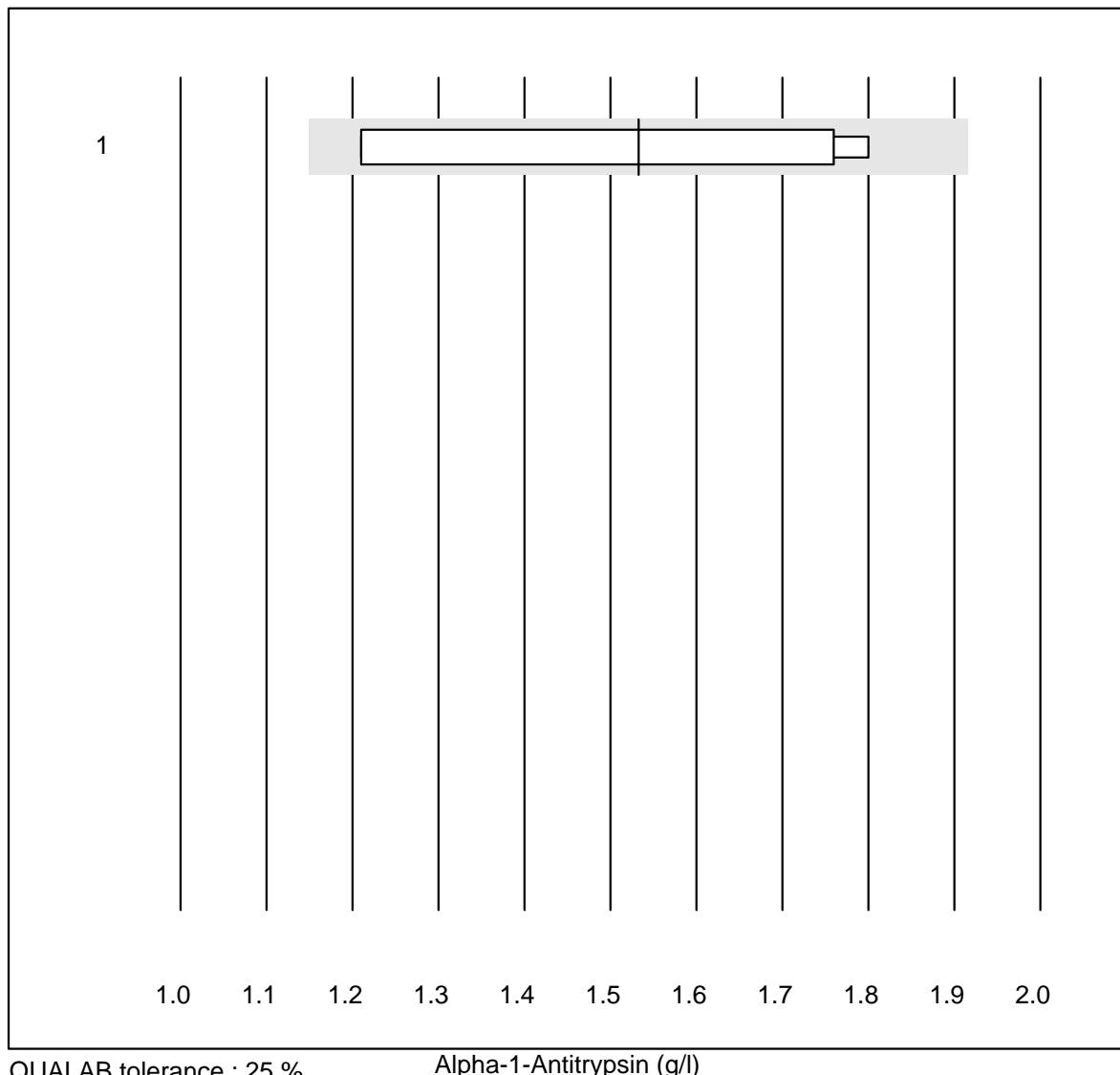
I2 Plasmaproteins

IgE



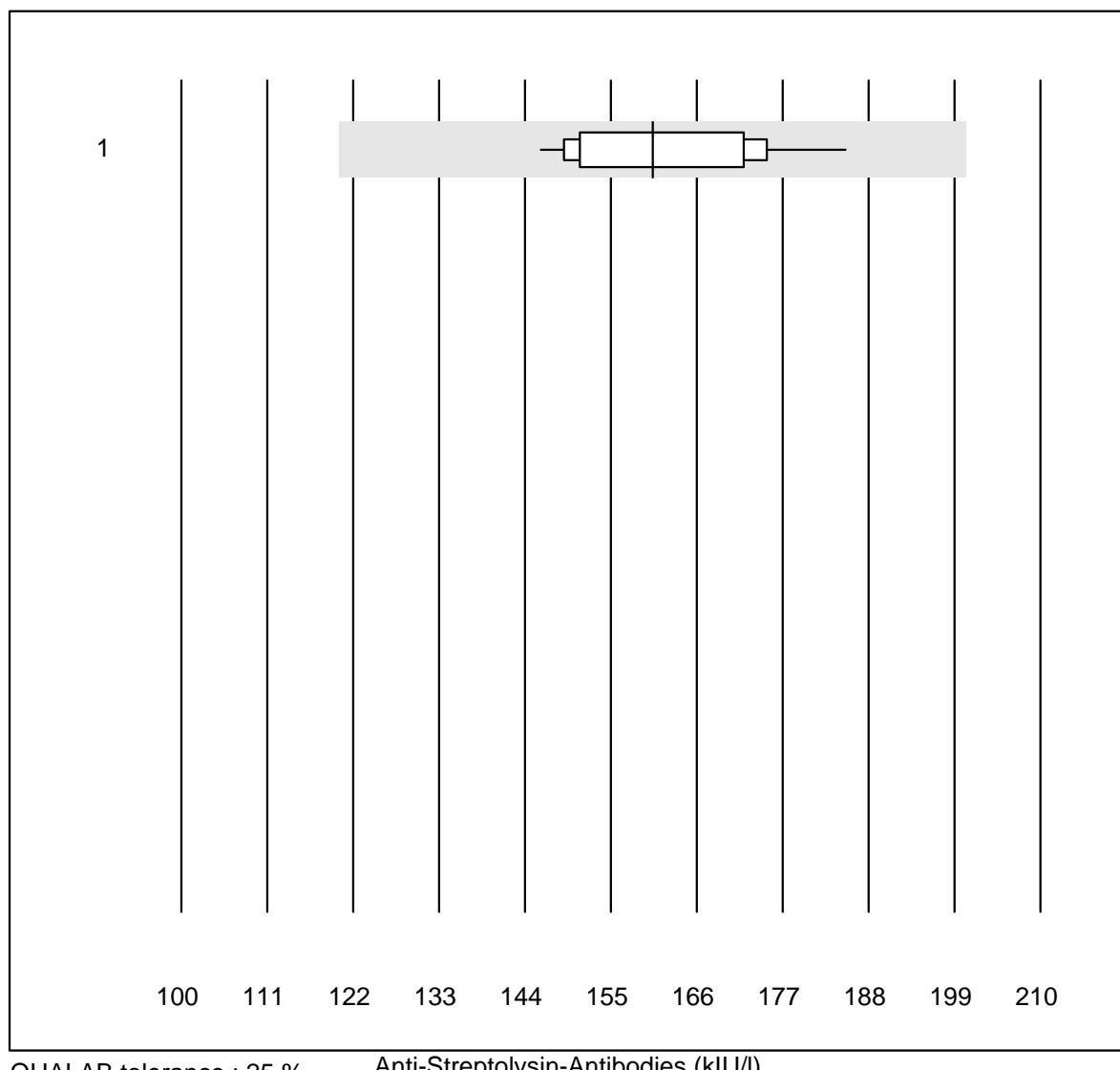
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 all Participants | 9 | 77.8 | 0.0 | 22.2 | 161 | 8.2 | e* |

Alpha-1-Antitrypsin

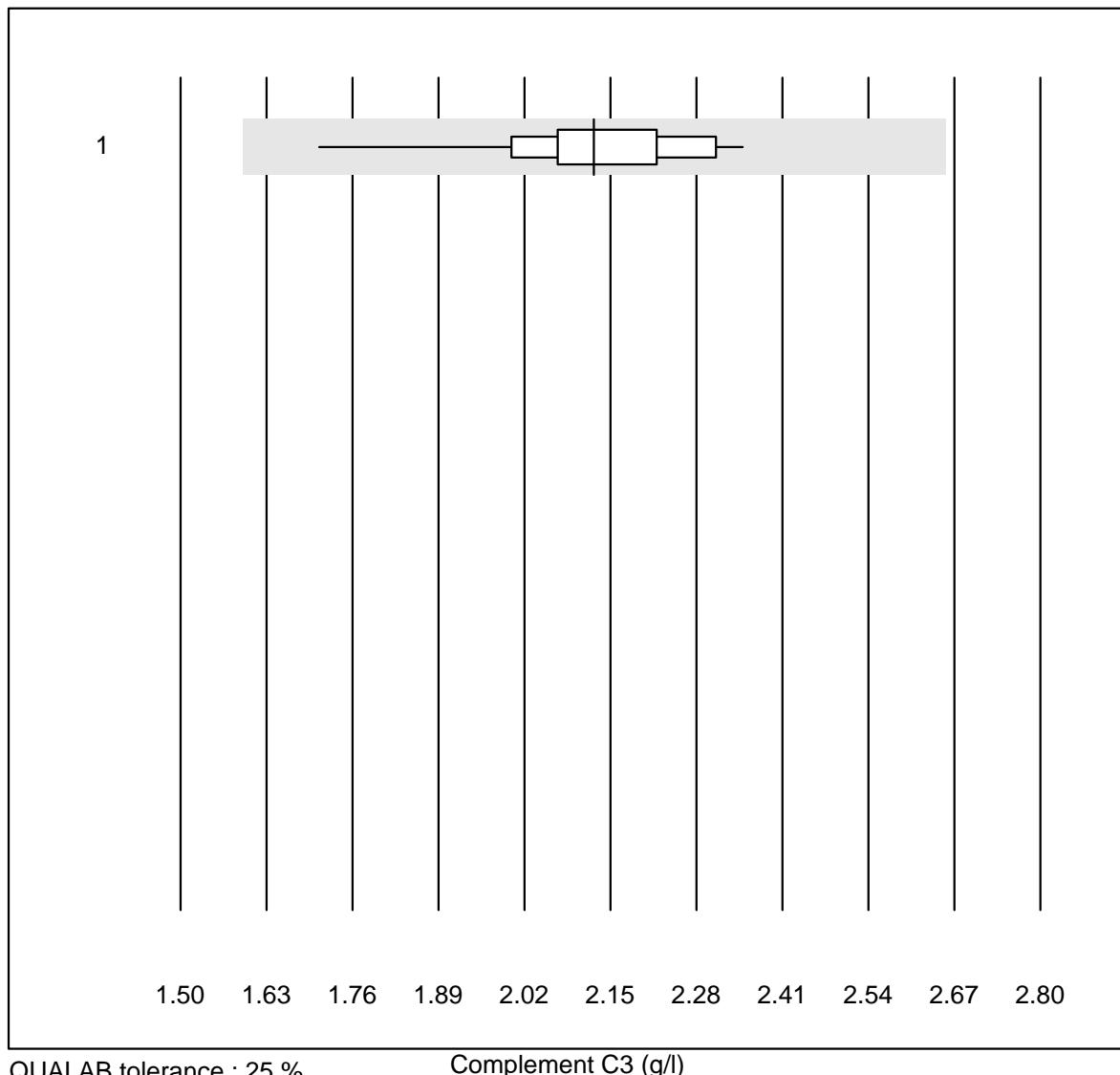


| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|--------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 | Nephelometry | 4 | 100.0 | 0.0 | 0.0 | 1.53 | 19.1 | a |

Anti-Streptolysin-Antibodies

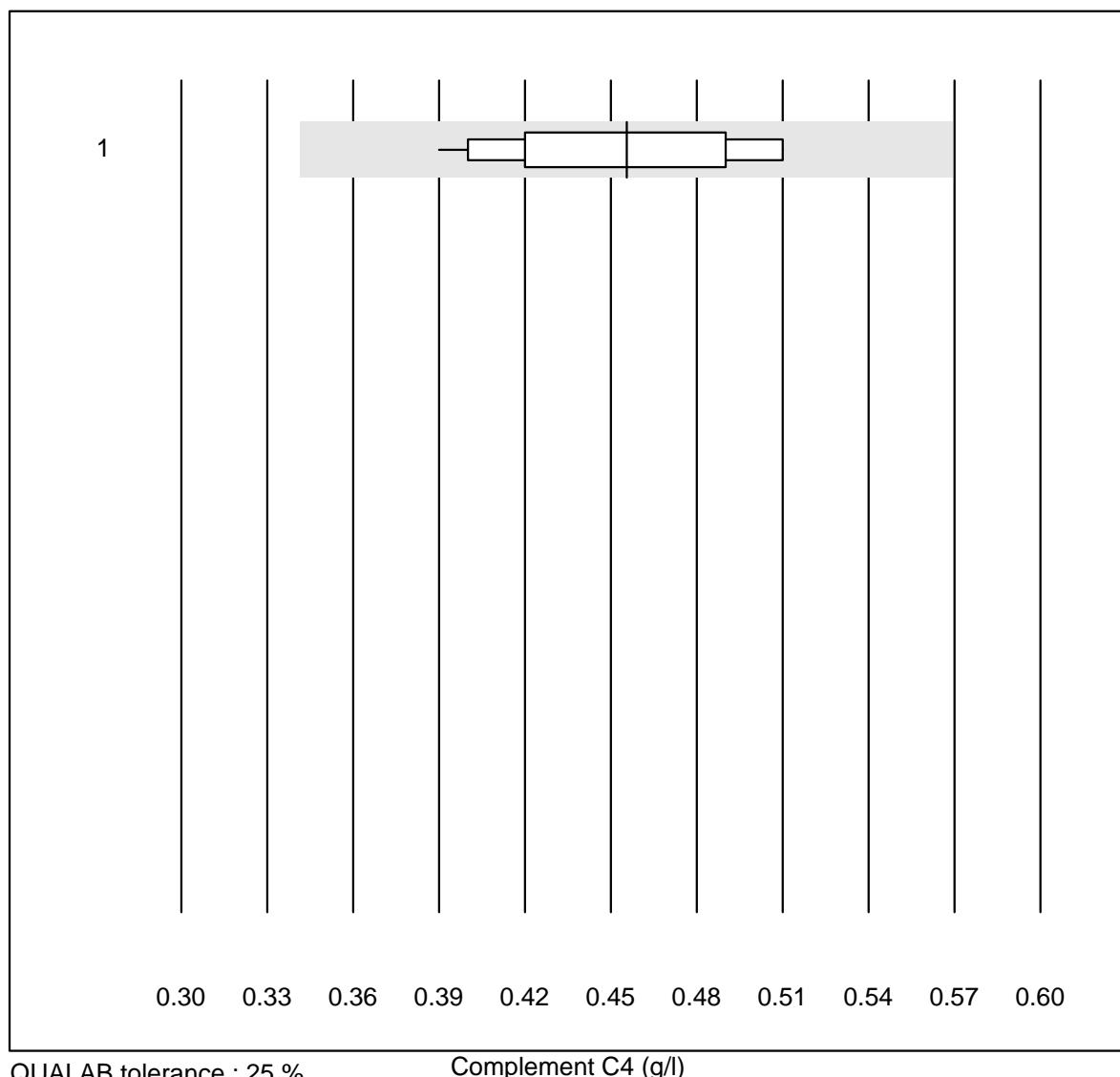


Complement C3



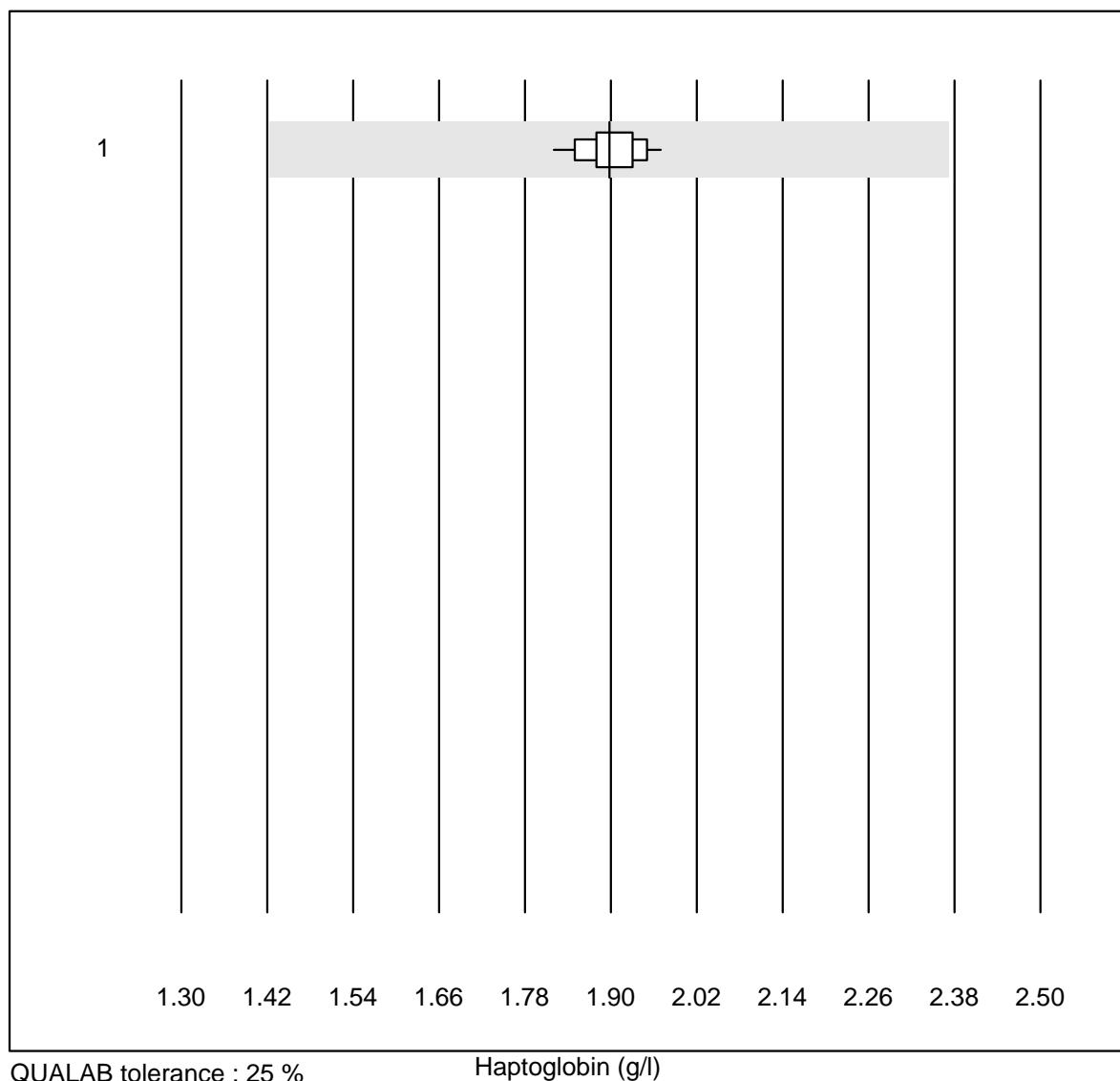
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 all Participants | 13 | 100.0 | 0.0 | 0.0 | 2.13 | 7.7 | e |

Complement C4



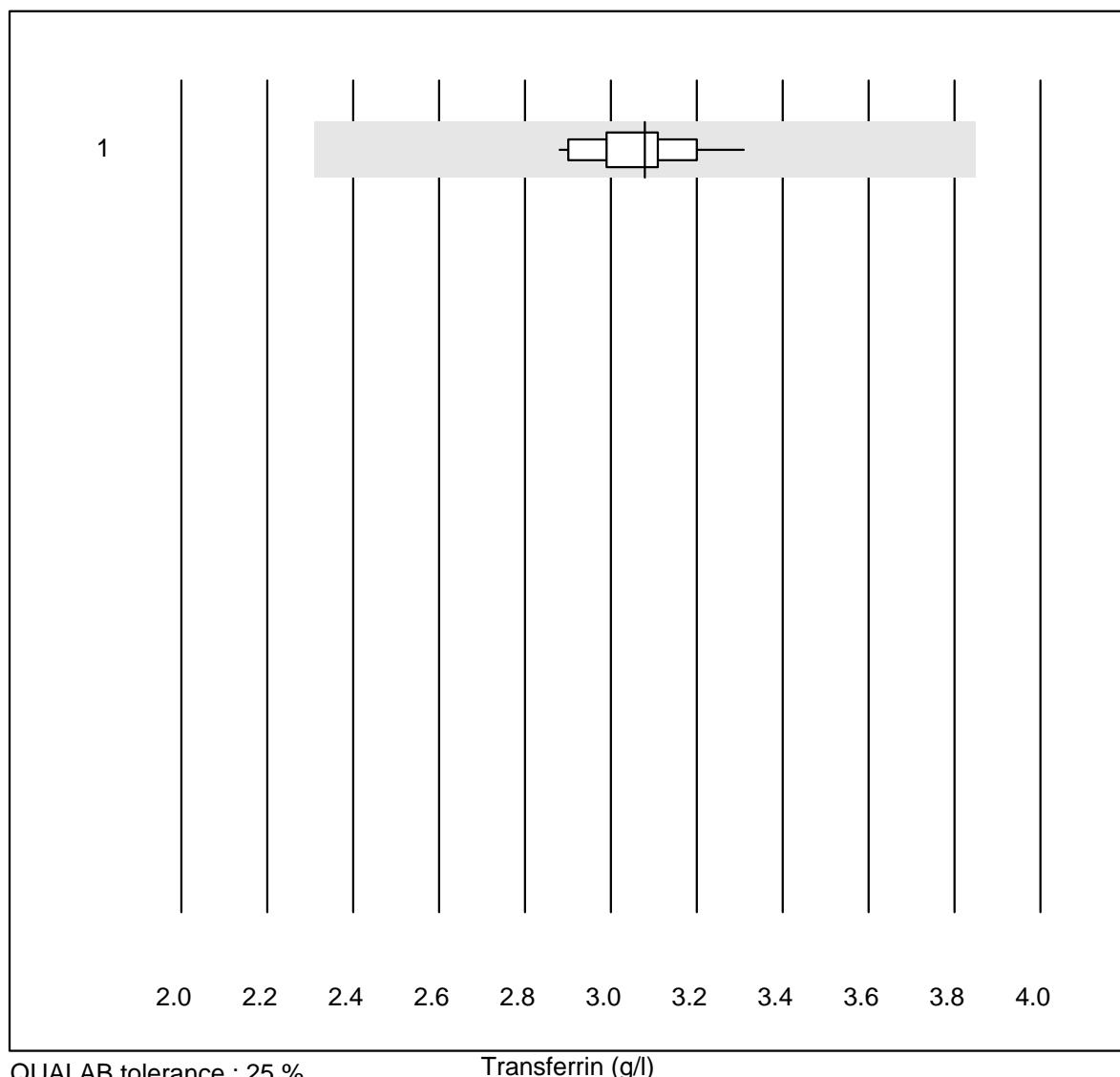
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | all Participants | 12 | 100.0 | 0.0 | 0.0 | 0.46 | 9.5 | e |

Haptoglobin

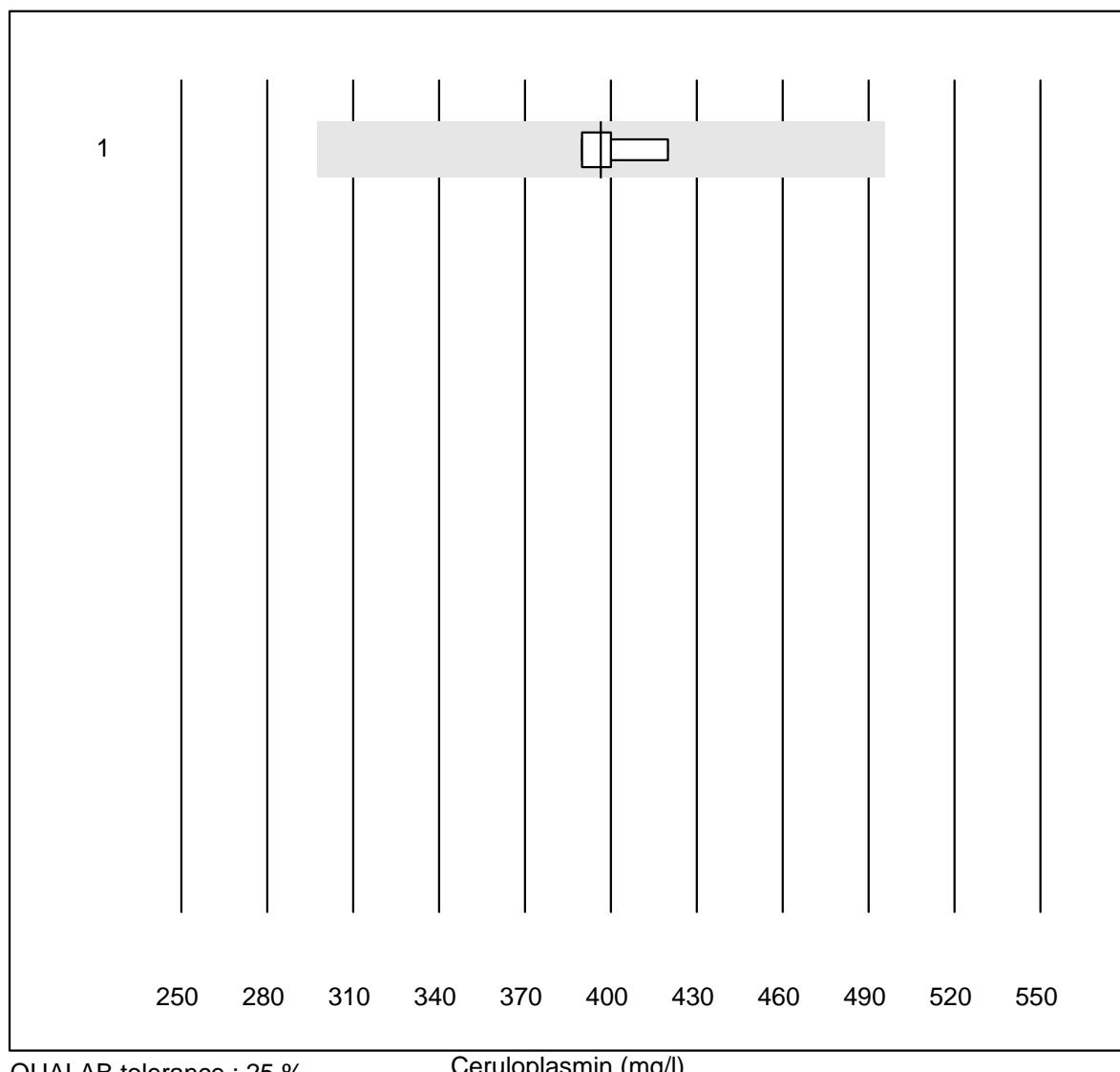


| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 all Participants | 13 | 100.0 | 0.0 | 0.0 | 1.90 | 2.2 | e |

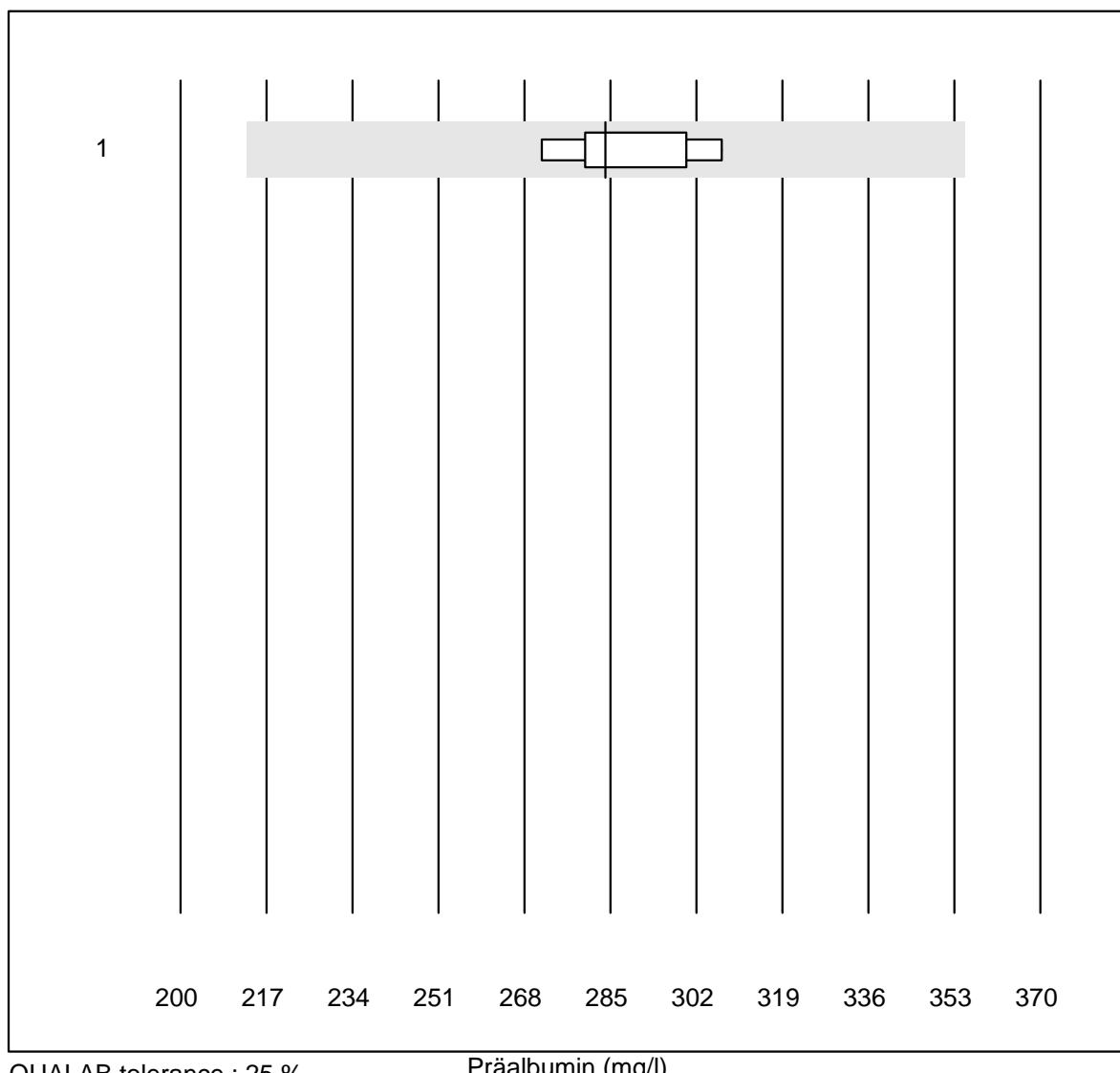
Transferrin



Ceruloplasmin

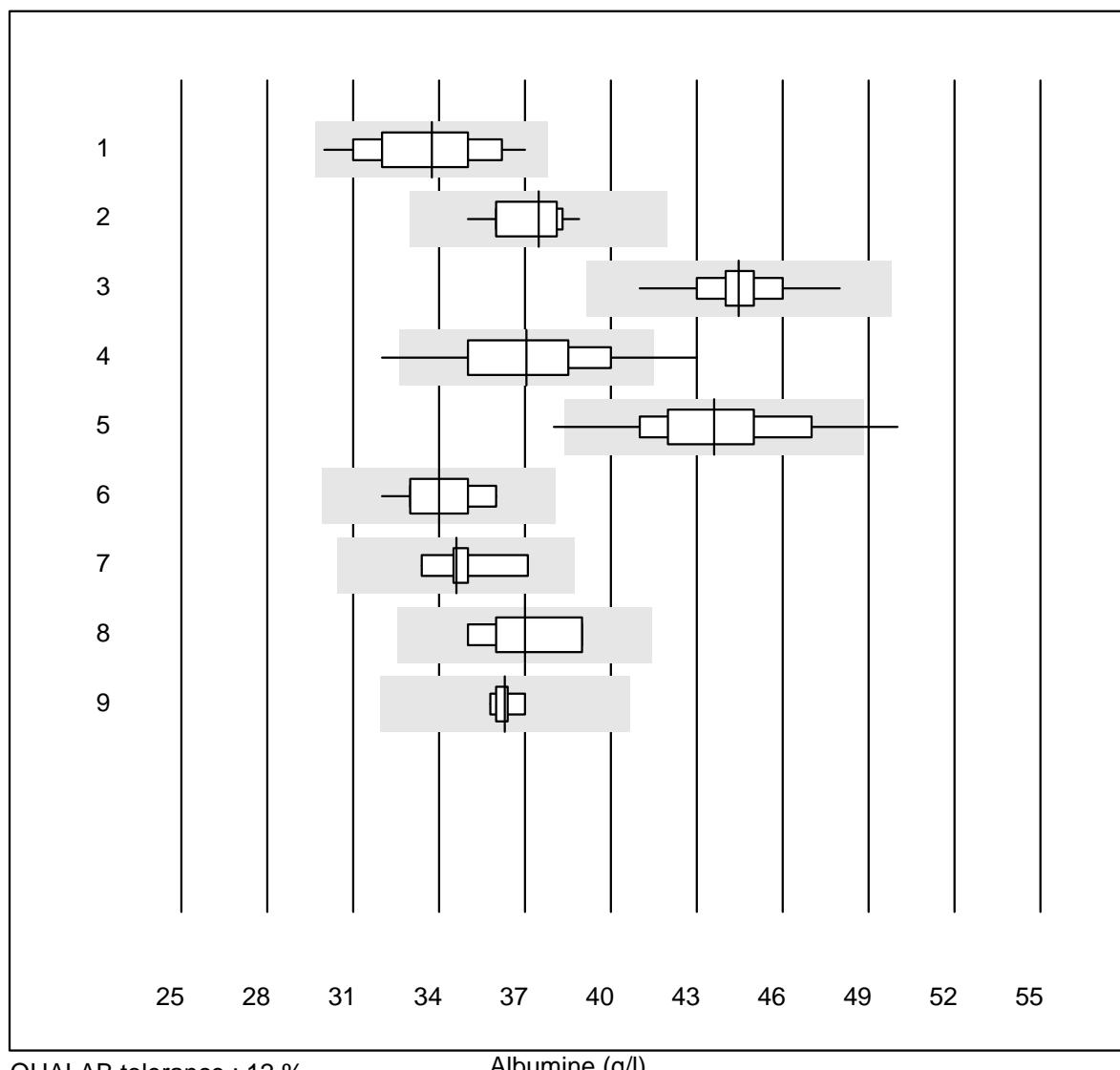


| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 all Participants | 4 | 100.0 | 0.0 | 0.0 | 396.5 | 3.4 | e |

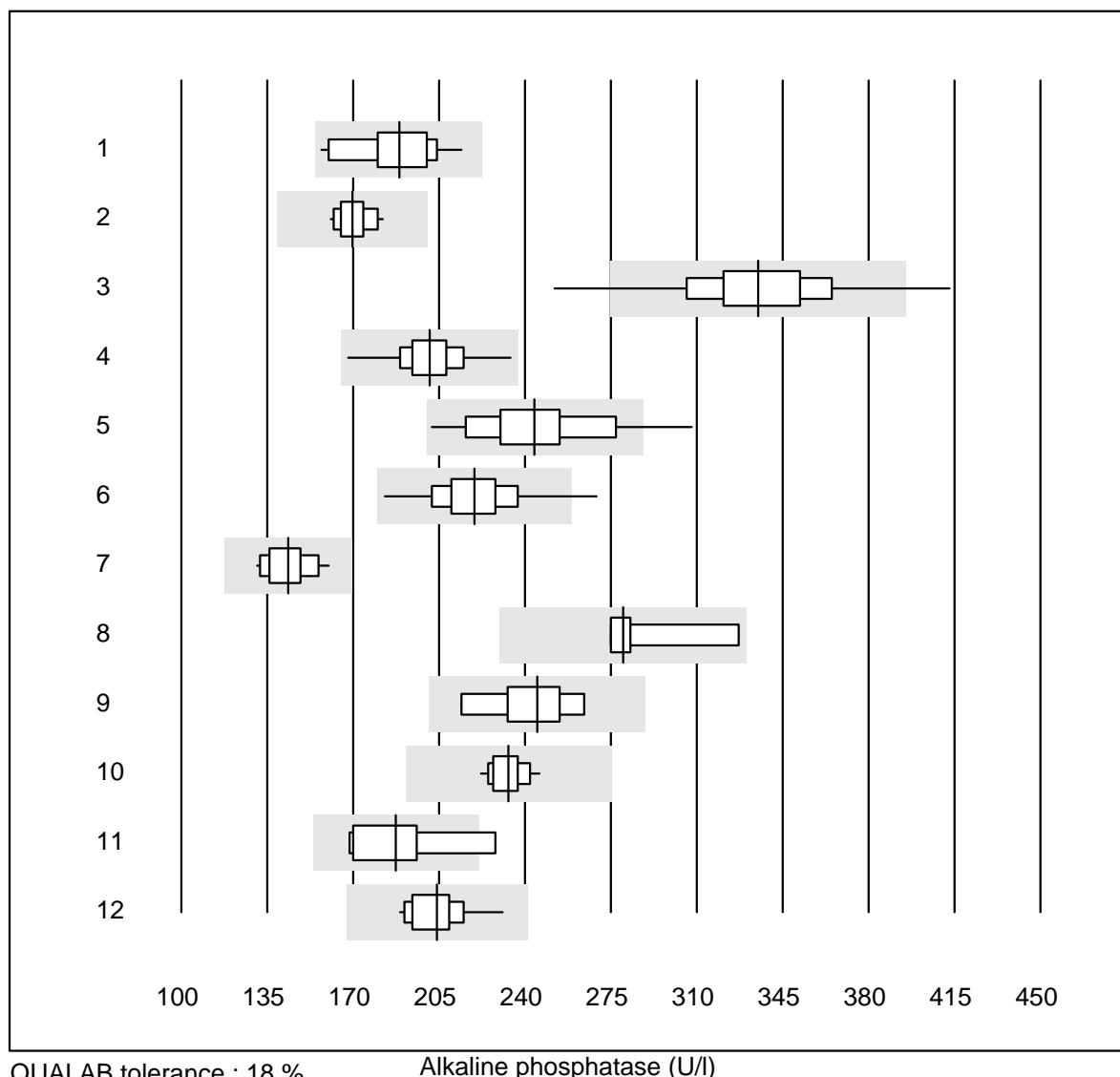
Präalbumin

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 all Participants | 7 | 100.0 | 0.0 | 0.0 | 284.0 | 4.3 | e |

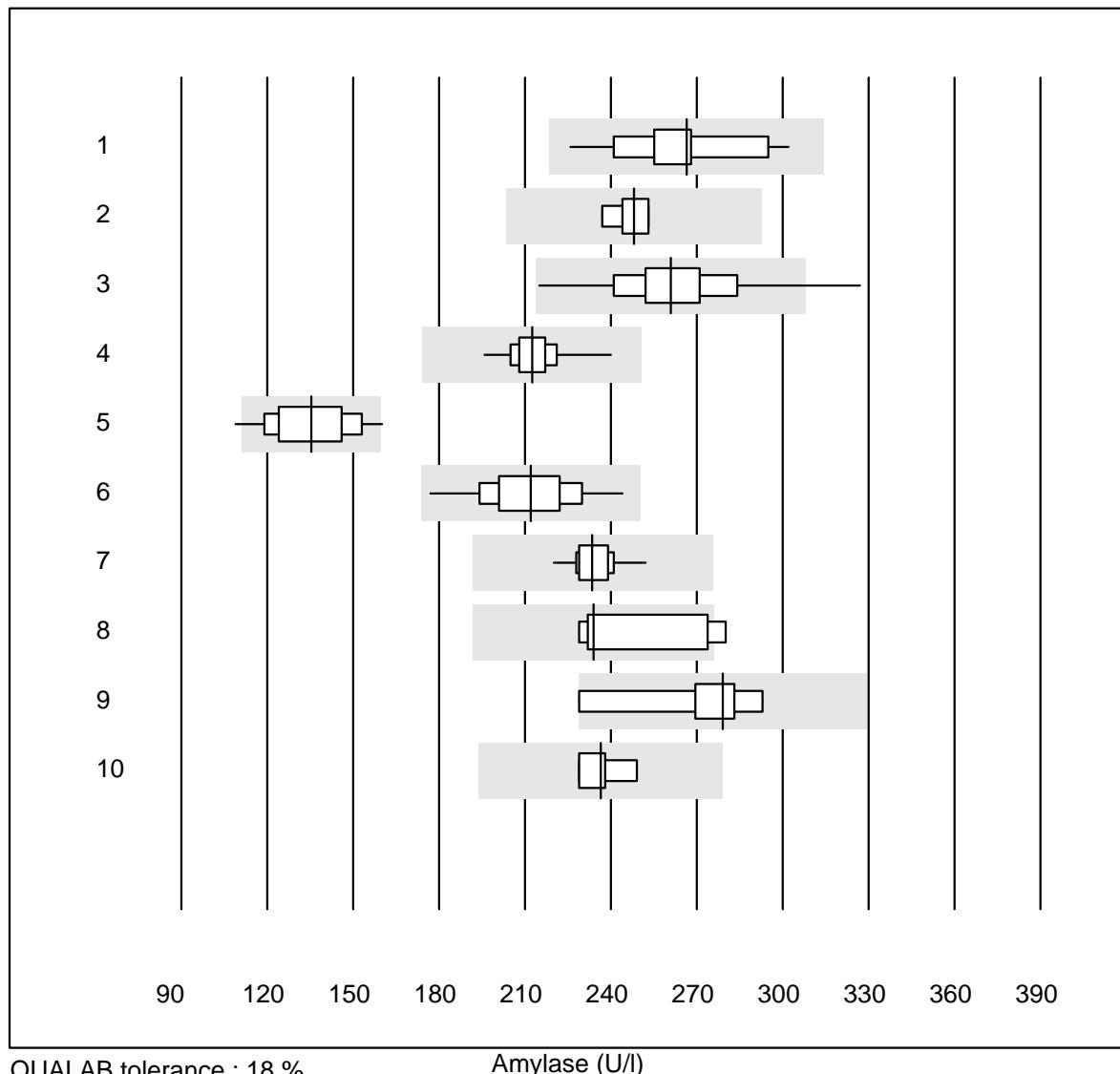
Albumine



Alkaline phosphatase

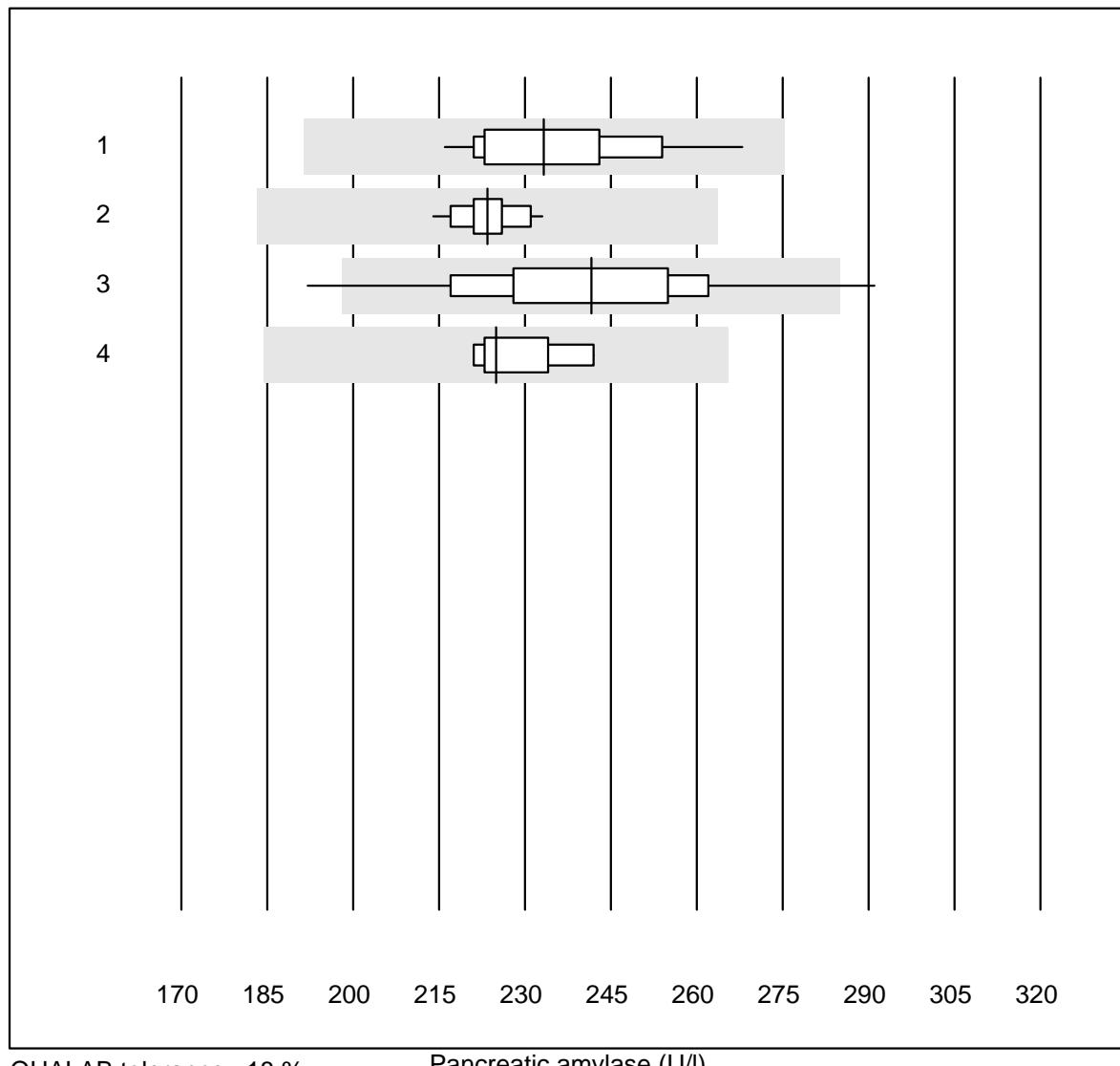


Amylase

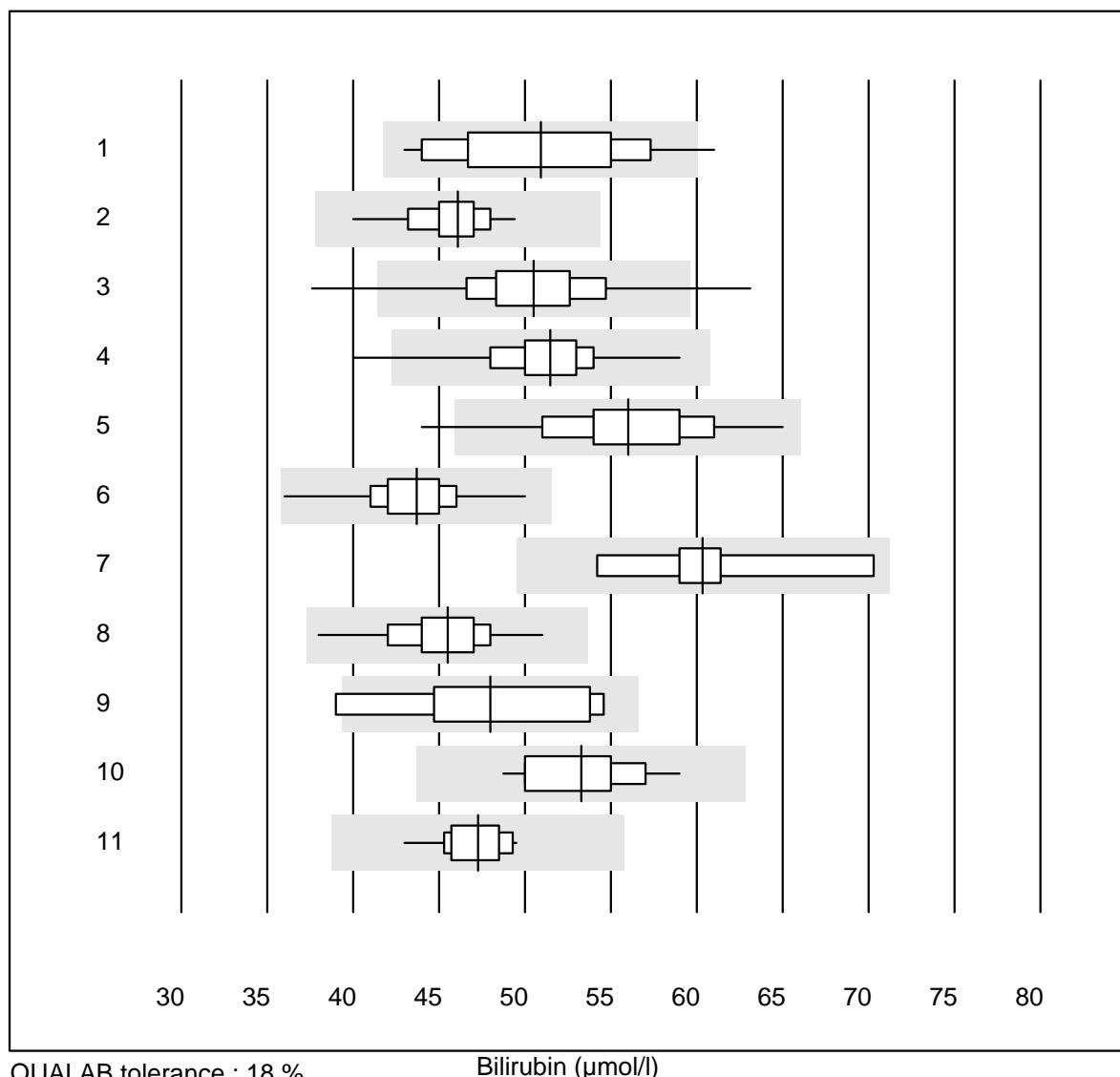


| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 IFCC | 12 | 100.0 | 0.0 | 0.0 | 267 | 8.2 | e* |
| 2 Cobas | 5 | 100.0 | 0.0 | 0.0 | 248 | 2.7 | e |
| 3 Reflotron | 179 | 98.3 | 1.1 | 0.6 | 261 | 6.5 | e |
| 4 Fuji Dri-Chem | 522 | 99.2 | 0.0 | 0.8 | 213 | 3.0 | e |
| 5 Spotchem/Ready | 73 | 84.9 | 4.1 | 11.0 | 135 | 9.7 | e |
| 6 Spotchem D-Concept | 115 | 99.1 | 0.0 | 0.9 | 212 | 6.4 | e |
| 7 Piccolo | 27 | 100.0 | 0.0 | 0.0 | 234 | 2.9 | e |
| 8 Abx Mira | 5 | 80.0 | 20.0 | 0.0 | 234 | 10.0 | e* |
| 9 Hitachi S40/M40 | 9 | 88.9 | 0.0 | 11.1 | 279 | 7.4 | e* |
| 10 Autolyser/DiaSys | 4 | 100.0 | 0.0 | 0.0 | 237 | 3.5 | e |

Pancreatic amylase

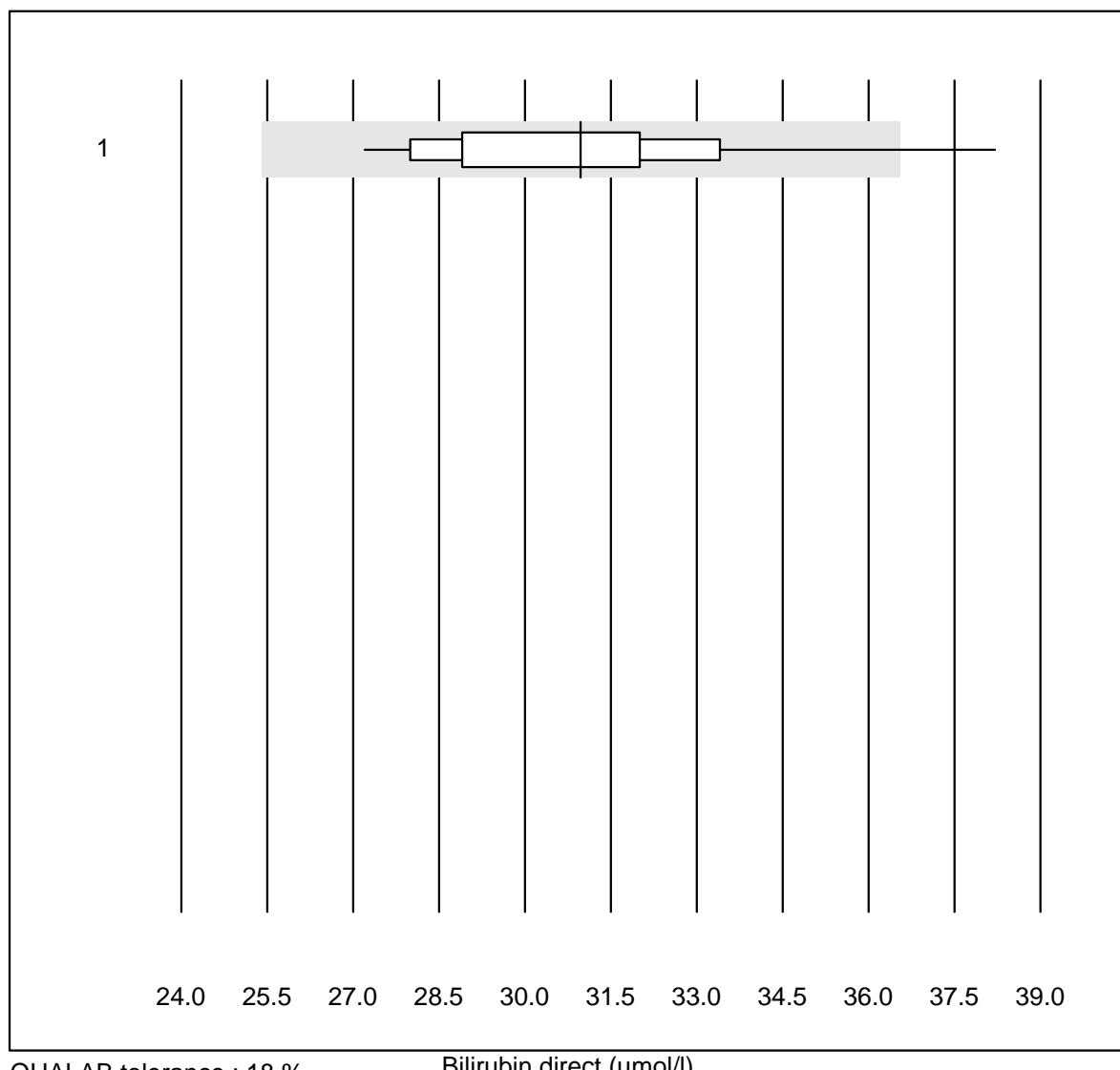


Bilirubin

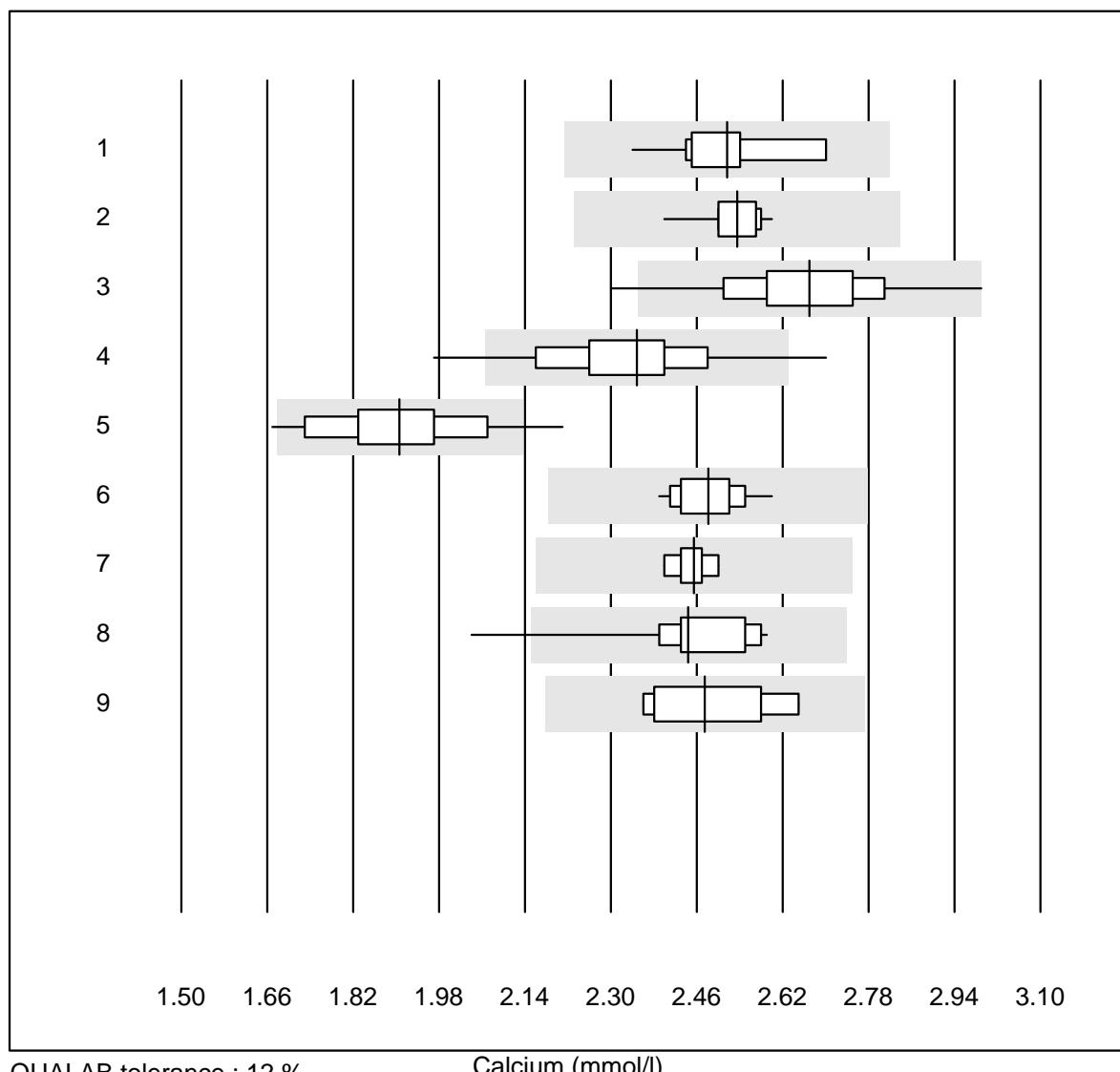


| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 Standard chemistry | 16 | 93.7 | 6.3 | 0.0 | 50.9 | 10.5 | e* |
| 2 Cobas | 15 | 100.0 | 0.0 | 0.0 | 46.1 | 4.8 | e |
| 3 Reflotron | 466 | 95.7 | 2.4 | 1.9 | 50.5 | 6.9 | e |
| 4 Fuji Dri-Chem | 534 | 98.8 | 0.6 | 0.6 | 51.5 | 4.7 | e |
| 5 Spotchem/Ready | 90 | 97.8 | 2.2 | 0.0 | 56.0 | 7.4 | e |
| 6 Spotchem D-Concept | 117 | 98.3 | 0.0 | 1.7 | 43.7 | 4.9 | e |
| 7 Beckman | 8 | 100.0 | 0.0 | 0.0 | 60.4 | 7.4 | e* |
| 8 Piccolo | 31 | 96.8 | 0.0 | 3.2 | 45.5 | 5.8 | e |
| 9 Abx Mira | 7 | 85.7 | 14.3 | 0.0 | 48.0 | 11.2 | e* |
| 10 Hitachi S40/M40 | 14 | 100.0 | 0.0 | 0.0 | 53.3 | 5.9 | e |
| 11 Autolyser/DiaSys | 14 | 92.9 | 0.0 | 7.1 | 47.3 | 4.1 | e |

Bilirubin direct

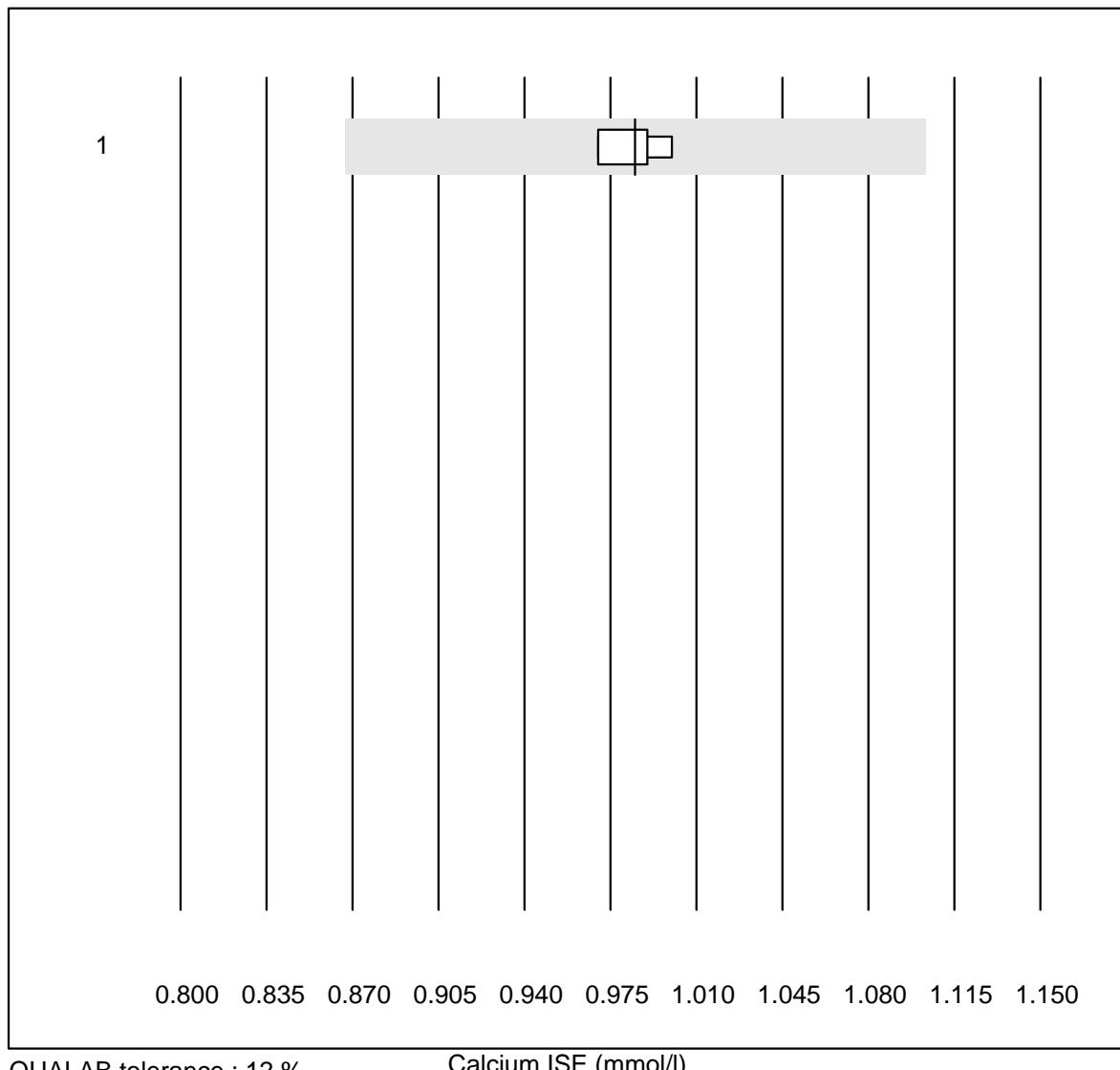


Calcium



| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Standard chemistry | 24 | 95.8 | 0.0 | 4.2 | 2.52 | 3.6 | e |
| 2 Cobas | 12 | 100.0 | 0.0 | 0.0 | 2.54 | 2.1 | e |
| 3 Fuji Dri-Chem | 361 | 97.7 | 0.6 | 1.7 | 2.67 | 4.4 | e |
| 4 Spotchem/Ready | 40 | 95.0 | 5.0 | 0.0 | 2.35 | 6.0 | e |
| 5 Spotchem D-Concept | 70 | 94.3 | 5.7 | 0.0 | 1.91 | 6.1 | e |
| 6 Piccolo | 30 | 100.0 | 0.0 | 0.0 | 2.48 | 2.3 | e |
| 7 Abx Mira | 6 | 100.0 | 0.0 | 0.0 | 2.46 | 1.4 | e |
| 8 Hitachi S40/M40 | 12 | 83.4 | 8.3 | 8.3 | 2.44 | 6.1 | e* |
| 9 Autolyser/DiaSys | 6 | 100.0 | 0.0 | 0.0 | 2.48 | 4.5 | e* |

Calcium ISE

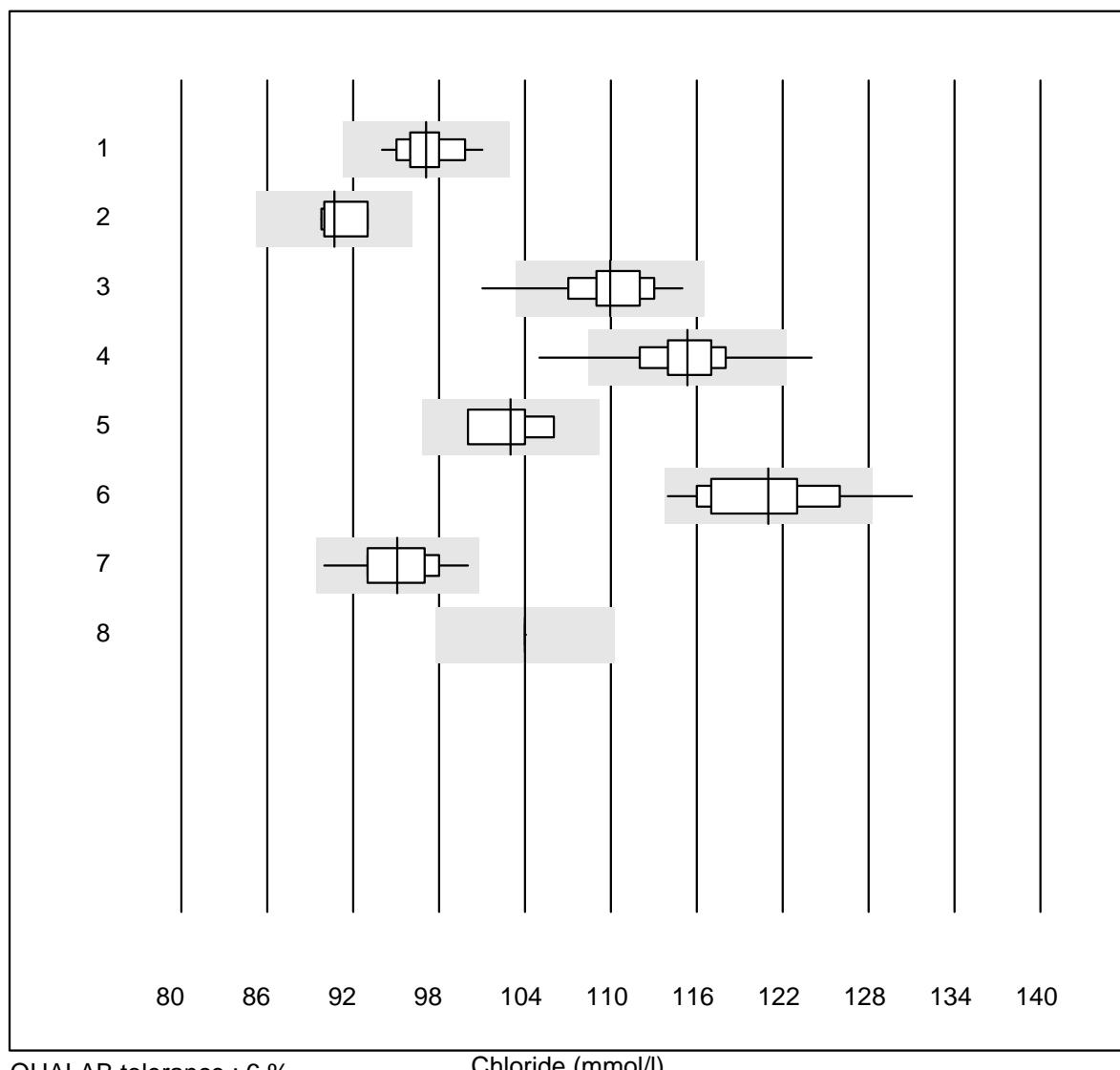


QUALAB tolerance : 12 %

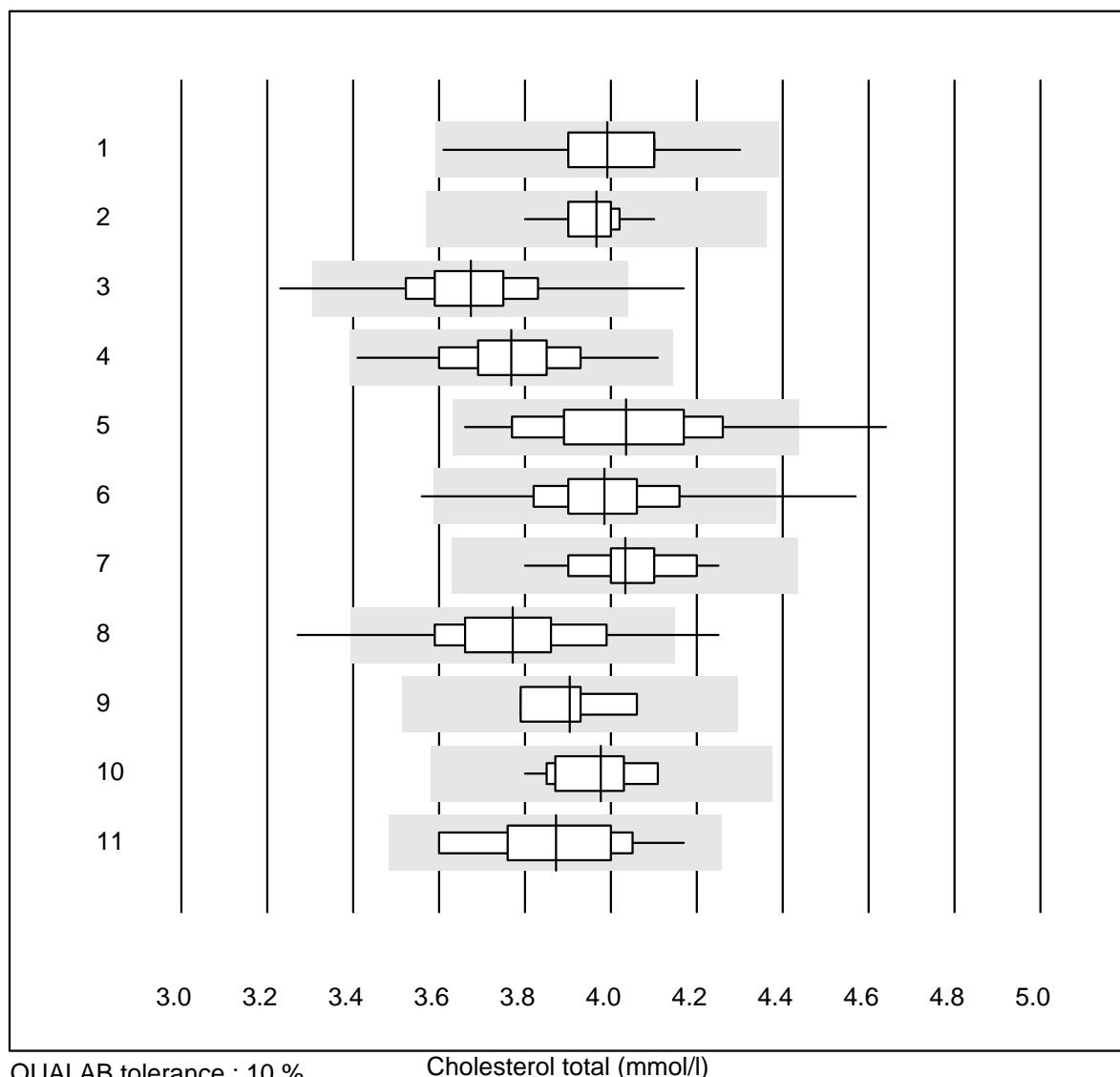
Calcium ISE (mmol/l)

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|---------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 iStat Chem8 | 4 | 100.0 | 0.0 | 0.0 | 0.99 | 1.3 | e |

Chloride

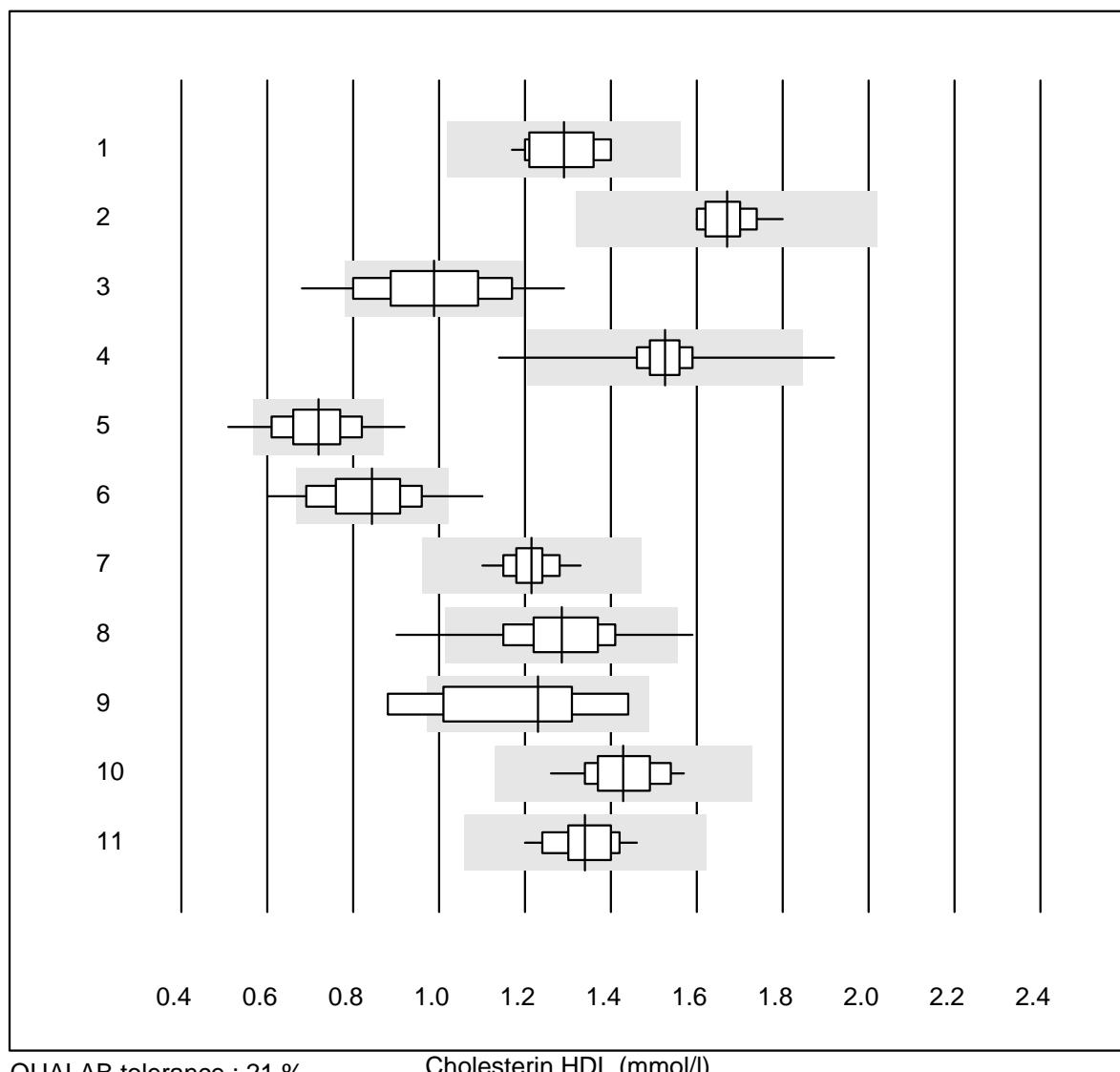


| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 ISE | 17 | 100.0 | 0.0 | 0.0 | 97 | 1.9 | e |
| 2 Cobas | 7 | 100.0 | 0.0 | 0.0 | 91 | 1.5 | e |
| 3 Fuji Dri-Chem | 637 | 96.9 | 2.2 | 0.9 | 110 | 2.2 | e |
| 4 Spotchem D-Concept | 139 | 95.0 | 1.4 | 3.6 | 115 | 2.3 | e |
| 5 Standard chemistry | 4 | 100.0 | 0.0 | 0.0 | 103 | 2.9 | a |
| 6 Spotchem EL-SE 1520 | 112 | 90.2 | 2.7 | 7.1 | 121 | 3.1 | e |
| 7 Piccolo | 16 | 100.0 | 0.0 | 0.0 | 95 | 2.6 | e |
| 8 iStat Chem8 | 4 | 100.0 | 0.0 | 0.0 | 104 | 0.0 | e |

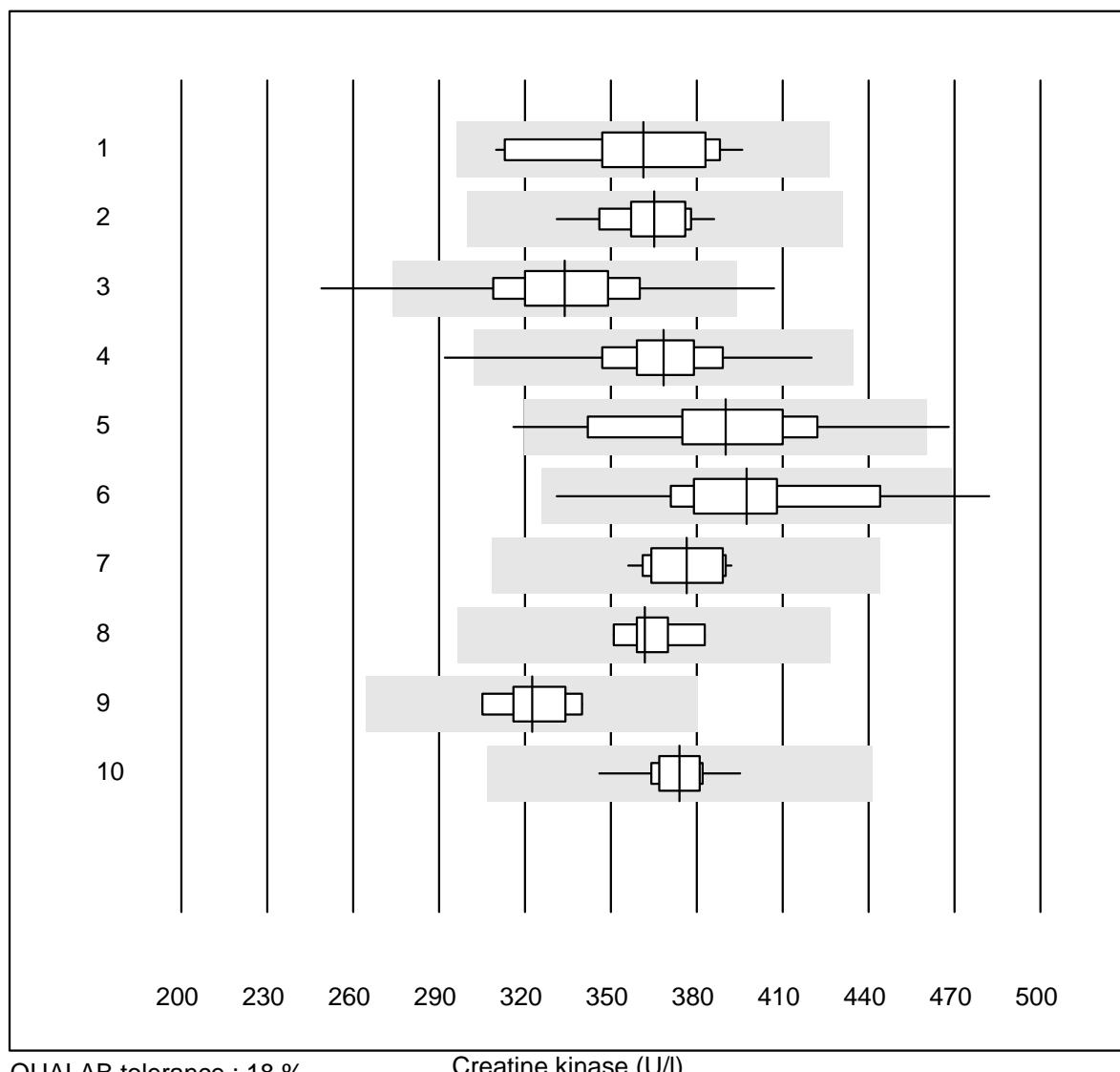
Cholesterol total

| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Standard chemistry | 22 | 100.0 | 0.0 | 0.0 | 3.99 | 3.6 | e |
| 2 | Cobas | 15 | 100.0 | 0.0 | 0.0 | 3.97 | 1.8 | e |
| 3 | Reflotron | 699 | 98.1 | 0.9 | 1.0 | 3.67 | 3.4 | e |
| 4 | Fuji Dri-Chem | 715 | 99.4 | 0.0 | 0.6 | 3.77 | 3.3 | e |
| 5 | Spotchem/Ready | 134 | 97.0 | 1.5 | 1.5 | 4.04 | 4.7 | e |
| 6 | Spotchem D-Concept | 156 | 97.5 | 1.9 | 0.6 | 3.98 | 3.6 | e |
| 7 | Piccolo | 19 | 100.0 | 0.0 | 0.0 | 4.03 | 3.0 | e |
| 8 | Cholestech LDX | 190 | 94.7 | 3.2 | 2.1 | 3.77 | 4.3 | e |
| 9 | Abx Mira | 8 | 100.0 | 0.0 | 0.0 | 3.91 | 2.6 | e |
| 10 | Hitachi S40/M40 | 16 | 100.0 | 0.0 | 0.0 | 3.98 | 2.4 | e |
| 11 | Autolyser/DiaSys | 13 | 100.0 | 0.0 | 0.0 | 3.87 | 4.5 | e* |

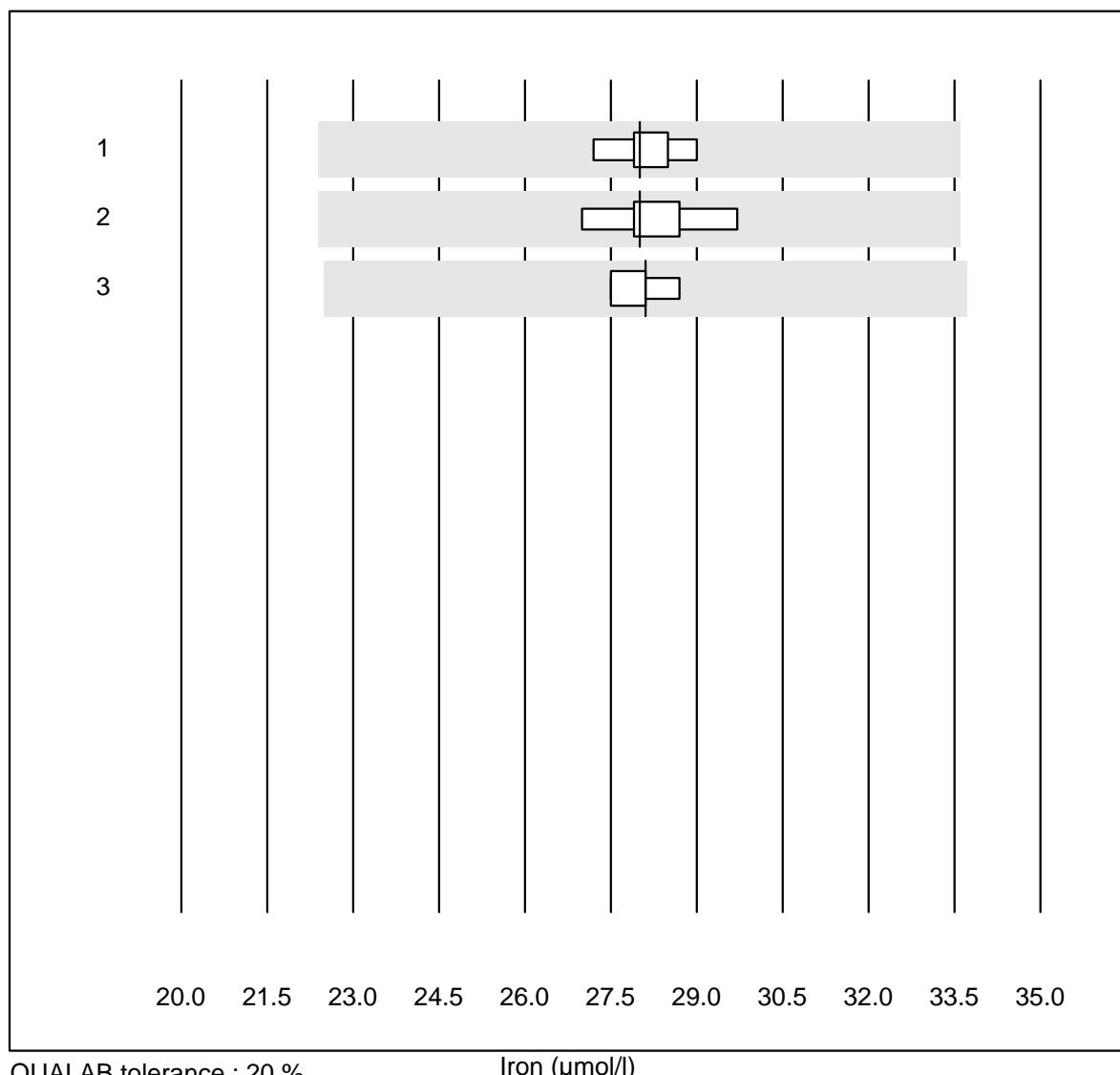
Cholesterin HDL



Creatine kinase



| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 IFCC | 19 | 100.0 | 0.0 | 0.0 | 361 | 7.1 | e |
| 2 Cobas | 14 | 100.0 | 0.0 | 0.0 | 365 | 4.1 | e |
| 3 Reflotron | 394 | 93.6 | 1.8 | 4.6 | 334 | 6.5 | e |
| 4 Fuji Dri-Chem | 451 | 99.4 | 0.2 | 0.4 | 368 | 4.5 | e |
| 5 Spotchem/Ready | 52 | 94.2 | 5.8 | 0.0 | 390 | 8.0 | e |
| 6 Spotchem D-Concept | 97 | 95.9 | 3.1 | 1.0 | 397 | 7.1 | e |
| 7 Piccolo | 11 | 100.0 | 0.0 | 0.0 | 376 | 3.3 | e |
| 8 Abx Mira | 6 | 100.0 | 0.0 | 0.0 | 362 | 3.0 | e |
| 9 Hitachi S40/M40 | 10 | 90.0 | 0.0 | 10.0 | 323 | 3.5 | e |
| 10 Autolyser/DiaSys | 12 | 100.0 | 0.0 | 0.0 | 374 | 3.2 | e |

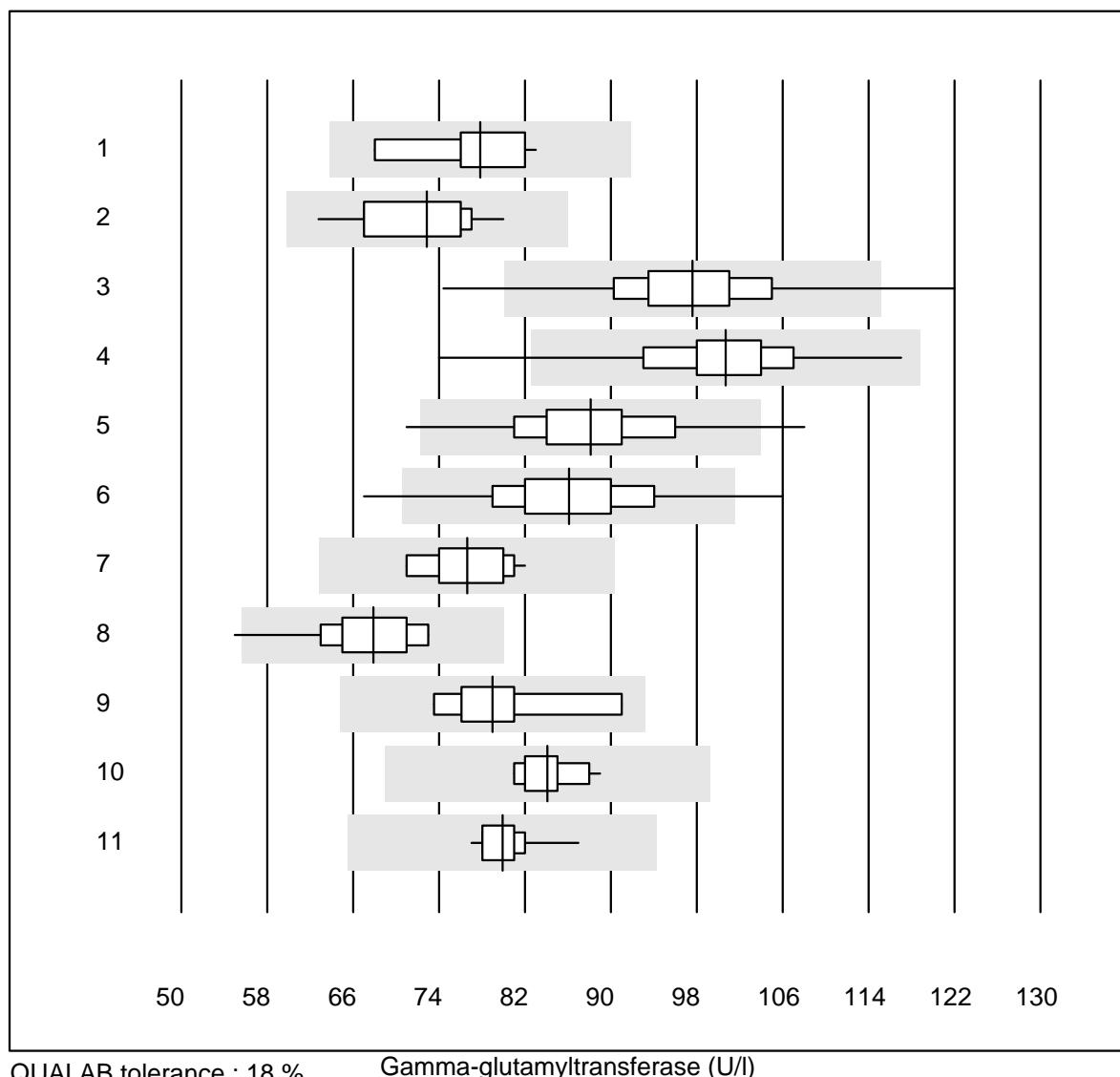
Iron

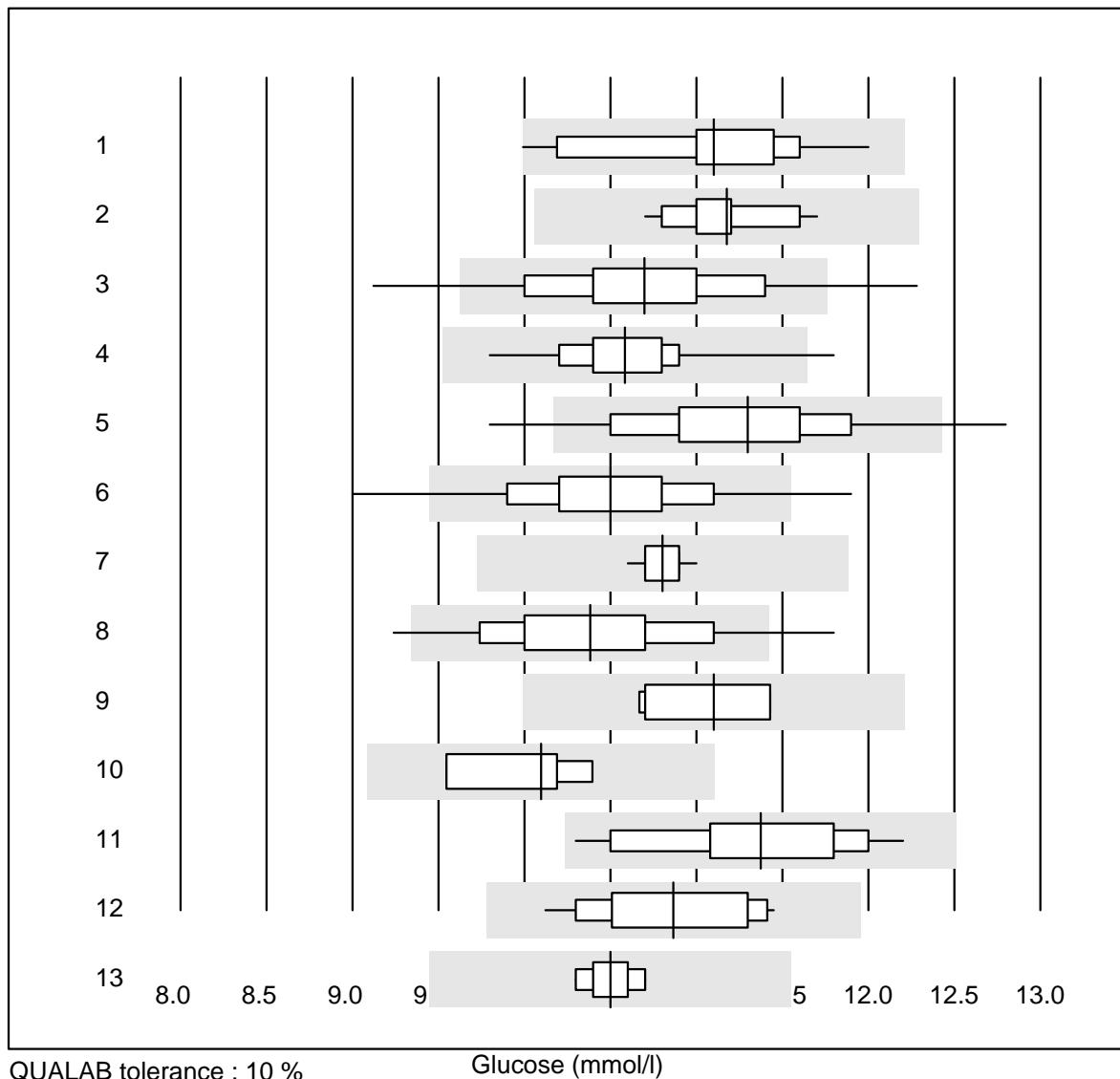
QUALAB tolerance : 20 %

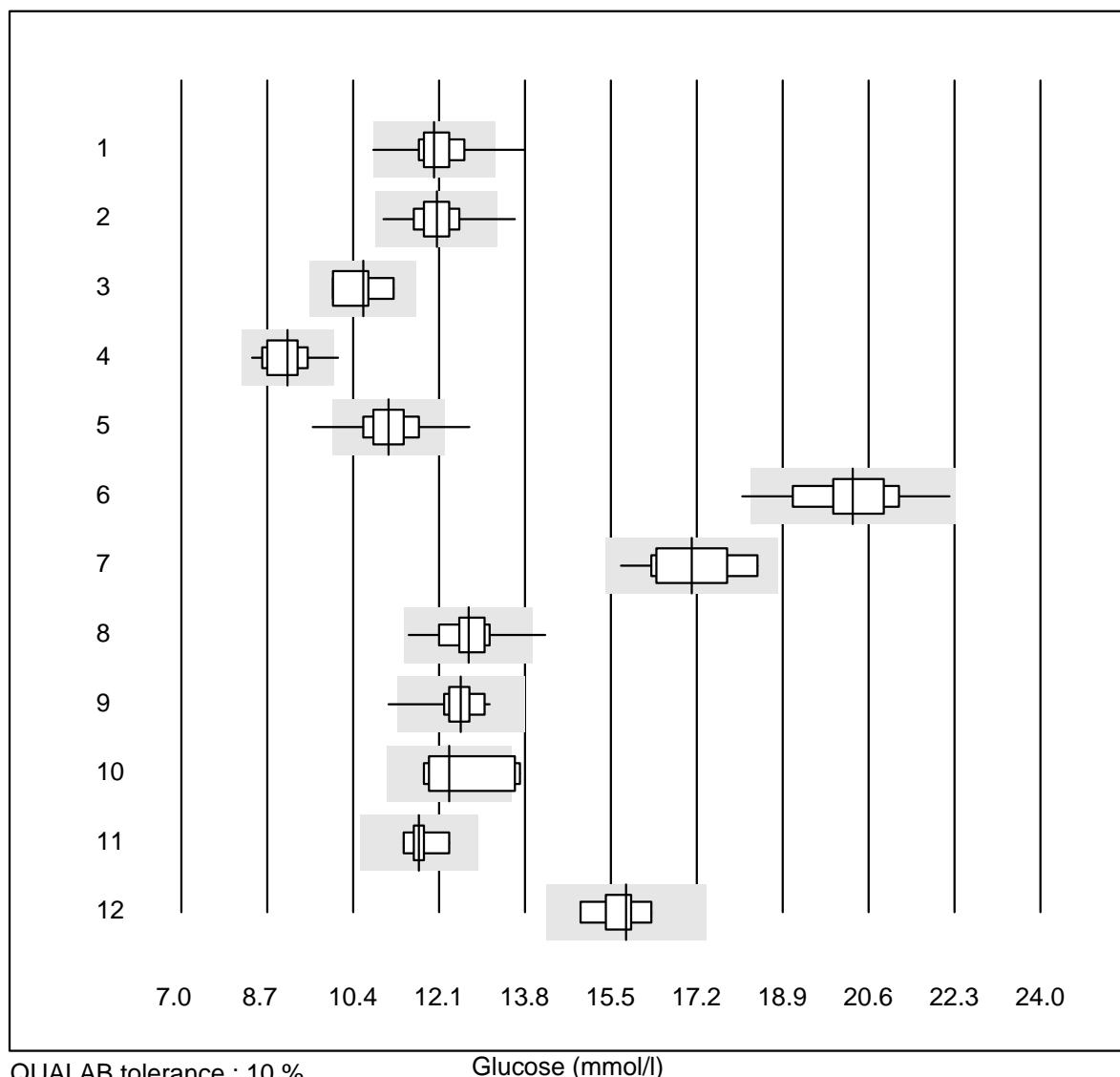
Iron (μmol/l)

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Standard chemistry | 7 | 100.0 | 0.0 | 0.0 | 28 | 2.0 | e |
| 2 Cobas | 9 | 100.0 | 0.0 | 0.0 | 28 | 2.8 | e |
| 3 Abx Mira | 5 | 80.0 | 0.0 | 20.0 | 28 | 1.8 | e |

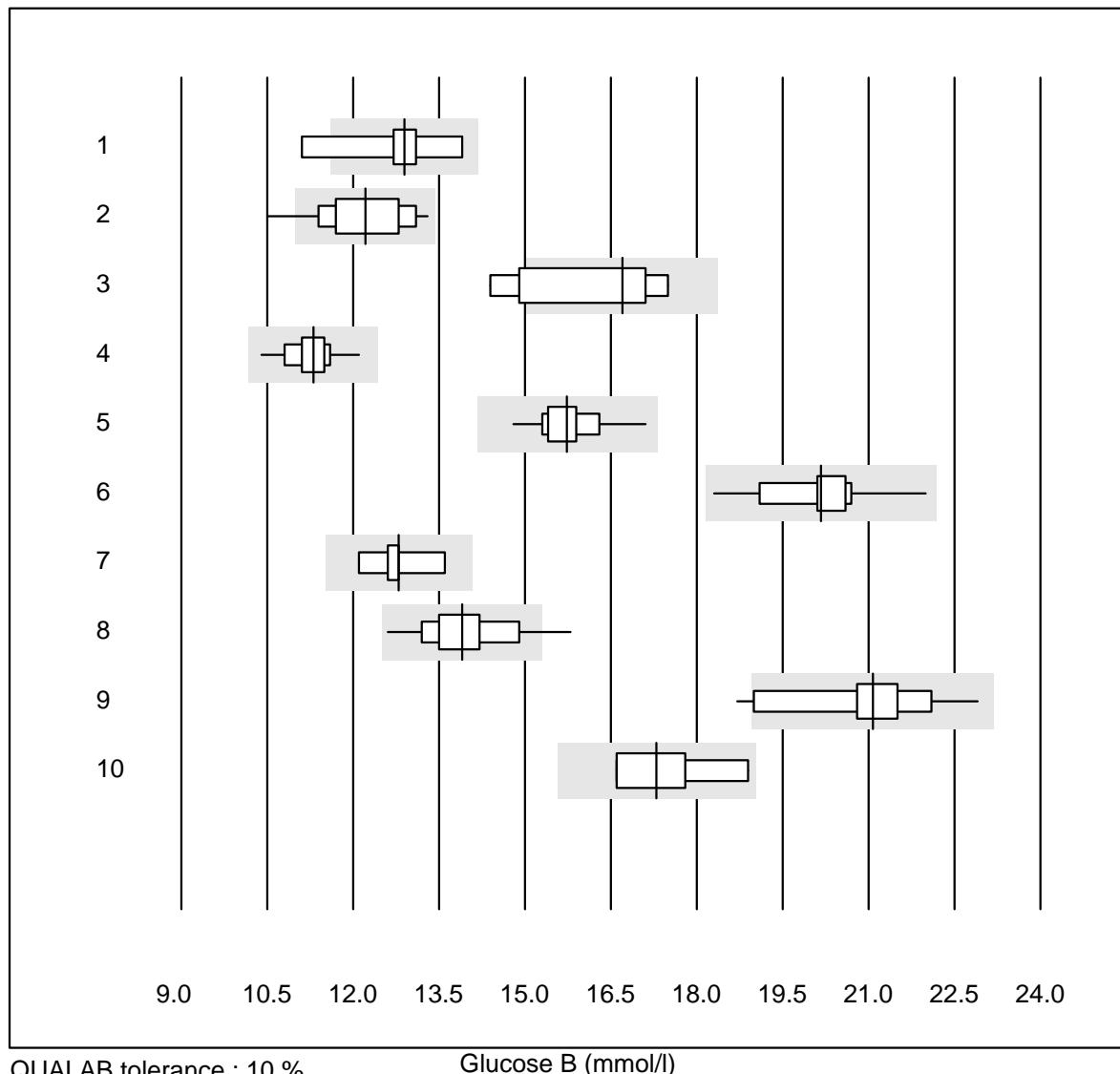
Gamma-glutamyltransferase

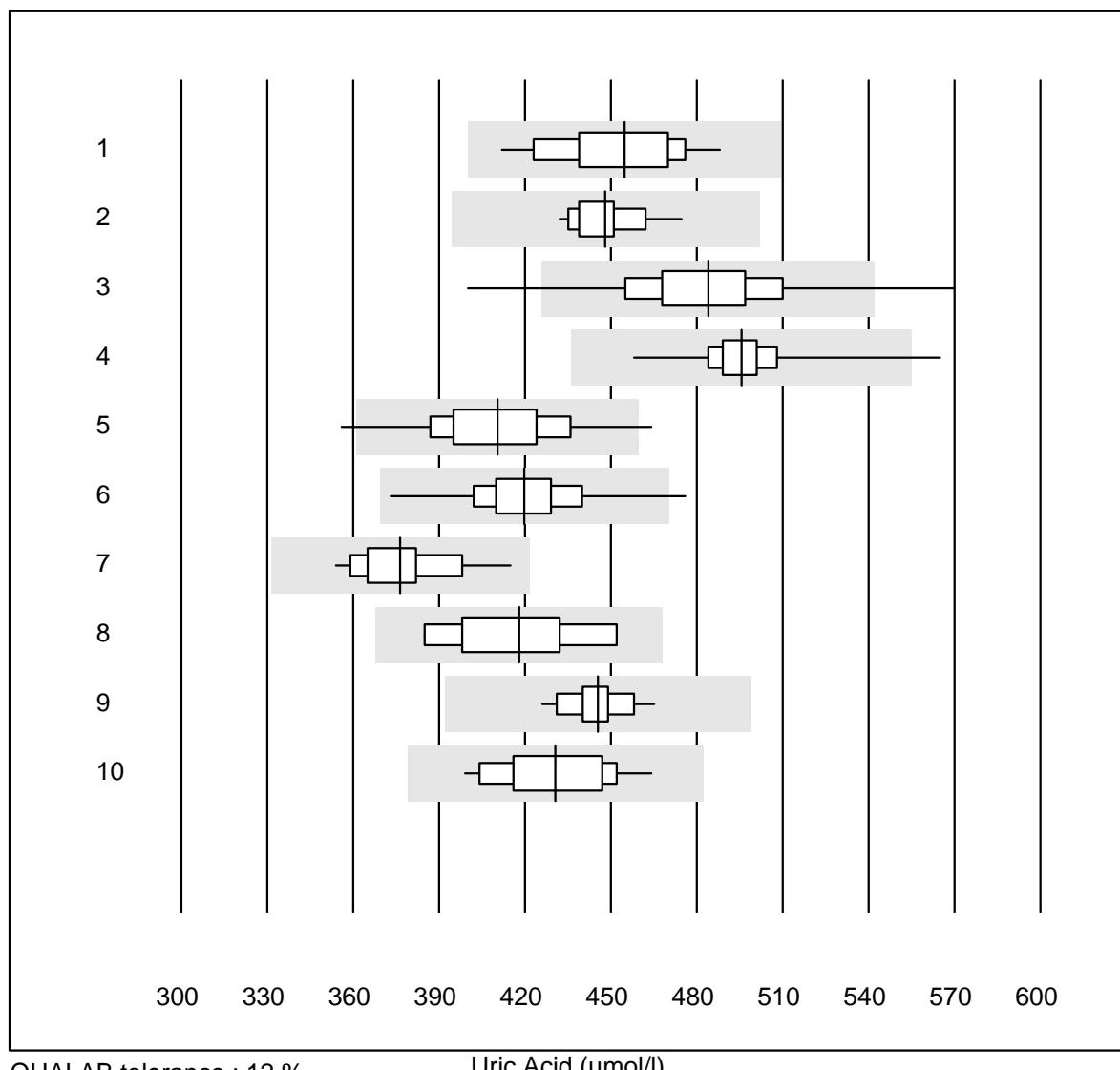


Glucose

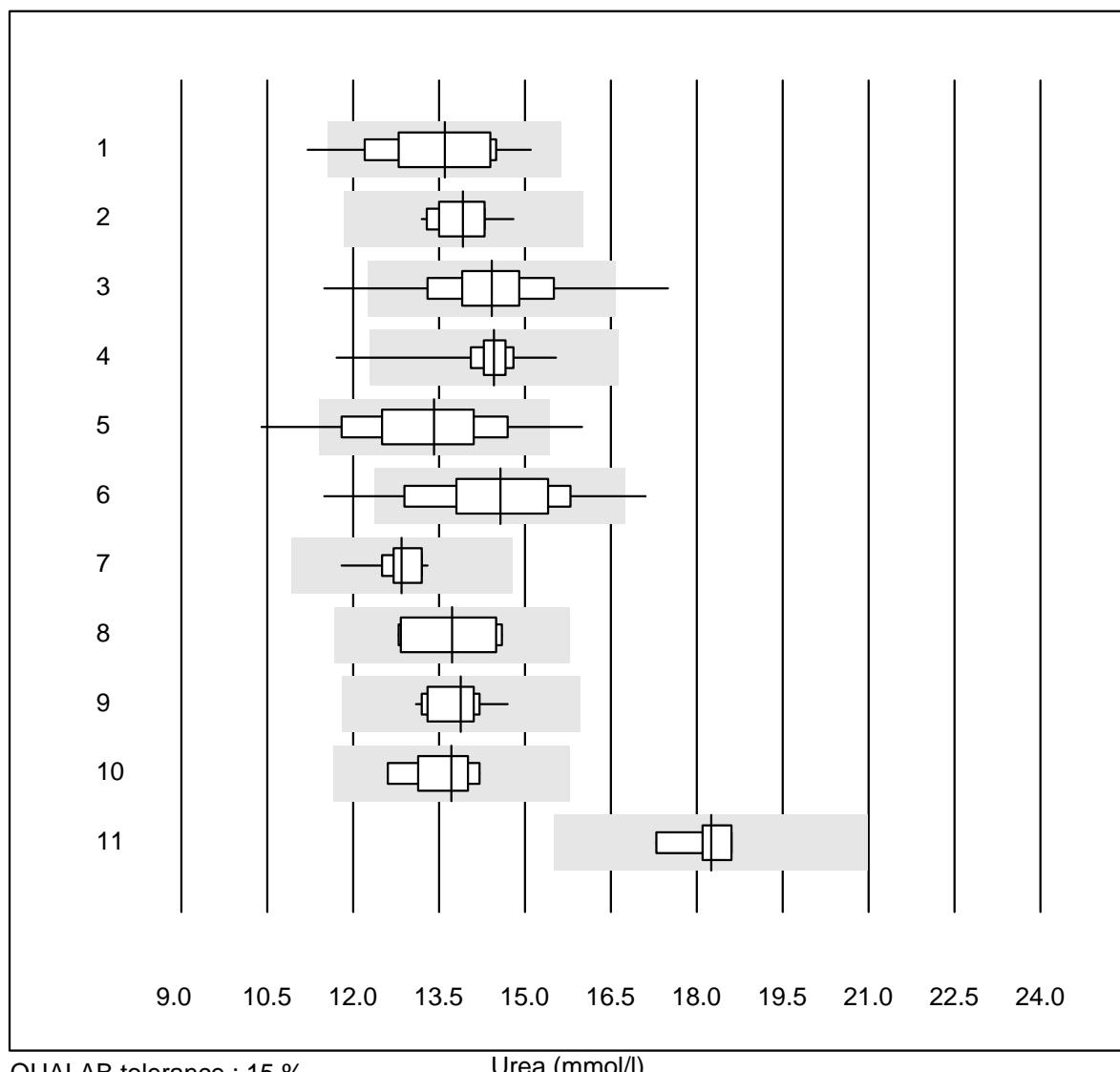
Glucose

Glucose B



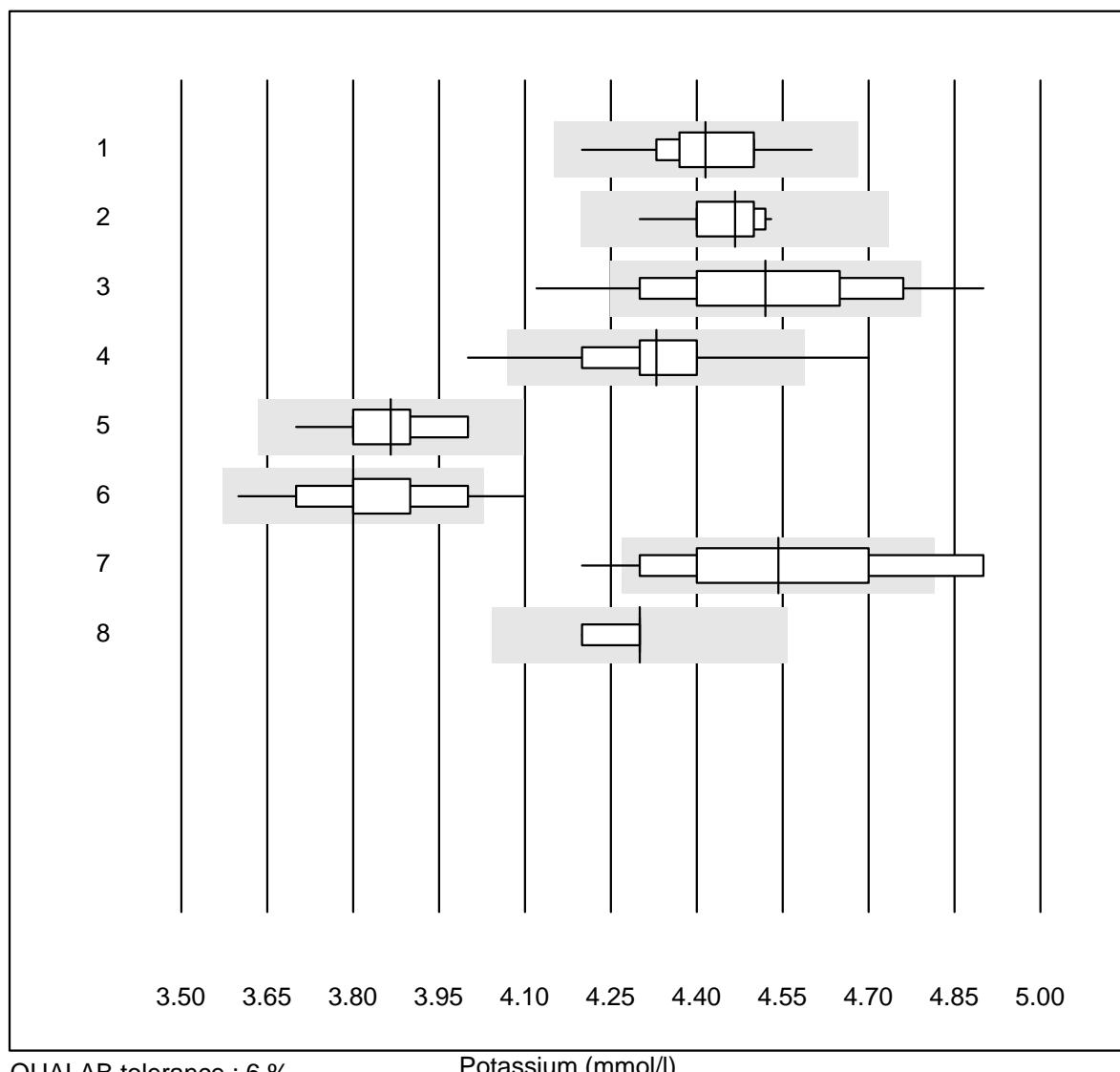
Uric Acid

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Standard chemistry | 25 | 100.0 | 0.0 | 0.0 | 455 | 4.5 | e |
| 2 Cobas | 11 | 100.0 | 0.0 | 0.0 | 448 | 2.7 | e |
| 3 Reflotron | 736 | 98.5 | 1.1 | 0.4 | 484 | 4.7 | e |
| 4 Fuji Dri-Chem | 729 | 99.6 | 0.1 | 0.3 | 496 | 2.0 | e |
| 5 Spotchem/Ready | 117 | 96.5 | 2.6 | 0.9 | 410 | 5.0 | e |
| 6 Spotchem D-Concept | 156 | 98.7 | 1.3 | 0.0 | 420 | 3.9 | e |
| 7 Piccolo | 23 | 100.0 | 0.0 | 0.0 | 376 | 4.2 | e |
| 8 Abx Mira | 8 | 100.0 | 0.0 | 0.0 | 418 | 5.6 | e* |
| 9 Hitachi S40/M40 | 17 | 100.0 | 0.0 | 0.0 | 445 | 2.2 | e |
| 10 Autolyser/DiaSys | 13 | 100.0 | 0.0 | 0.0 | 431 | 4.7 | e |

Urea

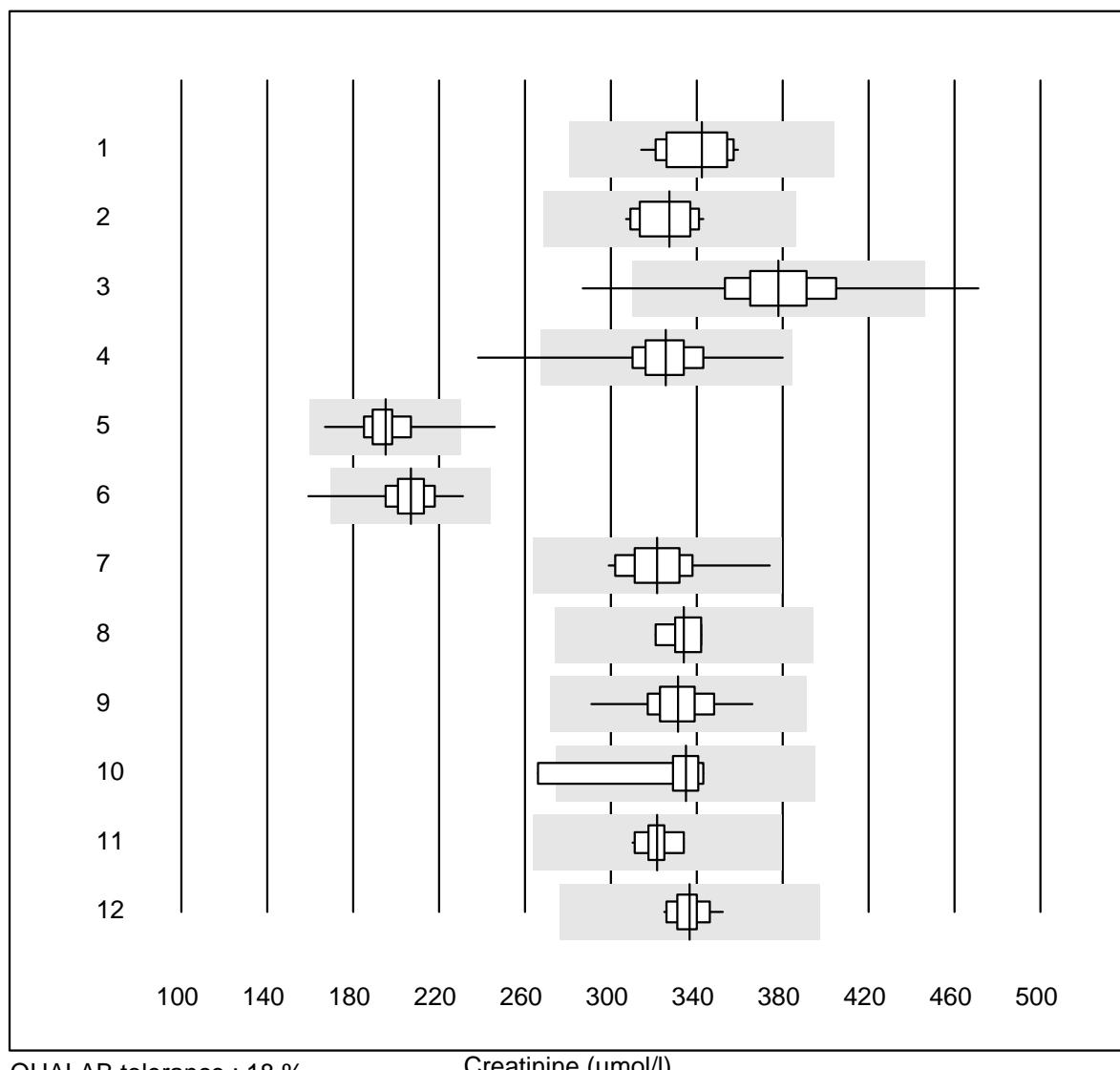
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Standard chemistry | 24 | 95.8 | 4.2 | 0.0 | 13.6 | 7.6 | e |
| 2 Cobas | 14 | 100.0 | 0.0 | 0.0 | 13.9 | 3.2 | e |
| 3 Reflotron | 329 | 96.4 | 3.0 | 0.6 | 14.4 | 6.0 | e |
| 4 Fuji Dri-Chem | 446 | 99.6 | 0.2 | 0.2 | 14.5 | 2.3 | e |
| 5 Spotchem/Ready | 80 | 93.7 | 6.3 | 0.0 | 13.4 | 8.5 | e |
| 6 Spotchem D-Concept | 95 | 90.5 | 6.3 | 3.2 | 14.6 | 8.2 | e |
| 7 Piccolo | 34 | 100.0 | 0.0 | 0.0 | 12.8 | 3.0 | e |
| 8 Abx Mira | 6 | 100.0 | 0.0 | 0.0 | 13.7 | 5.7 | e* |
| 9 Hitachi S40/M40 | 13 | 92.3 | 0.0 | 7.7 | 13.9 | 3.3 | e |
| 10 Autolyser/DiaSys | 7 | 100.0 | 0.0 | 0.0 | 13.7 | 4.0 | e |
| 11 iStat Chem8 | 6 | 100.0 | 0.0 | 0.0 | 18.3 | 2.7 | e |

Potassium

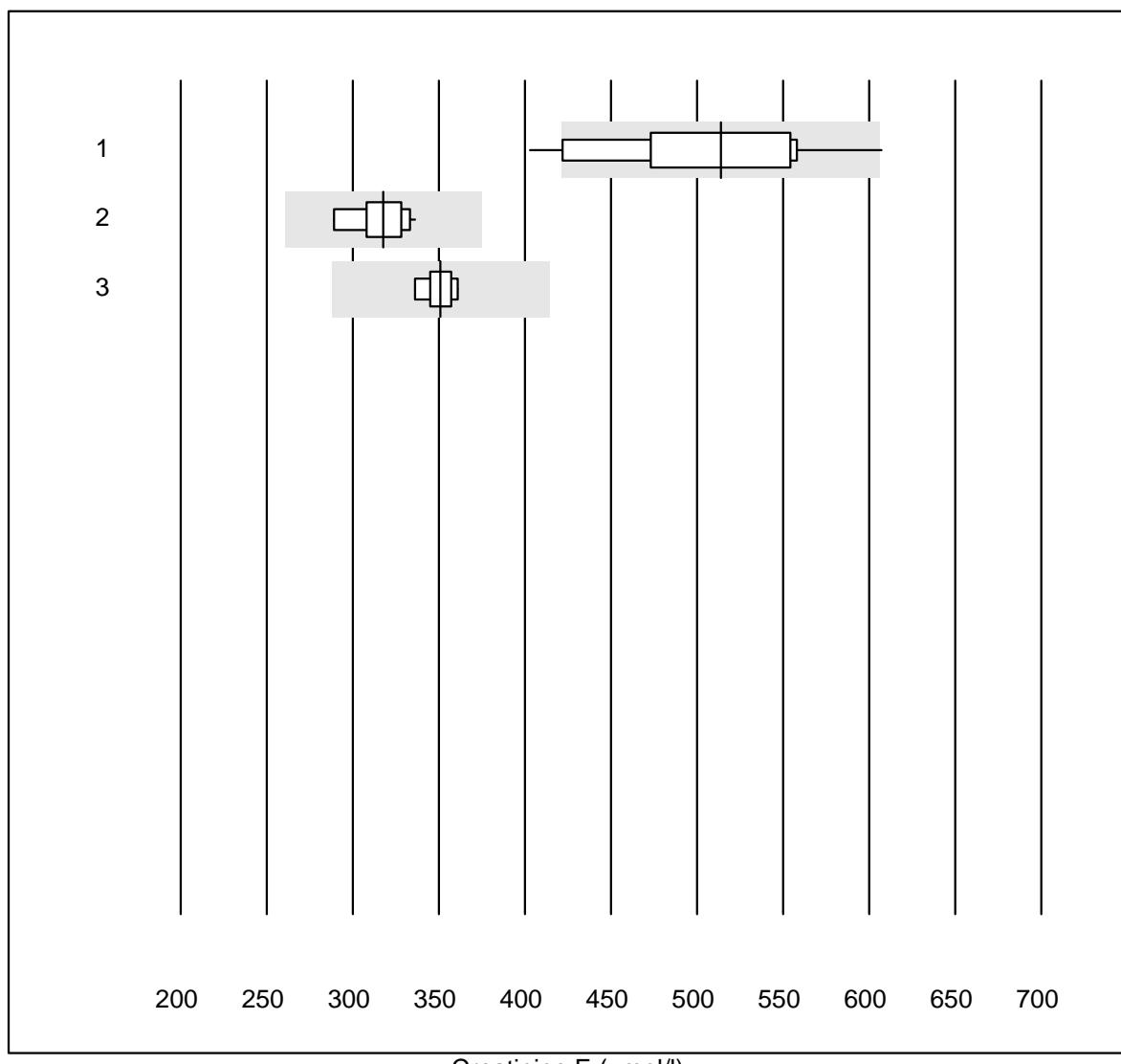


| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 ISE | 33 | 97.0 | 0.0 | 3.0 | 4.42 | 1.9 | e |
| 2 Cobas | 16 | 100.0 | 0.0 | 0.0 | 4.47 | 1.4 | e |
| 3 Reflotron | 752 | 85.5 | 9.6 | 4.9 | 4.52 | 3.7 | e |
| 4 Fuji Dri-Chem | 764 | 97.7 | 1.3 | 1.0 | 4.33 | 2.0 | e |
| 5 Spotchem D-Concept | 158 | 97.5 | 0.0 | 2.5 | 3.87 | 2.0 | e |
| 6 Spotchem EL-SE 1520 | 116 | 97.4 | 0.9 | 1.7 | 3.80 | 2.8 | e |
| 7 Piccolo | 22 | 72.8 | 13.6 | 13.6 | 4.54 | 4.4 | e* |
| 8 iStat Chem8 | 6 | 100.0 | 0.0 | 0.0 | 4.30 | 1.0 | e |

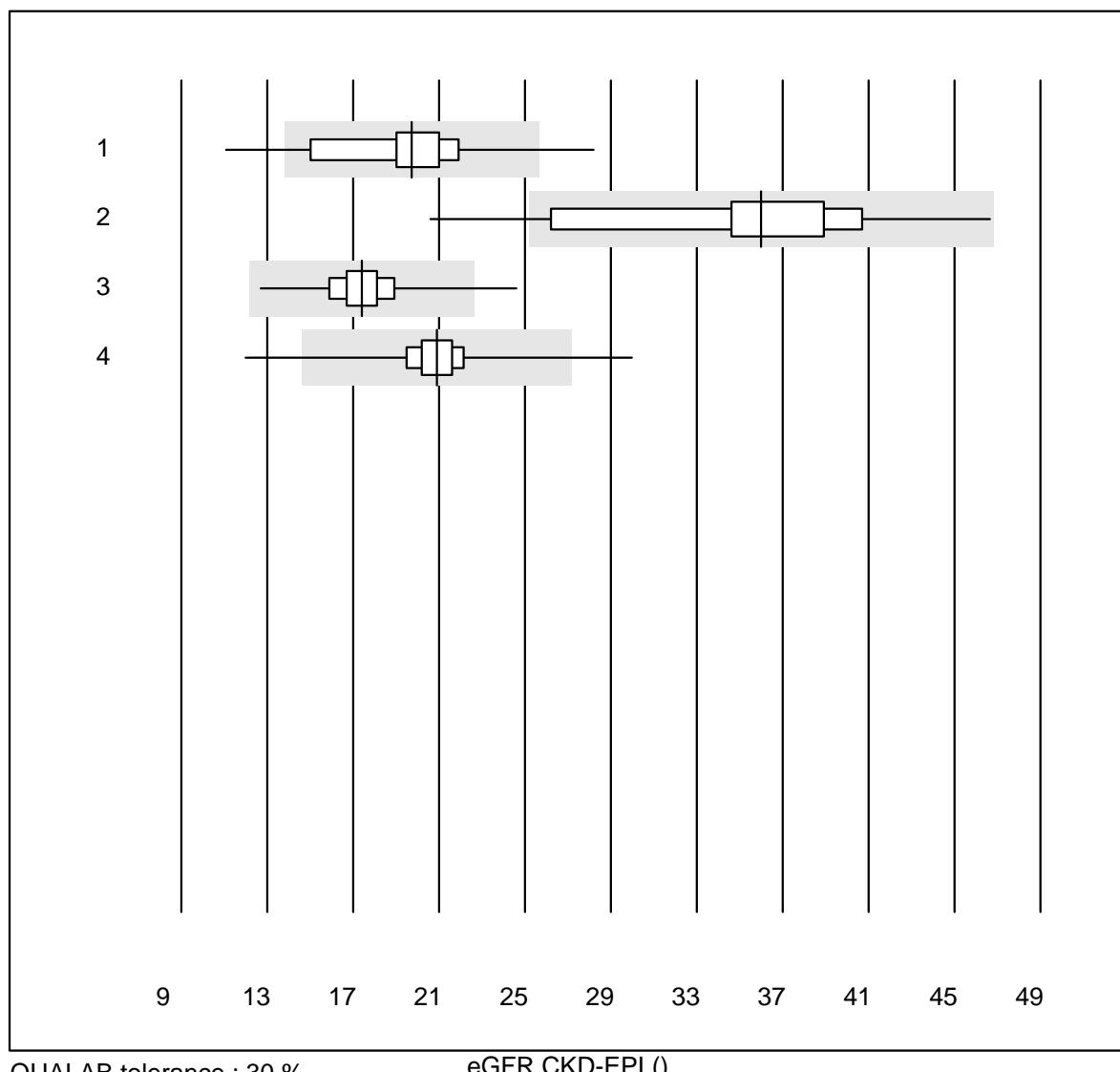
Creatinine



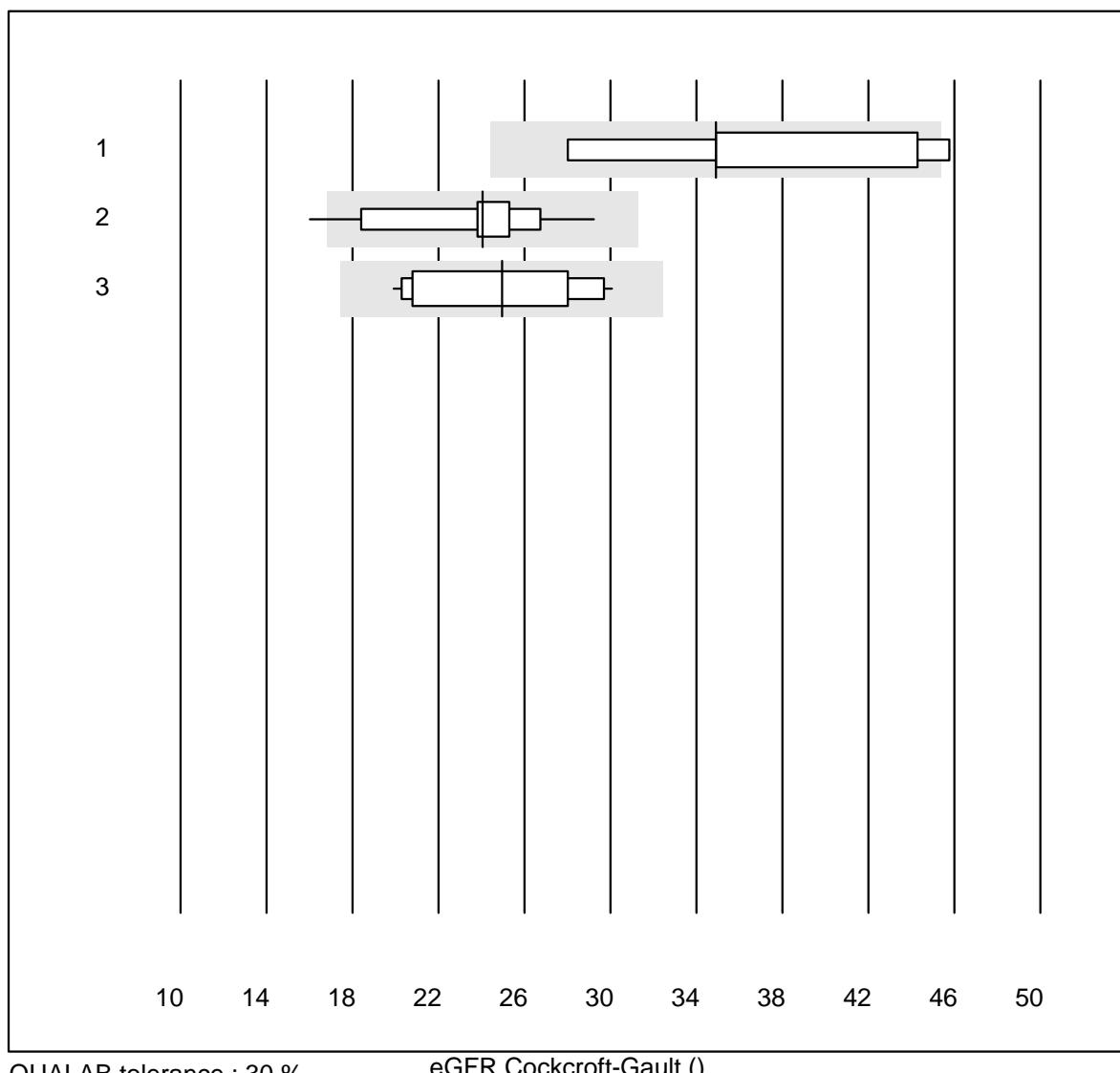
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Standard chemistry | 14 | 92.9 | 0.0 | 7.1 | 343 | 4.5 | e |
| 2 Cobas | 16 | 100.0 | 0.0 | 0.0 | 327 | 3.7 | e |
| 3 Reflotron | 937 | 98.3 | 1.0 | 0.7 | 378 | 5.7 | e |
| 4 Fuji Dri-Chem | 794 | 98.8 | 0.3 | 0.9 | 326 | 4.1 | e |
| 5 Spotchem/Ready | 135 | 81.5 | 1.5 | 17.0 | 195 | 6.1 | e |
| 6 Spotchem D-Concept | 157 | 96.9 | 0.6 | 2.5 | 207 | 5.2 | e |
| 7 Spotchem old test | 19 | 94.7 | 0.0 | 5.3 | 322 | 5.3 | e |
| 8 Enzymatic | 7 | 100.0 | 0.0 | 0.0 | 334 | 2.3 | e |
| 9 Piccolo | 34 | 100.0 | 0.0 | 0.0 | 331 | 4.3 | e |
| 10 Abx Mira | 10 | 80.0 | 10.0 | 10.0 | 335 | 7.9 | e* |
| 11 Hitachi S40/M40 | 18 | 100.0 | 0.0 | 0.0 | 321 | 2.1 | e |
| 12 Autolyser/DiaSys | 14 | 100.0 | 0.0 | 0.0 | 337 | 2.2 | e |

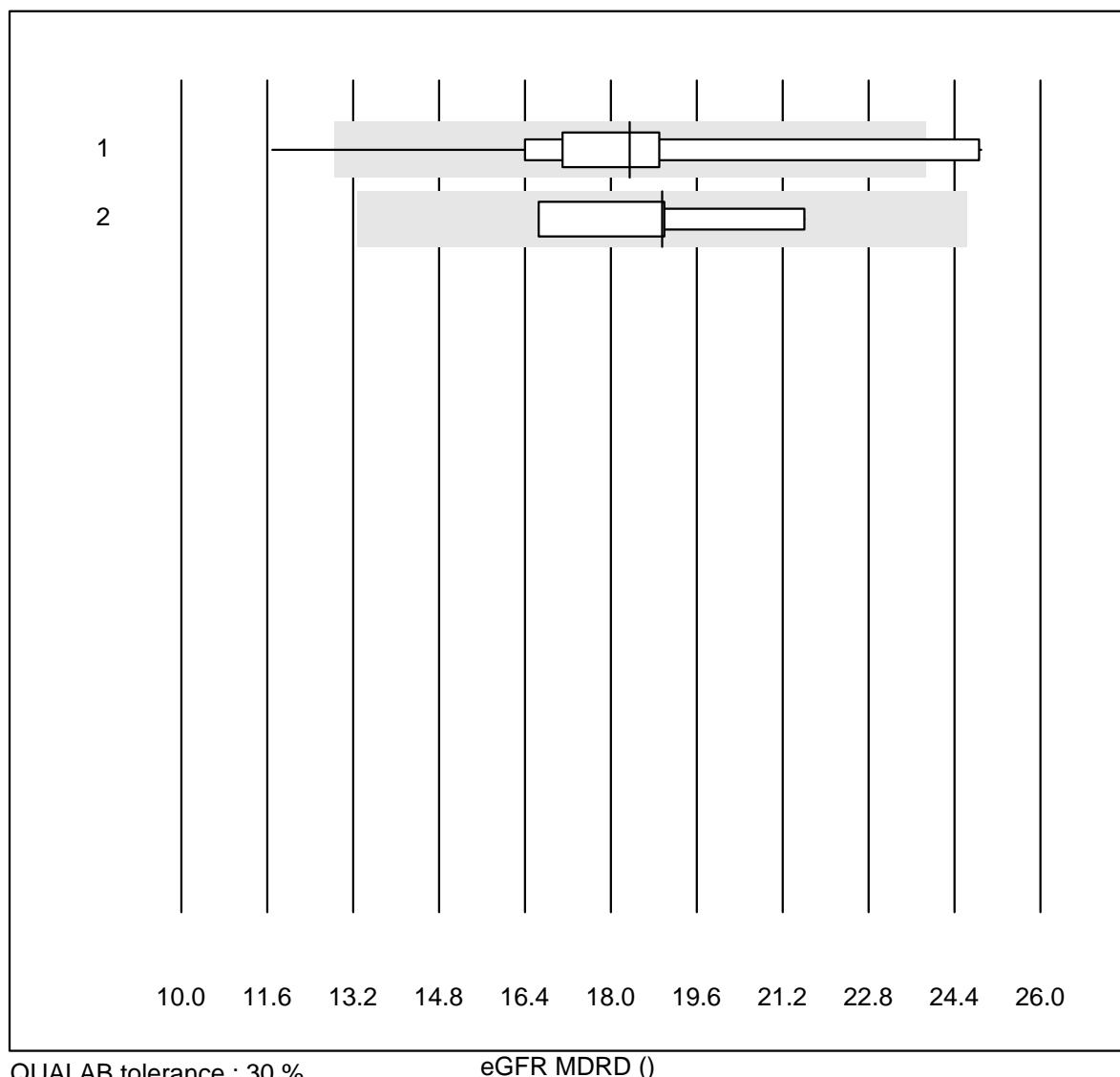
Creatinine E

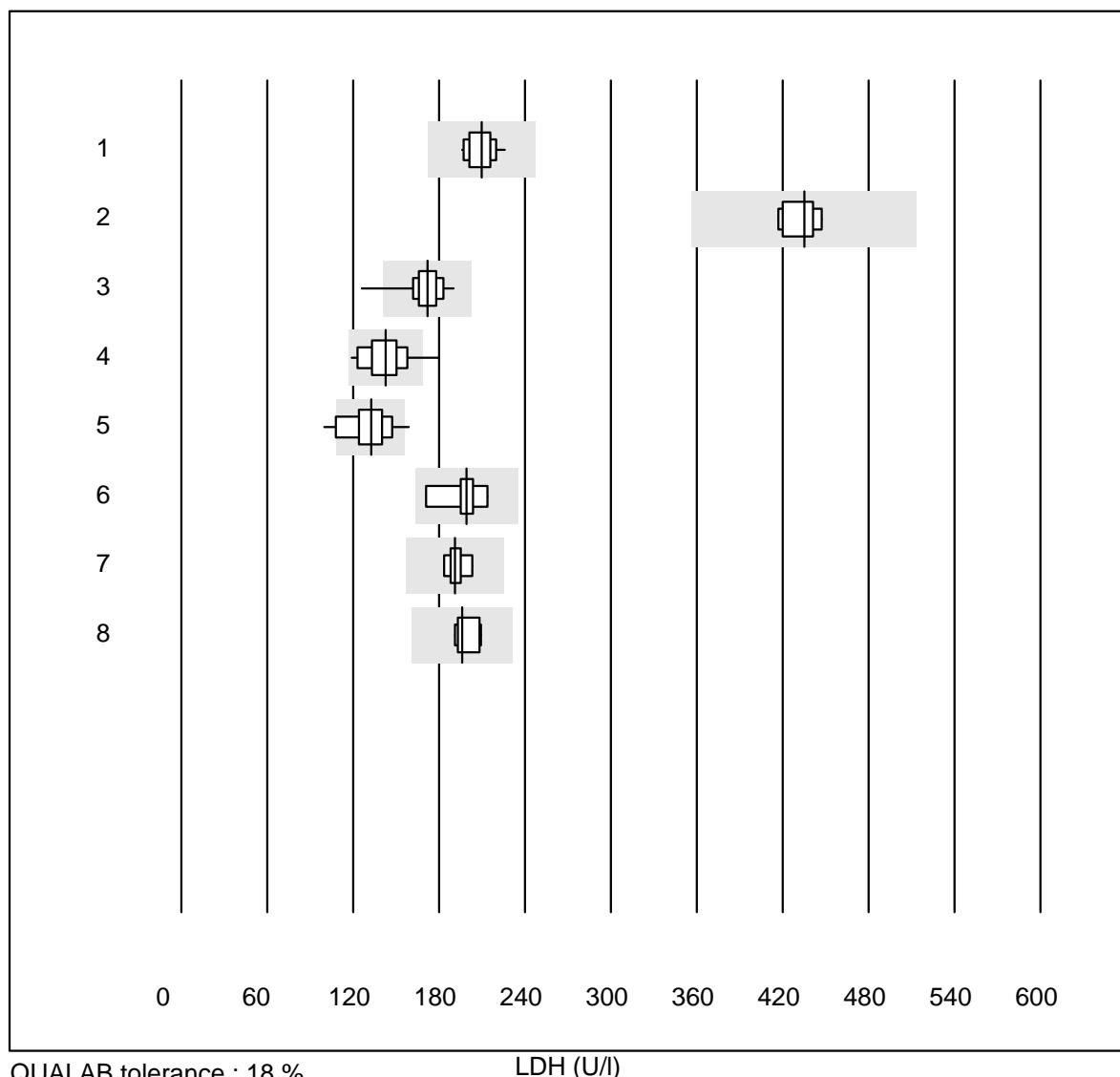
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 Statsensor i / Nova | 35 | 85.7 | 11.4 | 2.9 | 514 | 10.5 | e |
| 2 iStat Chem8 | 10 | 100.0 | 0.0 | 0.0 | 318 | 4.5 | e |
| 3 ABL700/800 | 9 | 100.0 | 0.0 | 0.0 | 351 | 2.5 | e |

eGFR CKD-EPI

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 Standard chemistry | 56 | 85.8 | 7.1 | 7.1 | 20 | 14.4 | e |
| 2 Spotchem/Ready | 111 | 86.5 | 8.1 | 5.4 | 36 | 14.6 | e |
| 3 Reflotron | 339 | 95.3 | 0.9 | 3.8 | 17 | 8.0 | e |
| 4 Fuji Dri-Chem | 327 | 94.1 | 2.8 | 3.1 | 21 | 8.7 | e |

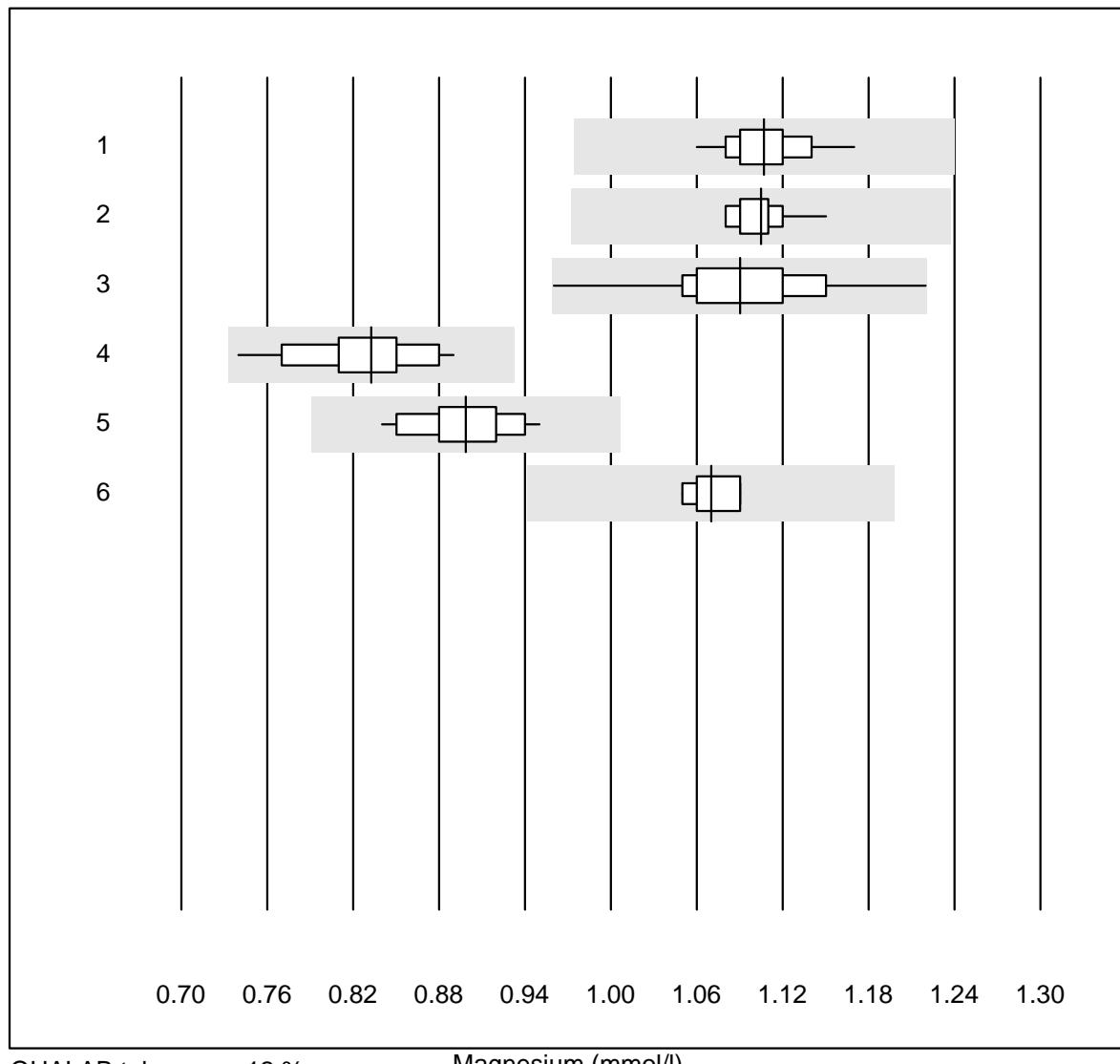
eGFR Cockcroft-Gault

eGFR MDRD

LDH

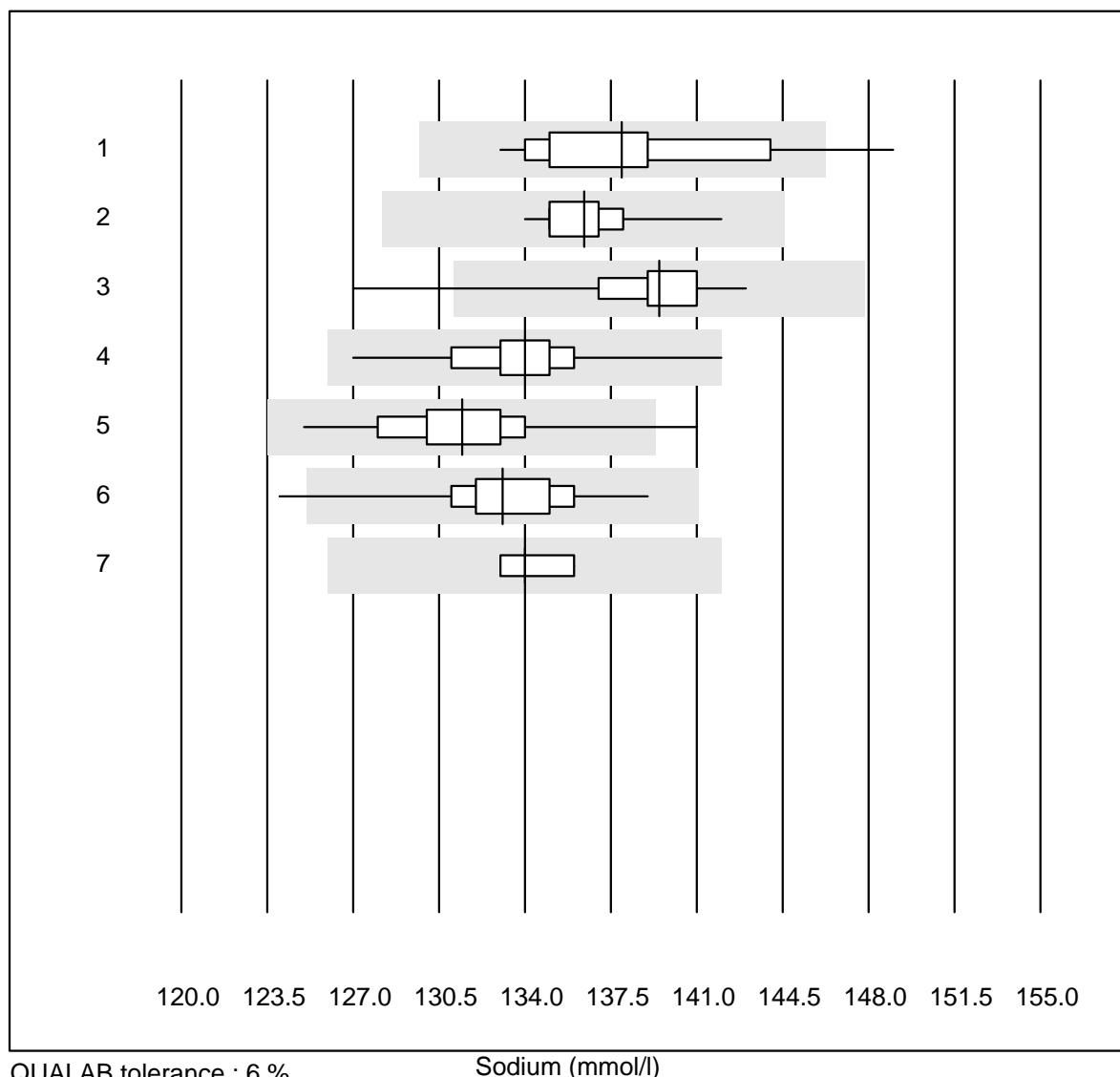
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 IFCC | 20 | 100.0 | 0.0 | 0.0 | 210 | 4.1 | e |
| 2 Cobas | 9 | 100.0 | 0.0 | 0.0 | 435 | 2.5 | e |
| 3 Fuji Dri-Chem | 144 | 97.9 | 0.7 | 1.4 | 172 | 5.1 | e |
| 4 Spotchem/Ready | 31 | 96.8 | 3.2 | 0.0 | 143 | 9.8 | e |
| 5 Spotchem D-Concept | 45 | 77.8 | 11.1 | 11.1 | 132 | 10.8 | e |
| 6 Abx Mira | 6 | 100.0 | 0.0 | 0.0 | 199 | 7.3 | e* |
| 7 Hitachi S40/M40 | 6 | 83.3 | 0.0 | 16.7 | 191 | 3.8 | e |
| 8 Autolyser/DiaSys | 7 | 100.0 | 0.0 | 0.0 | 196 | 3.8 | e |

Magnesium



| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Standard chemistry | 17 | 100.0 | 0.0 | 0.0 | 1.11 | 2.2 | e |
| 2 Cobas | 10 | 100.0 | 0.0 | 0.0 | 1.11 | 1.8 | e |
| 3 Fuji Dri-Chem | 119 | 98.3 | 0.0 | 1.7 | 1.09 | 3.8 | e |
| 4 Spotchem D-Concept | 33 | 100.0 | 0.0 | 0.0 | 0.83 | 4.4 | e |
| 5 Spotchem/Ready | 15 | 100.0 | 0.0 | 0.0 | 0.90 | 3.5 | e |
| 6 Piccolo | 5 | 100.0 | 0.0 | 0.0 | 1.07 | 1.7 | e |

Sodium

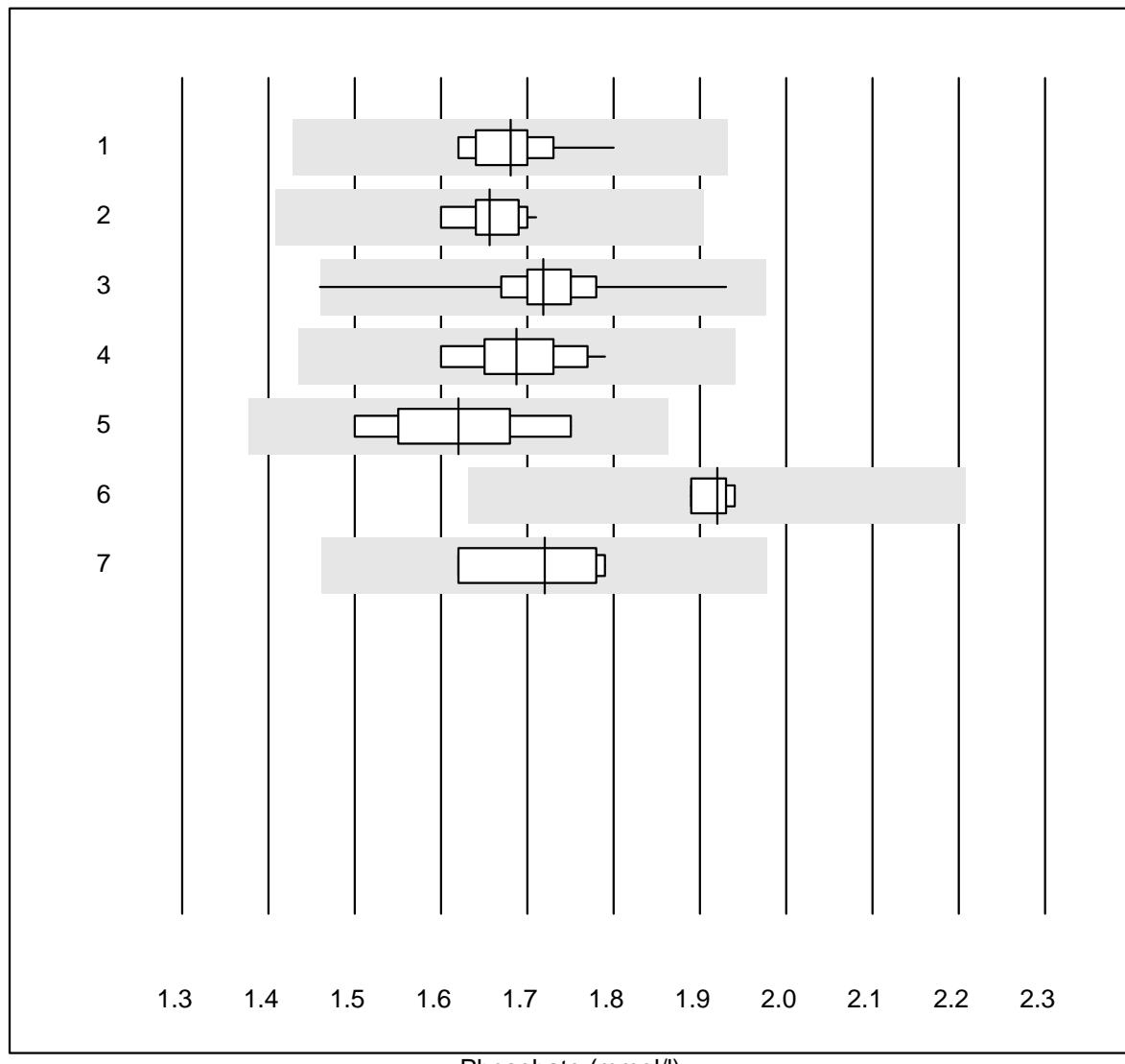


QUALAB tolerance : 6 %

Sodium (mmol/l)

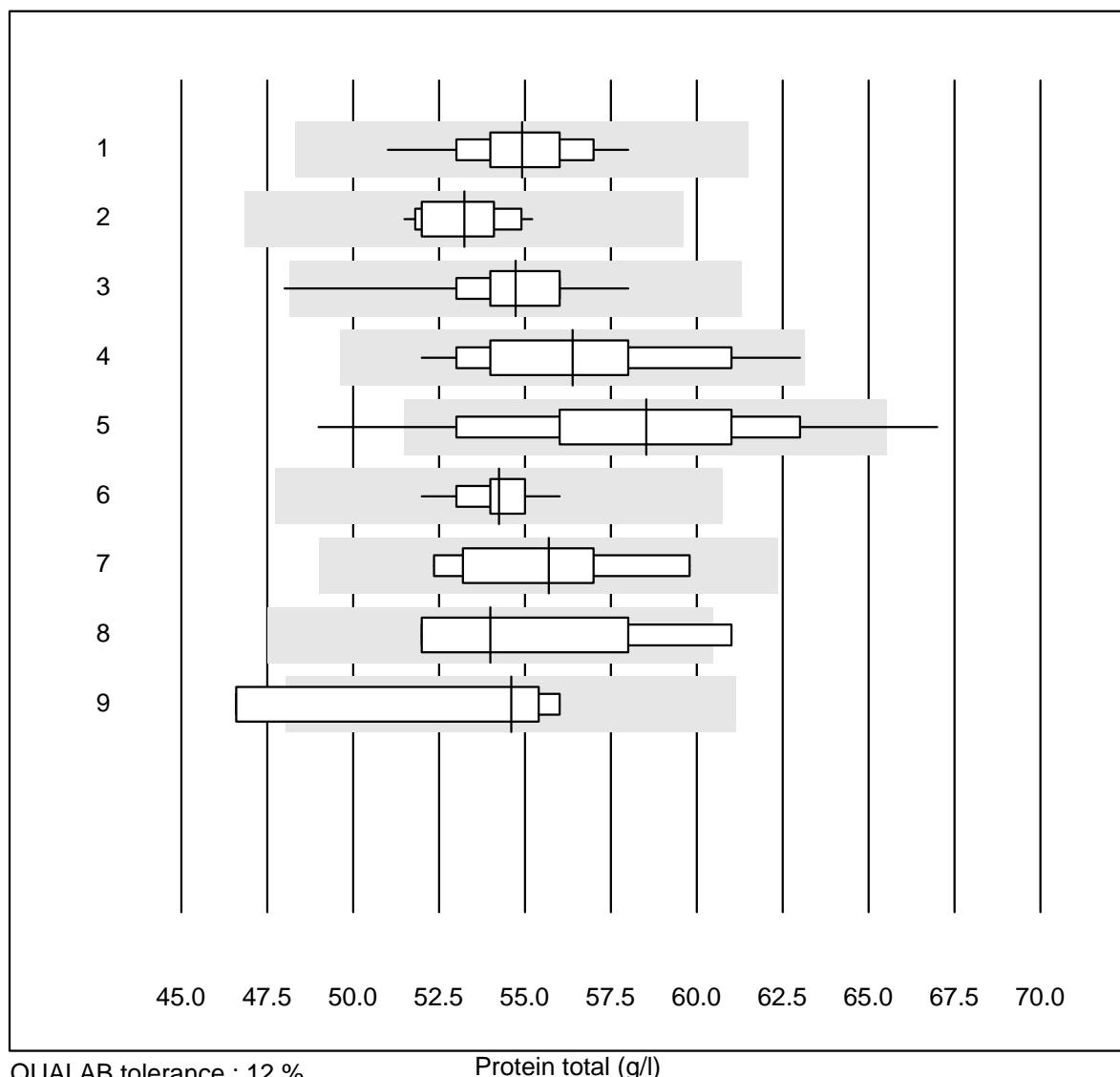
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 ISE | 32 | 90.6 | 9.4 | 0.0 | 138 | 3.0 | e |
| 2 Cobas | 15 | 100.0 | 0.0 | 0.0 | 136 | 1.4 | e |
| 3 Fuji Dri-Chem | 709 | 98.9 | 0.7 | 0.4 | 139 | 1.5 | e |
| 4 Spotchem D-Concept | 150 | 100.0 | 0.0 | 0.0 | 134 | 1.8 | e |
| 5 Spotchem EL-SE 1520 | 116 | 97.4 | 0.9 | 1.7 | 131 | 1.9 | e |
| 6 Piccolo | 23 | 95.7 | 4.3 | 0.0 | 133 | 2.3 | e |
| 7 iStat Chem8 | 5 | 100.0 | 0.0 | 0.0 | 134 | 0.8 | e |

Phosphate



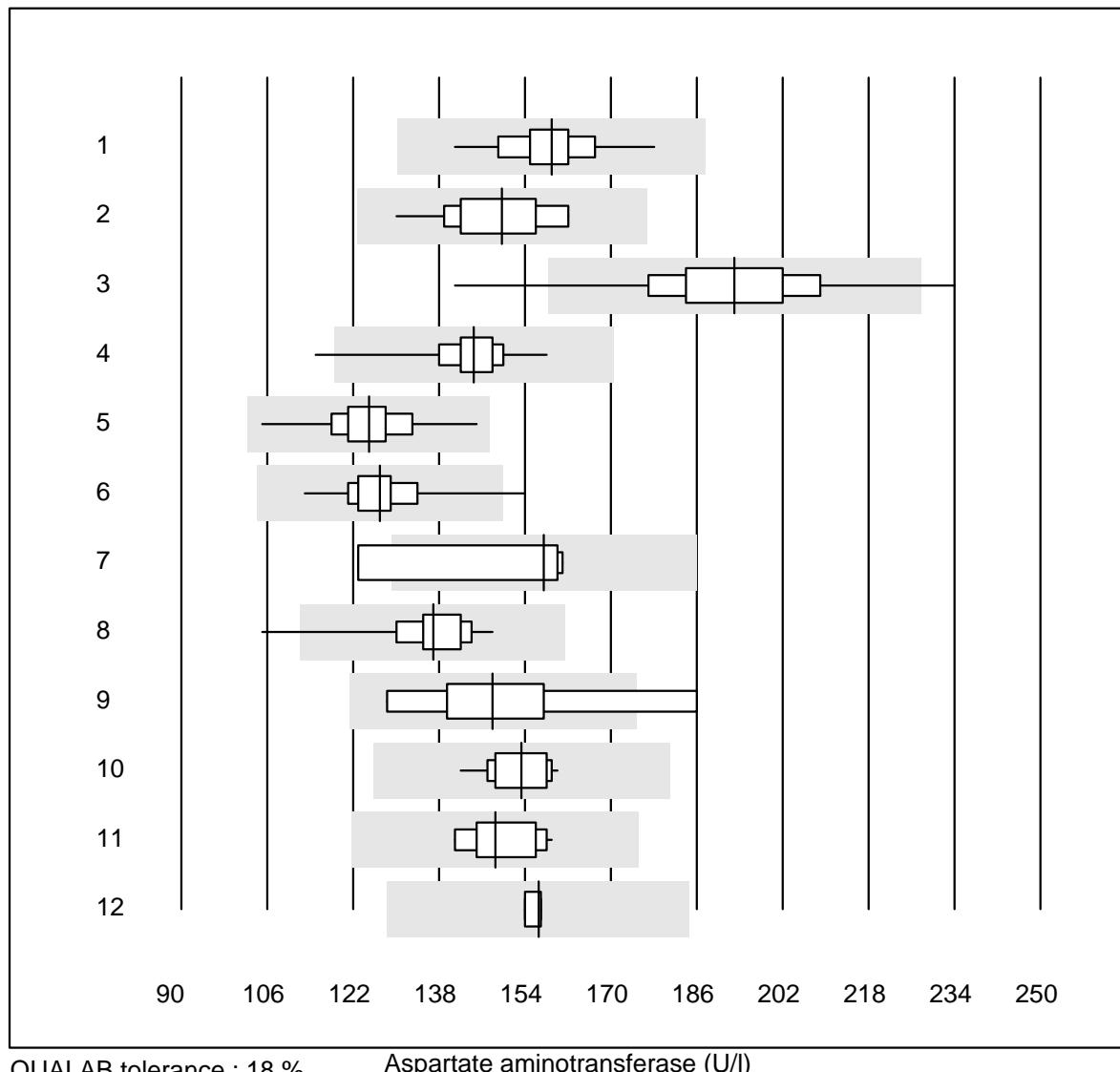
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Standard chemistry | 16 | 100.0 | 0.0 | 0.0 | 1.7 | 2.8 | e |
| 2 | Cobas | 10 | 100.0 | 0.0 | 0.0 | 1.7 | 2.2 | e |
| 3 | Fuji Dri-Chem | 81 | 98.8 | 1.2 | 0.0 | 1.7 | 3.7 | e |
| 4 | Spotchem D-Concept | 17 | 100.0 | 0.0 | 0.0 | 1.7 | 3.5 | e |
| 5 | Spotchem/Ready | 9 | 100.0 | 0.0 | 0.0 | 1.6 | 5.4 | e |
| 6 | Piccolo | 4 | 100.0 | 0.0 | 0.0 | 1.9 | 1.2 | e |
| 7 | Abx Mira | 4 | 100.0 | 0.0 | 0.0 | 1.7 | 5.0 | e* |

Protein total



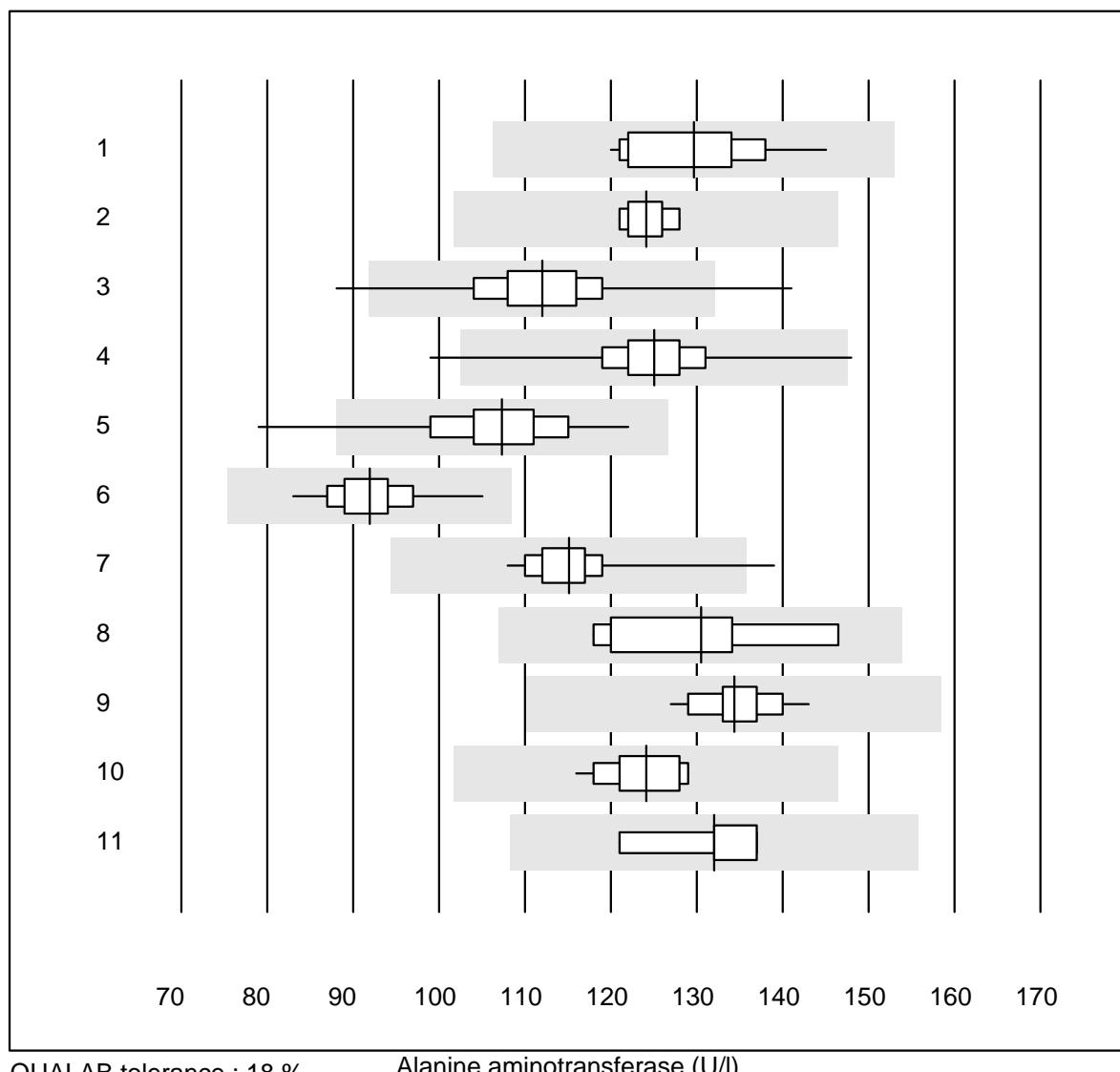
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Standard chemistry | 24 | 100.0 | 0.0 | 0.0 | 54.9 | 2.9 | e |
| 2 | Cobas | 11 | 100.0 | 0.0 | 0.0 | 53.2 | 2.4 | e |
| 3 | Fuji Dri-Chem | 184 | 99.5 | 0.5 | 0.0 | 54.7 | 2.3 | e |
| 4 | Spotchem/Ready | 38 | 97.4 | 0.0 | 2.6 | 56.4 | 5.7 | e |
| 5 | Spotchem D-Concept | 69 | 91.4 | 7.2 | 1.4 | 58.5 | 6.7 | e |
| 6 | Piccolo | 25 | 100.0 | 0.0 | 0.0 | 54.2 | 1.7 | e |
| 7 | Abx Mira | 5 | 100.0 | 0.0 | 0.0 | 55.7 | 5.4 | e* |
| 8 | Hitachi S40/M40 | 7 | 85.7 | 14.3 | 0.0 | 54.0 | 6.0 | e* |
| 9 | Autolyser/DiaSys | 4 | 75.0 | 25.0 | 0.0 | 54.6 | 8.2 | e* |

Aspartate aminotransferase

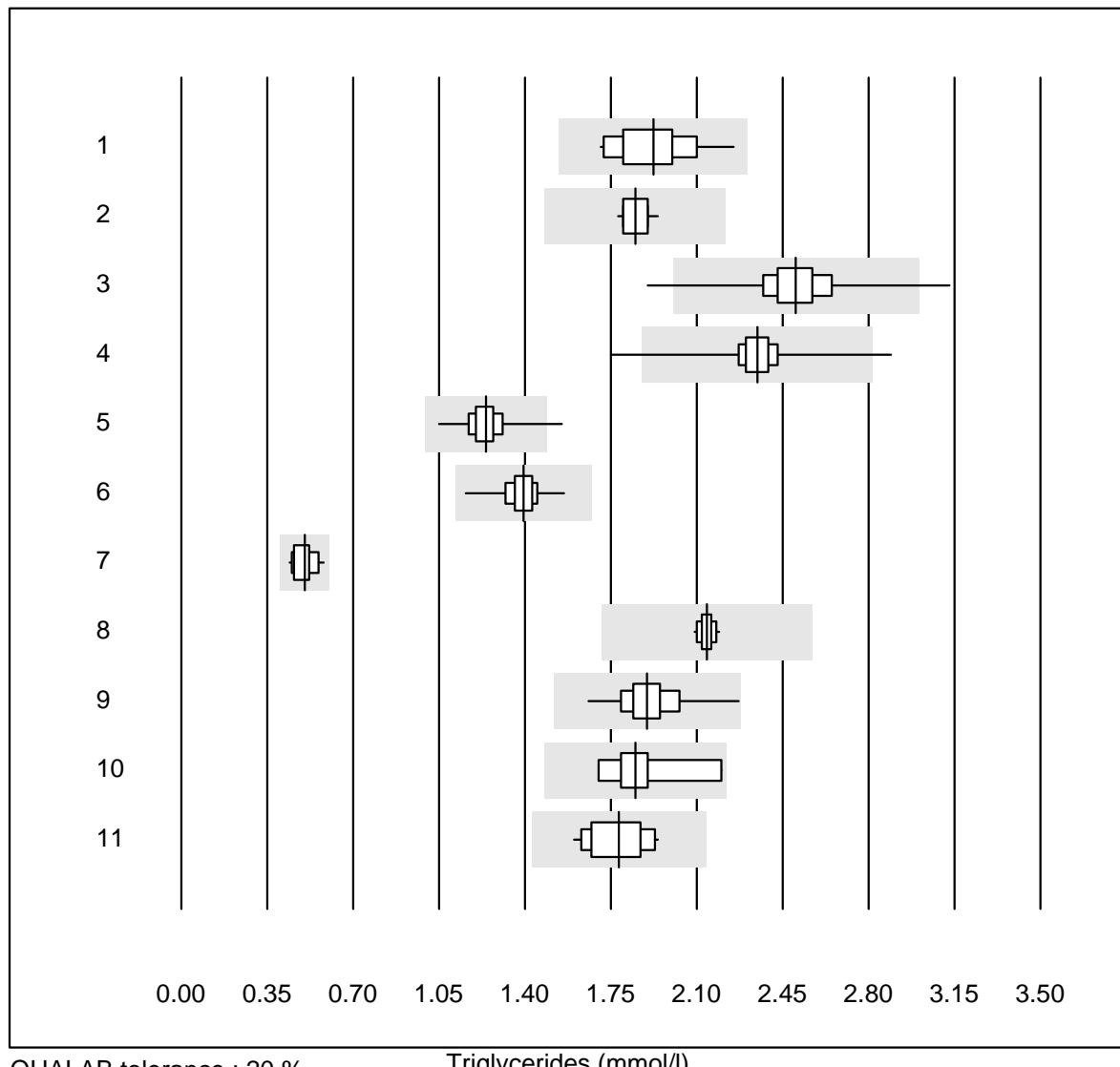


| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 IFCC with PP | 19 | 100.0 | 0.0 | 0.0 | 159 | 4.8 | e |
| 2 Cobas | 16 | 100.0 | 0.0 | 0.0 | 150 | 6.3 | e |
| 3 Reflotron | 838 | 97.1 | 2.1 | 0.8 | 193 | 7.0 | e |
| 4 Fuji Dri-Chem | 768 | 99.5 | 0.1 | 0.4 | 145 | 3.2 | e |
| 5 Spotchem/Ready | 147 | 99.3 | 0.0 | 0.7 | 125 | 4.8 | e |
| 6 Spotchem D-Concept | 166 | 98.8 | 1.2 | 0.0 | 127 | 4.6 | e |
| 7 IFCC without PP | 4 | 75.0 | 25.0 | 0.0 | 158 | 12.0 | e* |
| 8 Piccolo | 34 | 97.1 | 2.9 | 0.0 | 137 | 6.1 | e |
| 9 Abx Mira | 9 | 88.9 | 11.1 | 0.0 | 148 | 11.5 | e* |
| 10 Hitachi S40/M40 | 20 | 100.0 | 0.0 | 0.0 | 153 | 3.2 | e |
| 11 Autolyser/DiaSys | 14 | 100.0 | 0.0 | 0.0 | 148 | 4.4 | e |
| 12 Other methods | 4 | 100.0 | 0.0 | 0.0 | 157 | 0.9 | e |

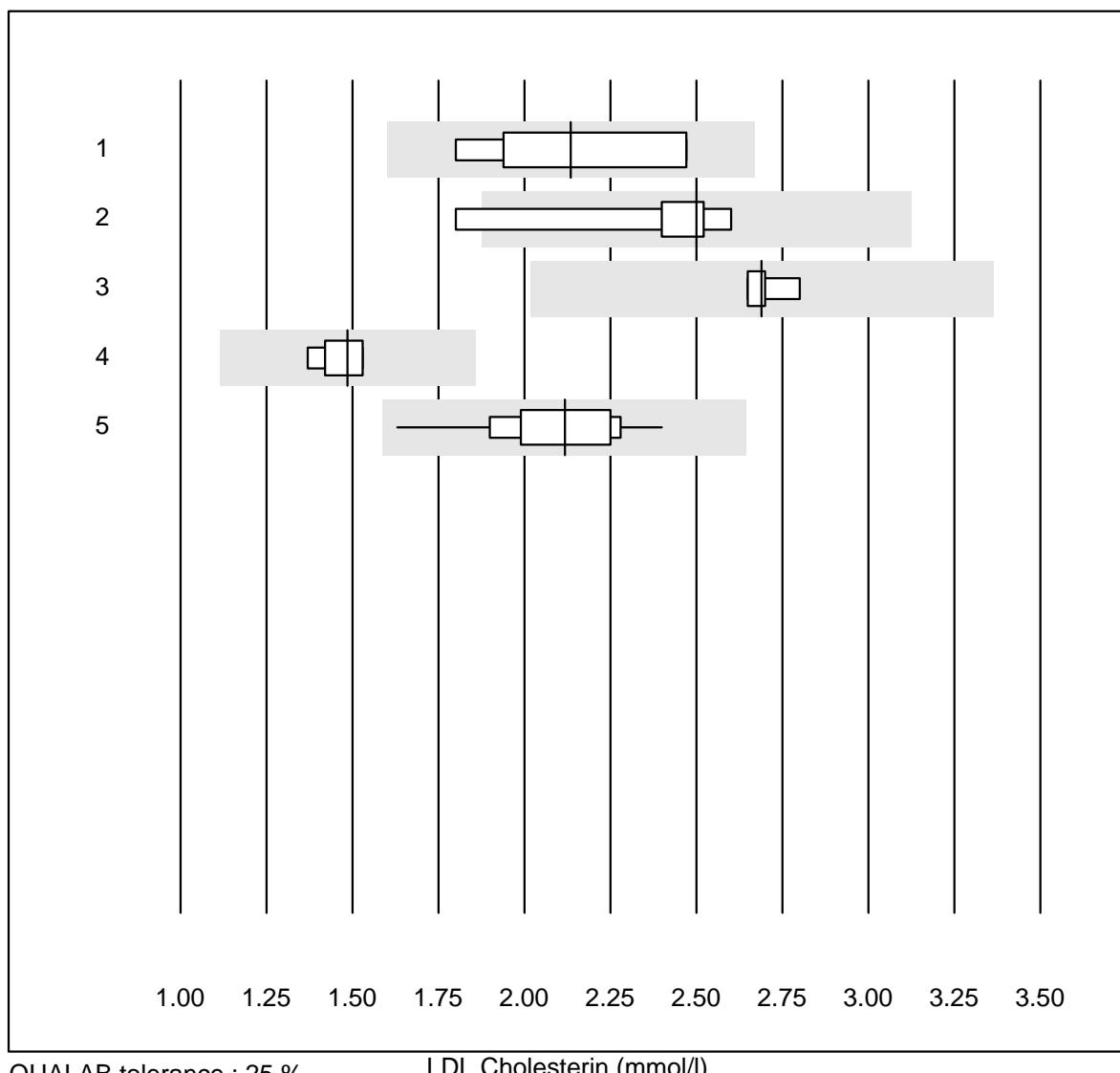
Alanine aminotransferase



Triglycerides



LDL Cholesterin

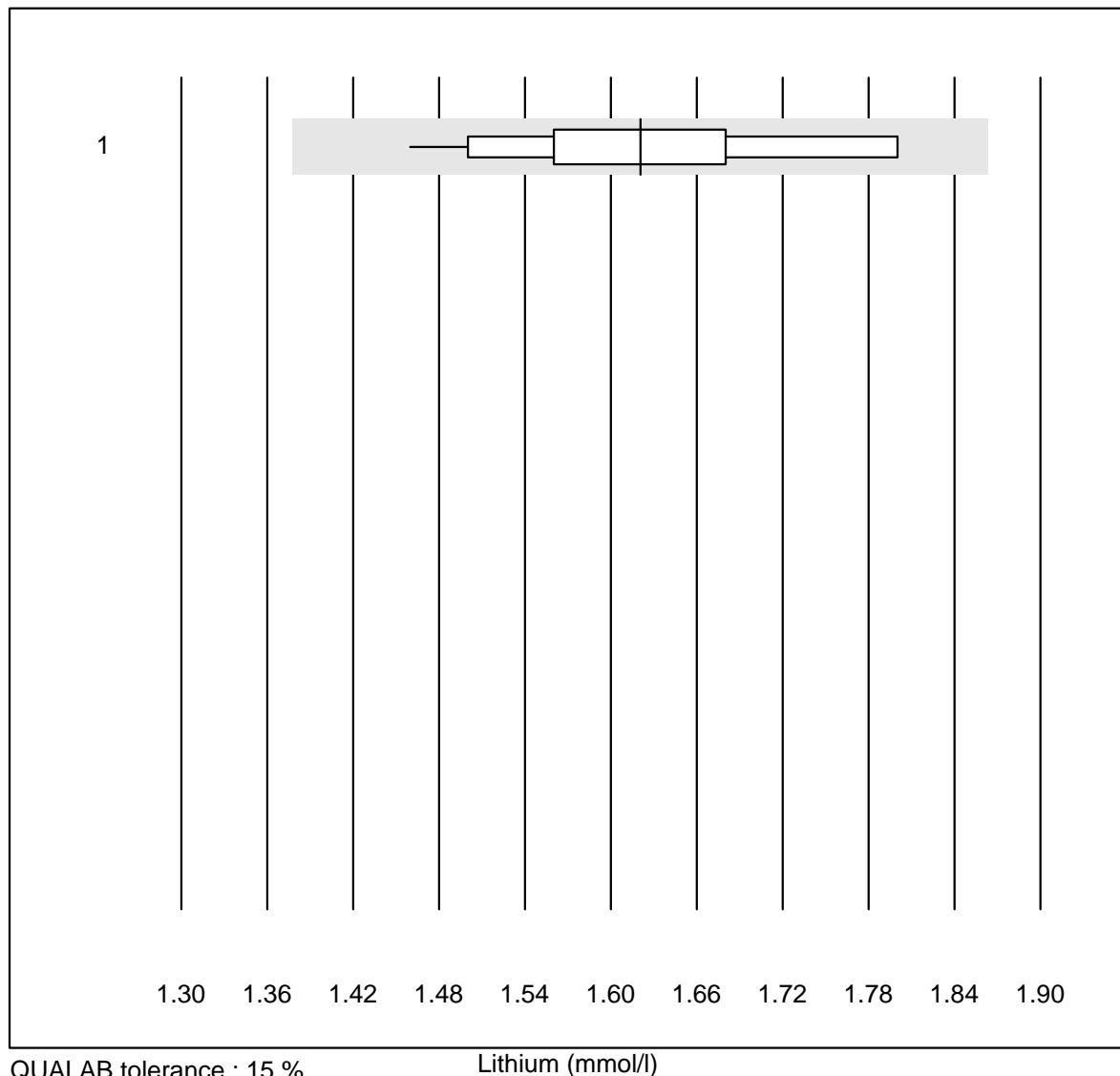


QUALAB tolerance : 25 %

LDL Cholesterin (mmol/l)

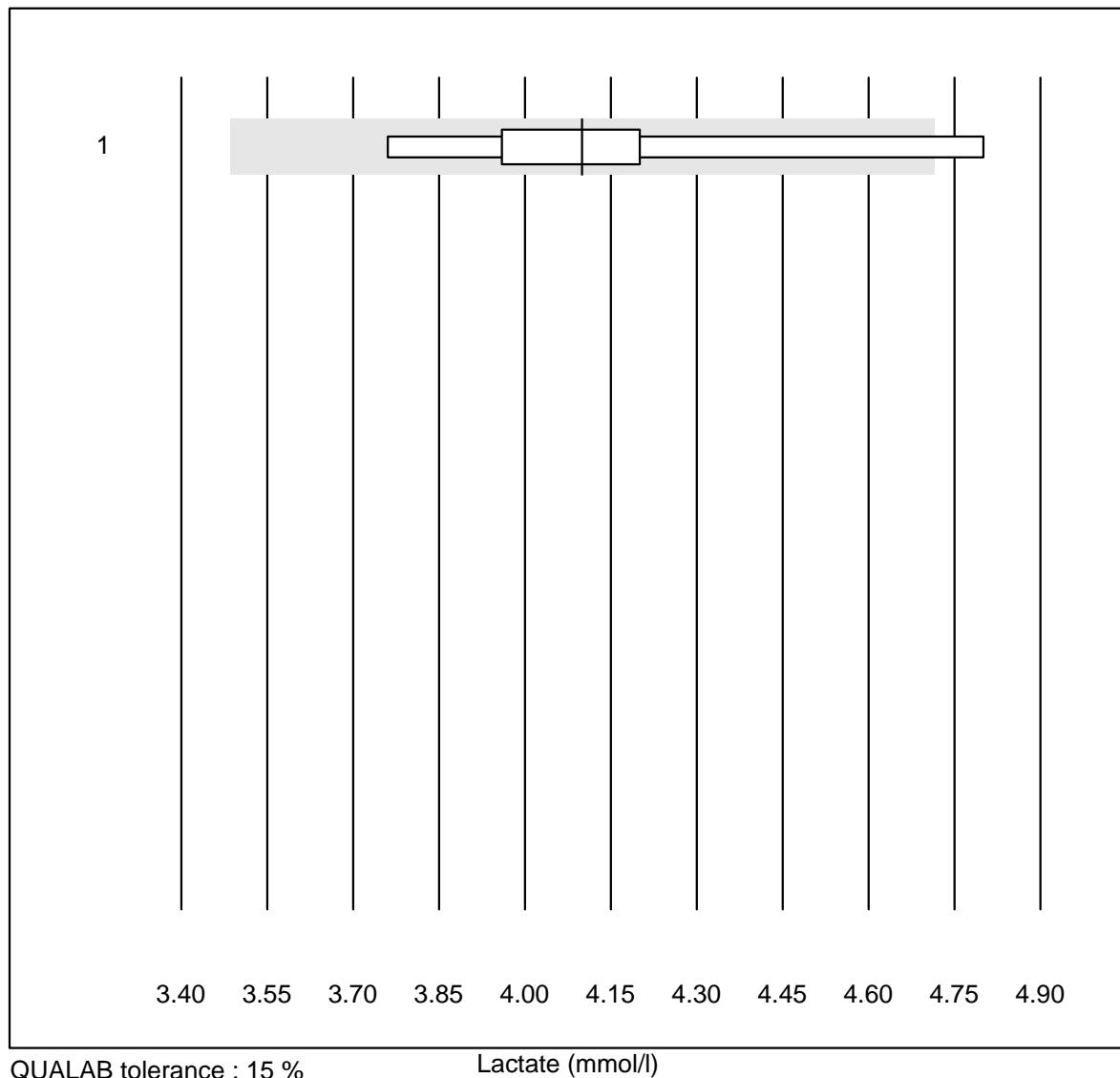
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 Abx Mira | 6 | 100.0 | 0.0 | 0.0 | 2.1 | 13.4 | e* |
| 2 Standard chemistry | 6 | 83.3 | 16.7 | 0.0 | 2.5 | 12.3 | e* |
| 3 Roche, Cobas | 4 | 100.0 | 0.0 | 0.0 | 2.7 | 2.4 | e |
| 4 Hitachi S40/M40 | 8 | 87.5 | 0.0 | 12.5 | 1.5 | 4.0 | e |
| 5 Autolyser/DiaSys | 12 | 100.0 | 0.0 | 0.0 | 2.1 | 9.7 | e |

Lithium

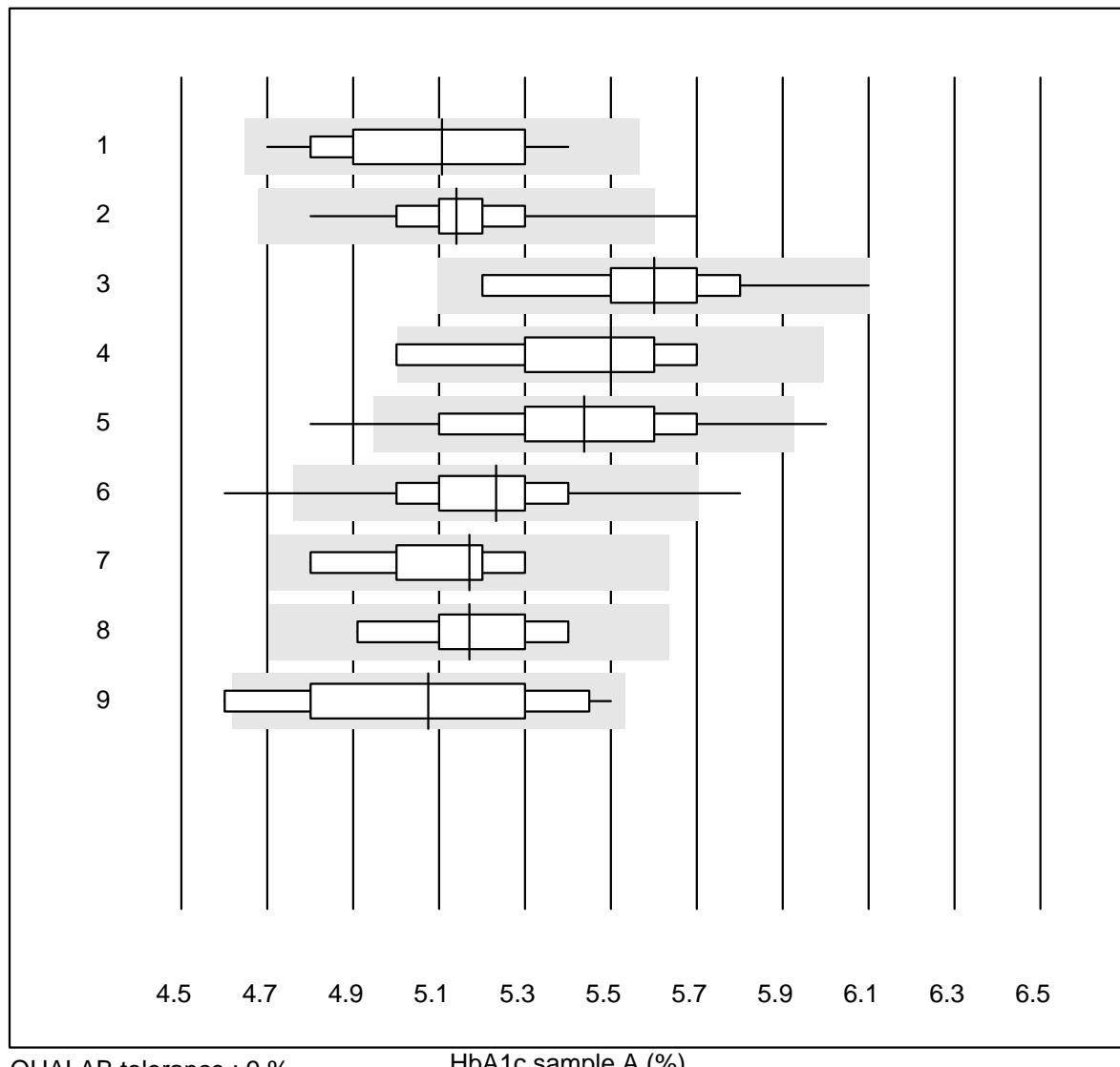


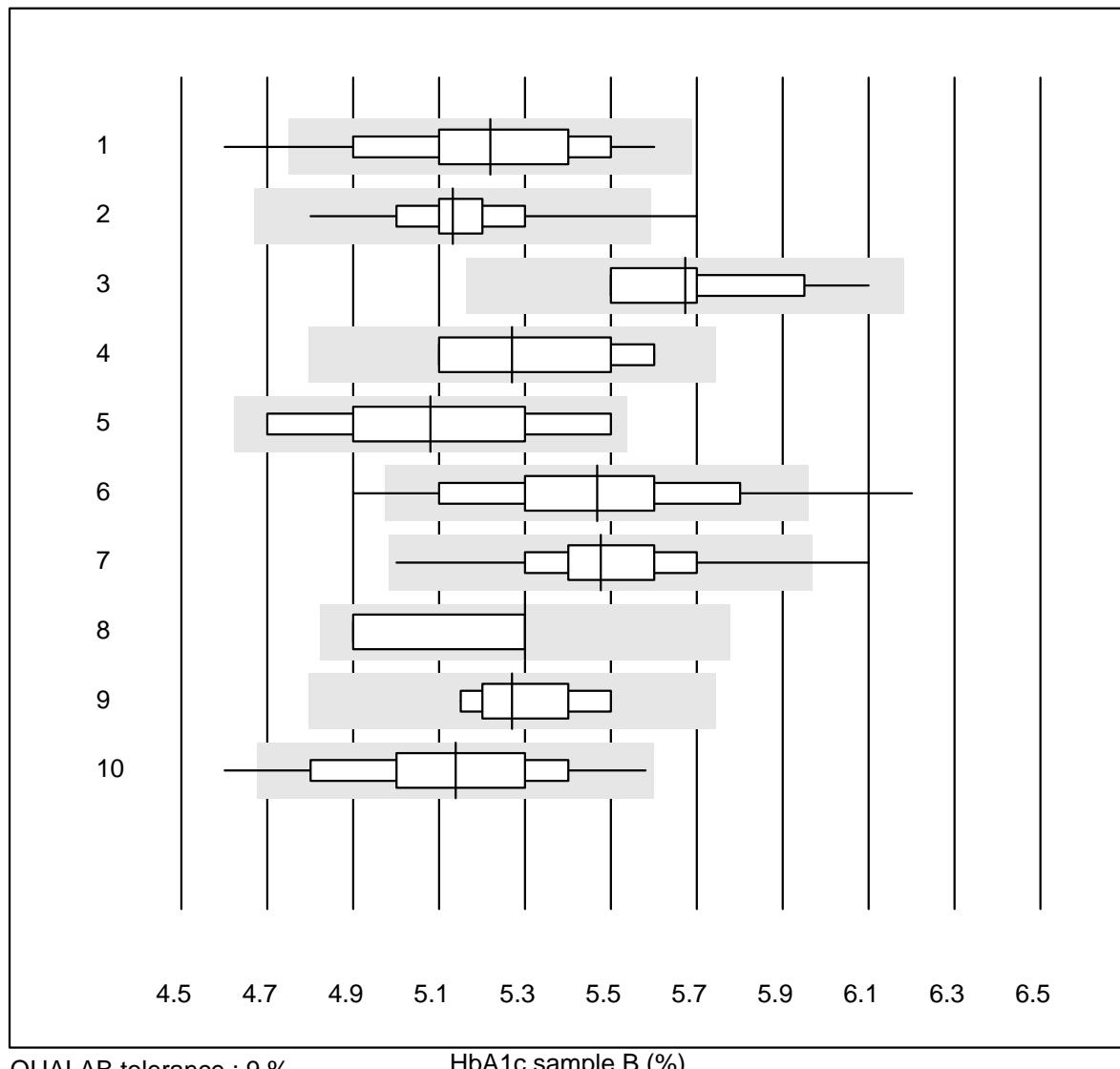
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 all Participants | 14 | 100.0 | 0.0 | 0.0 | 1.62 | 6.2 | e |

Lactate



| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | all Participants | 7 | 85.7 | 14.3 | 0.0 | 4.10 | 7.7 | e* |

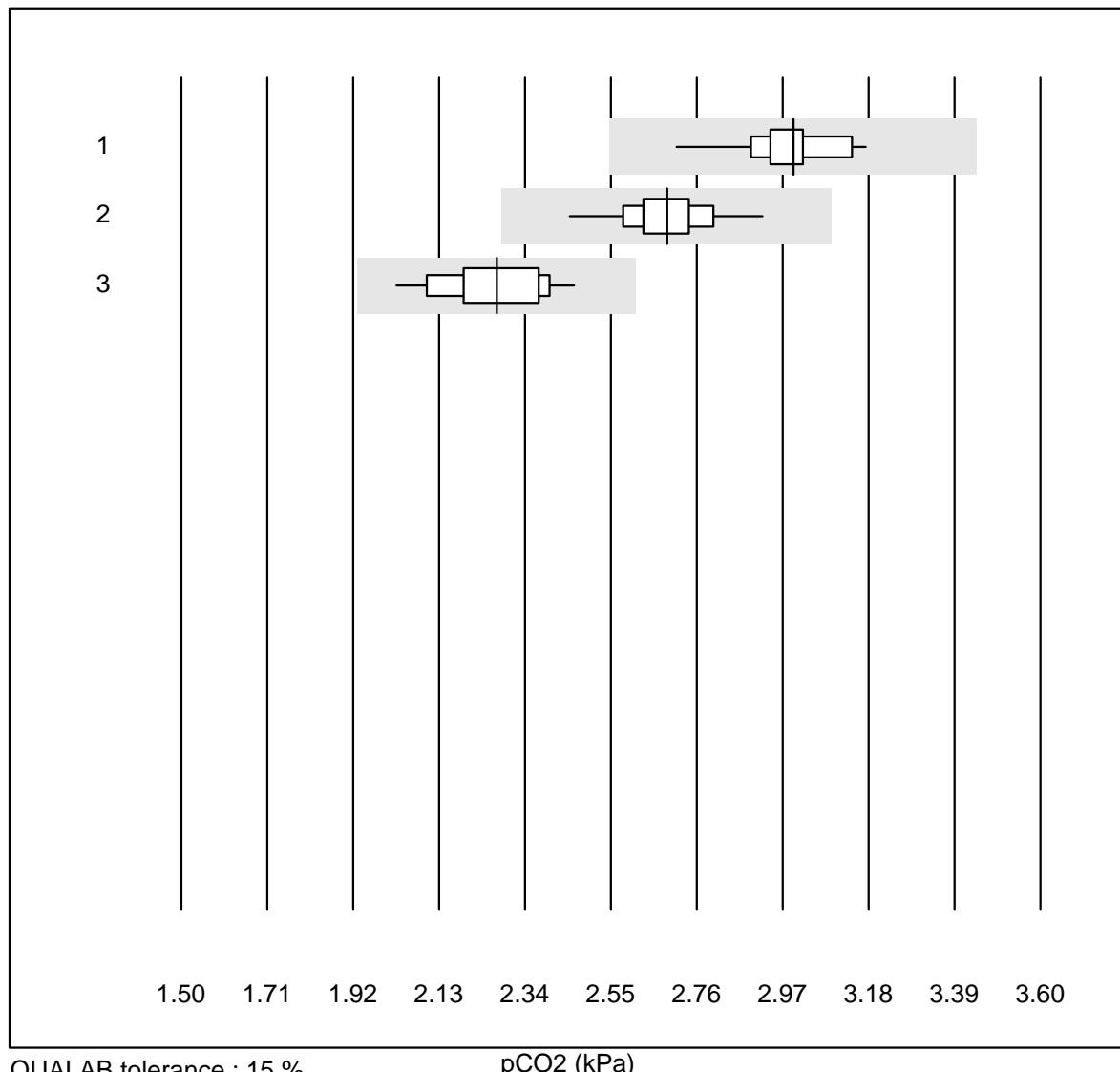
HbA1c sample A

HbA1c sample B

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|---------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Cobas b101 | 32 | 96.9 | 3.1 | 0.0 | 5.2 | 4.4 | e |
| 2 Afinion | 579 | 98.7 | 1.0 | 0.3 | 5.1 | 2.5 | e |
| 3 Eurolyser | 20 | 100.0 | 0.0 | 0.0 | 5.7 | 3.2 | e |
| 4 A1c Now | 4 | 100.0 | 0.0 | 0.0 | 5.3 | 4.9 | a |
| 5 Hemocue HbA1c 501 | 11 | 90.9 | 0.0 | 9.1 | 5.1 | 5.3 | e* |
| 6 NycoCard | 73 | 89.1 | 6.8 | 4.1 | 5.5 | 5.1 | e |
| 7 DCA2000/Vantage | 225 | 98.2 | 0.9 | 0.9 | 5.5 | 3.2 | e |
| 8 Others | 4 | 100.0 | 0.0 | 0.0 | 5.3 | 3.8 | e* |
| 9 HPLC | 5 | 100.0 | 0.0 | 0.0 | 5.3 | 2.8 | a |
| 10 Roche, Cobas | 13 | 92.3 | 7.7 | 0.0 | 5.1 | 5.2 | e* |

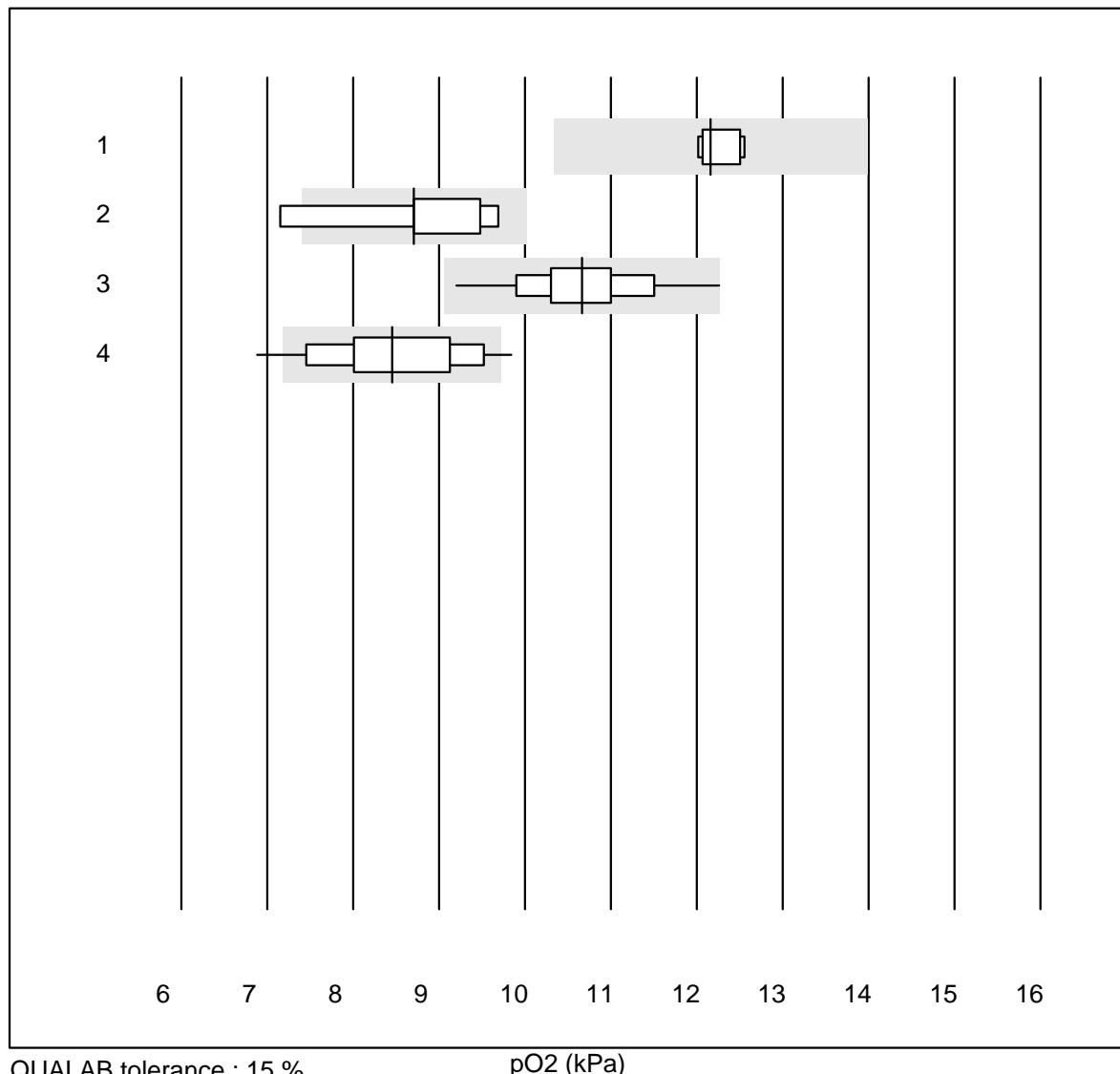
K4 Blood gases

pCO₂



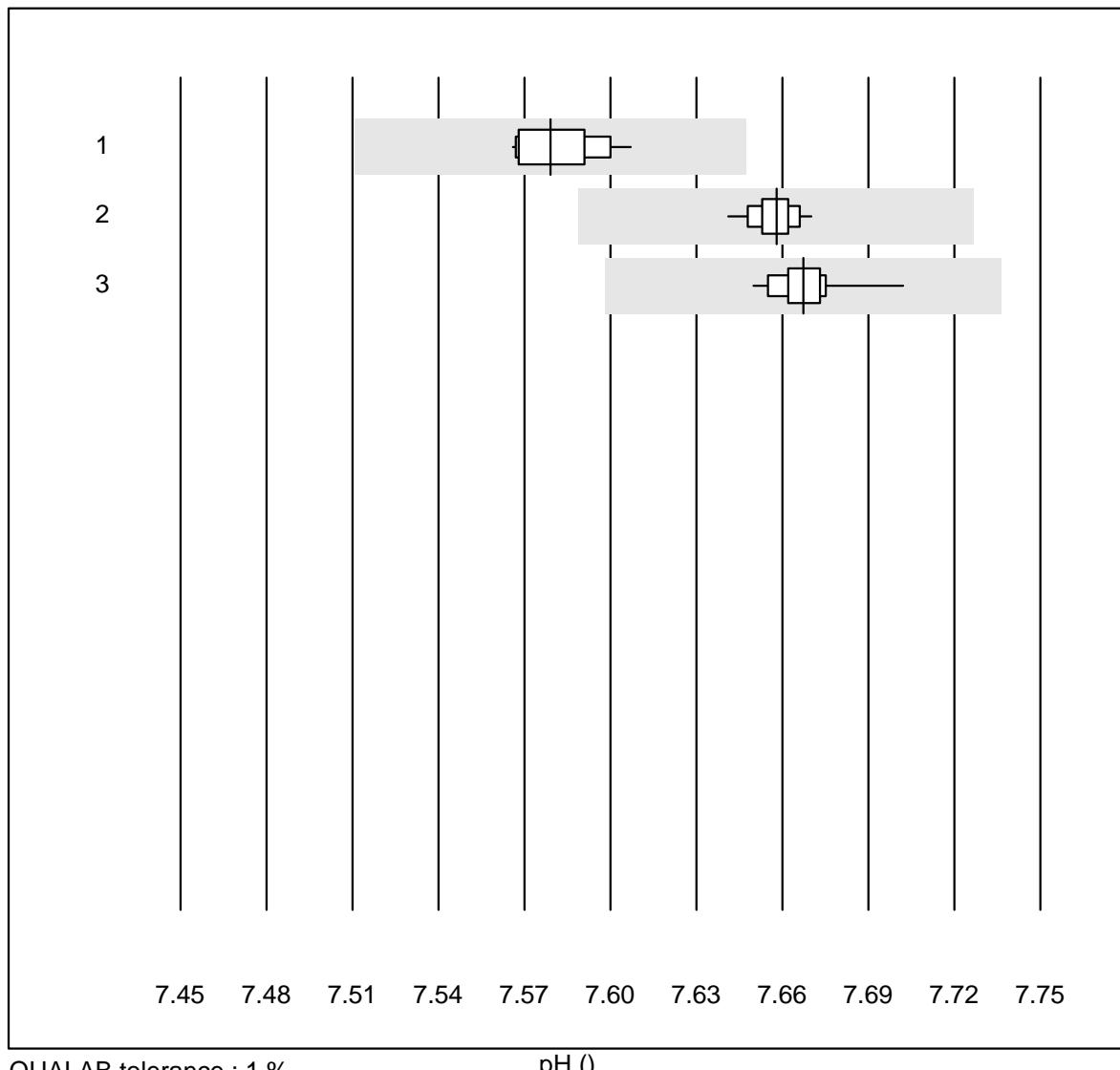
K4 Blood gases

pO₂



K4 Blood gases

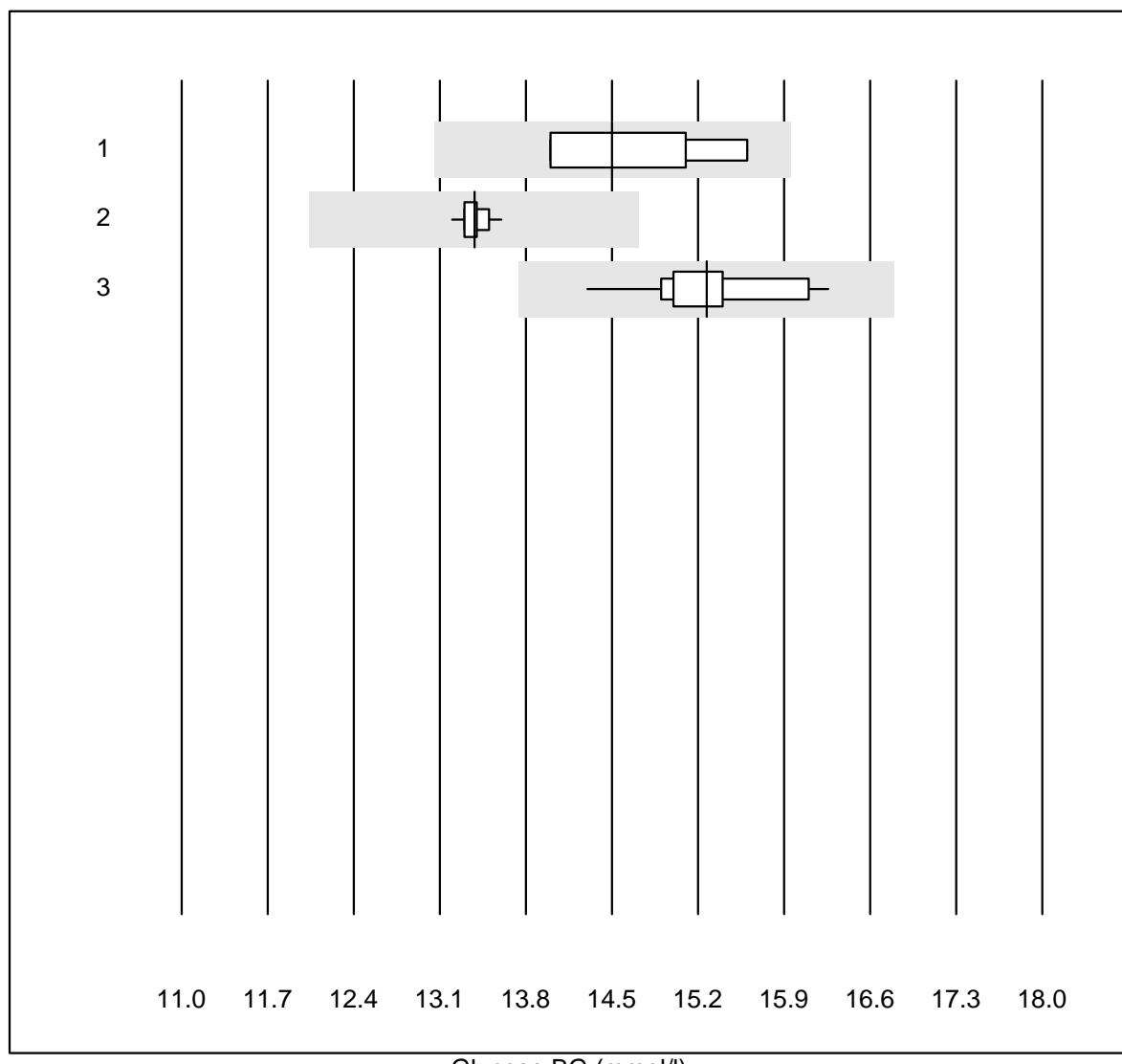
pH



QUALAB tolerance : 1 %

pH ()

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Cobas b121/123/221 | 15 | 100.0 | 0.0 | 0.0 | 7.58 | 0.2 | e |
| 2 iStat | 36 | 100.0 | 0.0 | 0.0 | 7.66 | 0.1 | e |
| 3 EPOC | 24 | 95.8 | 0.0 | 4.2 | 7.67 | 0.1 | e |

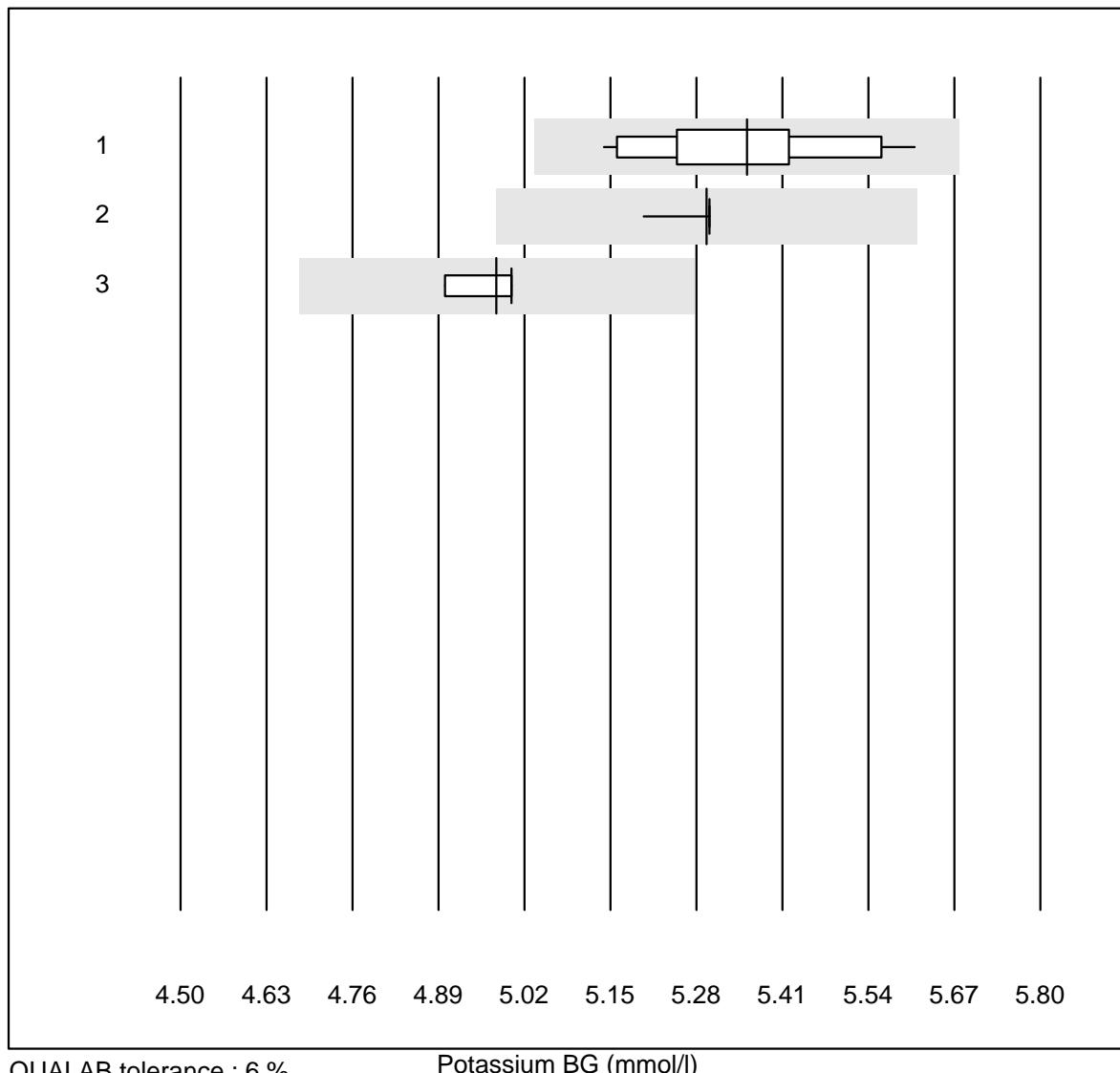
Glucose BG

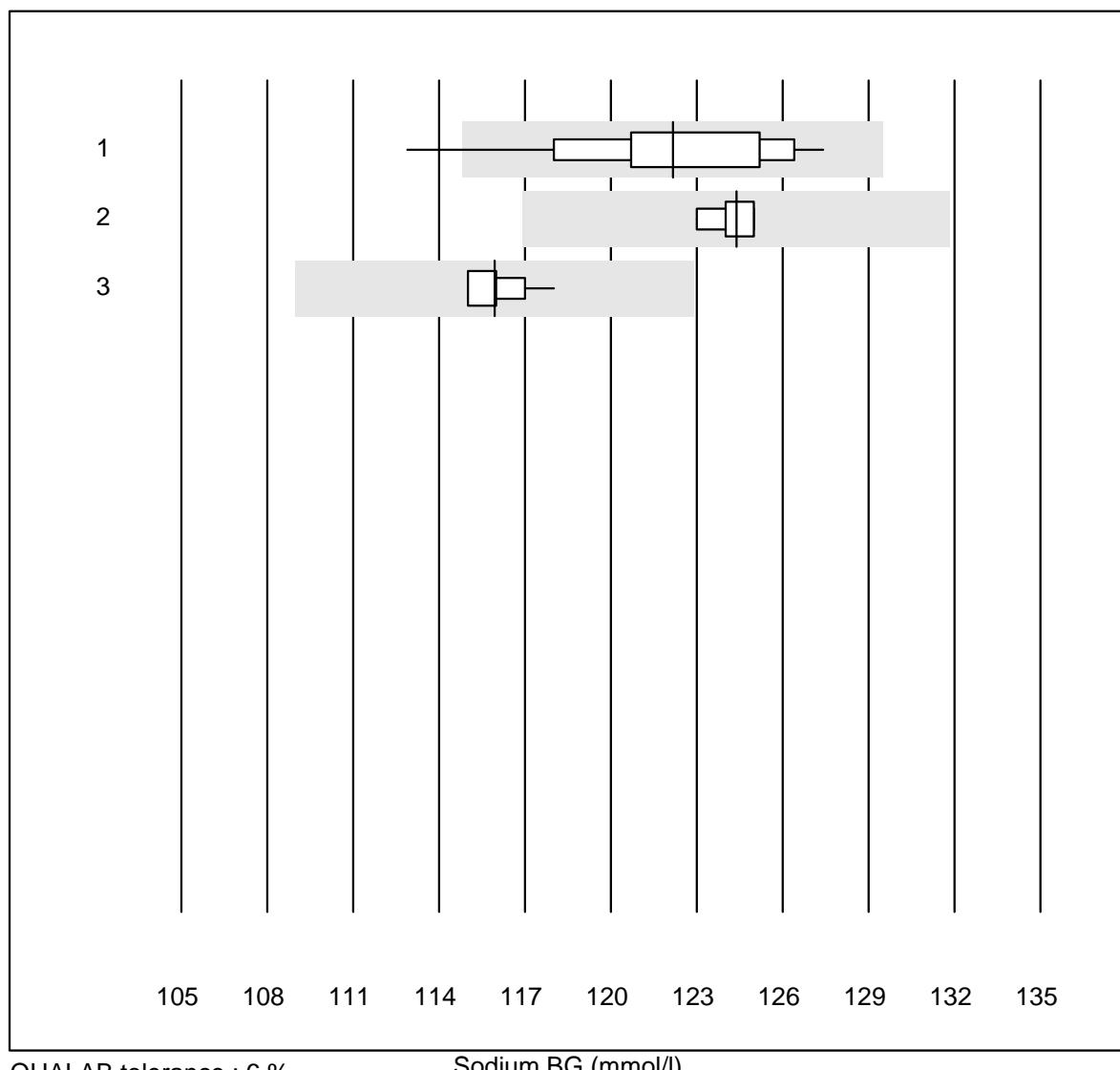
QUALAB tolerance : 10 %

Glucose BG (mmol/l)

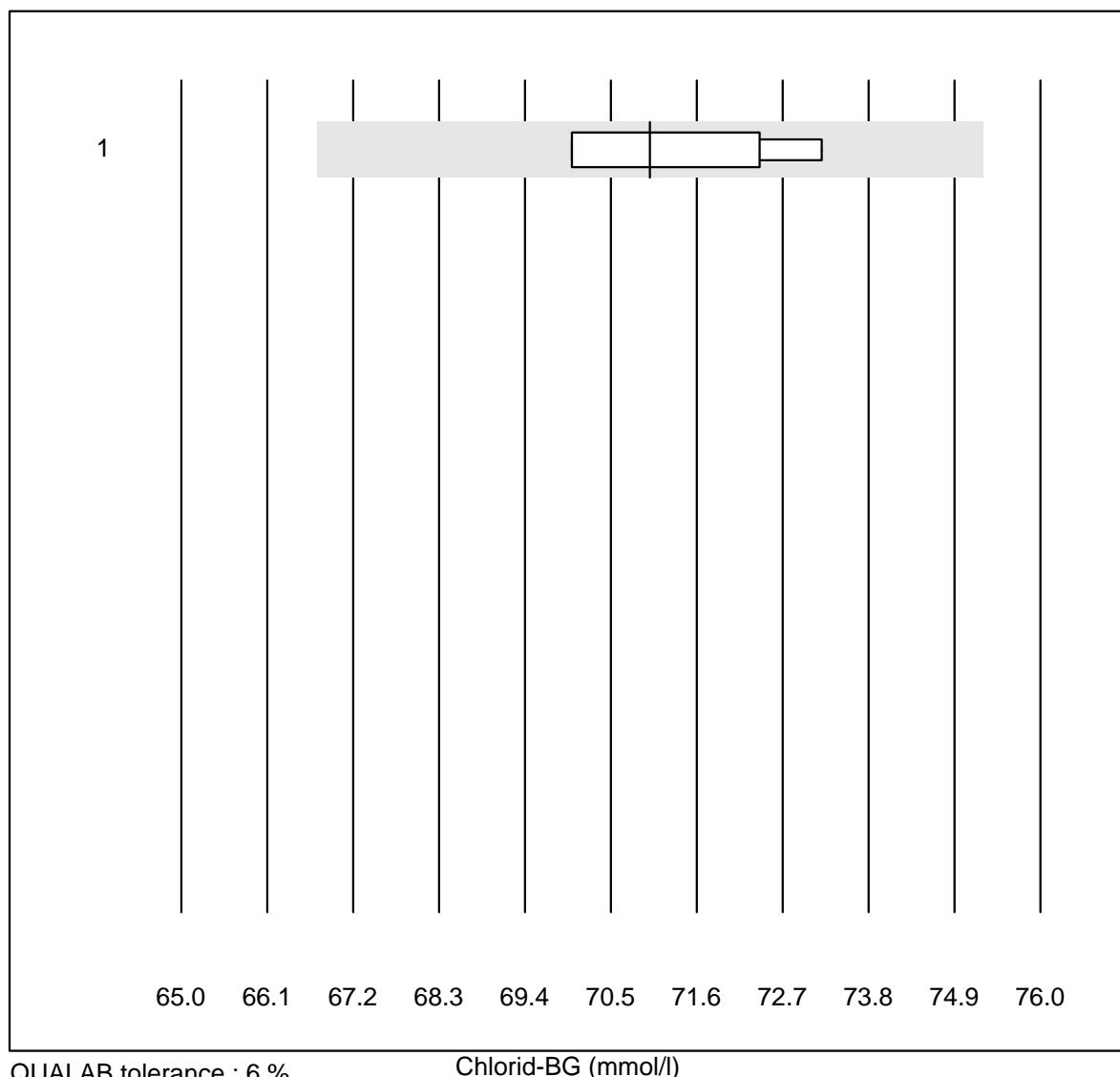
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Cobas b121/123/221 | 8 | 87.5 | 0.0 | 12.5 | 14.5 | 4.1 | e* |
| 2 iStat | 11 | 100.0 | 0.0 | 0.0 | 13.4 | 0.8 | e |
| 3 EPOC | 16 | 93.7 | 0.0 | 6.3 | 15.3 | 3.2 | e |

Potassium BG

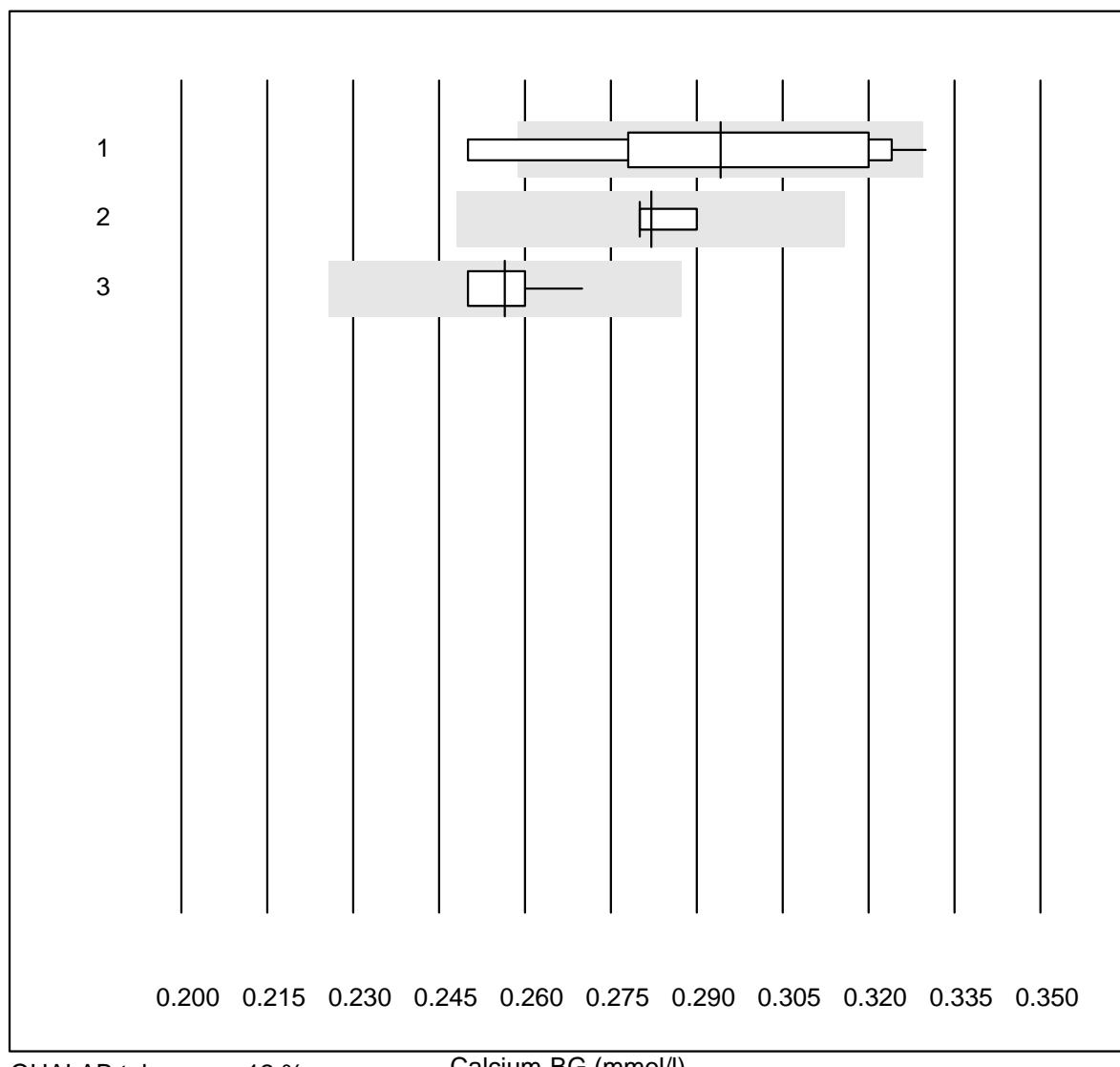


Sodium BG

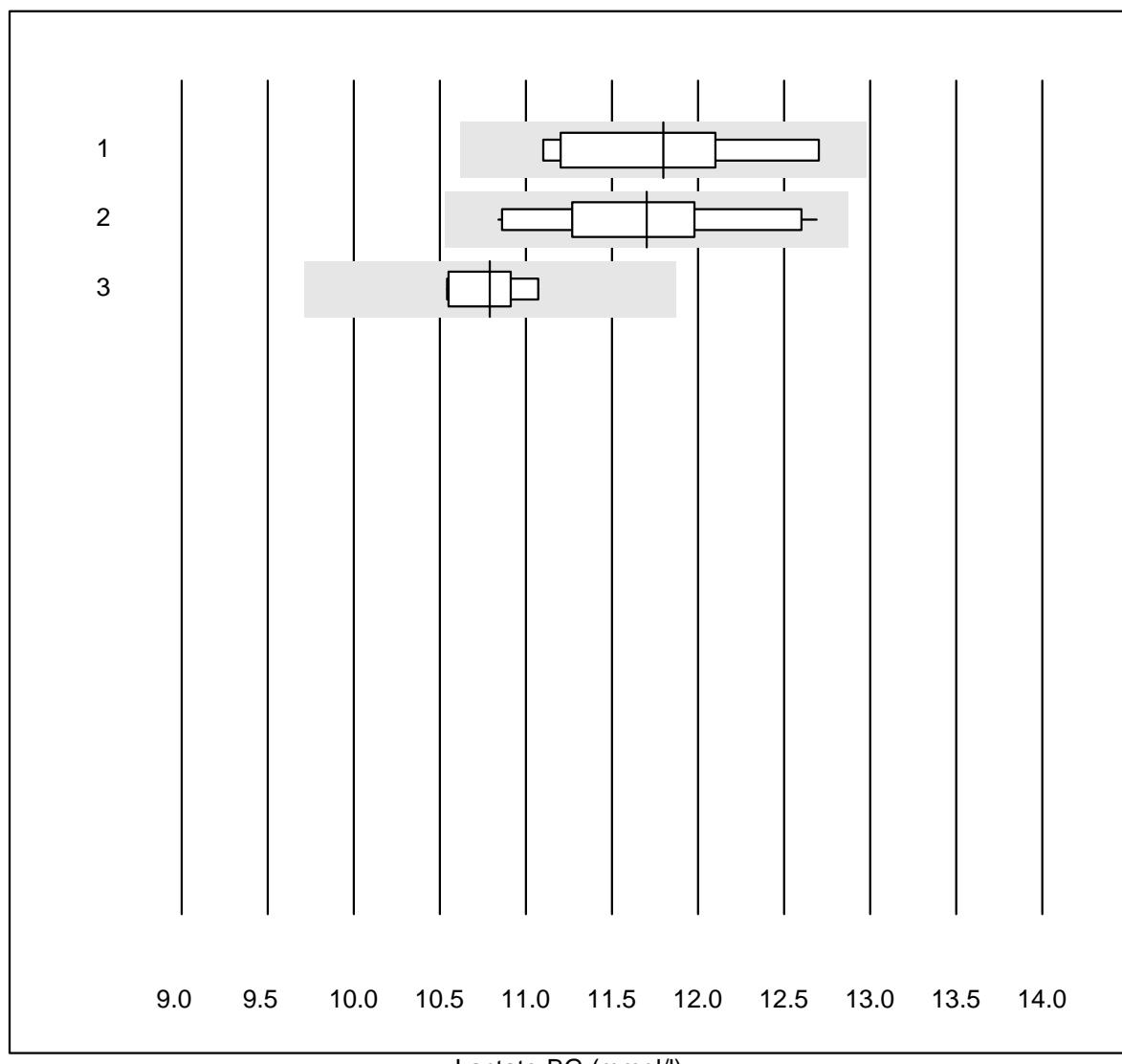
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Cobas b121/123/221 | 16 | 93.7 | 6.3 | 0.0 | 122.2 | 3.0 | e* |
| 2 iStat | 21 | 100.0 | 0.0 | 0.0 | 124.4 | 0.6 | e |
| 3 EPOC | 19 | 94.7 | 0.0 | 5.3 | 115.9 | 0.8 | e |

Chlorid-BG

| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Cobas b121/123/221 | 6 | 83.3 | 0.0 | 16.7 | 71.0 | 2.0 | e* |

Calcium-BG

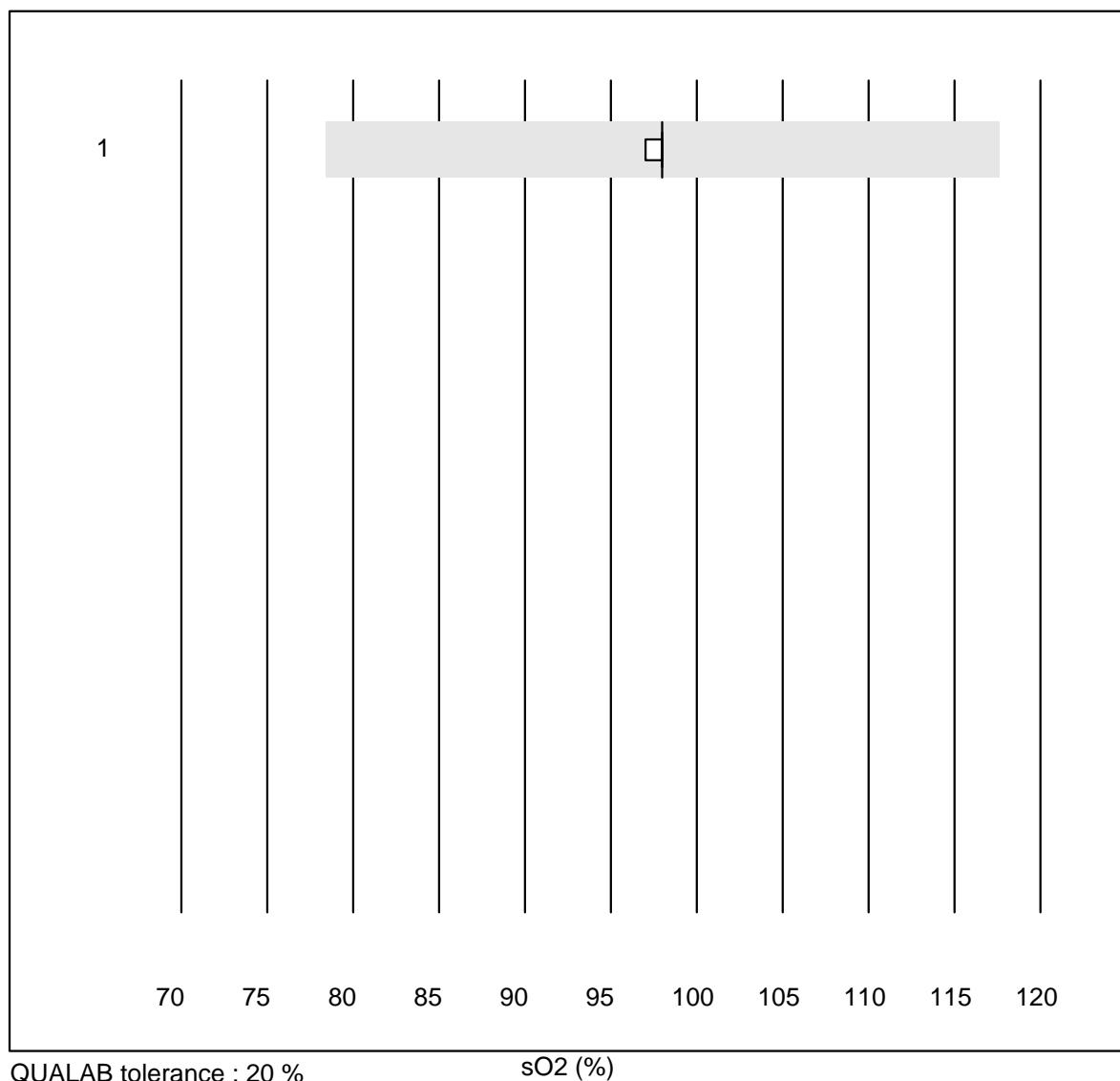
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Cobas b121/123/221 | 10 | 80.0 | 20.0 | 0.0 | 0.29 | 8.7 | e* |
| 2 | iStat | 10 | 100.0 | 0.0 | 0.0 | 0.28 | 1.5 | e |
| 3 | EPOC | 18 | 94.4 | 0.0 | 5.6 | 0.26 | 2.4 | e |

Lactate-BG

QUALAB tolerance : 10 %

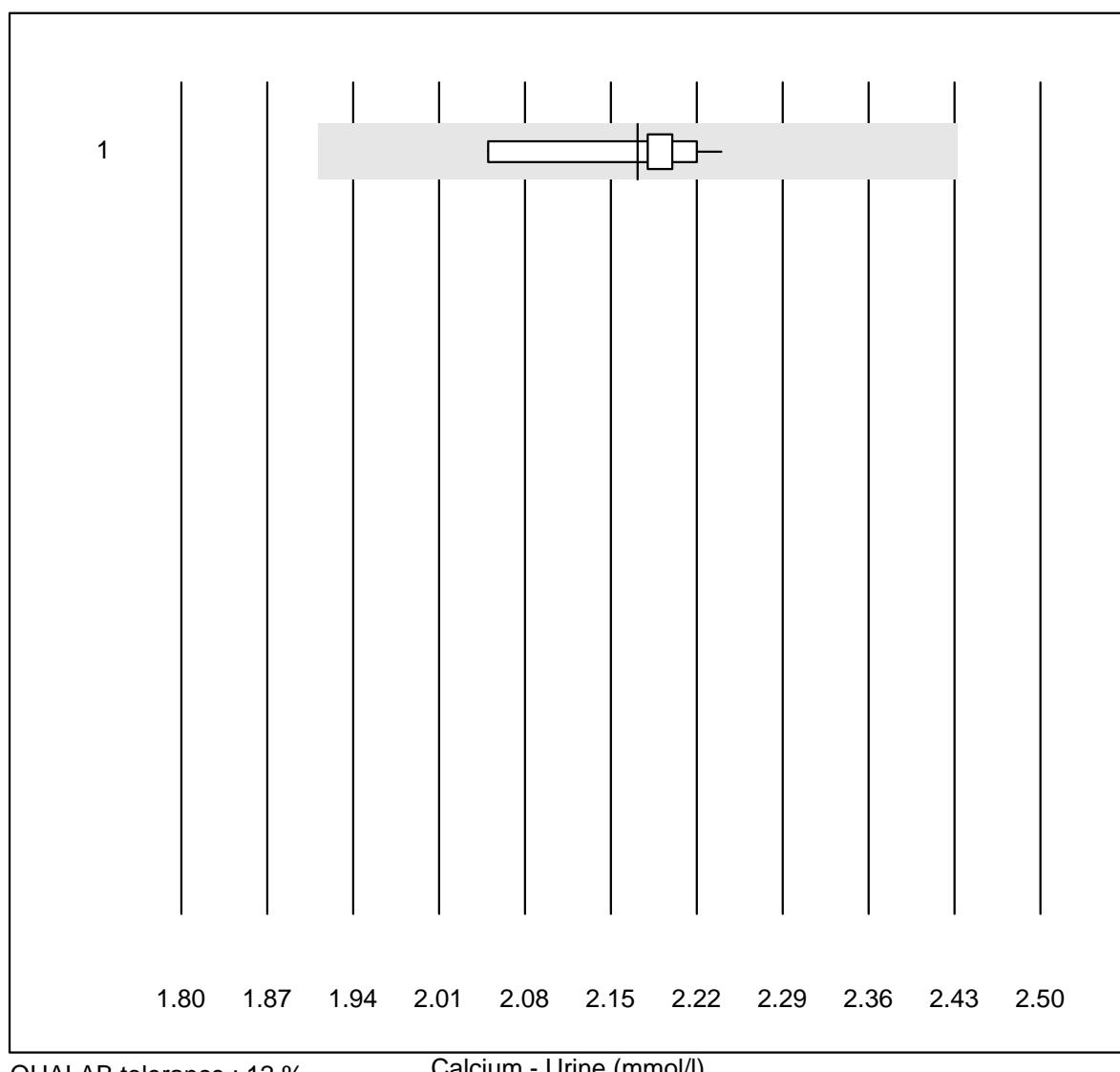
Lactate-BG (mmol/l)

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Cobas b121/123/221 | 6 | 83.3 | 0.0 | 16.7 | 11.80 | 5.7 | e* |
| 2 EPOC | 20 | 95.0 | 0.0 | 5.0 | 11.70 | 4.8 | e |
| 3 iStat | 8 | 100.0 | 0.0 | 0.0 | 10.79 | 1.7 | e |

sO2

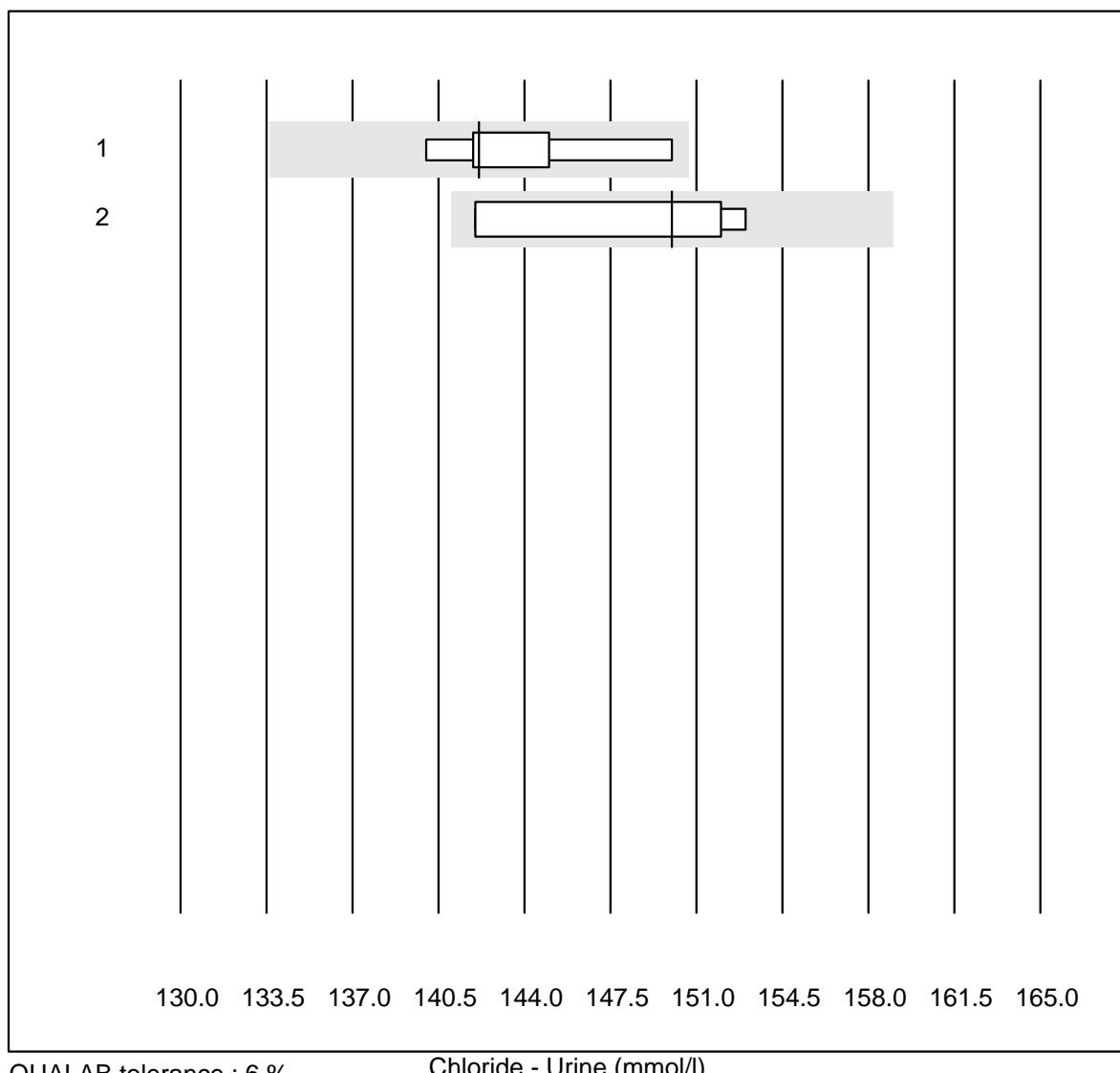
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | iStat | 6 | 100.0 | 0.0 | 0.0 | 98.000 | 0.4 | e |

Calcium - Urine



| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Standard chemistry | 13 | 100.0 | 0.0 | 0.0 | 2.17 | 2.9 | e |

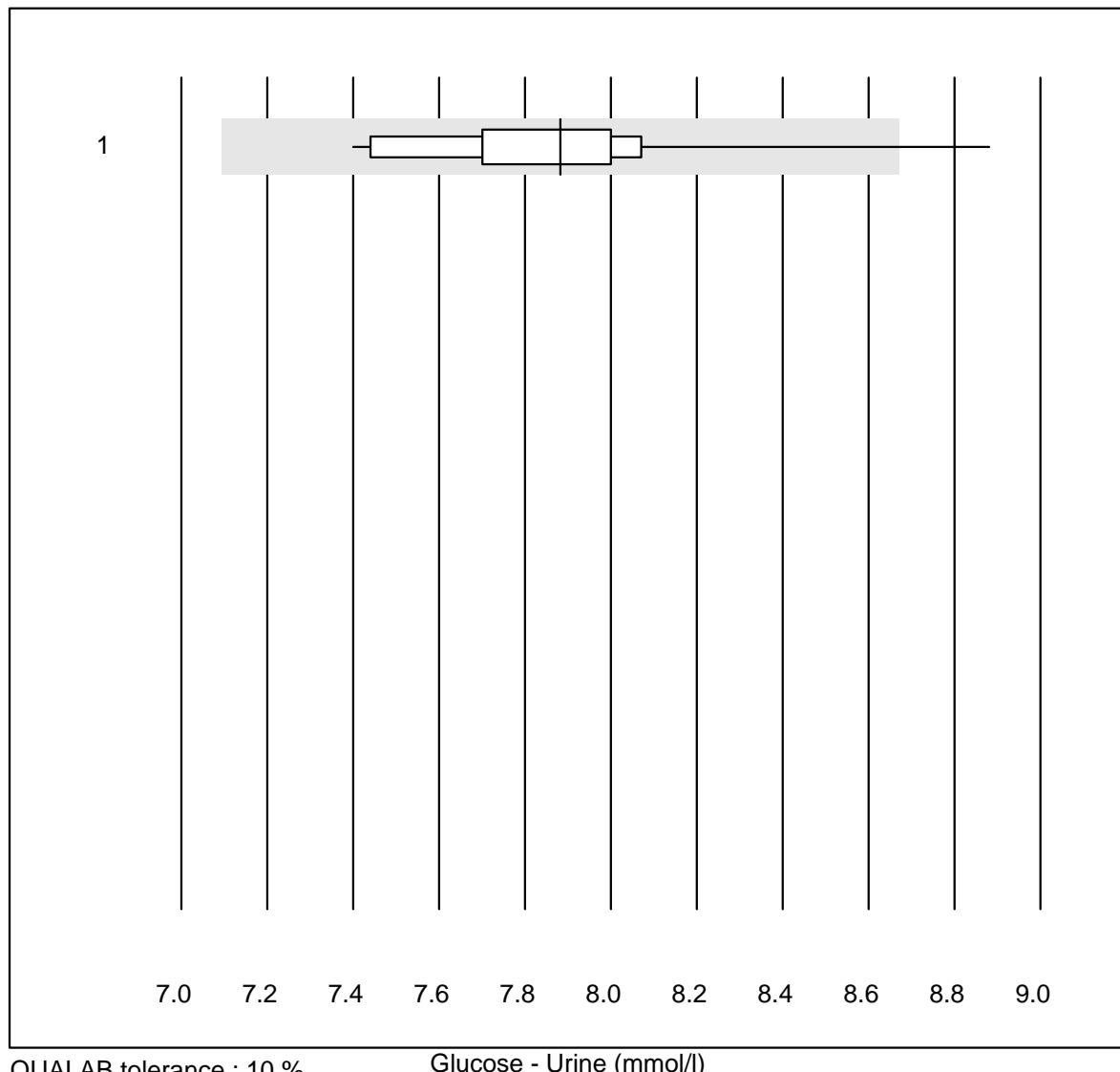
Chloride - Urine



QUALAB tolerance : 6 %

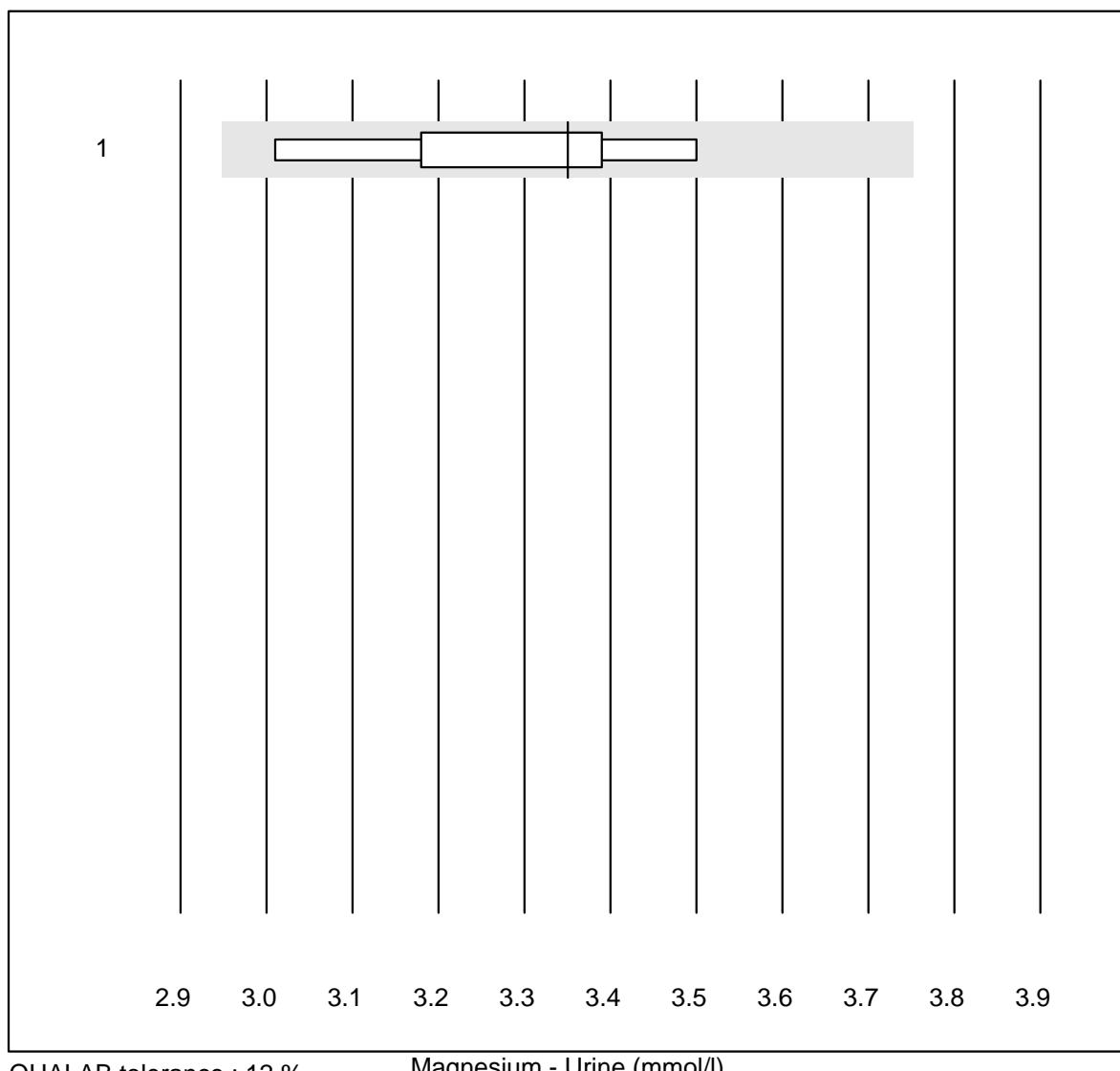
Chloride - Urine (mmol/l)

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Standard chemistry | 6 | 83.3 | 0.0 | 16.7 | 142 | 2.7 | e* |
| 2 ISE | 4 | 100.0 | 0.0 | 0.0 | 150 | 3.4 | e* |

Glucose - Urine

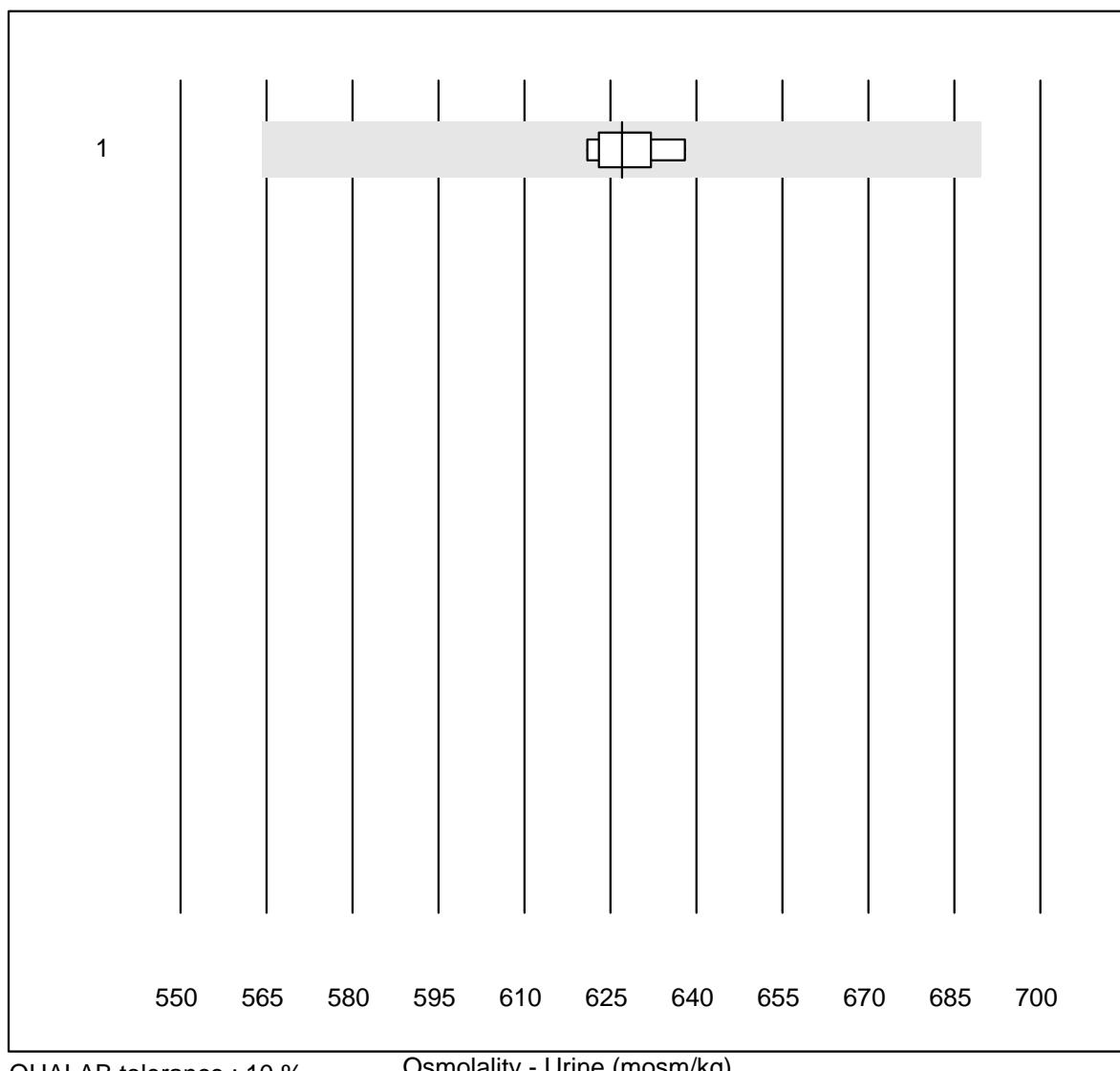
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Standard chemistry | 13 | 92.3 | 7.7 | 0.0 | 7.9 | 4.6 | e* |

Magnesium - Urine



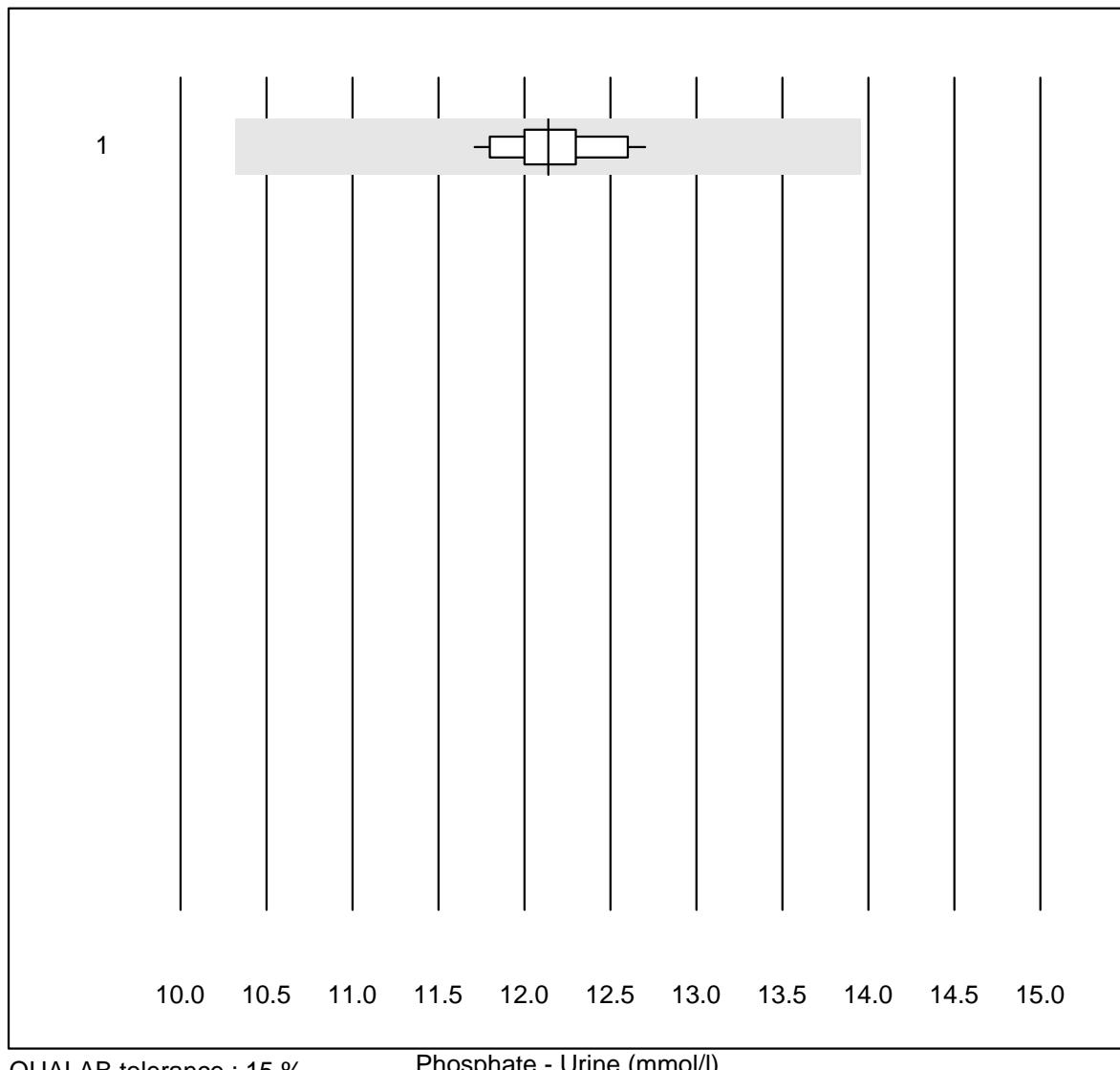
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Standard chemistry | 8 | 100.0 | 0.0 | 0.0 | 3.4 | 4.6 | e* |

Osmolality - Urine



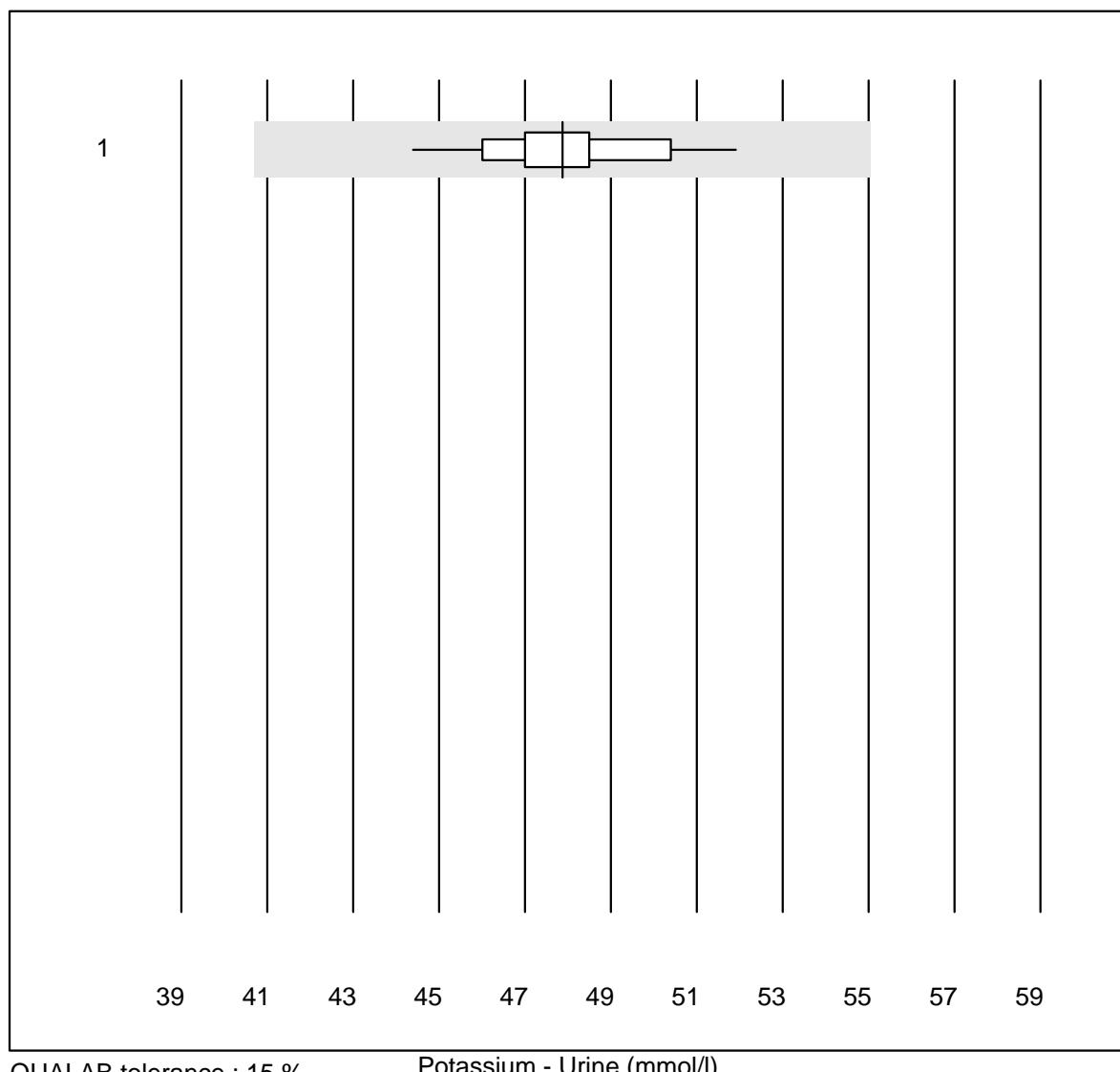
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|-----------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Cryoskopy | 7 | 100.0 | 0.0 | 0.0 | 627 | 0.9 | e |

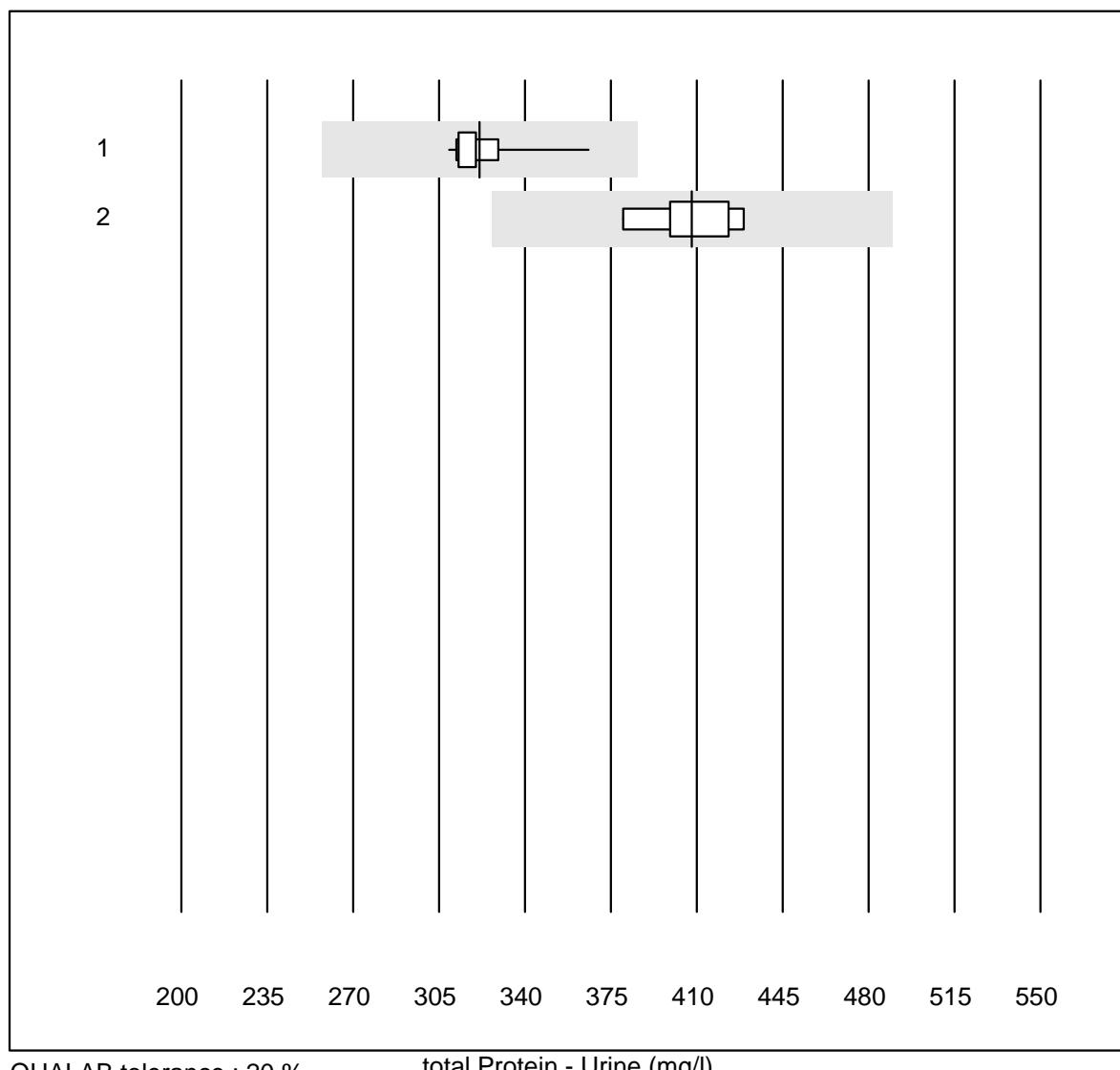
Phosphate - Urine



| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Standard chemistry | 13 | 100.0 | 0.0 | 0.0 | 12.1 | 2.4 | e |

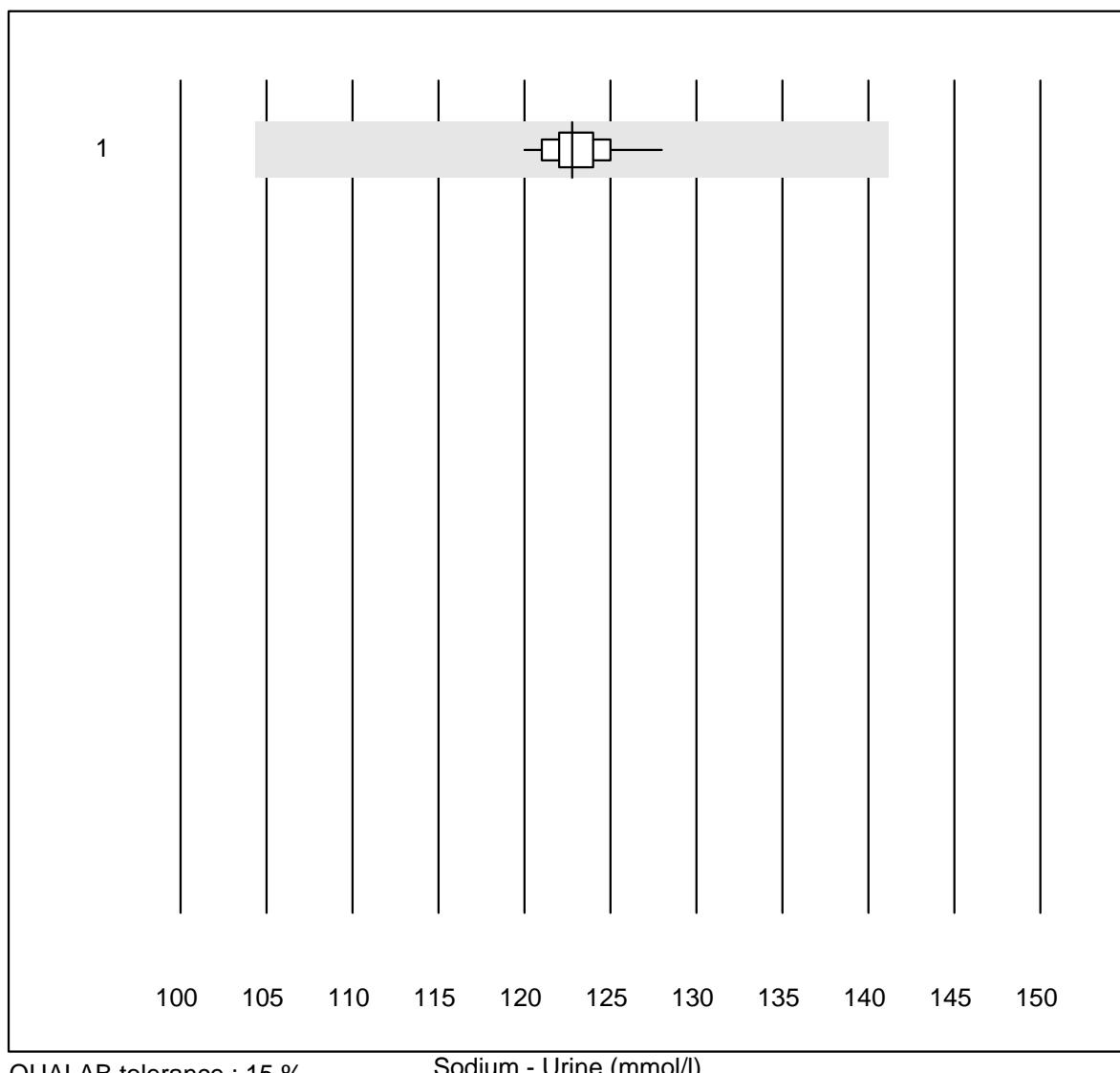
Potassium - Urine



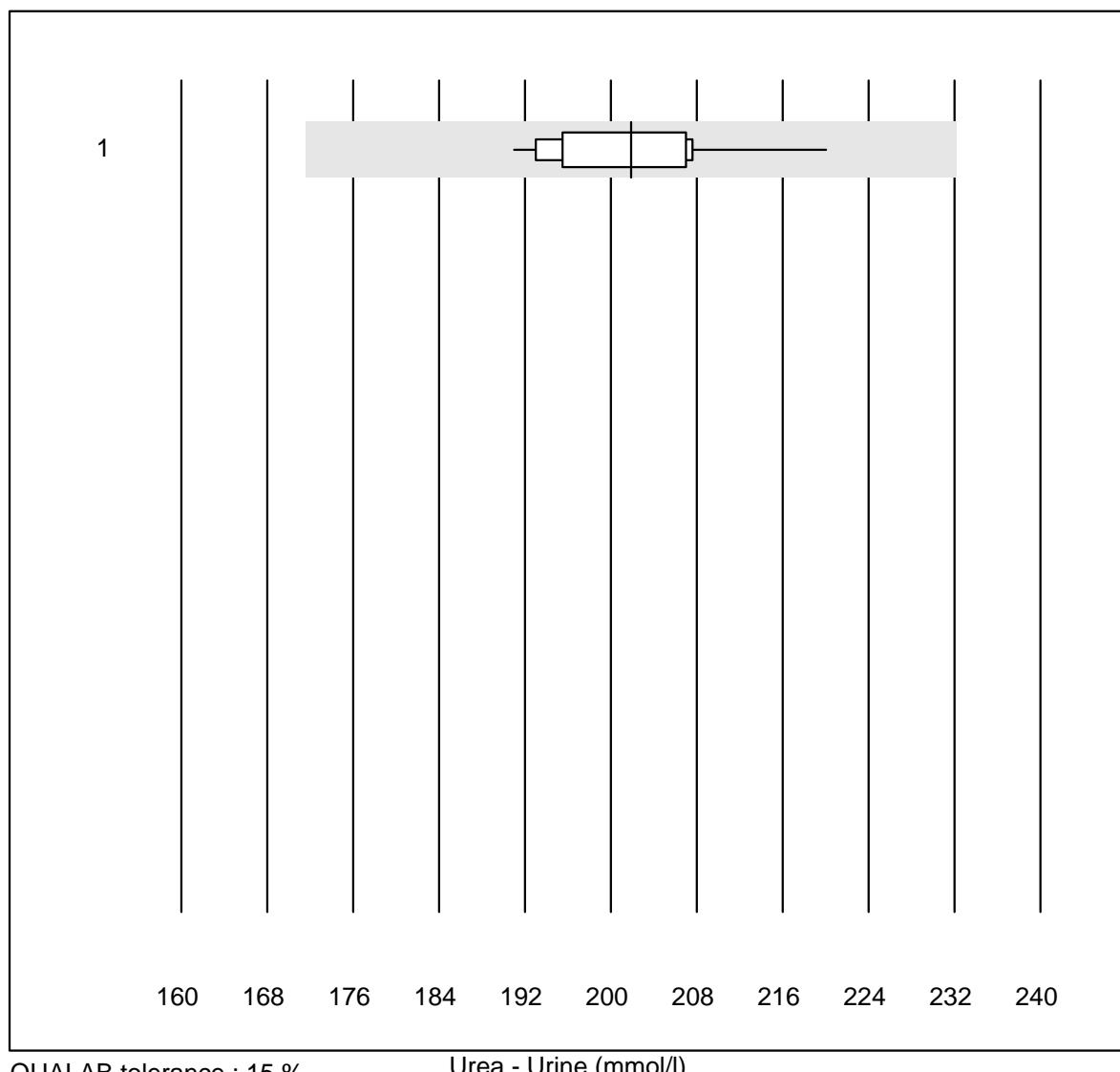
total Protein - Urine

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Cobas/Roche | 13 | 100.0 | 0.0 | 0.0 | 321.6 | 4.5 | e |
| 2 Standard chemistry | 5 | 100.0 | 0.0 | 0.0 | 408.0 | 4.8 | e |

Sodium - Urine

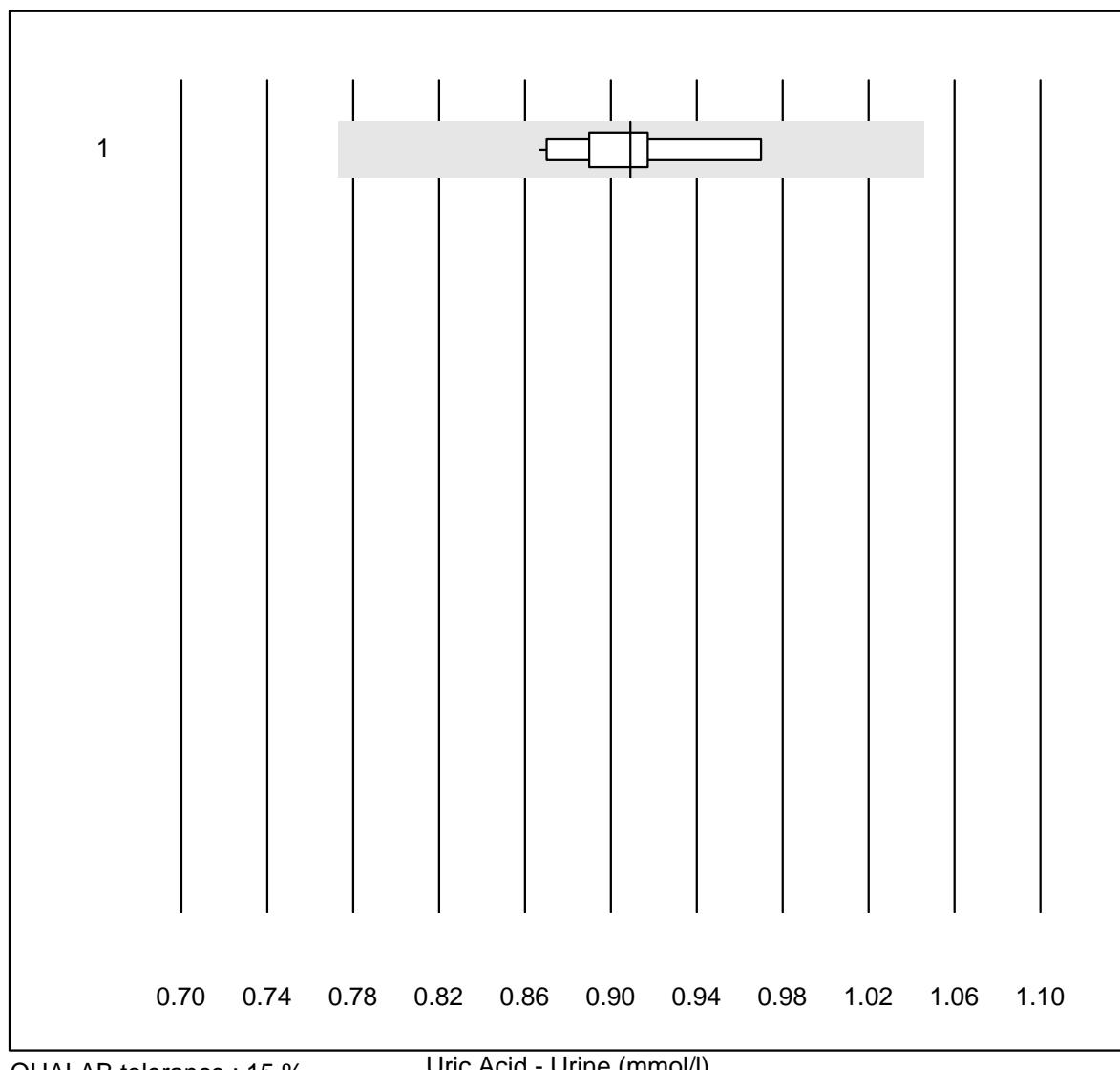


| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 all Participants | 19 | 94.7 | 0.0 | 5.3 | 123 | 1.5 | e |

Urea - Urine

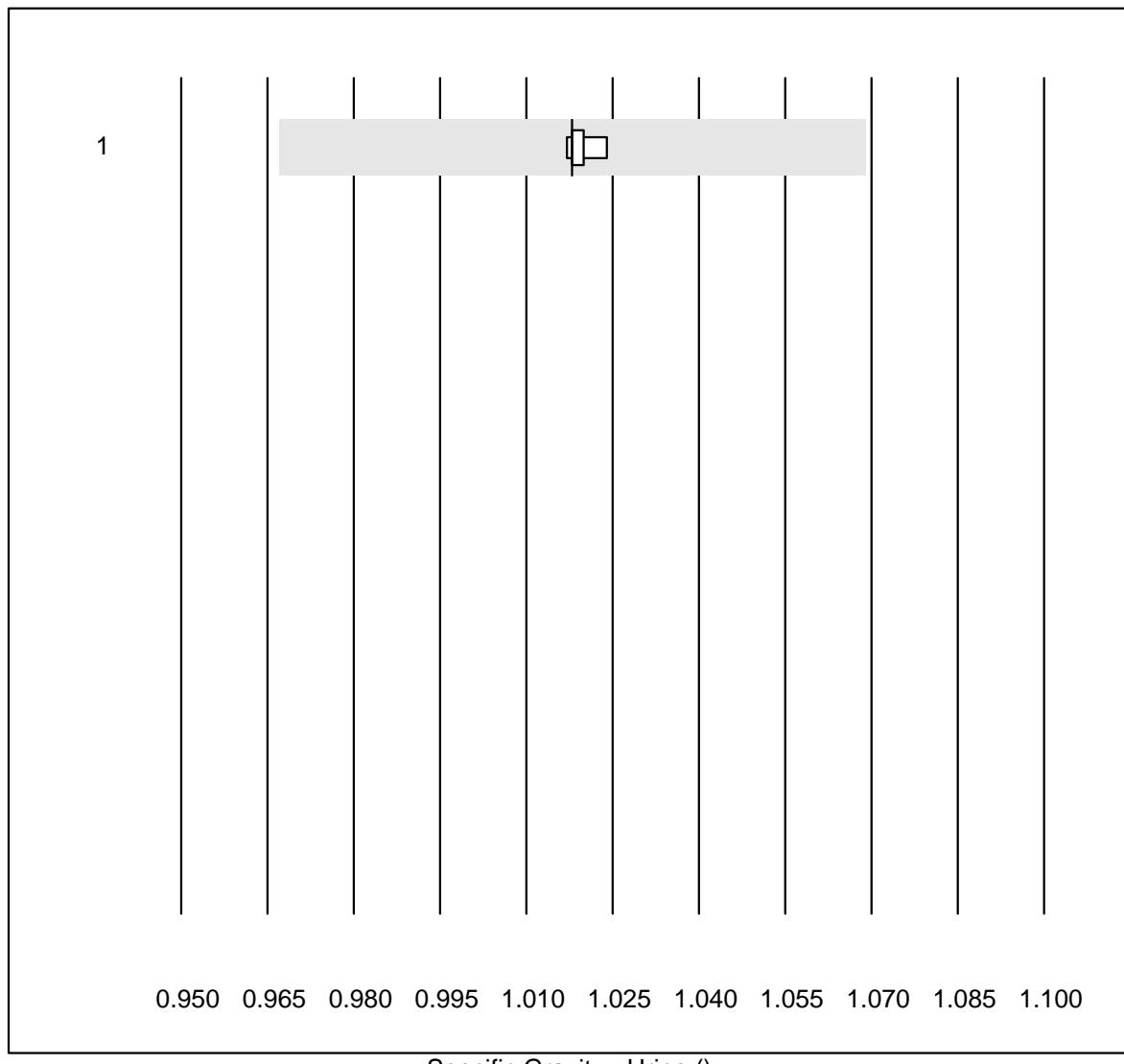
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Standard chemistry | 14 | 100.0 | 0.0 | 0.0 | 202 | 3.7 | e |

Uric Acid - Urine



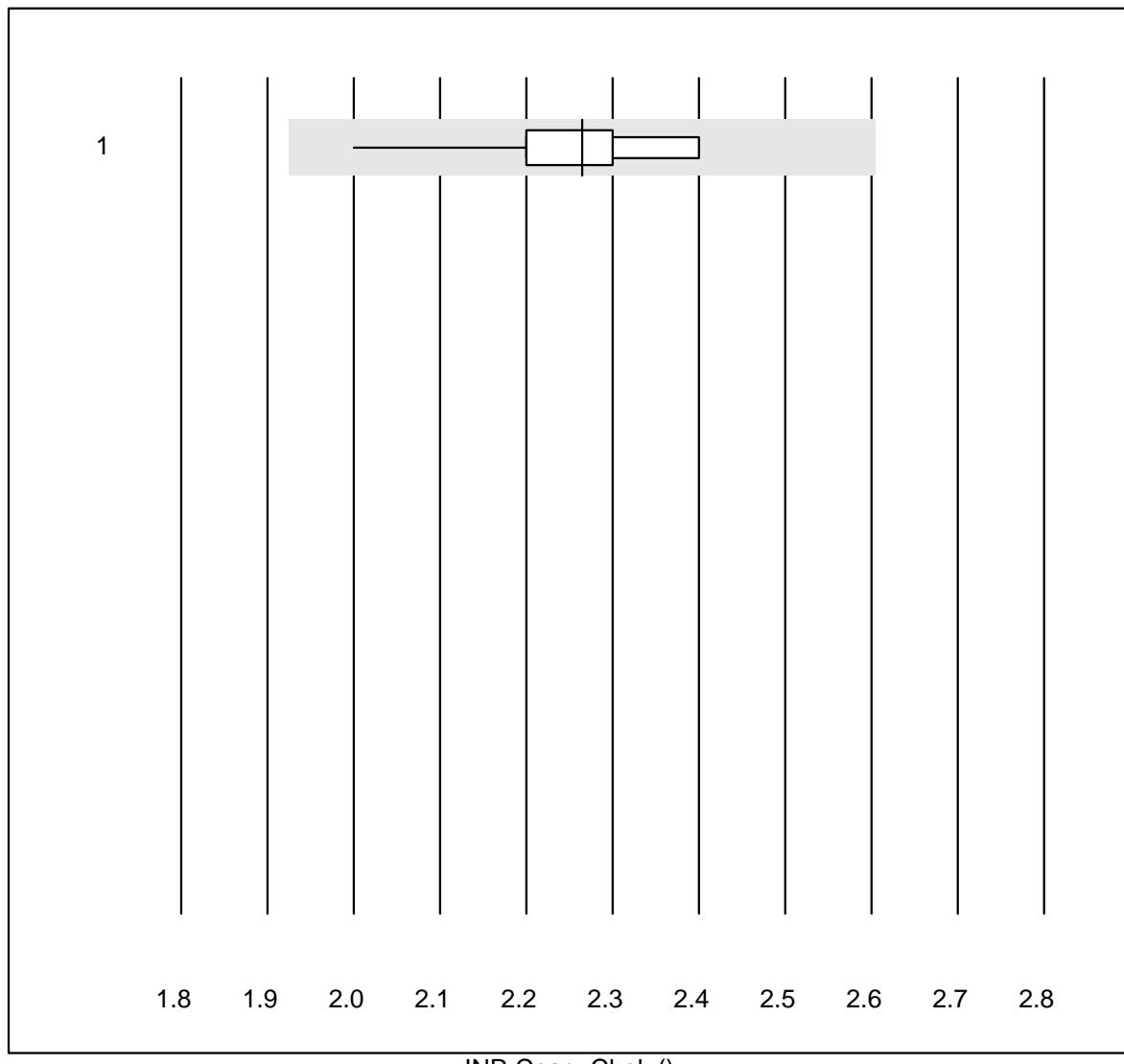
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Standard chemistry | 13 | 100.0 | 0.0 | 0.0 | 0.91 | 3.6 | e |

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 |

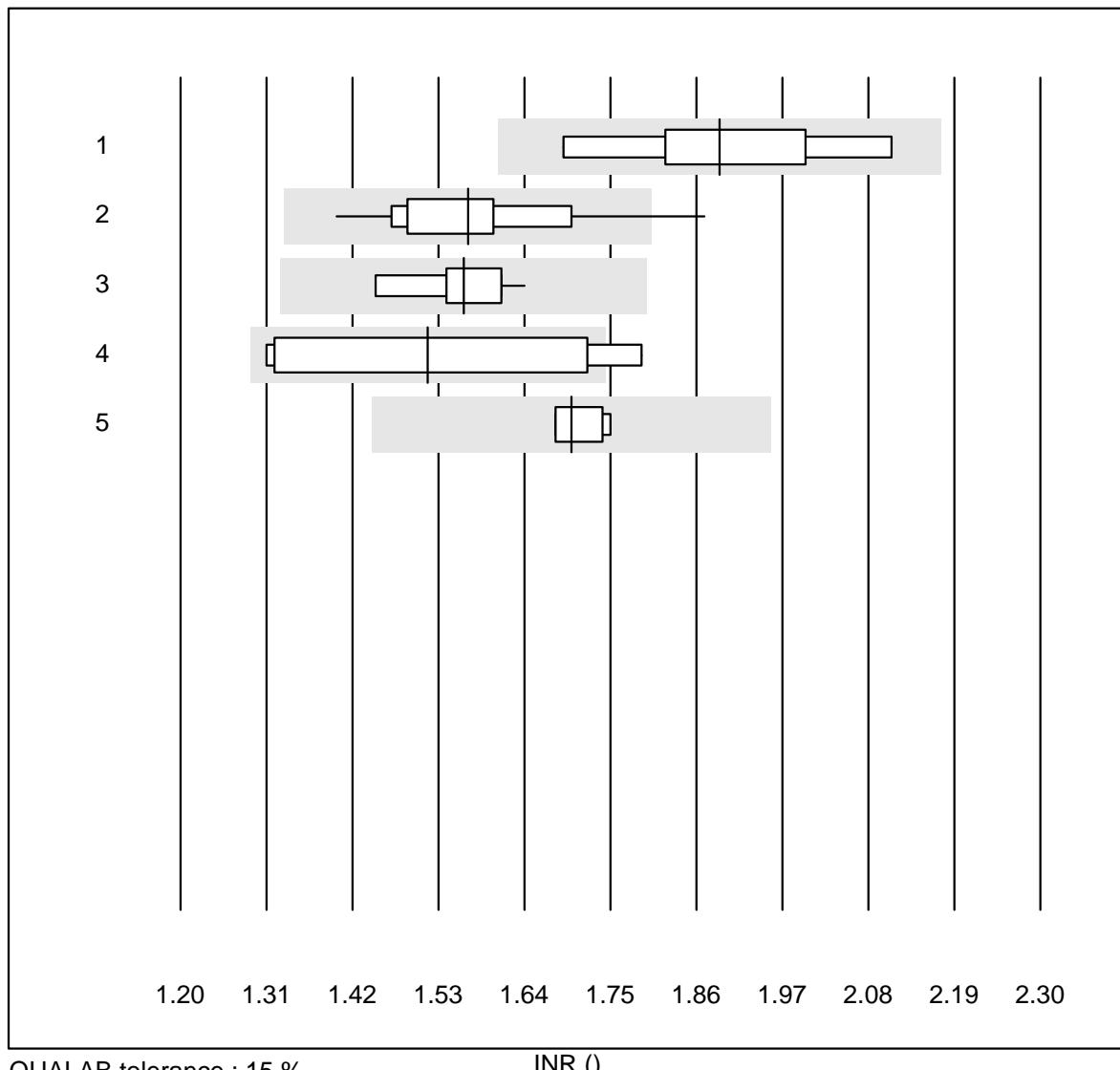
INR CoaguChek



| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | CoaguChek Pro II | 15 | 93.3 | 0.0 | 6.7 | 2.3 | 4.5 | e |

G1 Coagulation INR

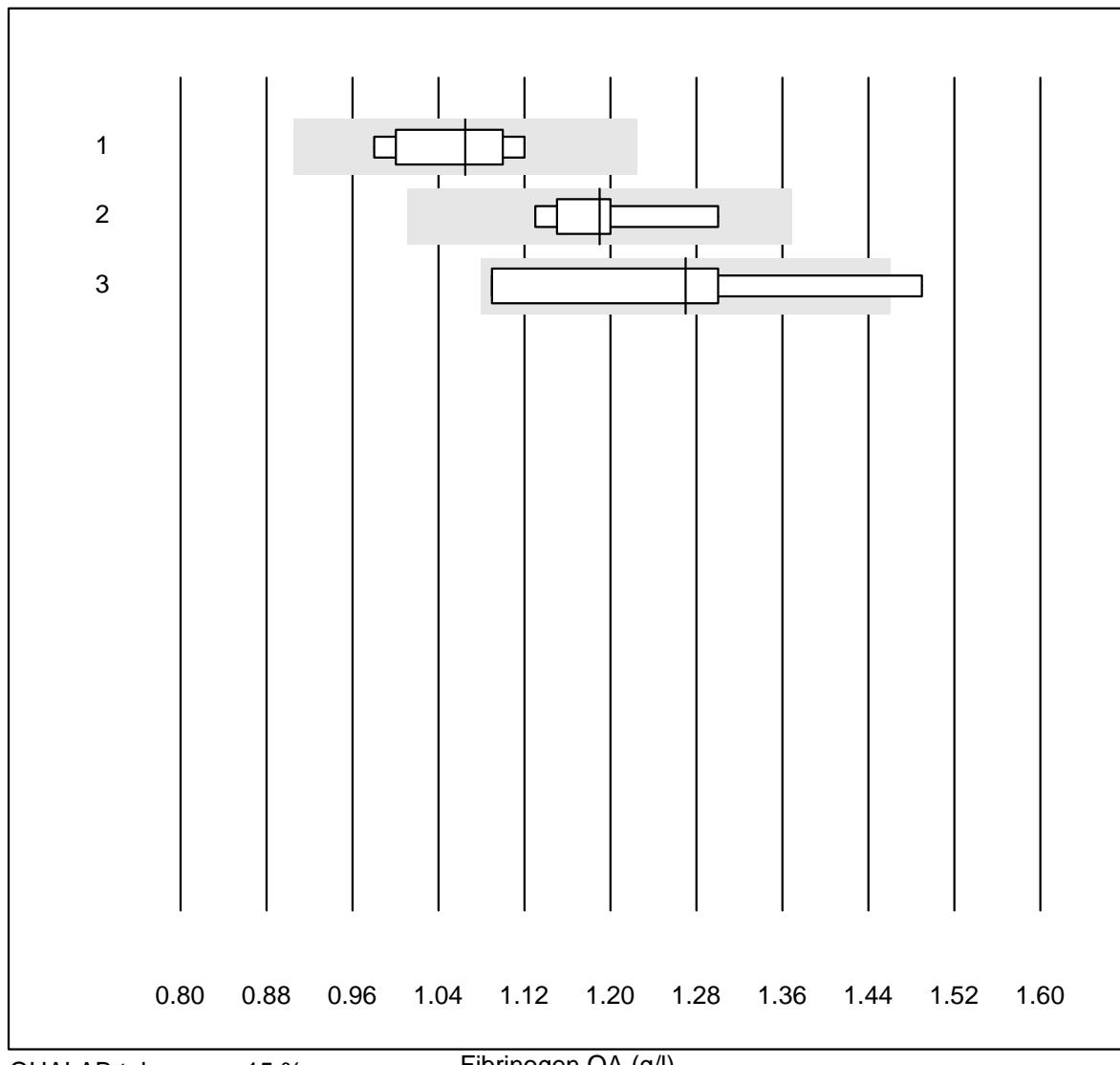
INR



QUALAB tolerance : 15 %

INR ()

| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|-------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 | Neoplastin Plus | 6 | 100.0 | 0.0 | 0.0 | 1.89 | 7.6 | e* |
| 2 | Innovin | 17 | 94.1 | 5.9 | 0.0 | 1.57 | 7.0 | e |
| 3 | Recombiplastin 2G | 10 | 100.0 | 0.0 | 0.0 | 1.56 | 3.7 | e |
| 4 | Eurolyser | 6 | 83.3 | 16.7 | 0.0 | 1.52 | 13.8 | a |
| 5 | Neoplastin R | 7 | 100.0 | 0.0 | 0.0 | 1.70 | 1.7 | e |

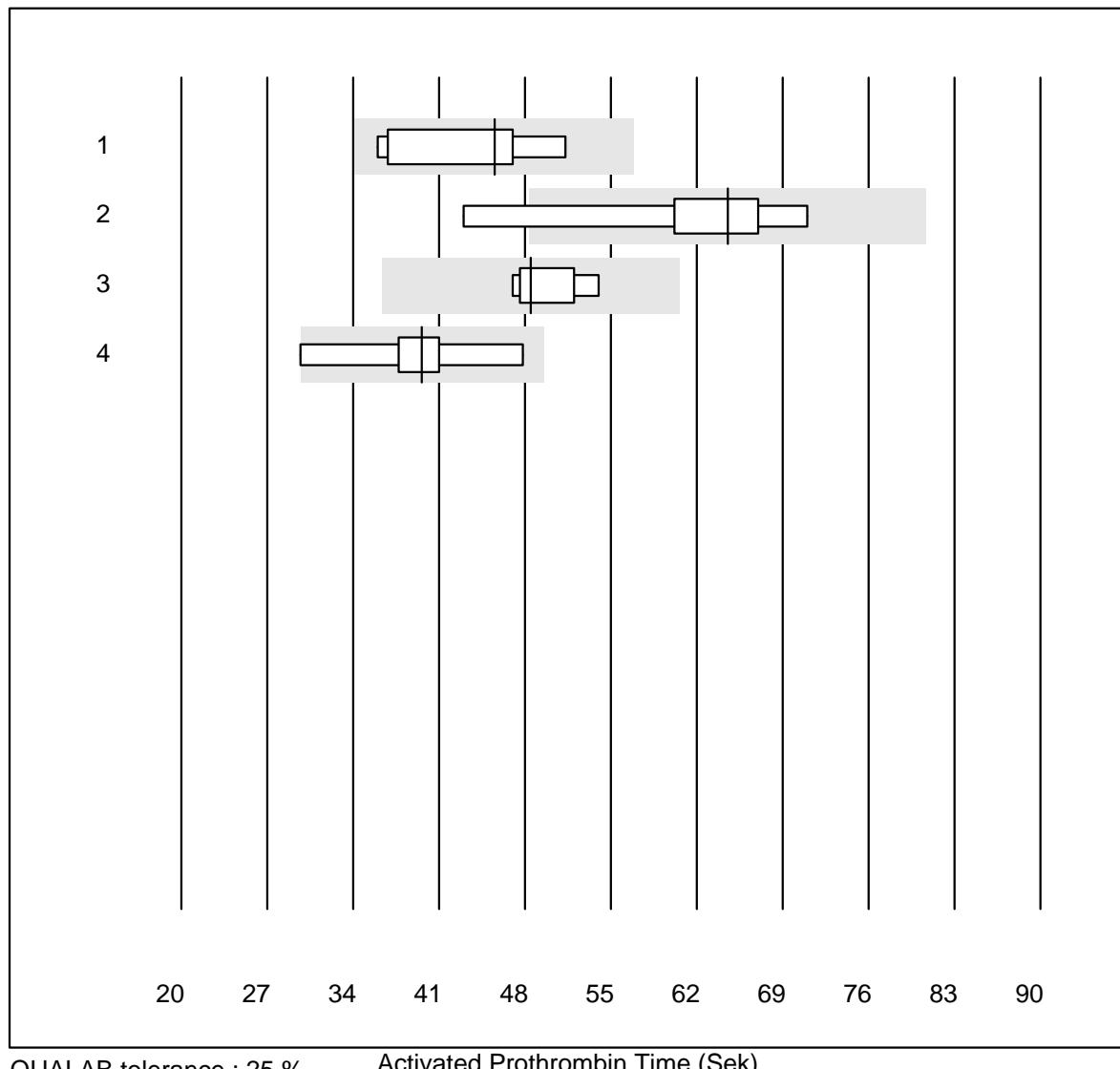
Fibrinogen OA

QUALAB tolerance : 15 %

Fibrinogen OA (g/l)

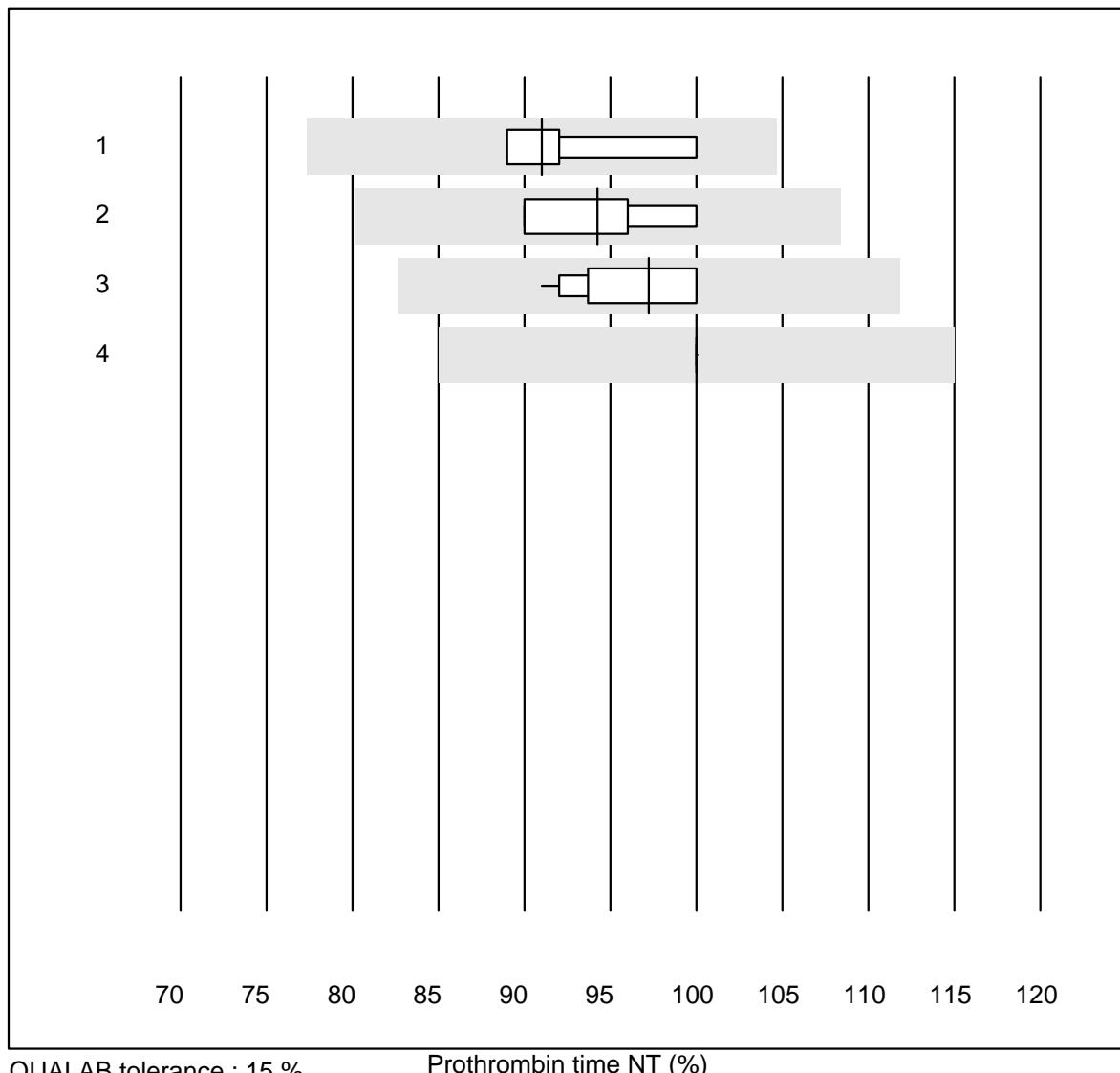
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|---------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 Siemens Thrombin | 6 | 100.0 | 0.0 | 0.0 | 1.07 | 5.6 | e* |
| 2 Stago/STA | 9 | 100.0 | 0.0 | 0.0 | 1.19 | 5.0 | e |
| 3 Fibrinogen Q.F.A. | 4 | 75.0 | 25.0 | 0.0 | 1.27 | 12.6 | a |

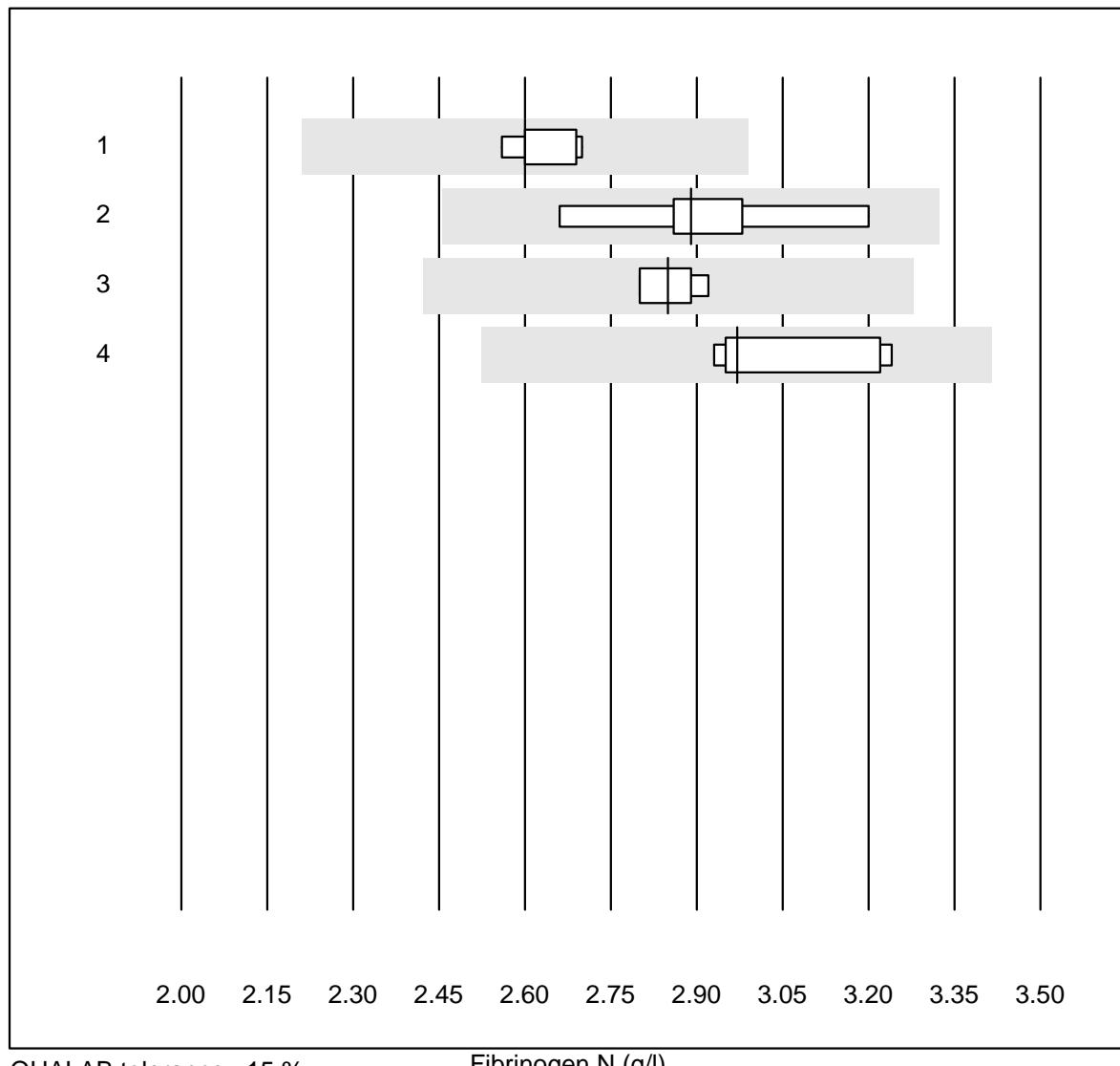
Activated Prothrombin Time



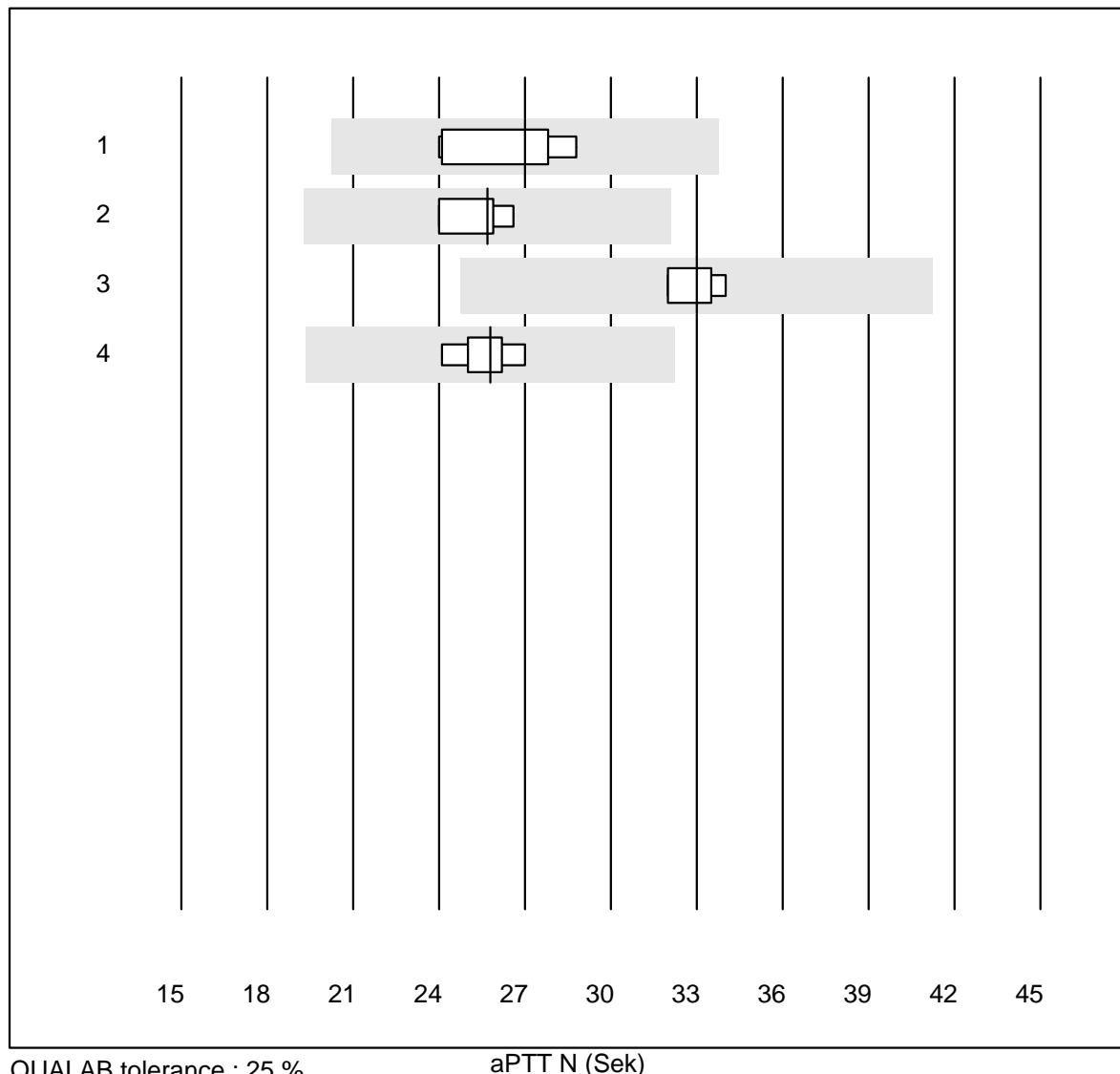
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 Actin FS | 8 | 100.0 | 0.0 | 0.0 | 45.5 | 12.1 | e* |
| 2 Pathromtin SL | 5 | 80.0 | 20.0 | 0.0 | 64.5 | 17.8 | e* |
| 3 Stago/STA | 6 | 100.0 | 0.0 | 0.0 | 48.5 | 5.6 | e |
| 4 aPTT-SP | 8 | 87.5 | 12.5 | 0.0 | 39.6 | 13.2 | e* |

Prothrombin time NT

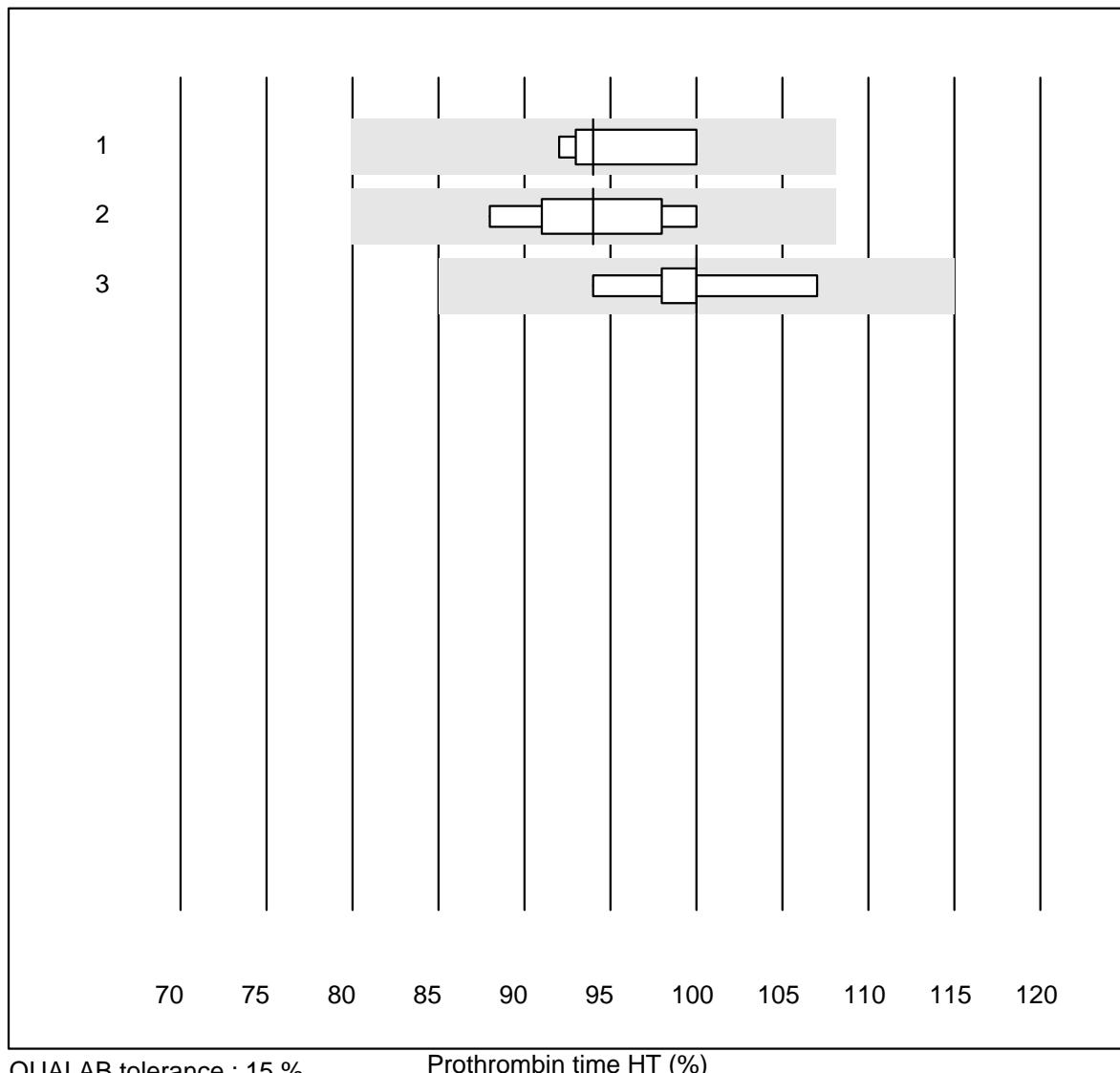


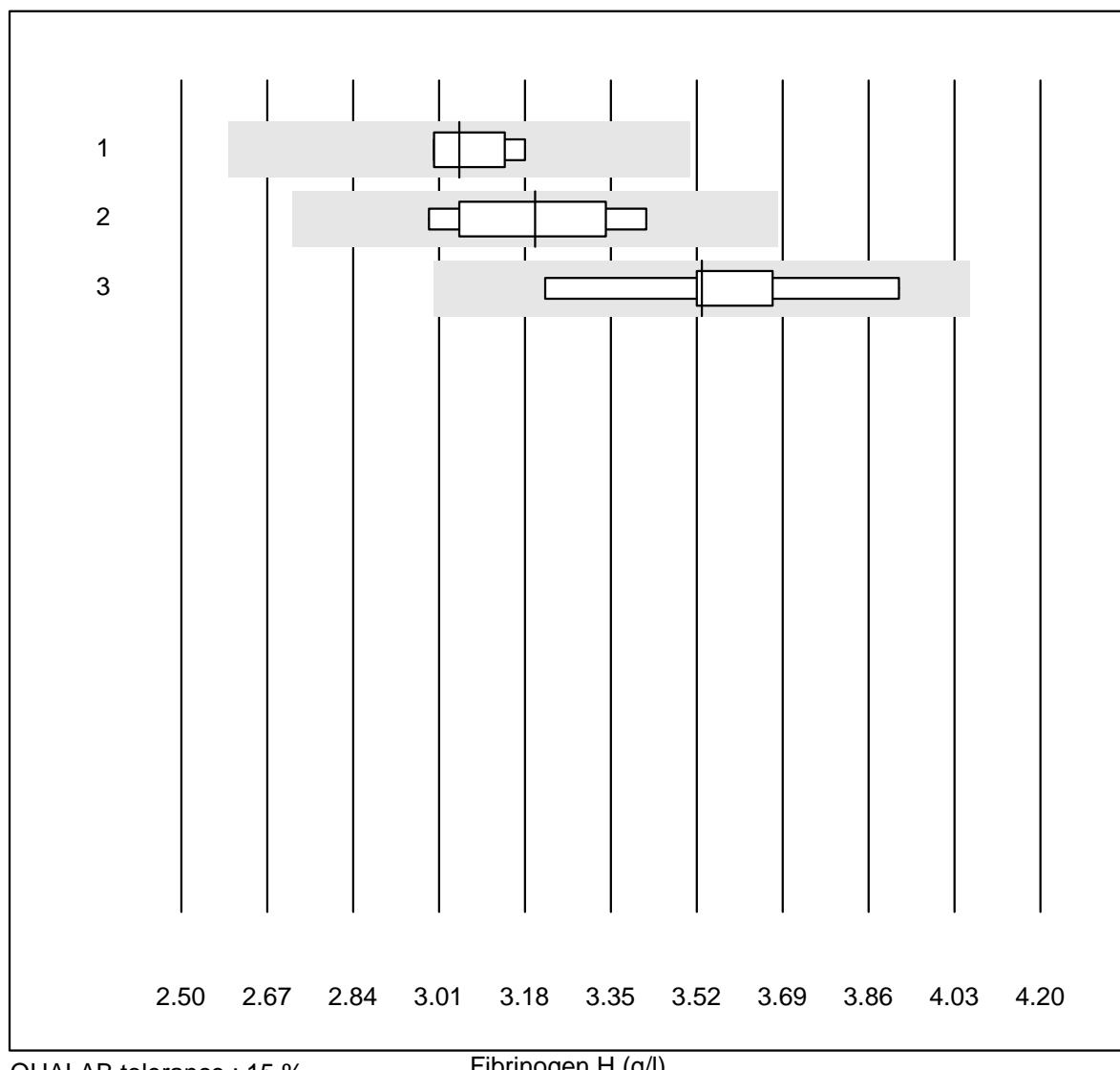
Fibrinogen N

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|---------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Siemens Thrombin | 7 | 100.0 | 0.0 | 0.0 | 2.60 | 2.1 | e |
| 2 Stago/STA | 9 | 100.0 | 0.0 | 0.0 | 2.89 | 5.3 | e |
| 3 Fibrinogen Q.F.A. | 4 | 100.0 | 0.0 | 0.0 | 2.85 | 2.1 | e |
| 4 Fib Clauss (IL) | 5 | 100.0 | 0.0 | 0.0 | 2.97 | 5.0 | e* |

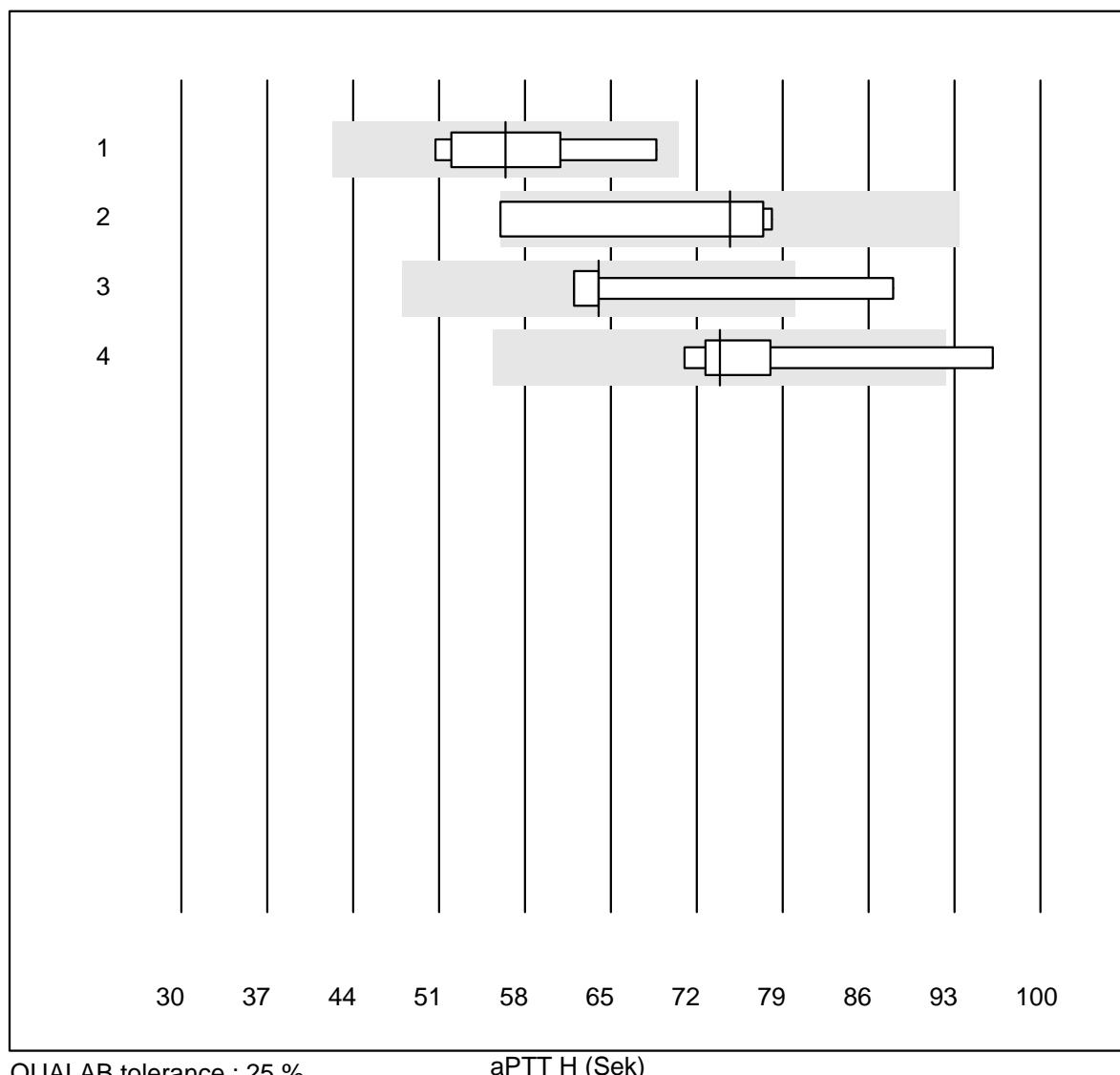
aPTT N

Prothrombin time HT

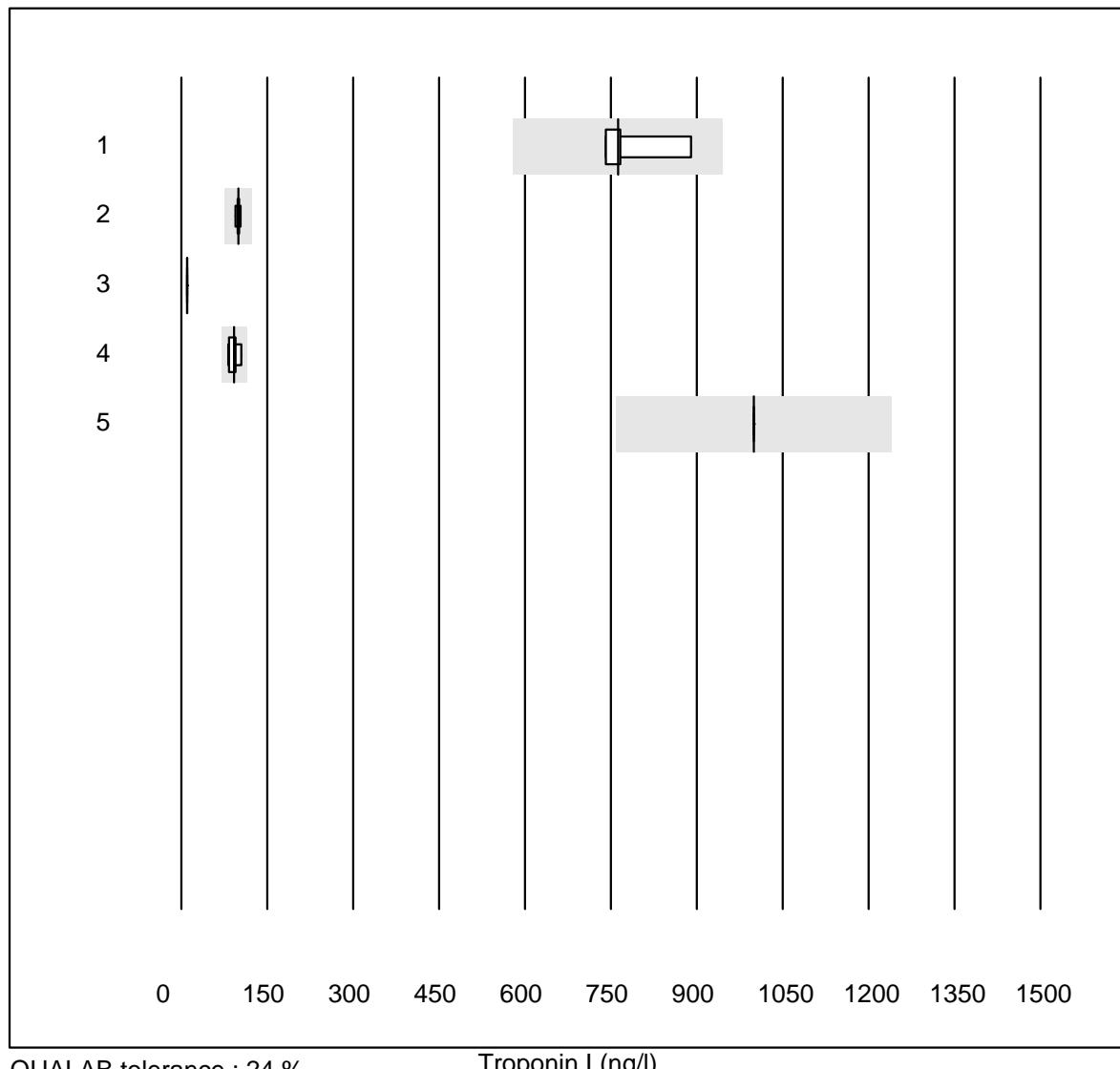


Fibrinogen H

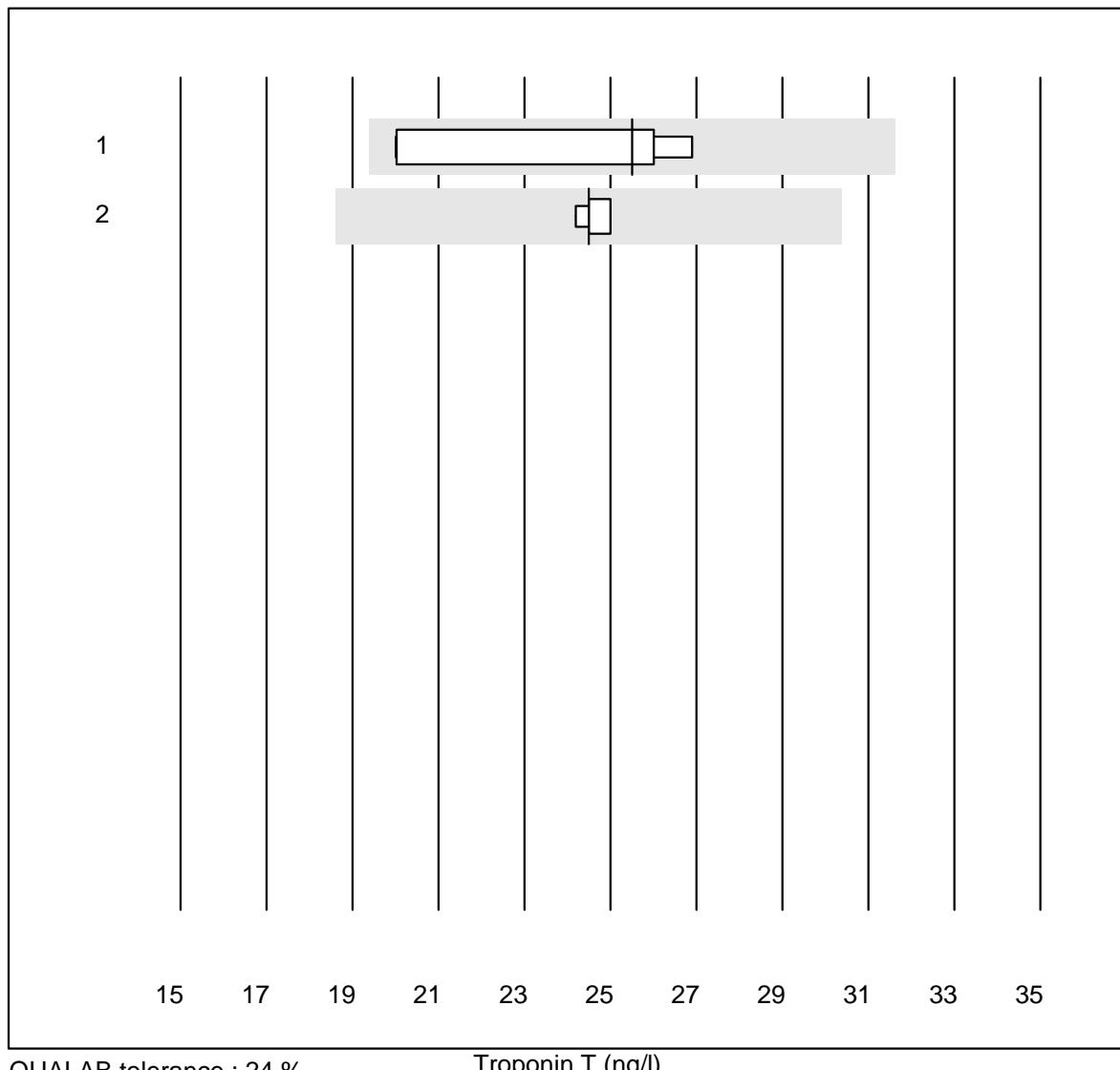
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Siemens Thrombin | 6 | 100.0 | 0.0 | 0.0 | 3.05 | 2.6 | e |
| 2 Stago/STA | 7 | 100.0 | 0.0 | 0.0 | 3.20 | 4.9 | e* |
| 3 Fib Clauss (IL) | 5 | 100.0 | 0.0 | 0.0 | 3.53 | 7.1 | e* |

aPTT H

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 Actin FS | 6 | 100.0 | 0.0 | 0.0 | 56.4 | 11.7 | e* |
| 2 Other methods | 4 | 75.0 | 25.0 | 0.0 | 74.7 | 14.5 | e* |
| 3 Stago/STA | 5 | 60.0 | 20.0 | 20.0 | 64.0 | 17.8 | e* |
| 4 aPTT-SP | 6 | 83.3 | 16.7 | 0.0 | 73.9 | 12.1 | e* |

Troponin I

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|------------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Vidas hs | 4 | 100.0 | 0.0 | 0.0 | 762.2 | 8.6 | e* |
| 2 Architect High Sensi | 6 | 100.0 | 0.0 | 0.0 | 99.6 | 3.0 | e |
| 3 AQT 90 FLEX | 6 | 100.0 | 0.0 | 0.0 | 10.0 | 0.0 | e |
| 4 ADVIA Centaur XP/CP | 5 | 100.0 | 0.0 | 0.0 | 92.4 | 9.8 | e* |
| 5 Eurolyser | 14 | 100.0 | 0.0 | 0.0 | 1000.0 | 0.0 | e |

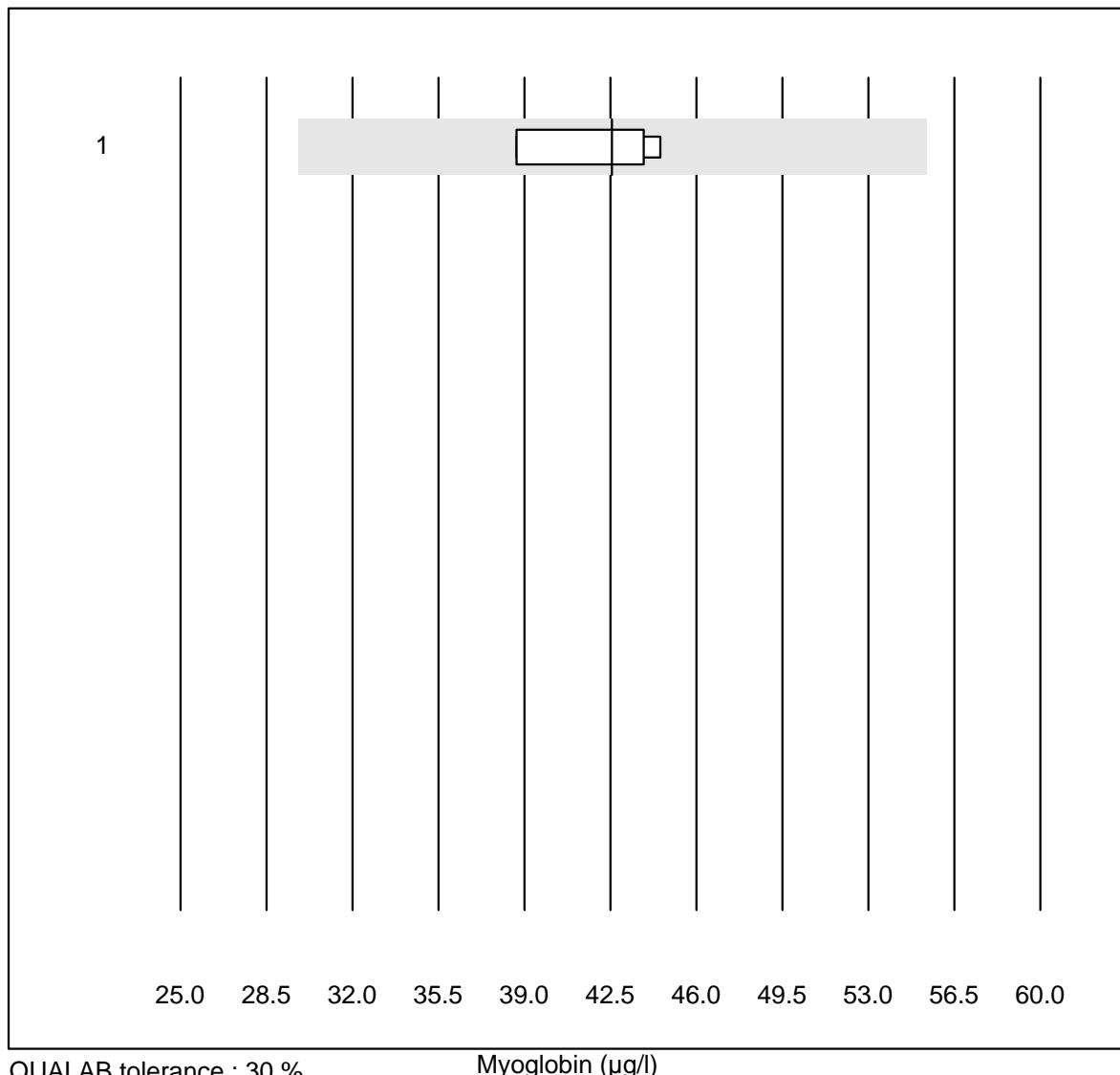
Troponin T

QUALAB tolerance : 24 %

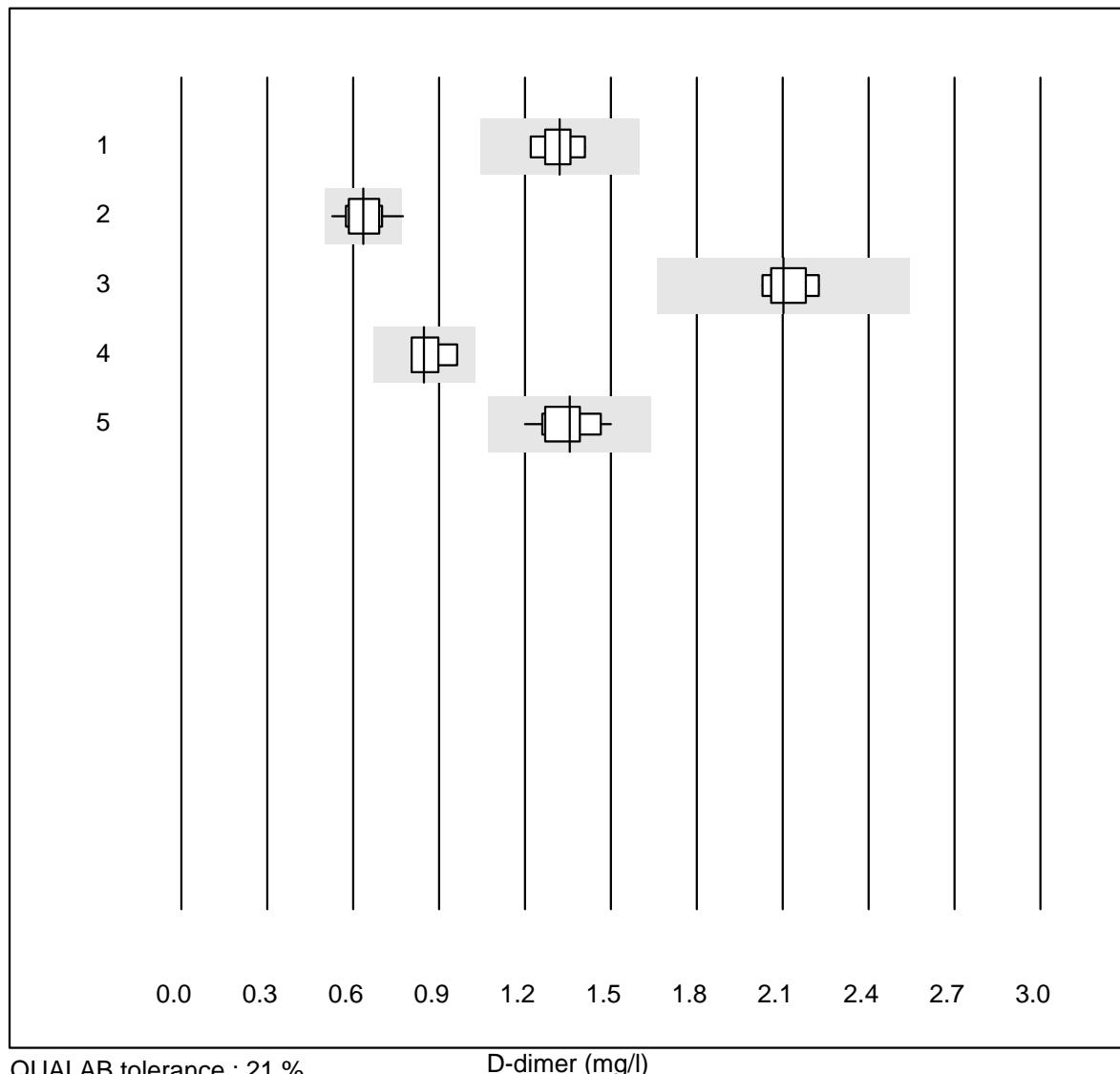
Troponin T (ng/l)

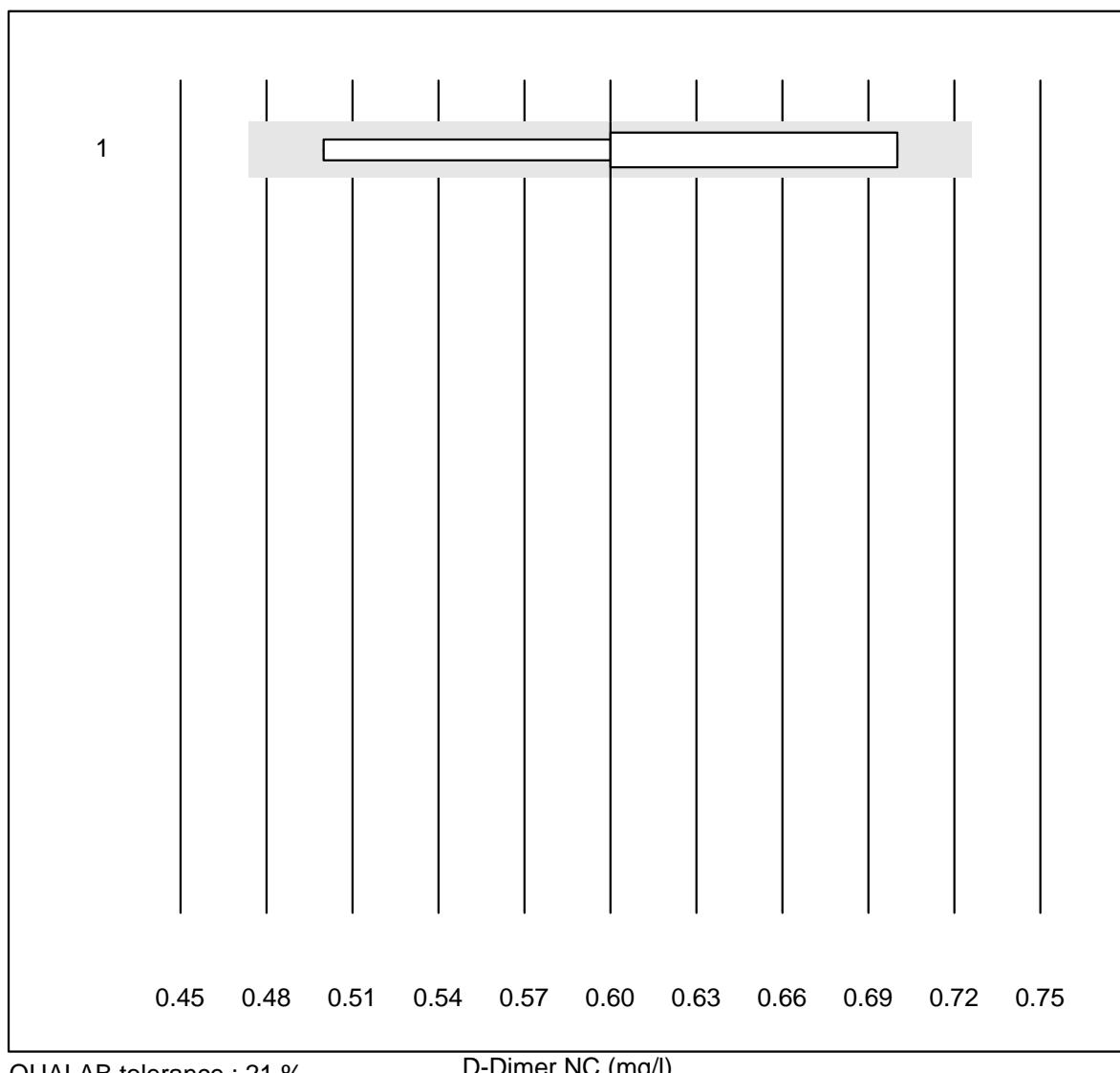
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 Cobas hs | 5 | 100.0 | 0.0 | 0.0 | 25.50 | 14.3 | e* |
| 2 Cobas hs STAT | 6 | 100.0 | 0.0 | 0.0 | 24.50 | 1.3 | e |

Myoglobin



| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|-------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Cobas E / Elecsys | 4 | 100.0 | 0.0 | 0.0 | 42.6 | 6.3 | e |

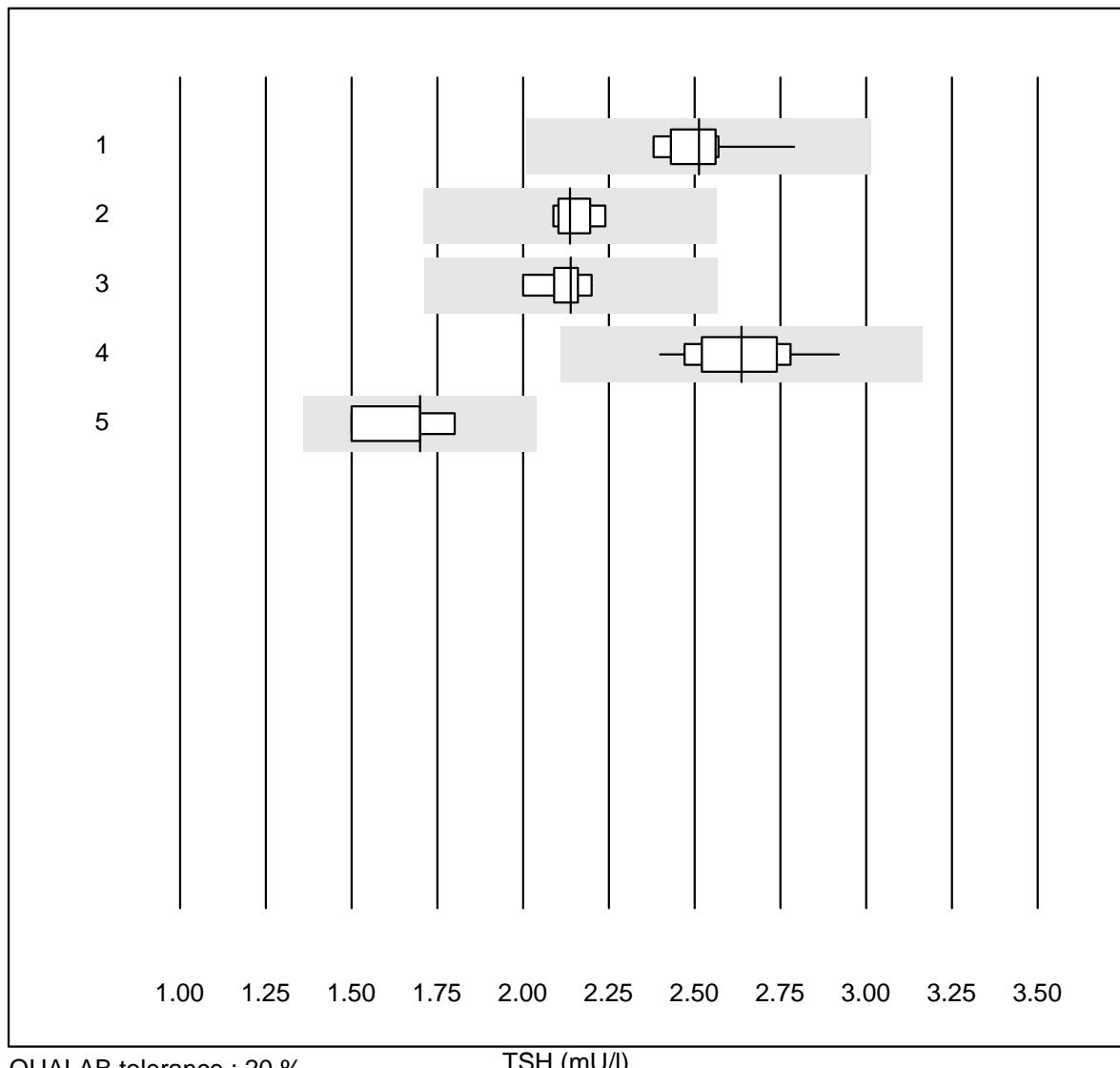
D-dimer

D-Dimer NC

| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|----------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 | NycoCard | 25 | 64.0 | 0.0 | 36.0 | 0.60 | 11.7 | e |

K6 Hormones

TSH



| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Cobas E / Elecsys | 10 | 100.0 | 0.0 | 0.0 | 2.5 | 4.7 | e |
| 2 ADVIA Centaur XP/CP | 6 | 100.0 | 0.0 | 0.0 | 2.1 | 2.7 | e |
| 3 Architect | 8 | 100.0 | 0.0 | 0.0 | 2.1 | 2.9 | e |
| 4 Vidas | 13 | 100.0 | 0.0 | 0.0 | 2.6 | 5.6 | e |
| 5 Qualigen | 5 | 80.0 | 0.0 | 20.0 | 1.7 | 7.8 | e* |

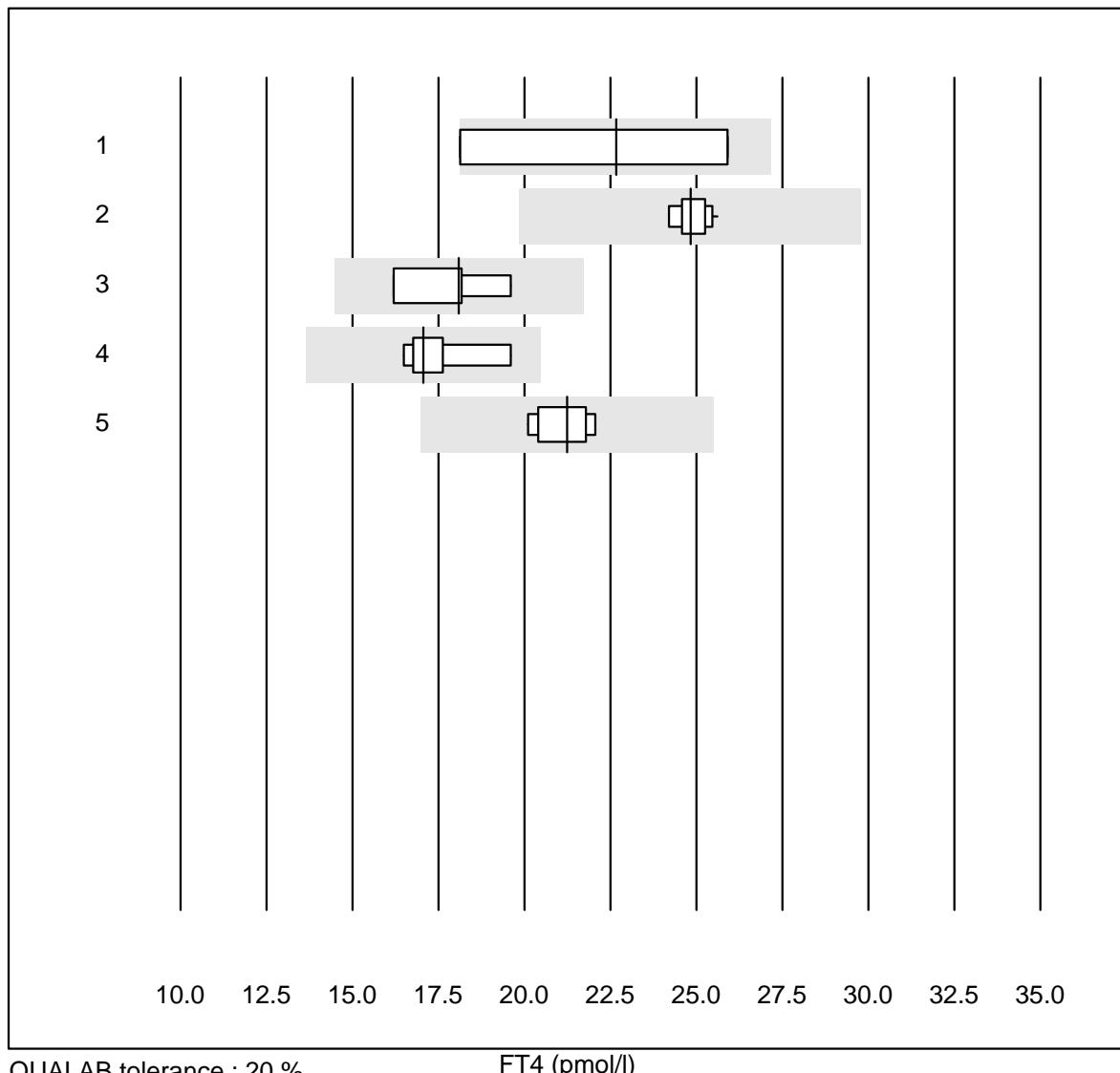
K6 Hormones

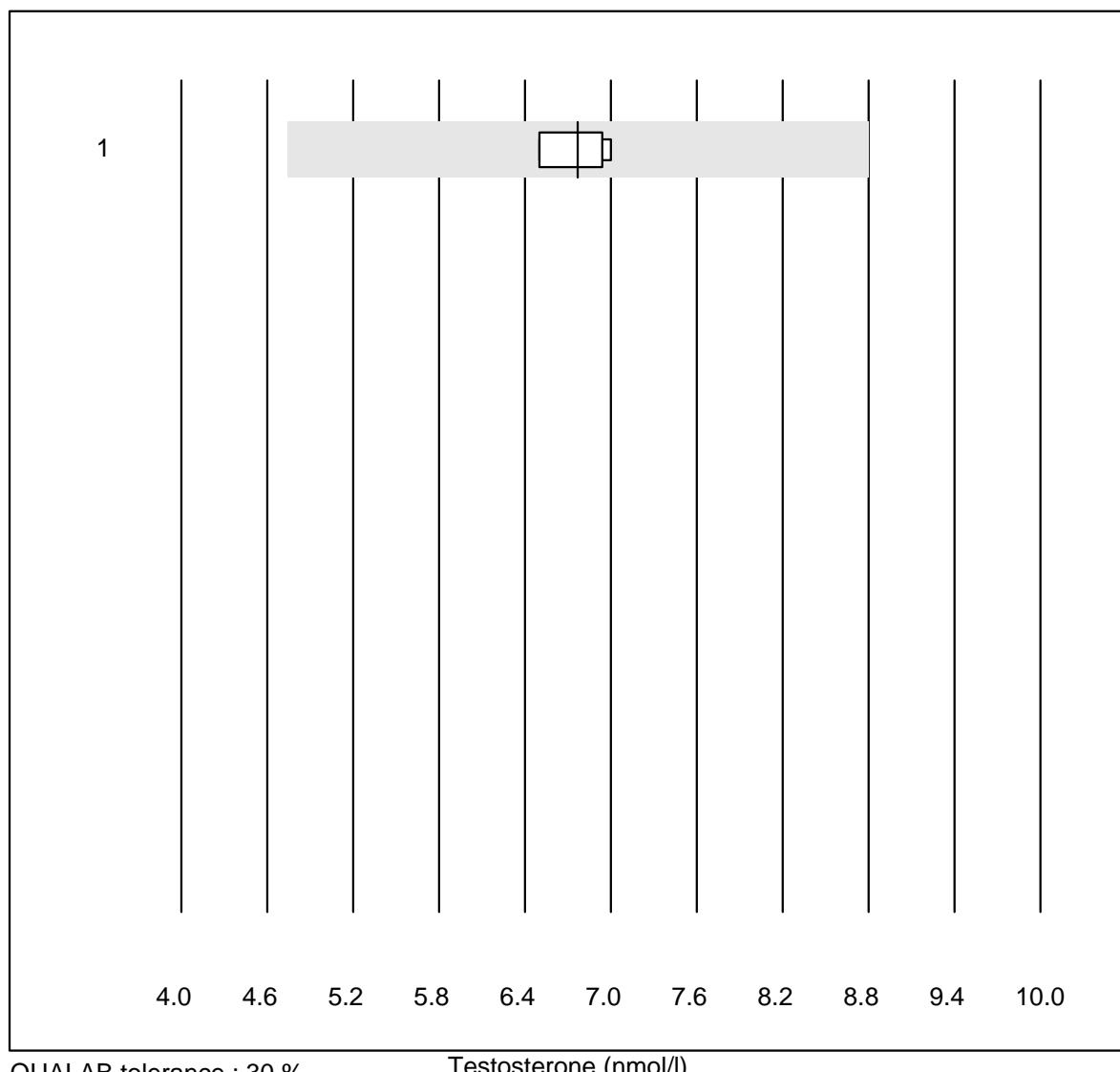
FT3



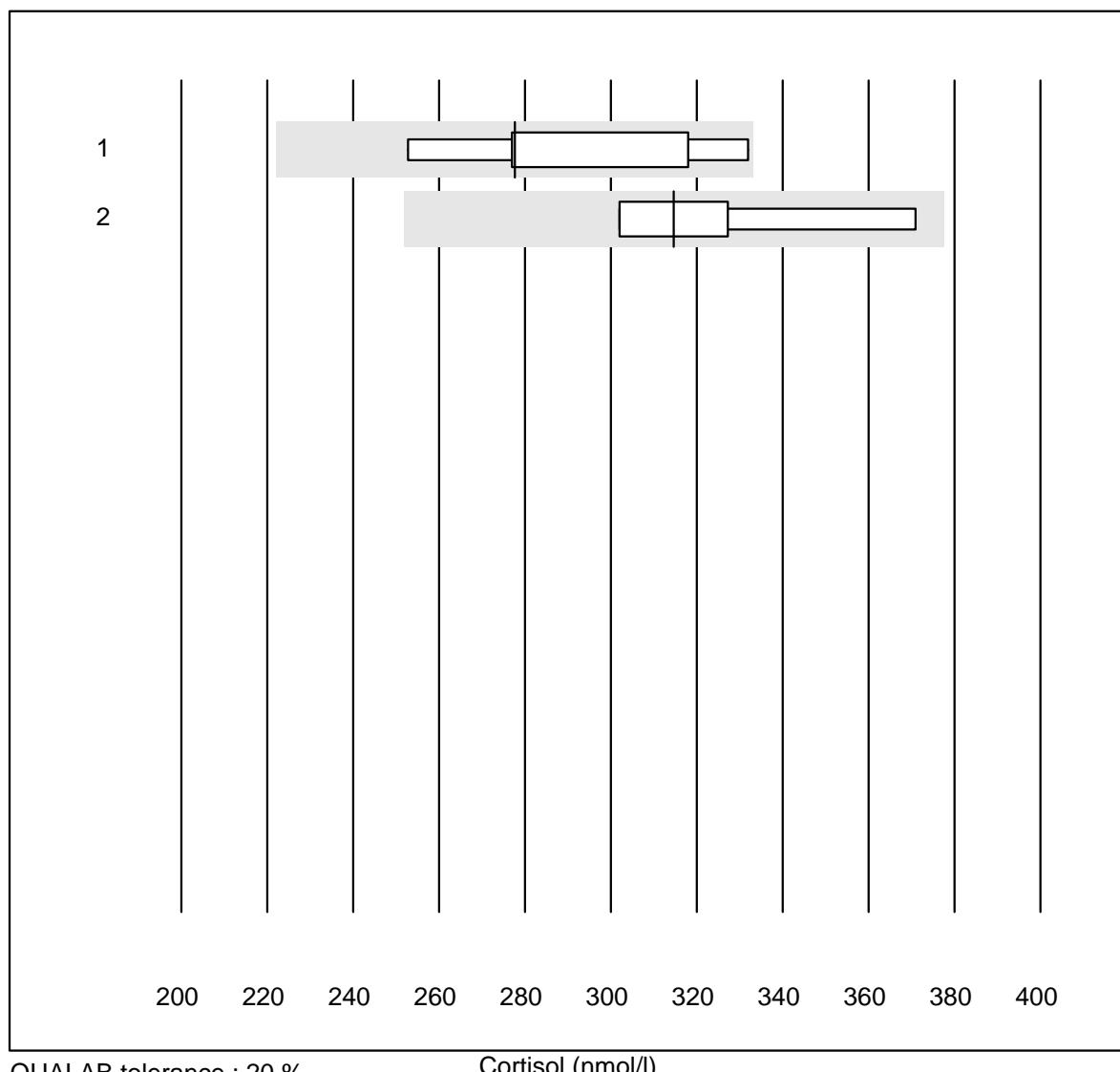
K6 Hormones

FT4



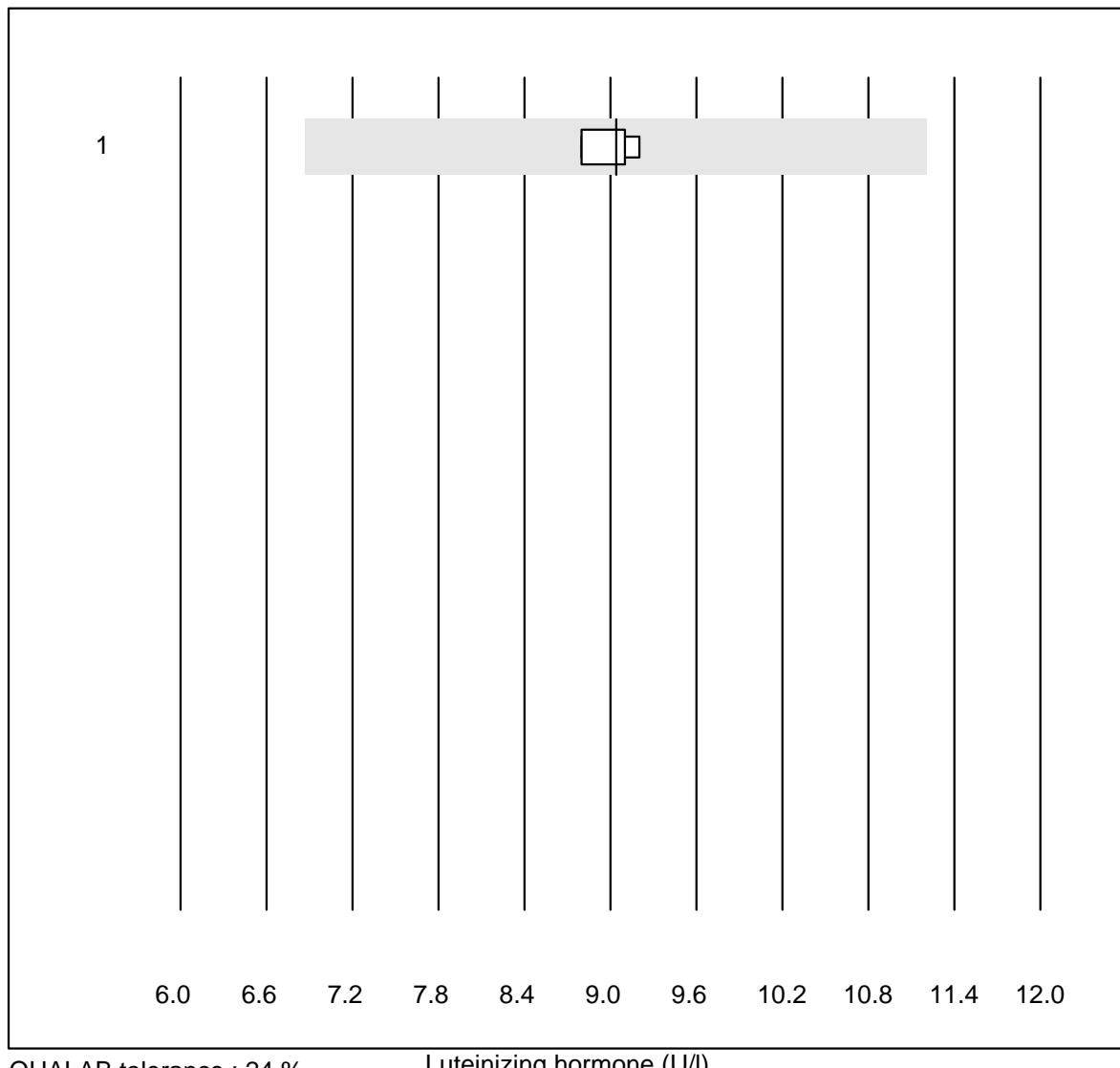
Testosterone

| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | ADVIA Centaur XP/CP | 4 | 100.0 | 0.0 | 0.0 | 7 | 3.7 | e |

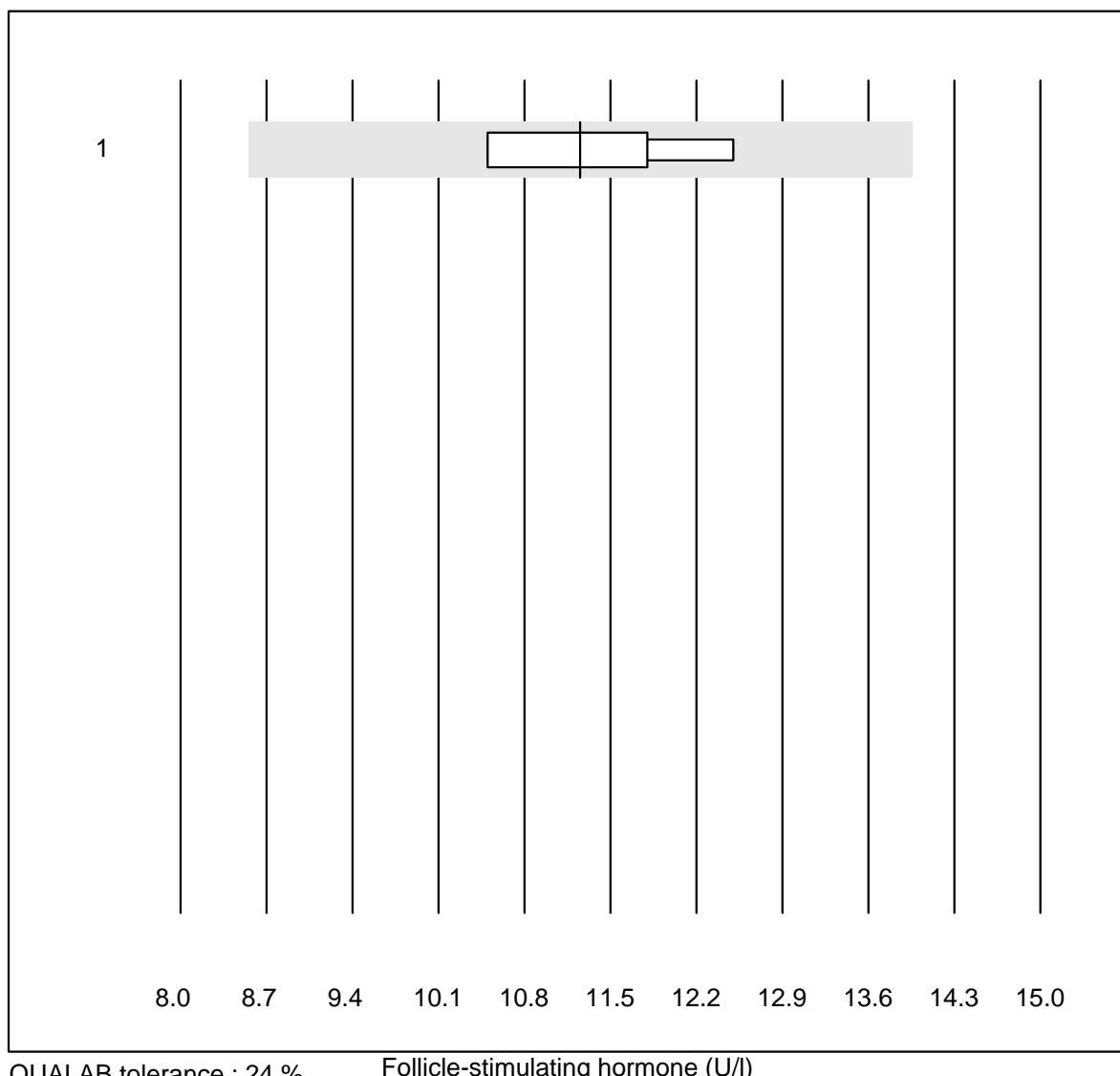
Cortisol

| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 | Cobas E / Elecsys | 6 | 100.0 | 0.0 | 0.0 | 278 | 10.3 | e* |
| 2 | ADVIA Centaur XP/CP | 4 | 100.0 | 0.0 | 0.0 | 315 | 10.0 | e* |

Luteinizing hormone

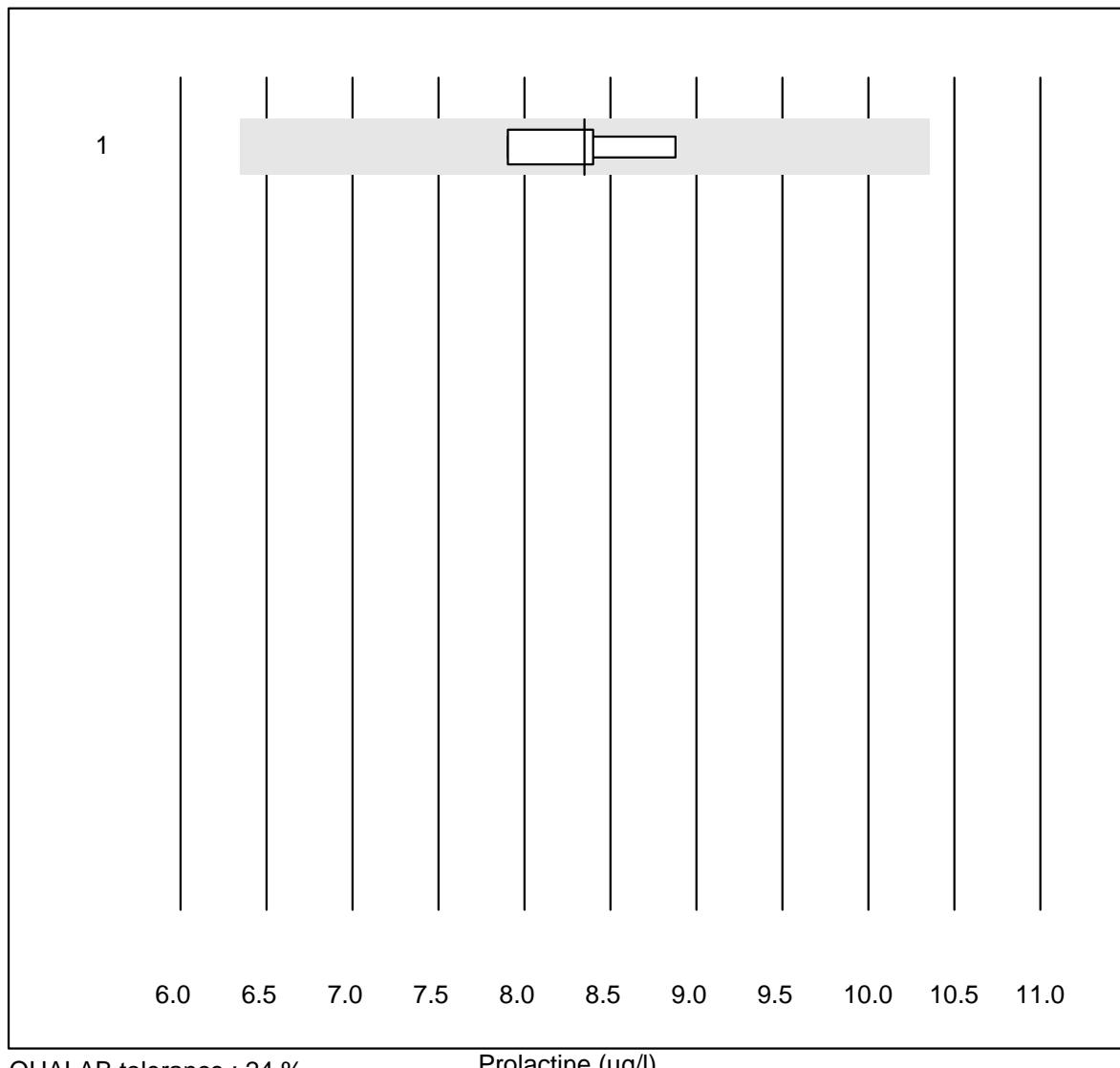


Follicle-stimulating hormone



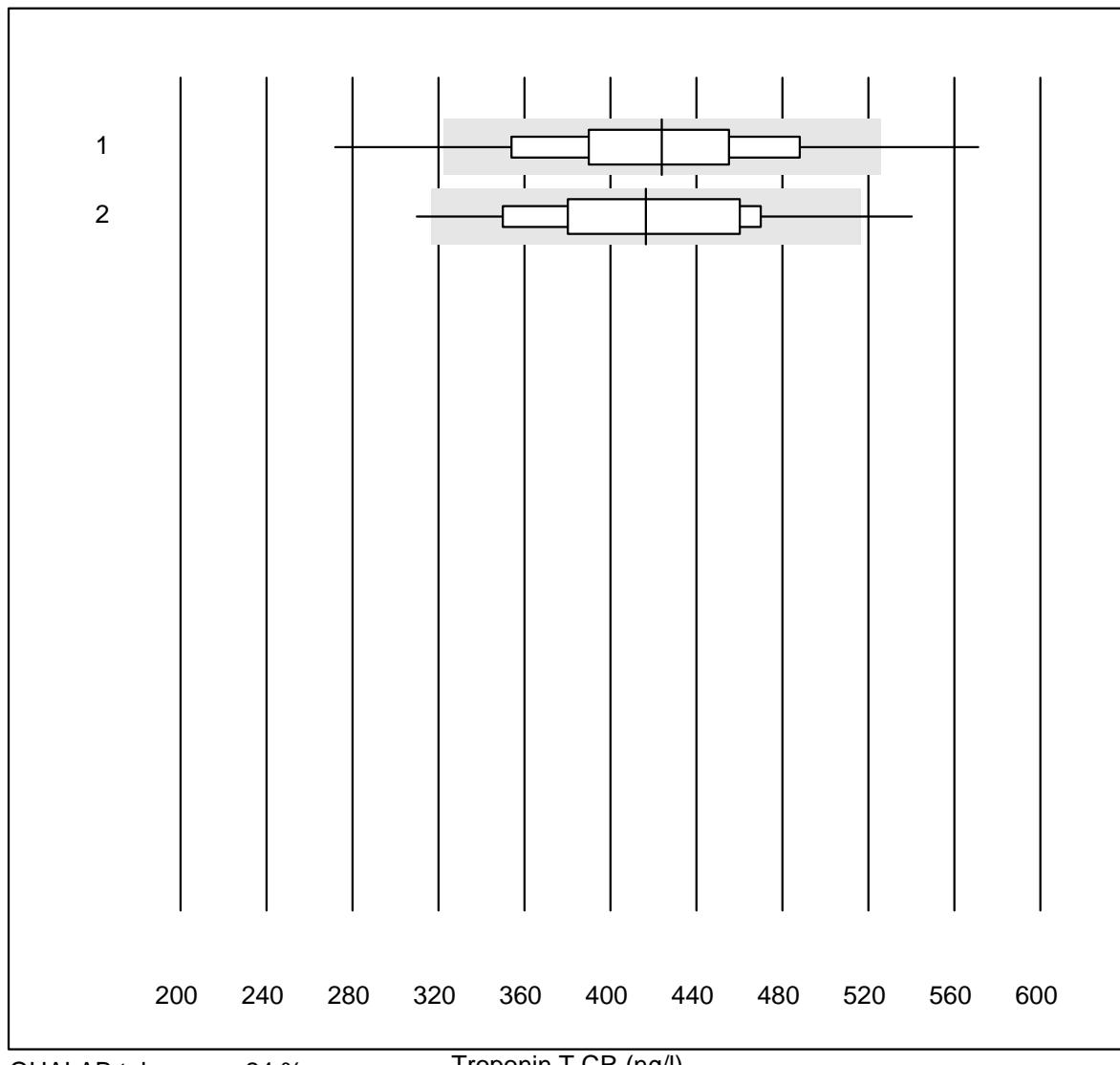
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | ADVIA Centaur XP/CP | 4 | 100.0 | 0.0 | 0.0 | 11.3 | 8.3 | e* |

Prolactine



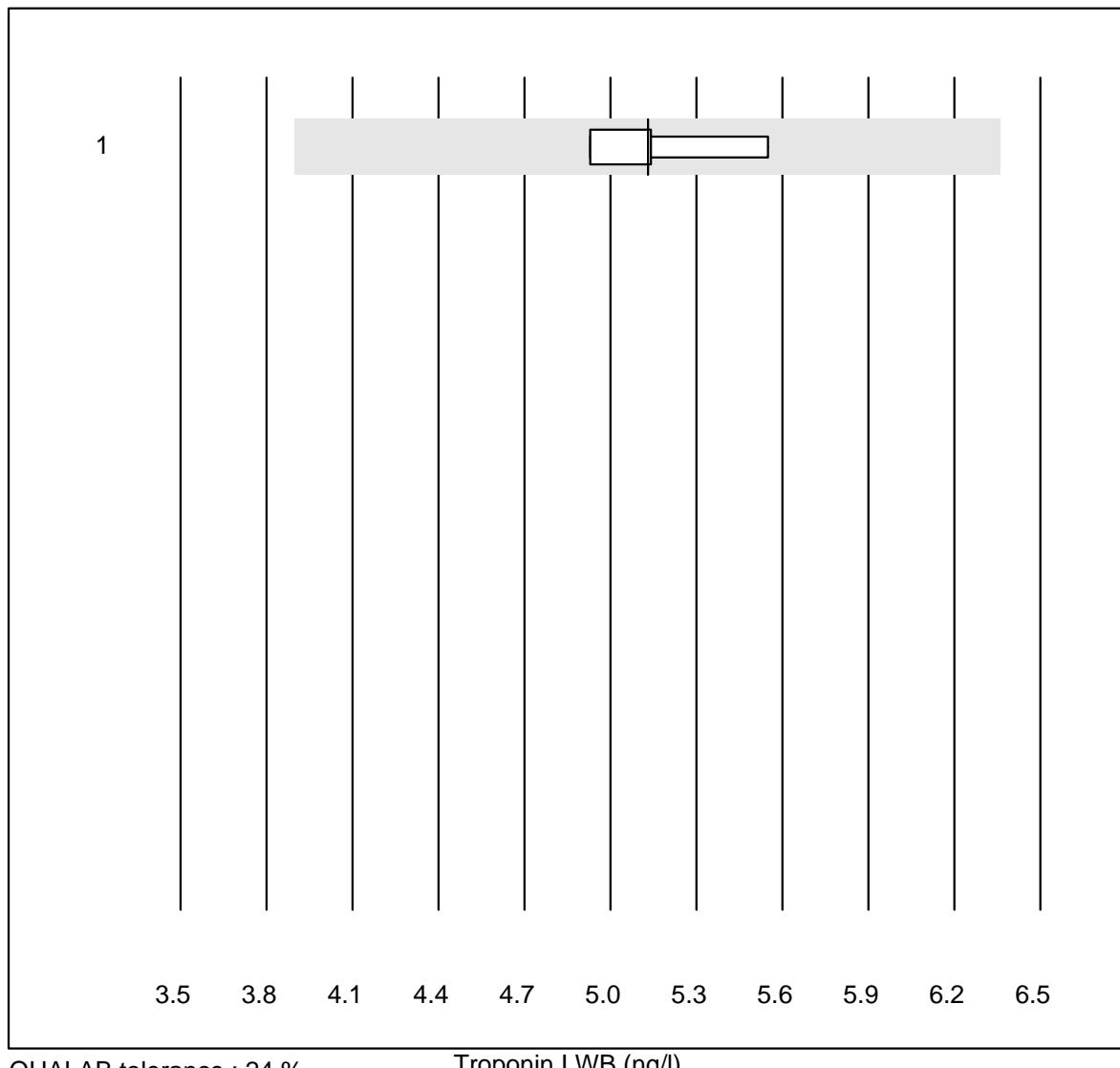
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | ADVIA Centaur XP/CP | 4 | 100.0 | 0.0 | 0.0 | 8.3 | 4.8 | e |

Troponin T CR

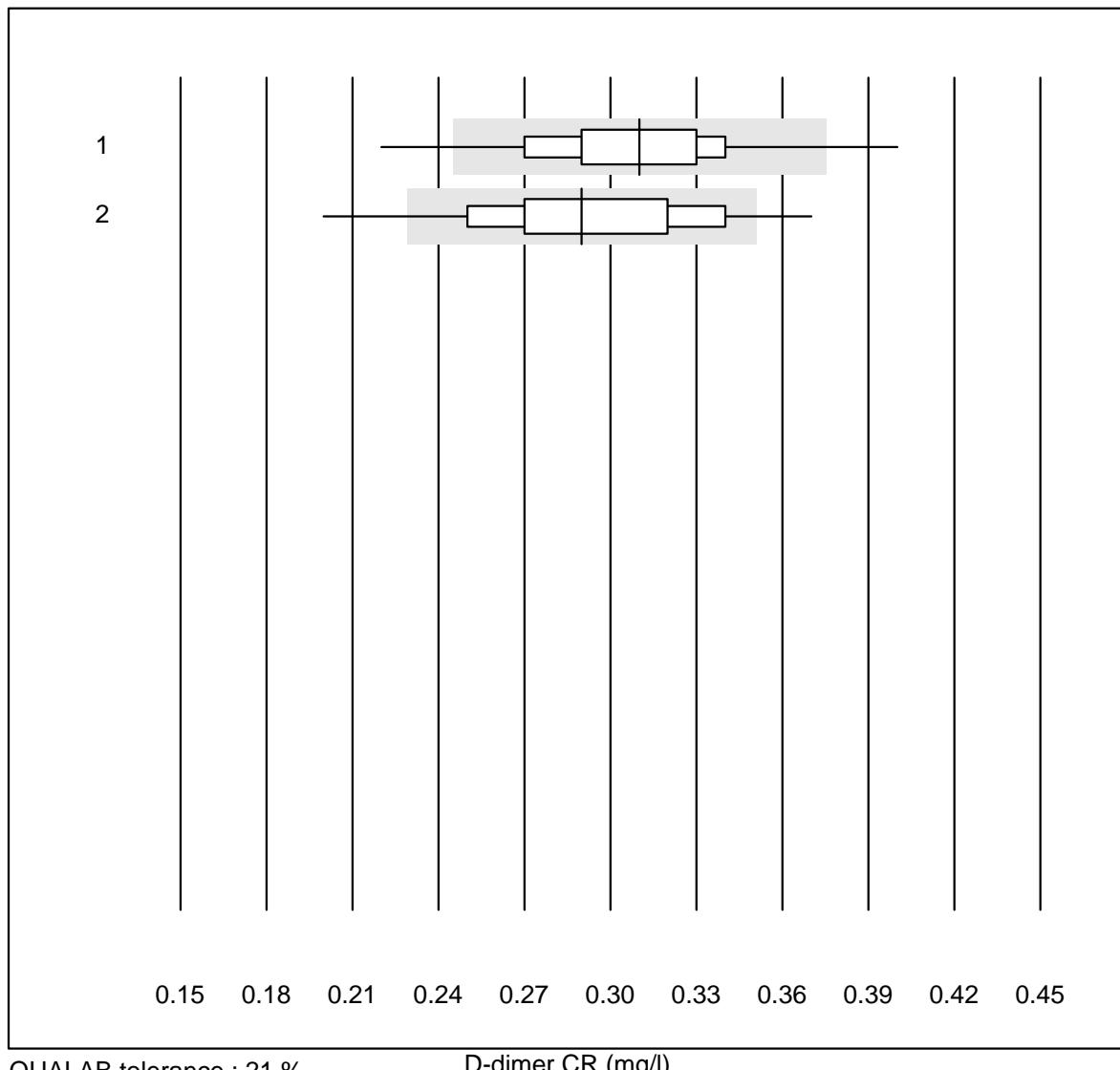


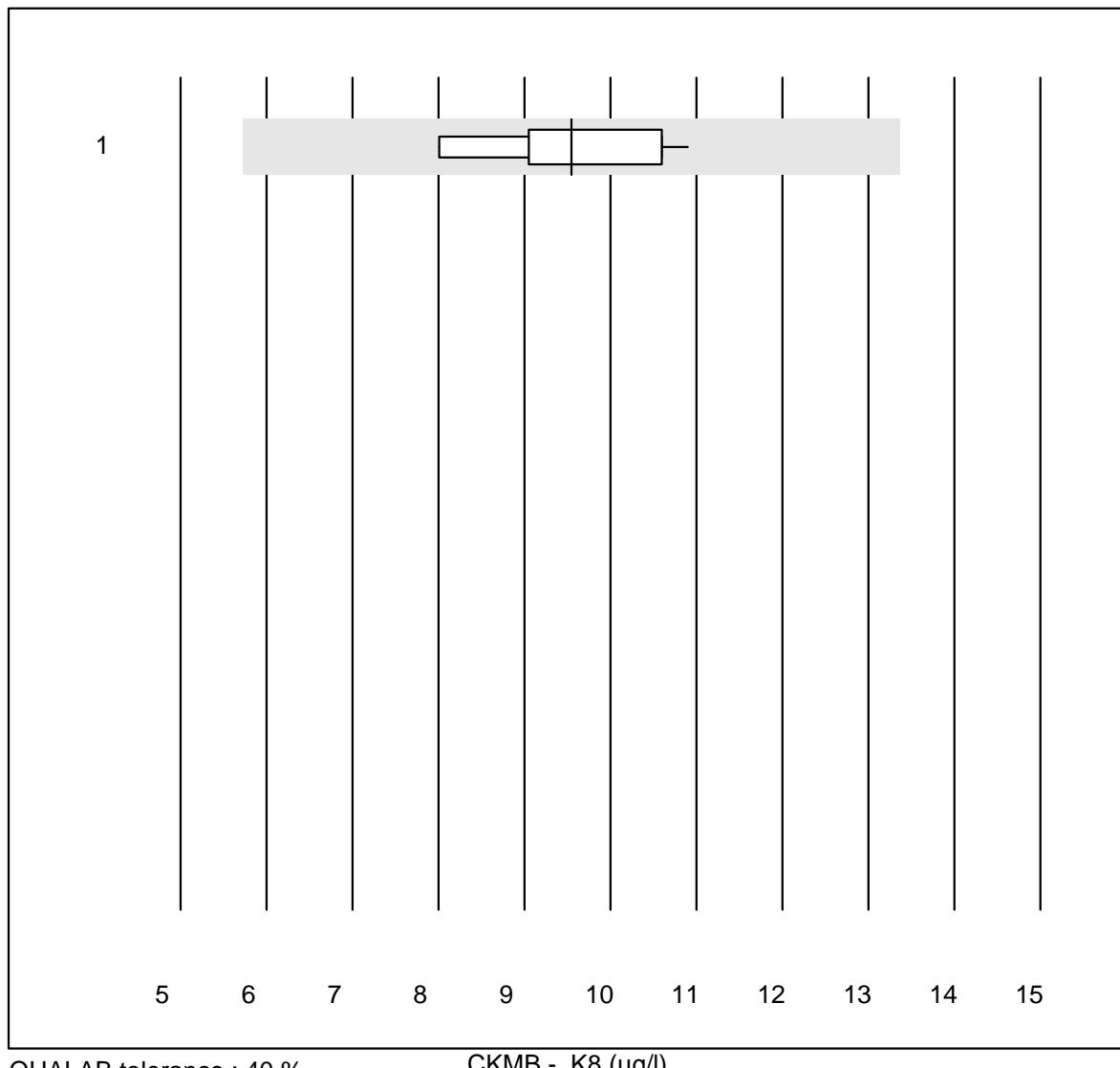
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 Cobas h 232 | 869 | 93.2 | 5.5 | 1.3 | 424.00 | 12.5 | e |
| 2 Cardiac Reader | 51 | 88.3 | 7.8 | 3.9 | 416.45 | 13.1 | e |

Troponin I WB



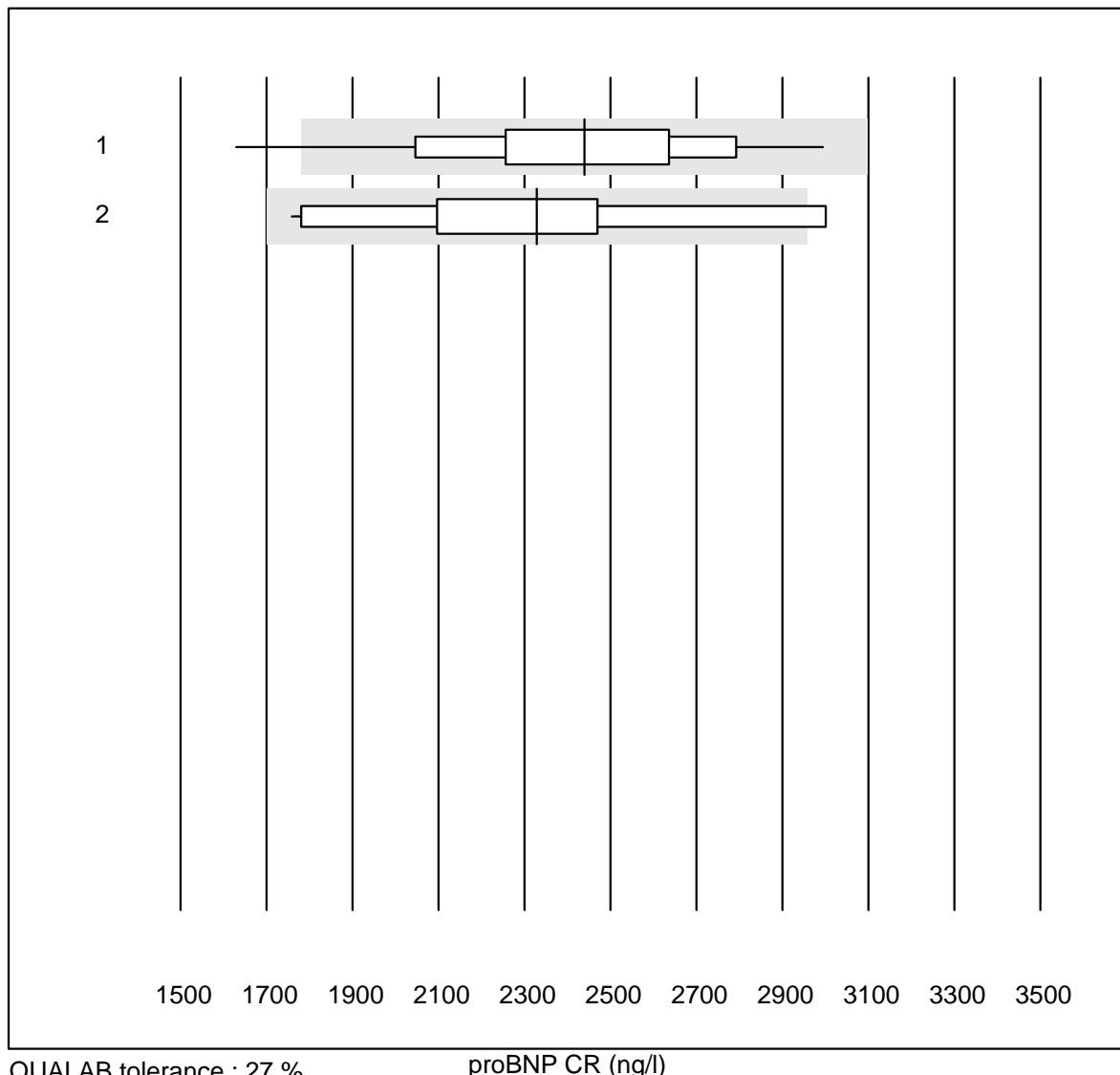
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | iStat | 4 | 100.0 | 0.0 | 0.0 | 5.13 | 5.0 | e |

D-dimer CR

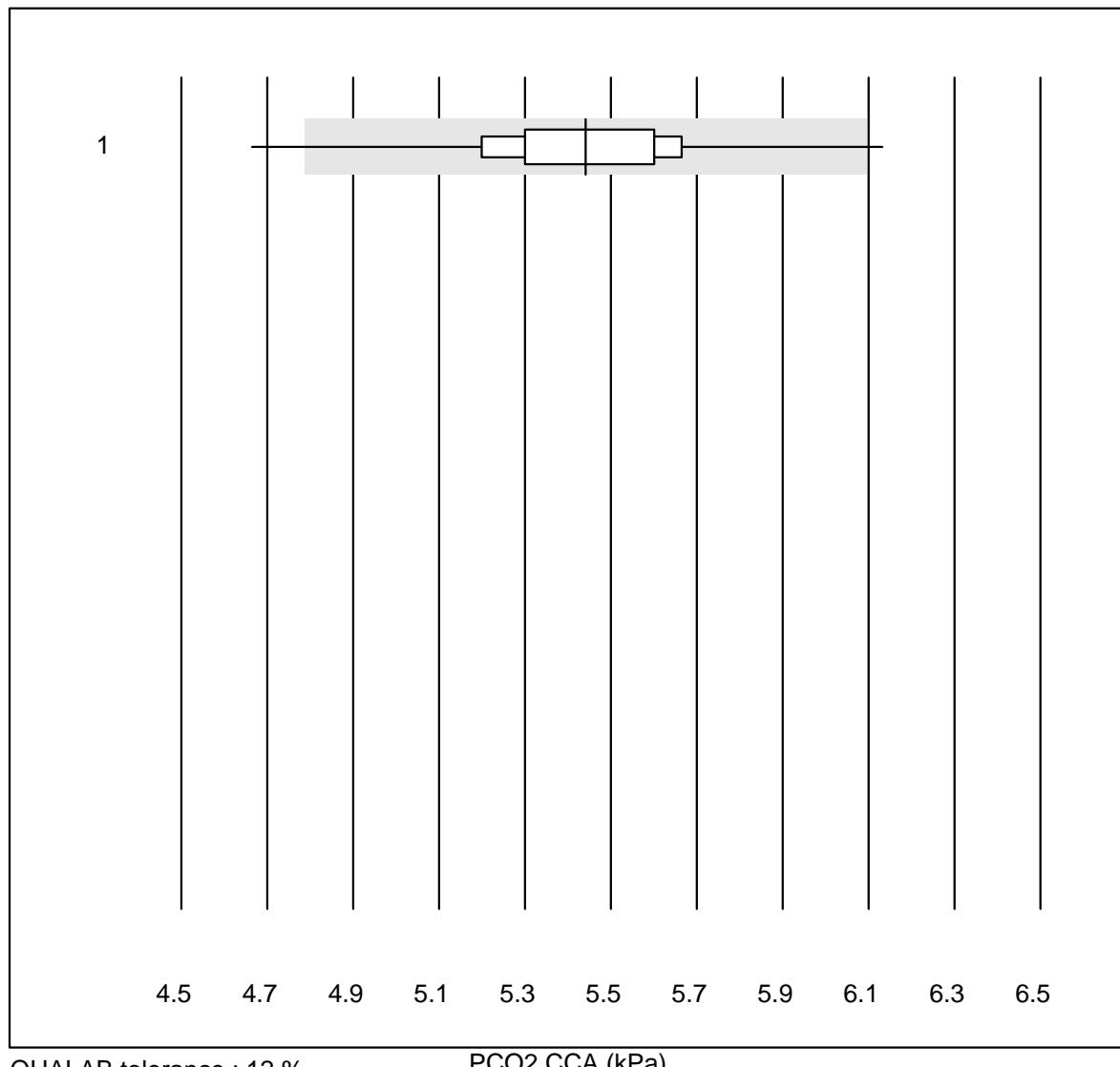
CKMB - K8

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|---------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 Cobas h 232 | 10 | 100.0 | 0.0 | 0.0 | 9.5 | 10.1 | e |

proBNP CR



| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 Cobas h 232 | 543 | 95.2 | 1.7 | 3.1 | 2439 | 11.5 | e |
| 2 Cardiac Reader | 13 | 76.9 | 23.1 | 0.0 | 2329 | 19.0 | e* |

PCO₂ CCA

No. Methode

Total

% good

% insuff.

% outlier

Target value

CV% Typ

1 OPTI CCA

13

84.6

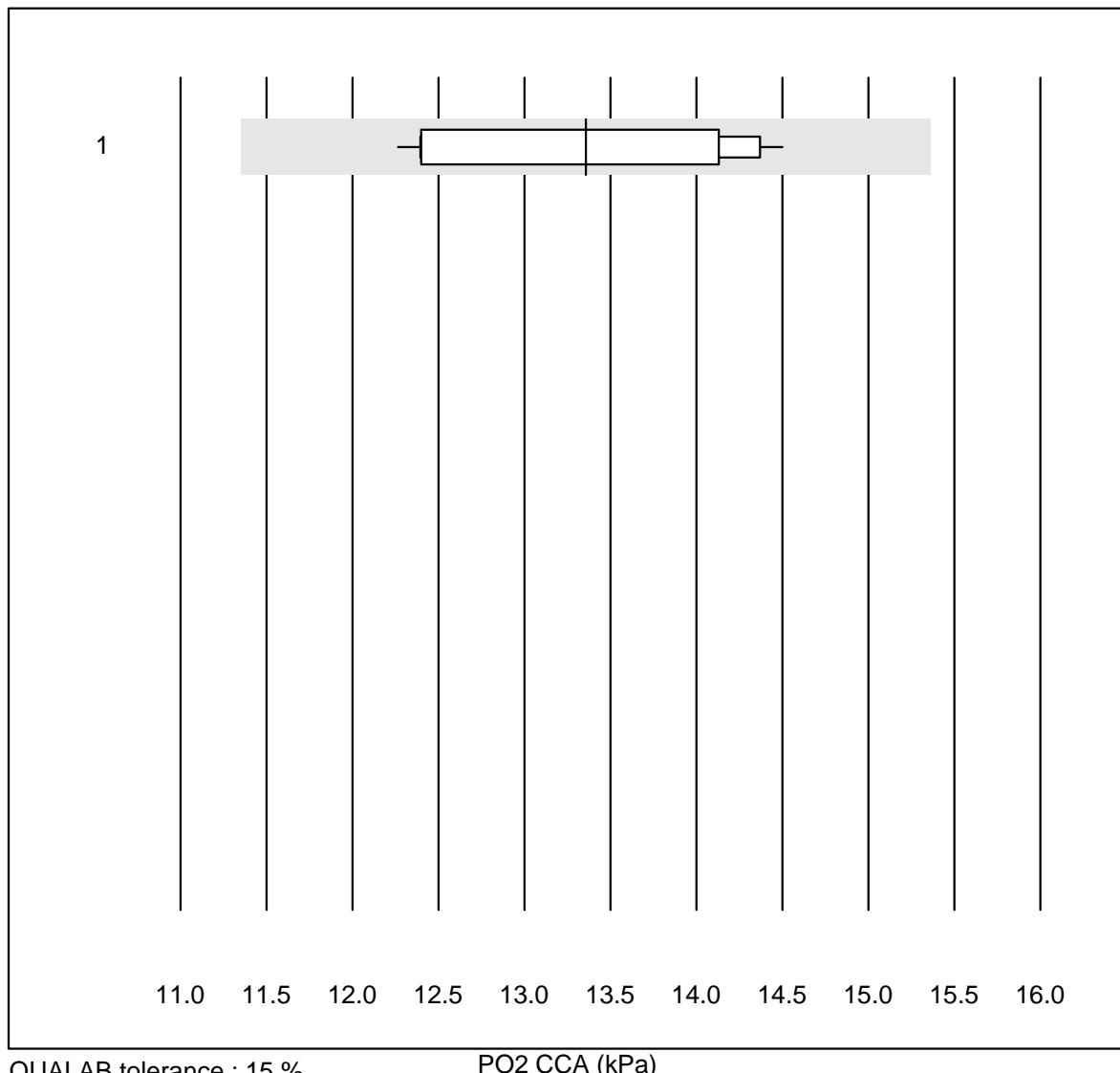
15.4

0.0

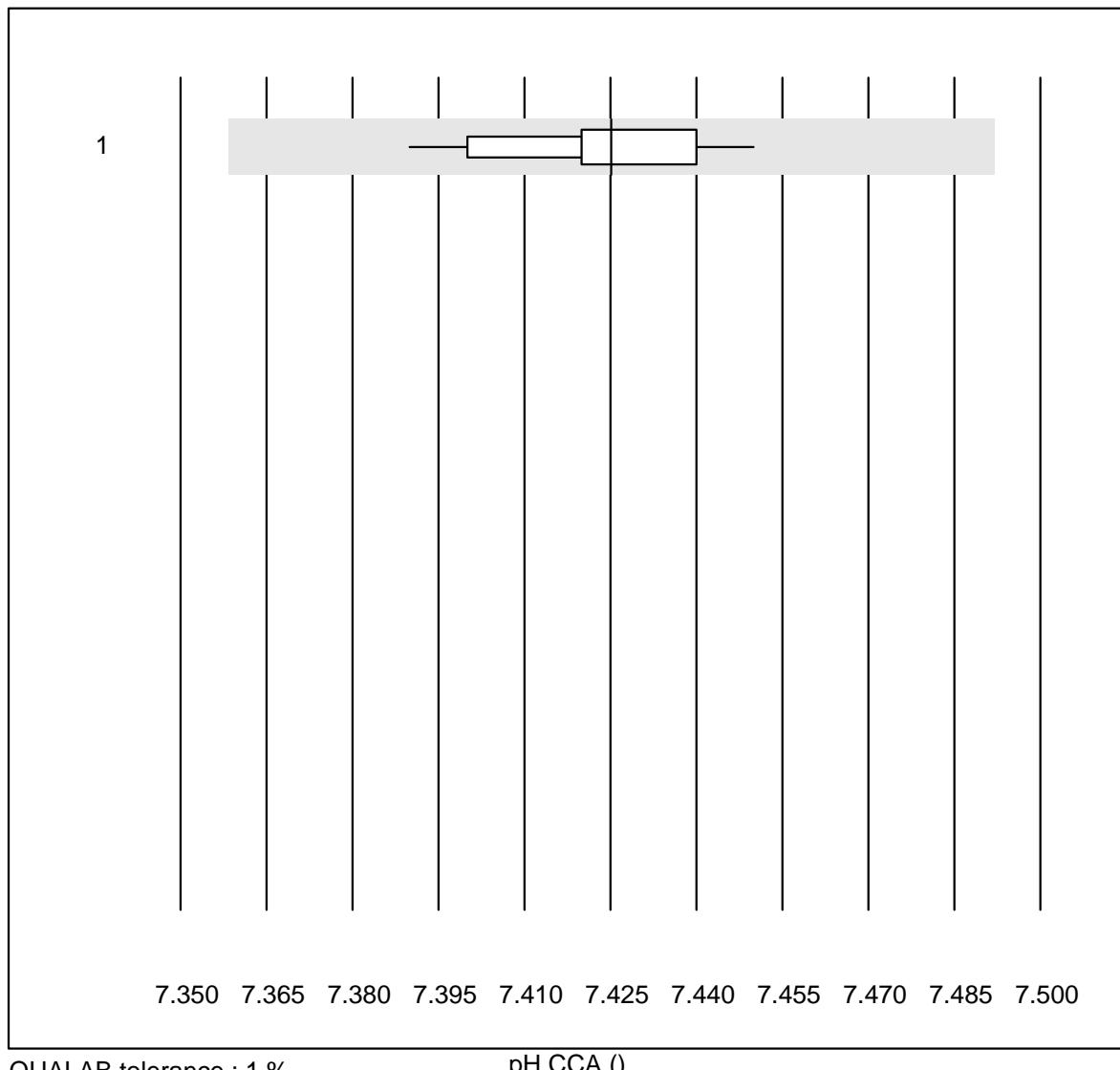
5.44

6.2

e*

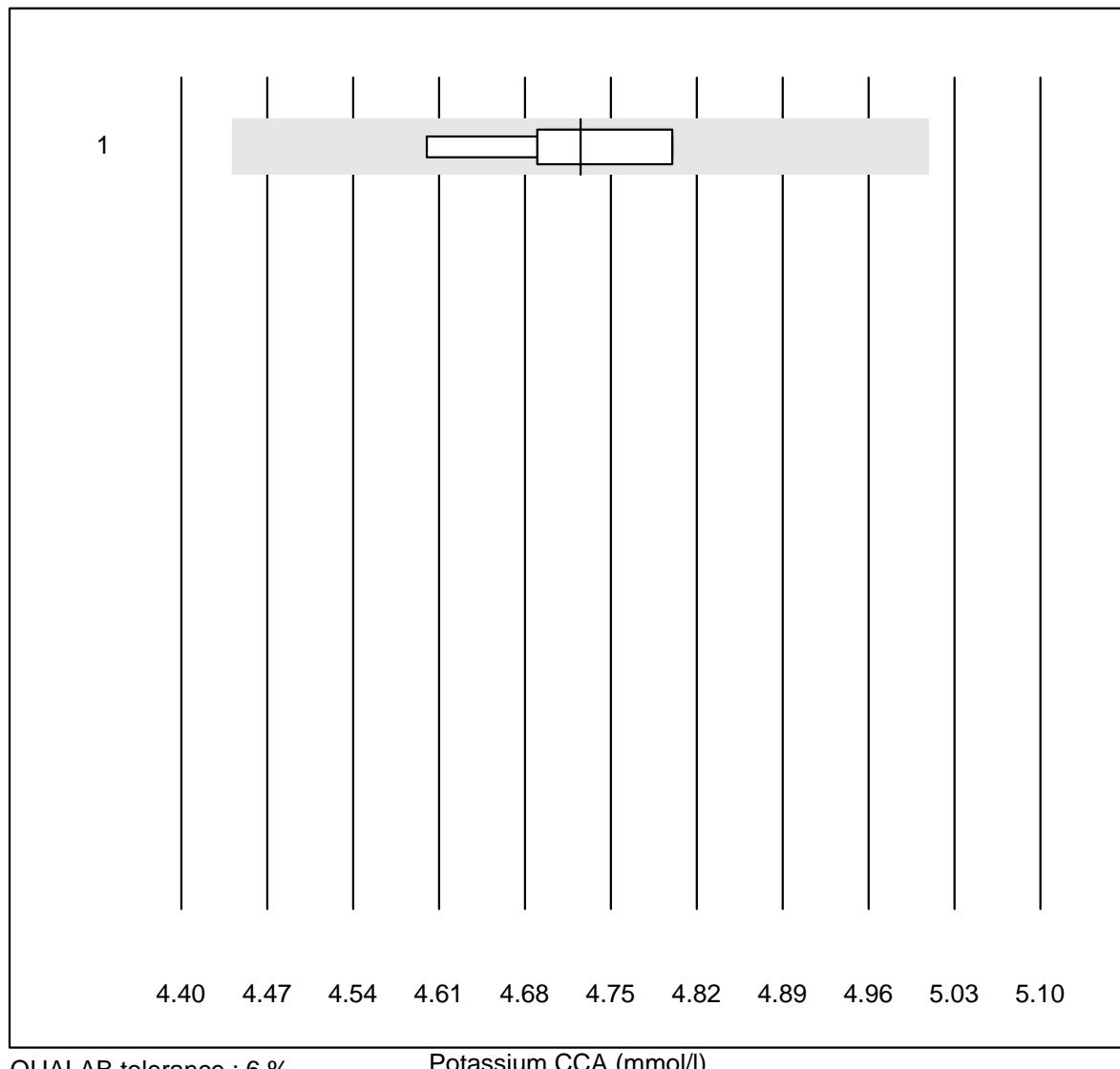
PO2 CCA

| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|----------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | OPTI CCA | 13 | 92.3 | 0.0 | 7.7 | 13.36 | 6.7 | e* |

pH CCA

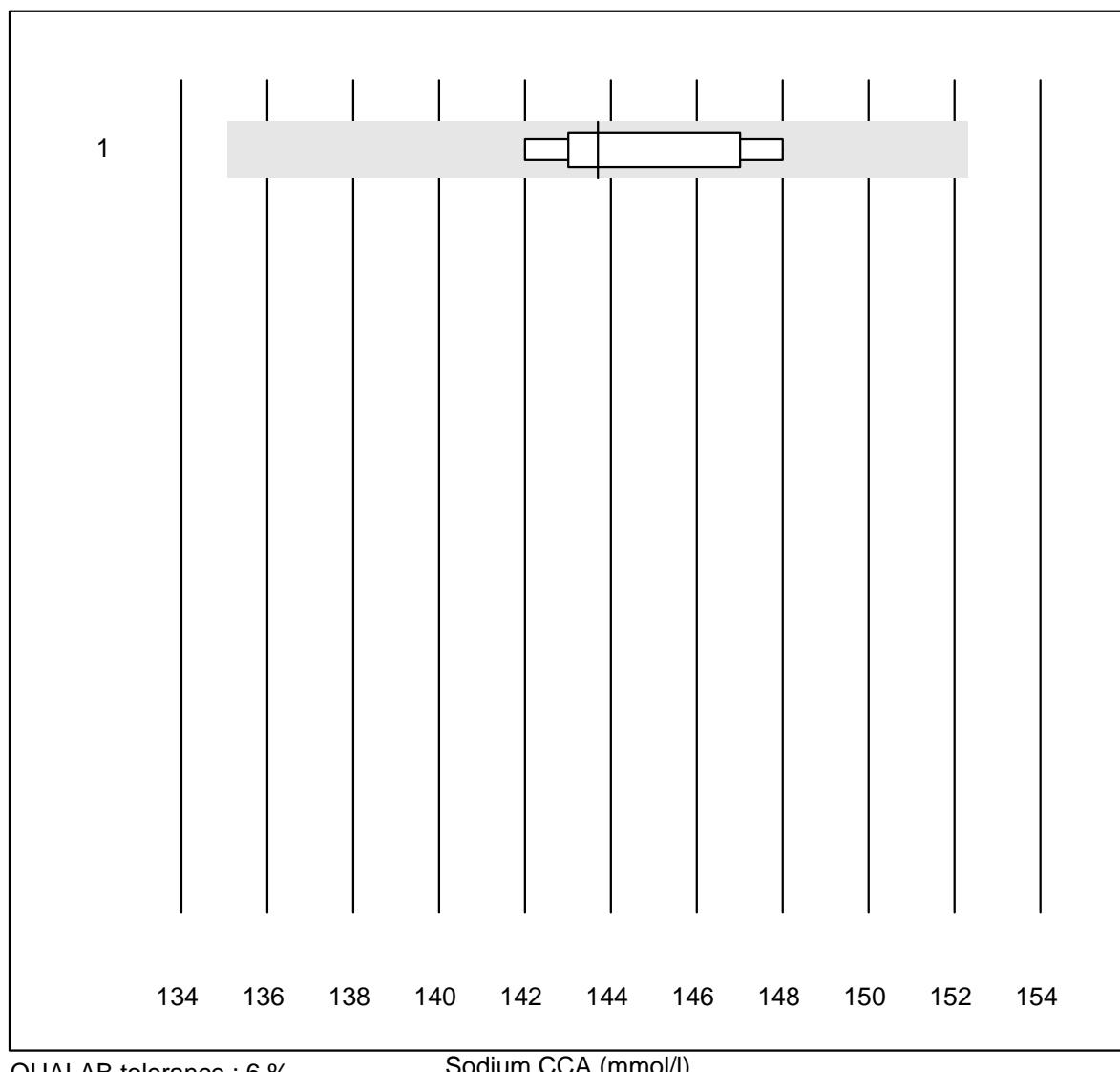
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 OPTI CCA | 13 | 100.0 | 0.0 | 0.0 | 7.43 | 0.2 | e |

Potassium CCA

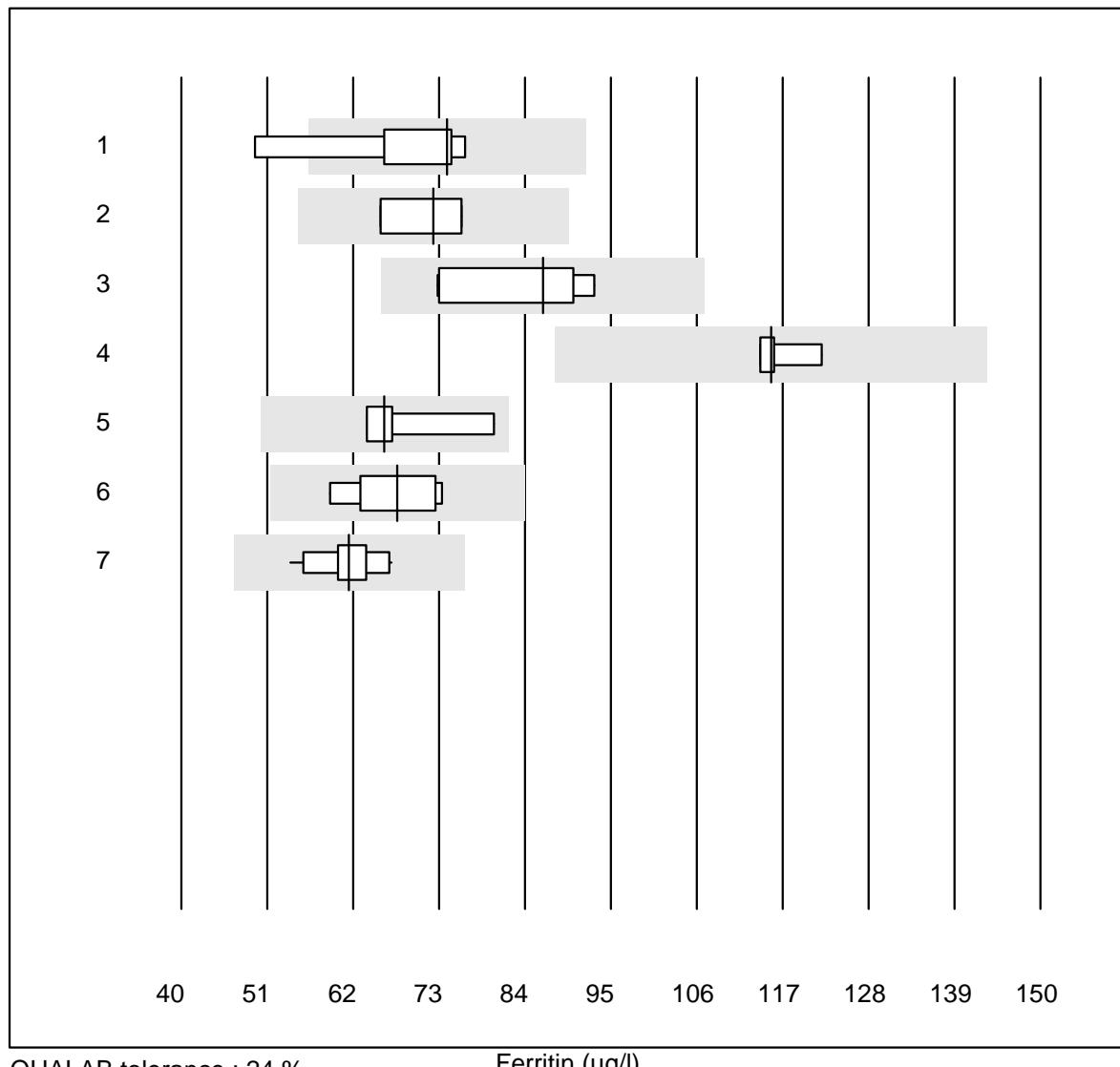


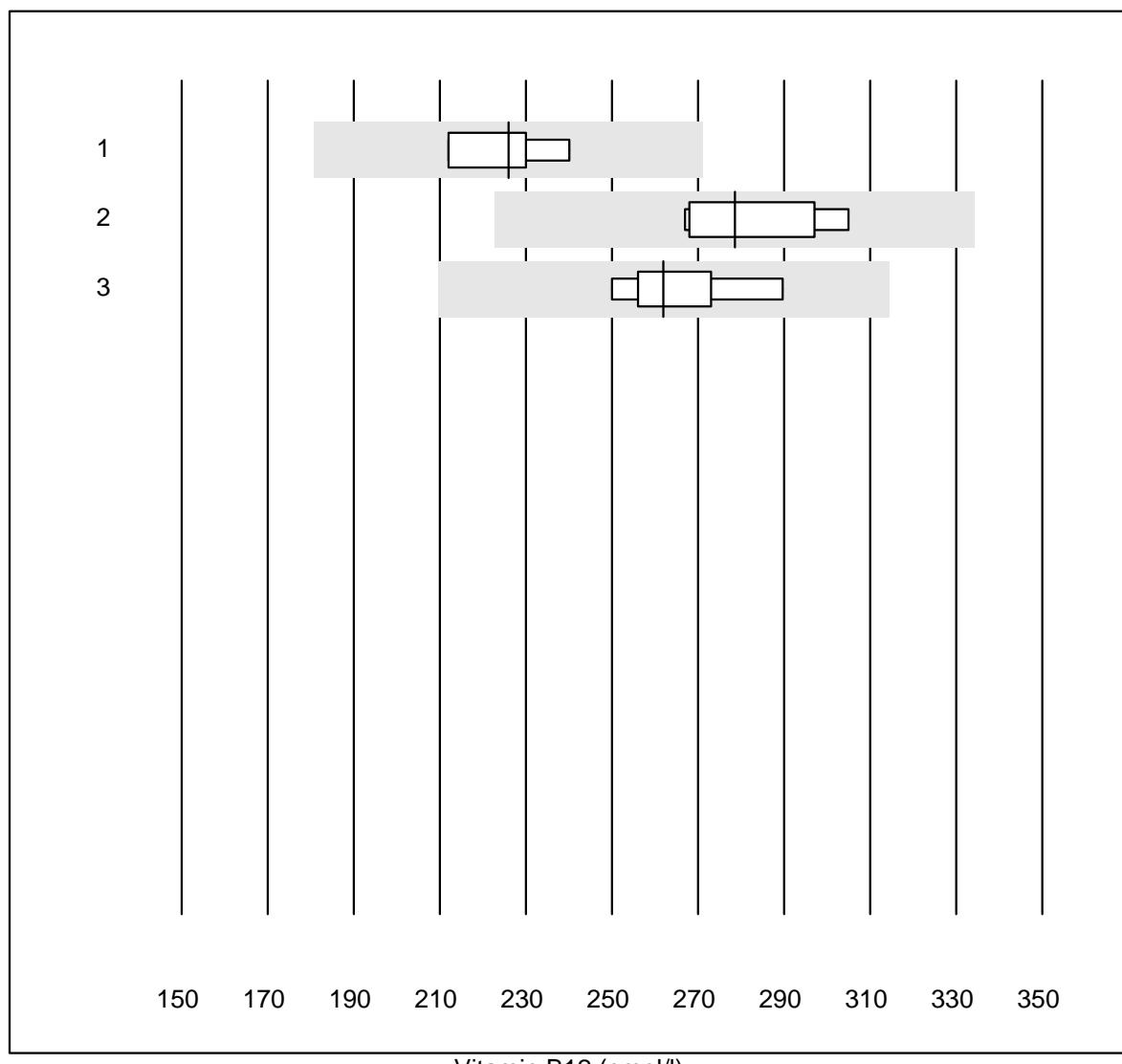
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|----------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | OPTI CCA | 8 | 100.0 | 0.0 | 0.0 | 4.7 | 1.5 | e |

Sodium CCA

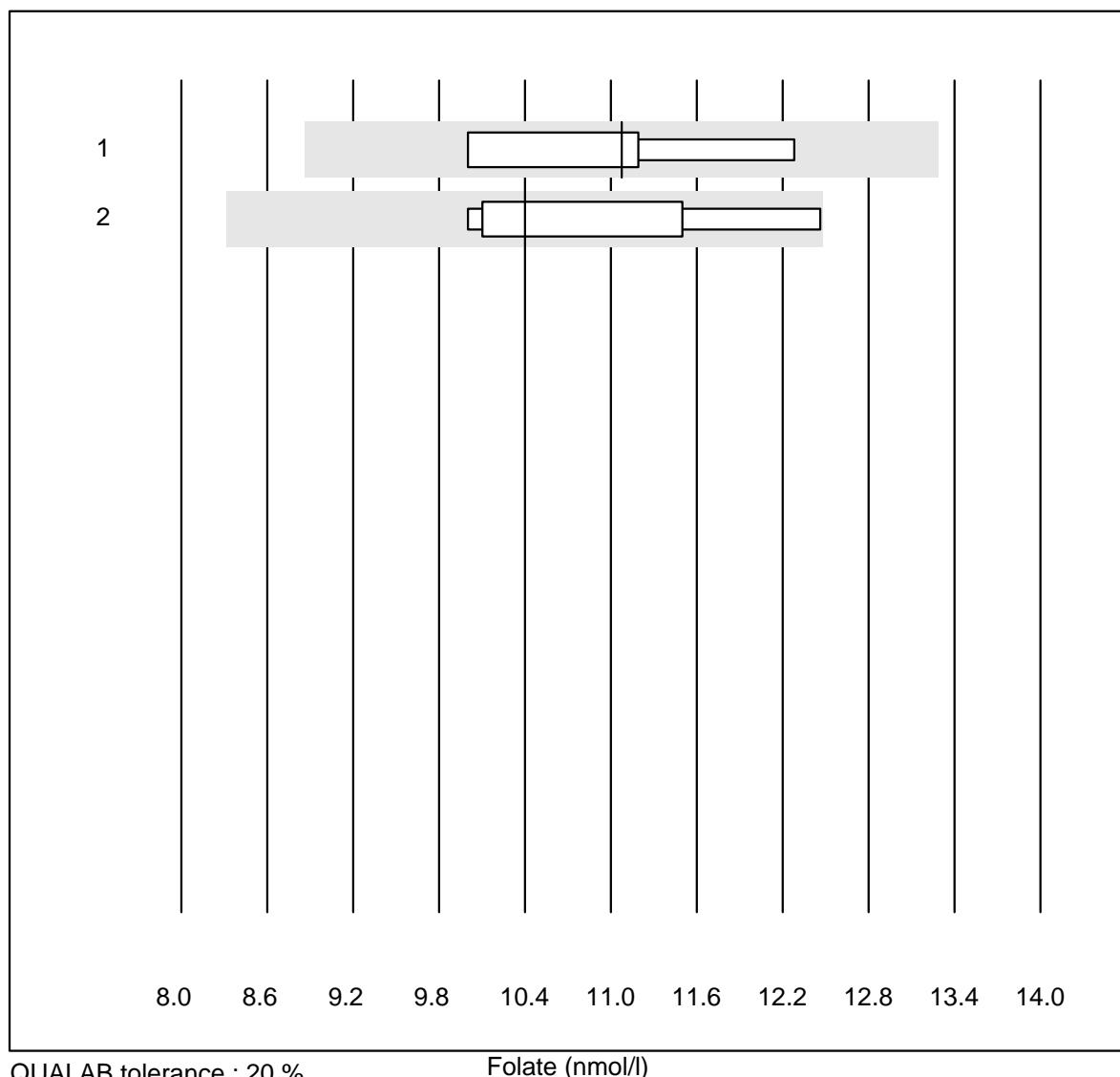


| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|----------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | OPTI CCA | 7 | 100.0 | 0.0 | 0.0 | 143.7 | 1.6 | e |

Ferritin

Vitamin B12

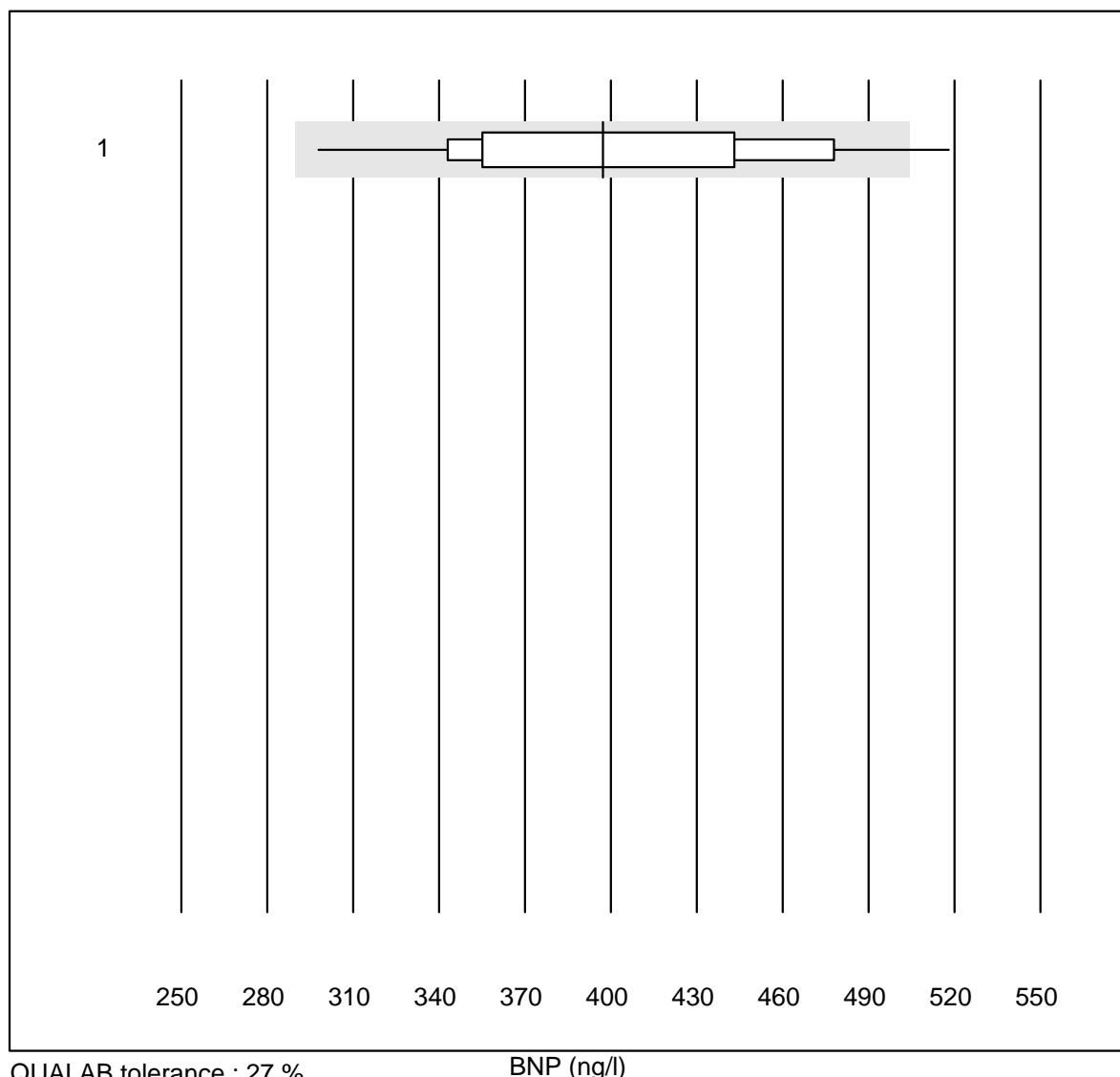
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 ADVIA Centaur XP/CP | 4 | 100.0 | 0.0 | 0.0 | 226.00 | 5.3 | e* |
| 2 Cobas E / Elecsys | 7 | 100.0 | 0.0 | 0.0 | 278.50 | 5.4 | e |
| 3 Architect | 6 | 100.0 | 0.0 | 0.0 | 262.00 | 5.4 | e |

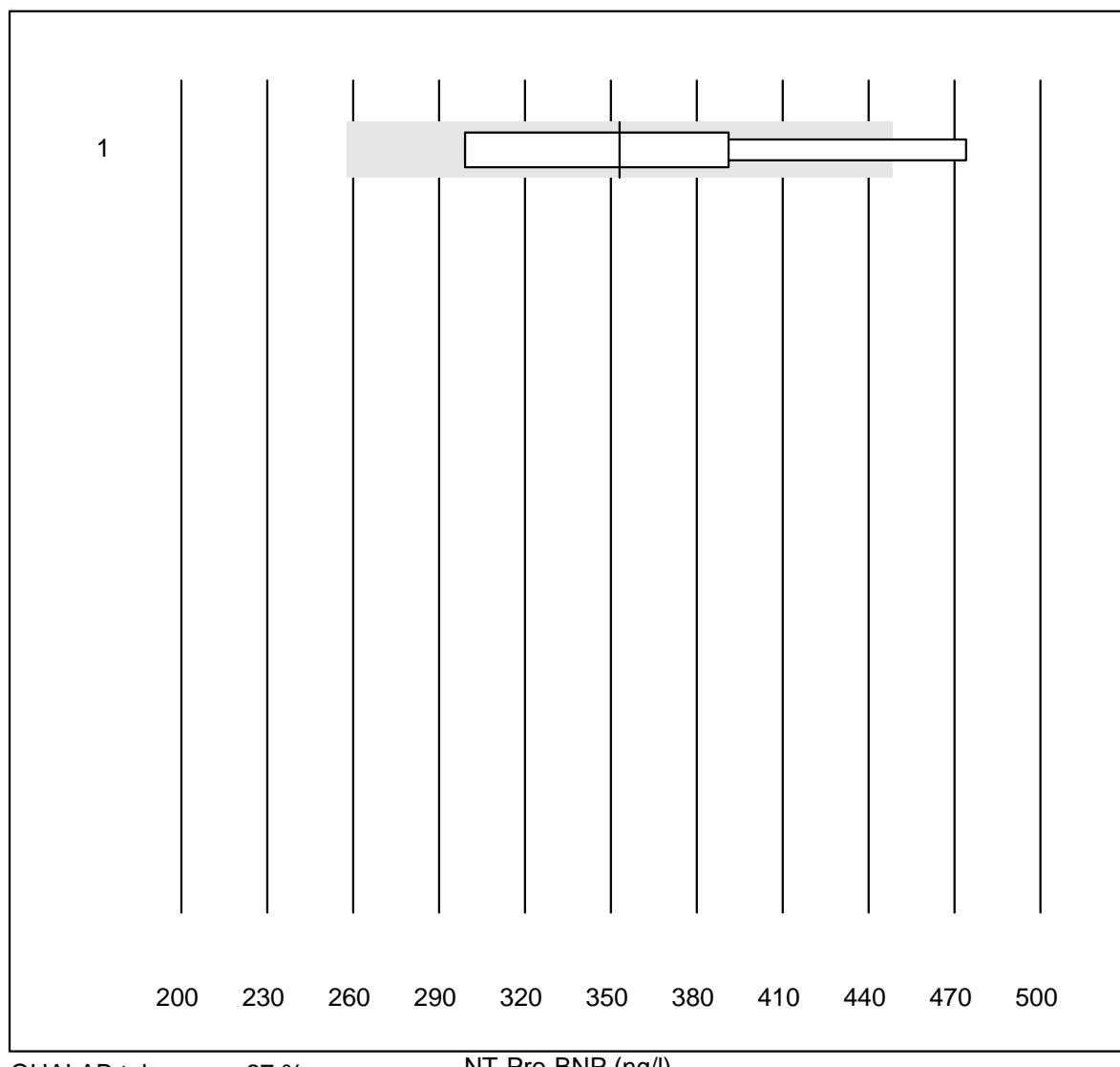
Folate

QUALAB tolerance : 20 %

Folate (nmol/l)

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Cobas alt | 4 | 100.0 | 0.0 | 0.0 | 11.08 | 8.4 | a |
| 2 Architect | 5 | 100.0 | 0.0 | 0.0 | 10.40 | 9.7 | e* |

BNP

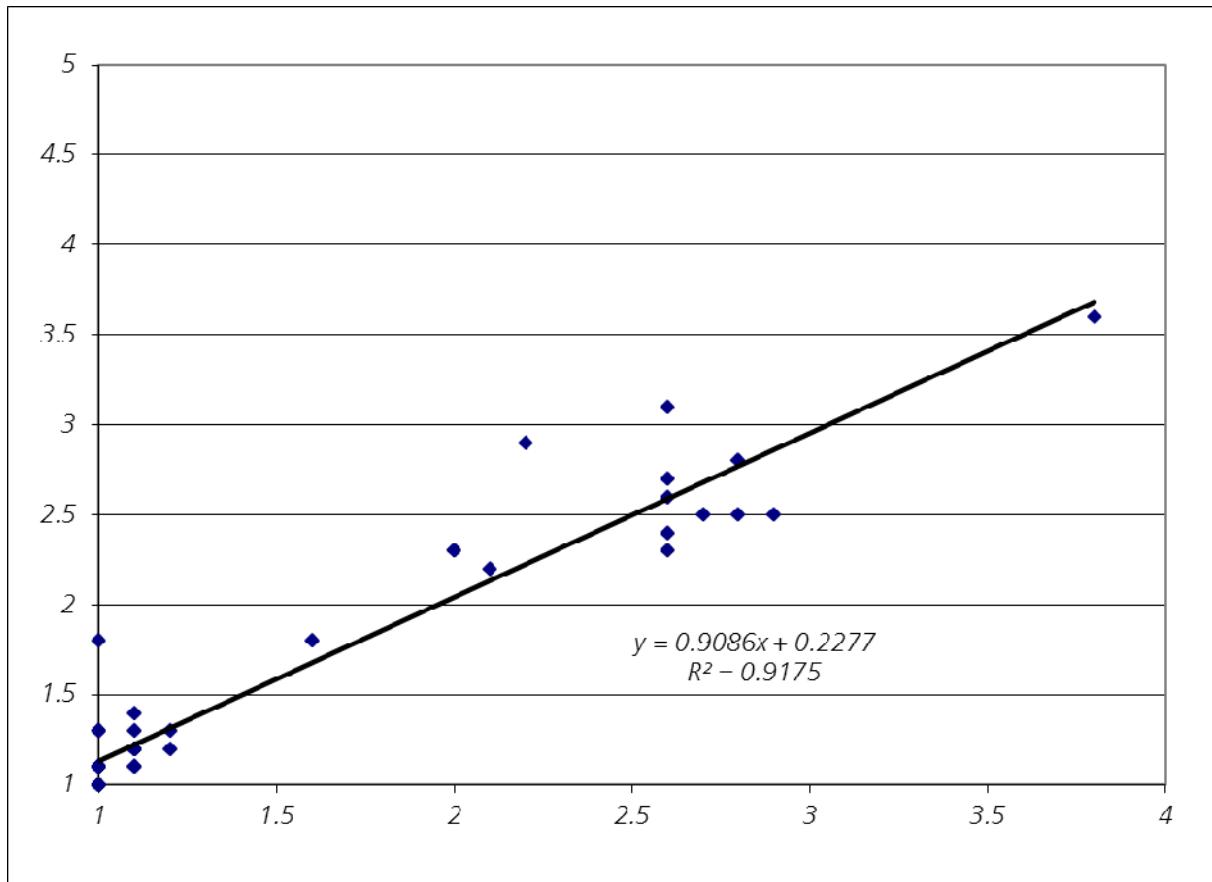
NT-Pro-BNP

| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 | Triage | 9 | 33.3 | 11.1 | 55.6 | 353 | 19.4 | e* |

G10 INR INRatio

INR INRatio

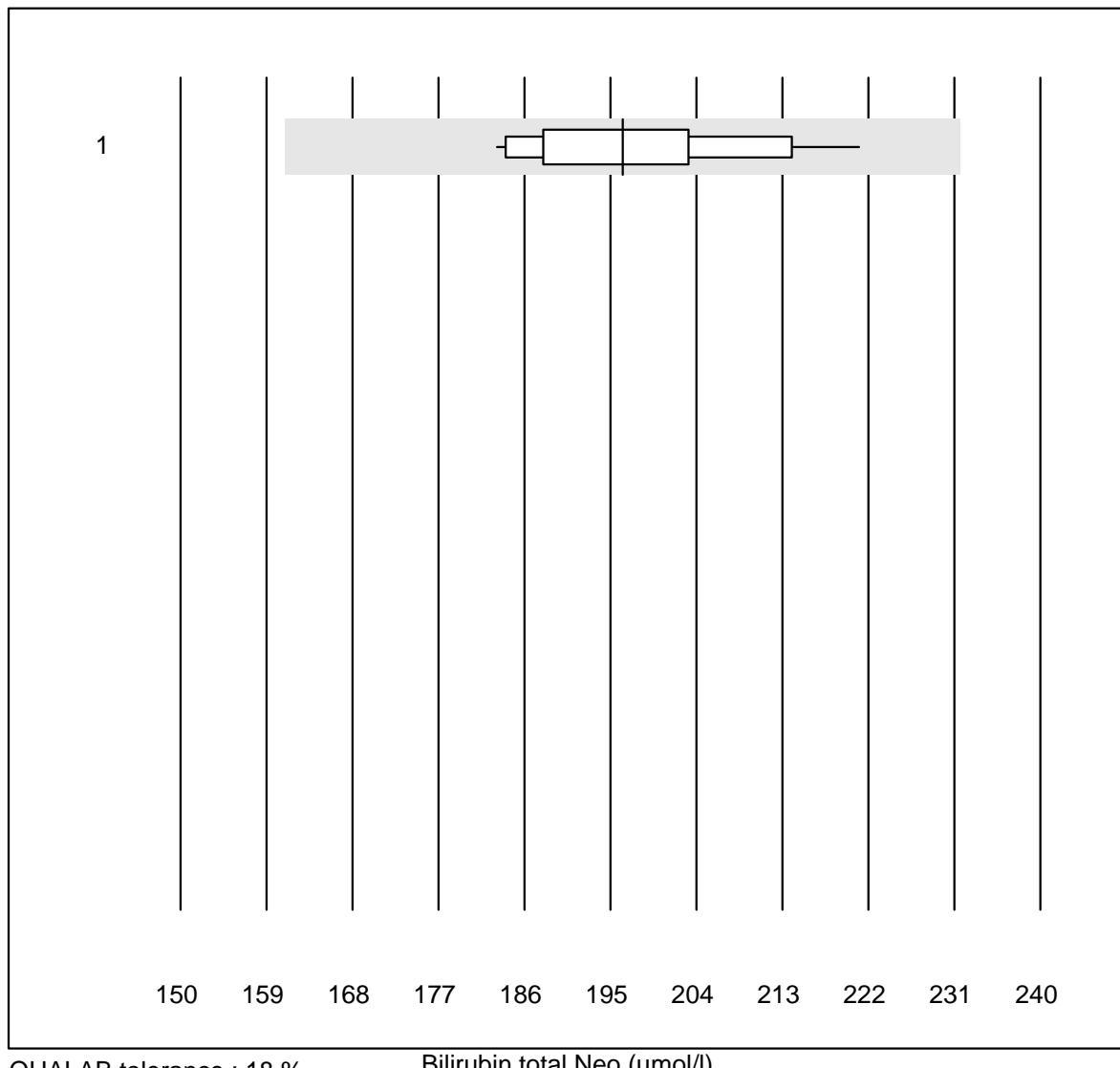
University Hospital Zuerich



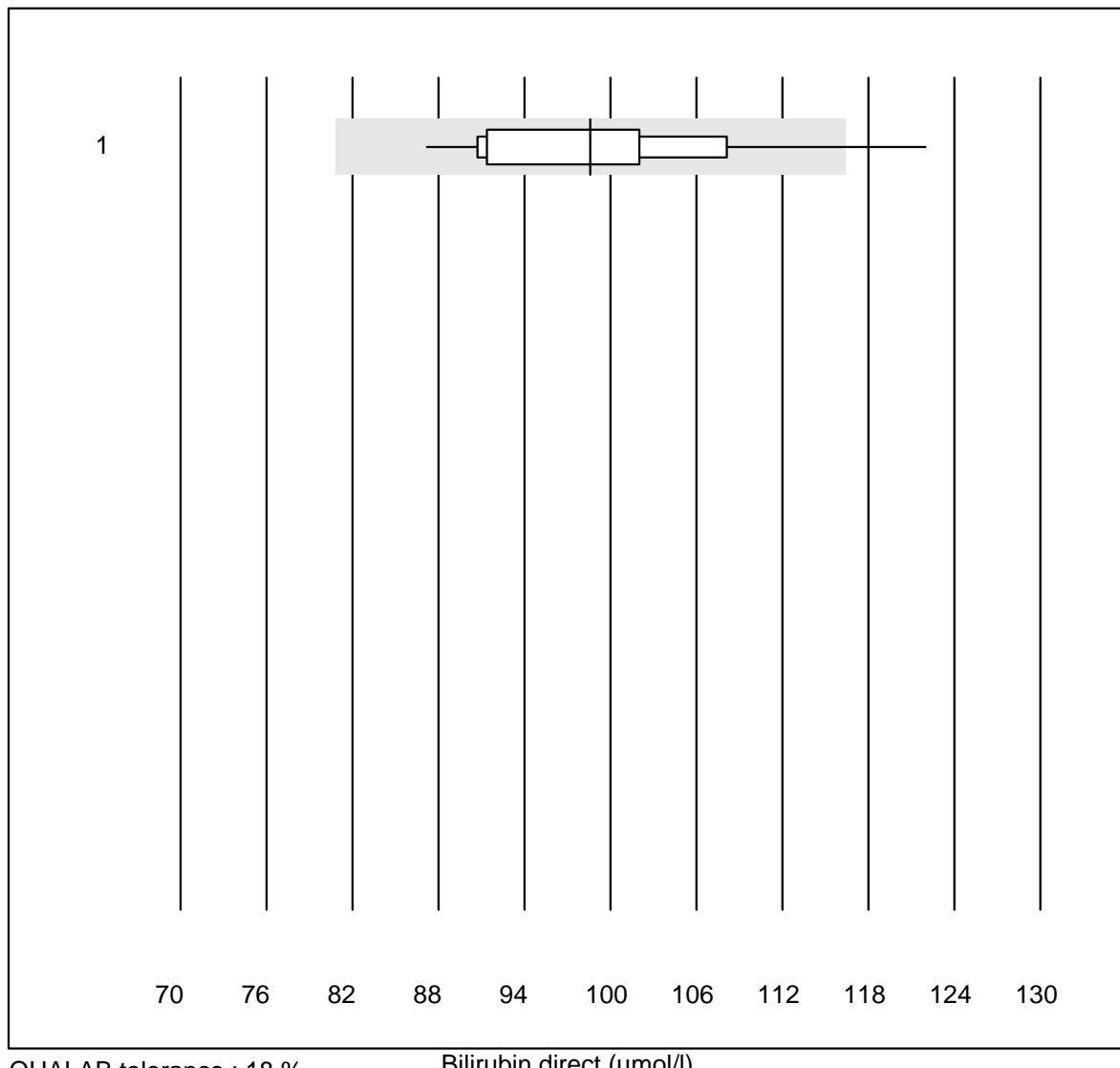
INRatio Participants

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 | INRatio | 50 | 82.00 | 8.00 | 10.00 |

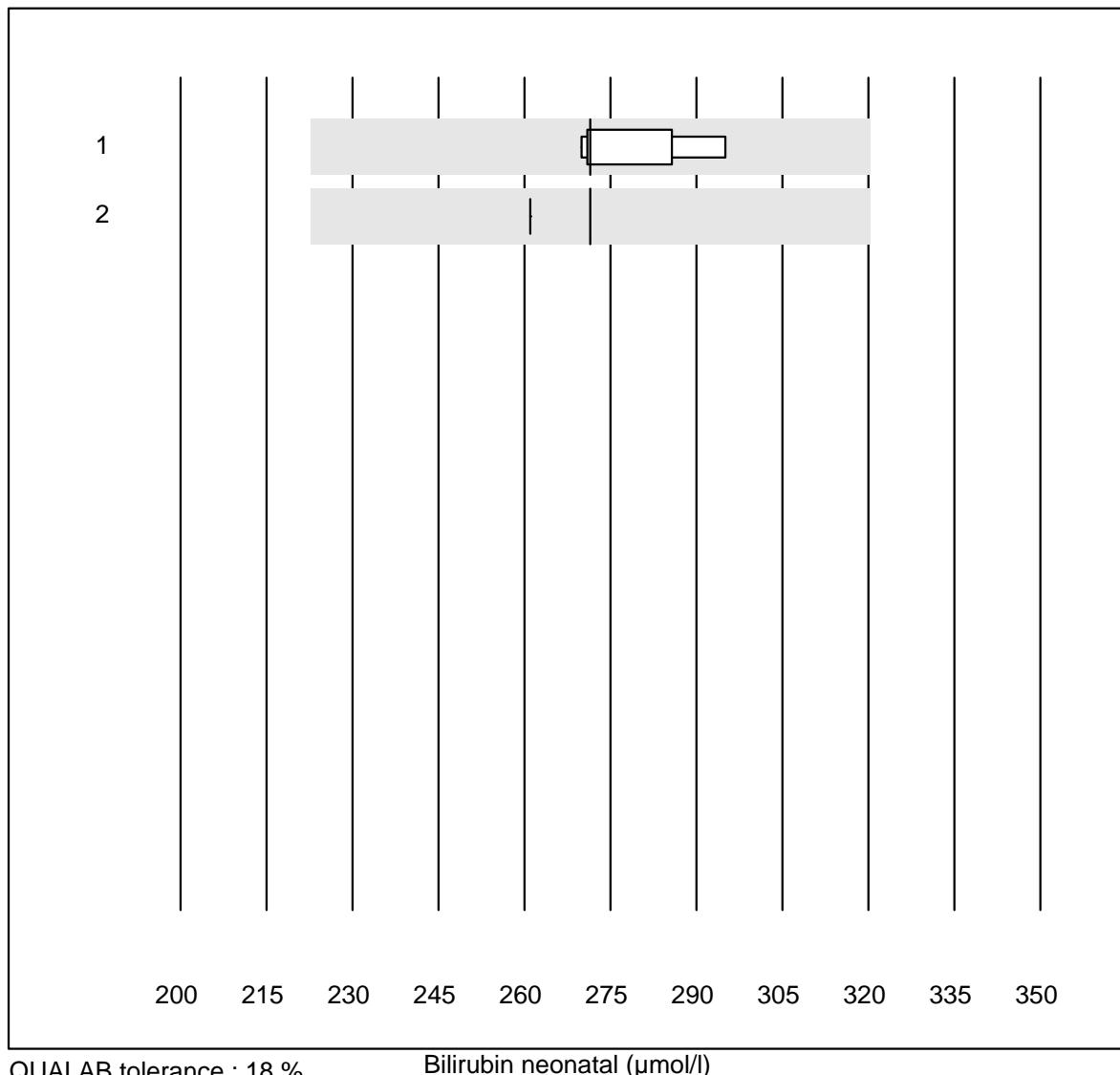
Bilirubin total Neo

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 all Participants | 14 | 100.0 | 0.0 | 0.0 | 196 | 5.8 | e |

Bilirubin direct

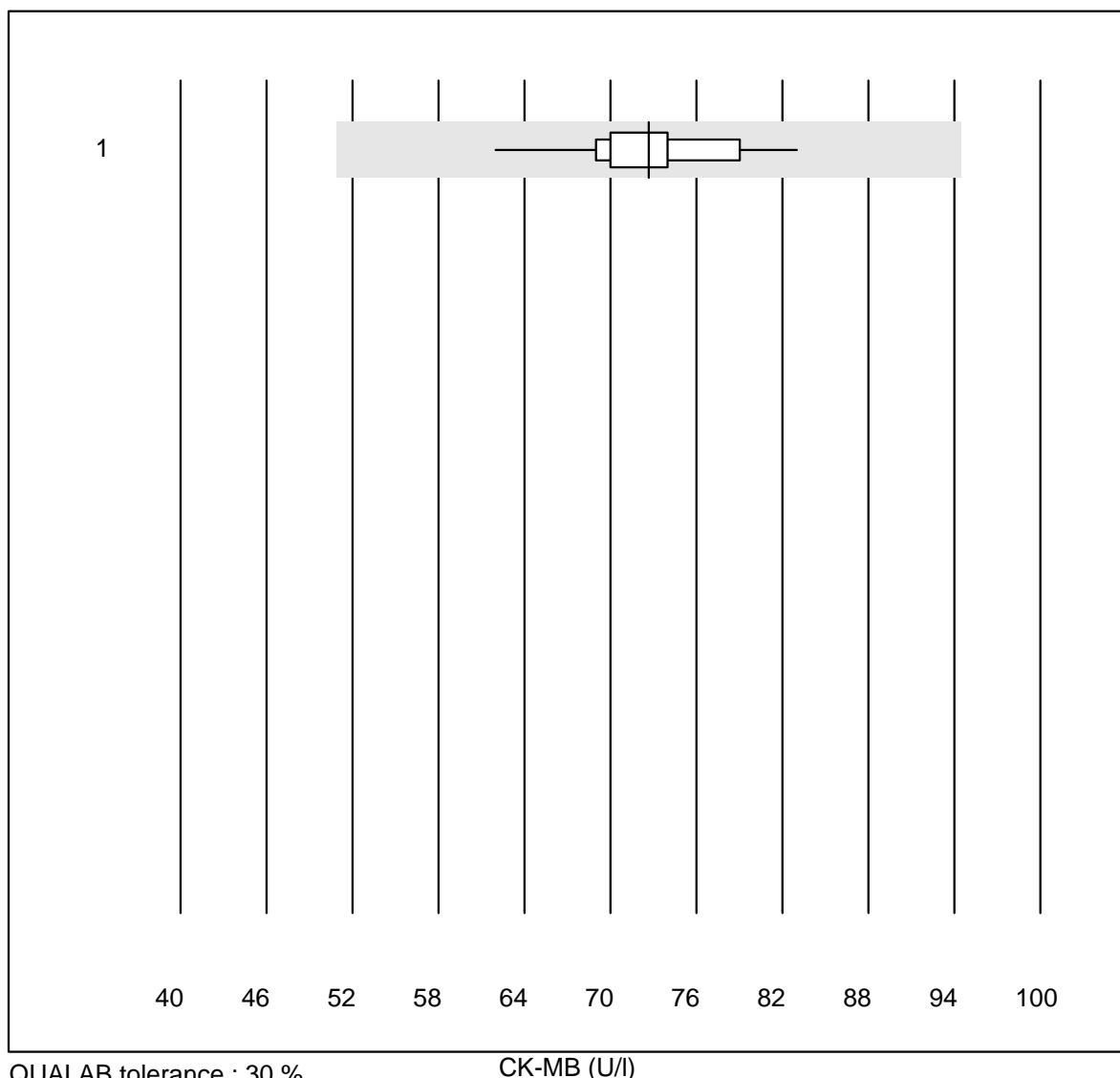
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 all Participants | 12 | 91.7 | 8.3 | 0.0 | 99 | 9.7 | e* |

Bilirubin neonatal



K15 Creatinkinase Activity

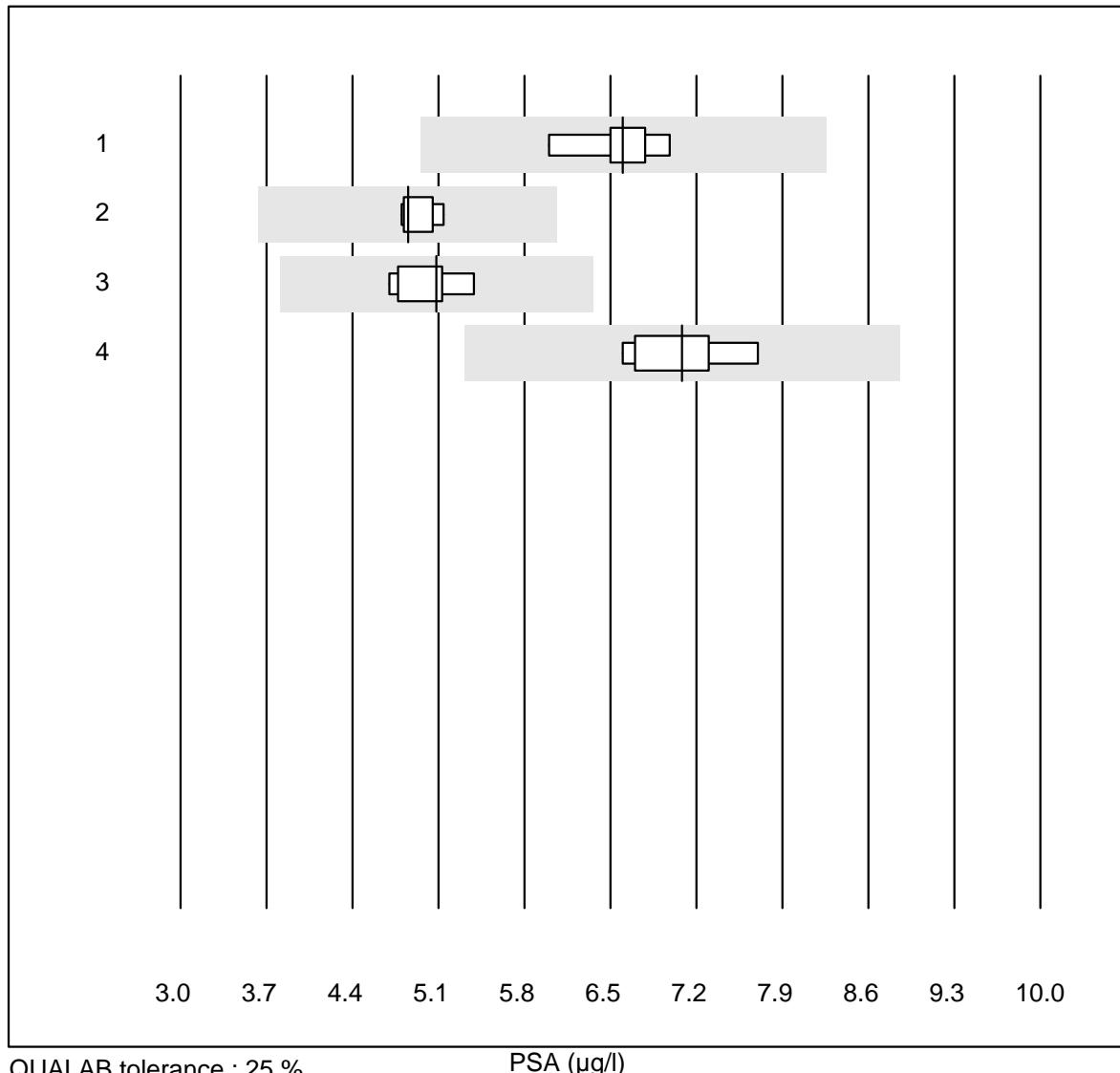
CK-MB



| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Fuji Dri-Chem | 44 | 100.0 | 0.0 | 0.0 | 72.7 | 6.0 | e |

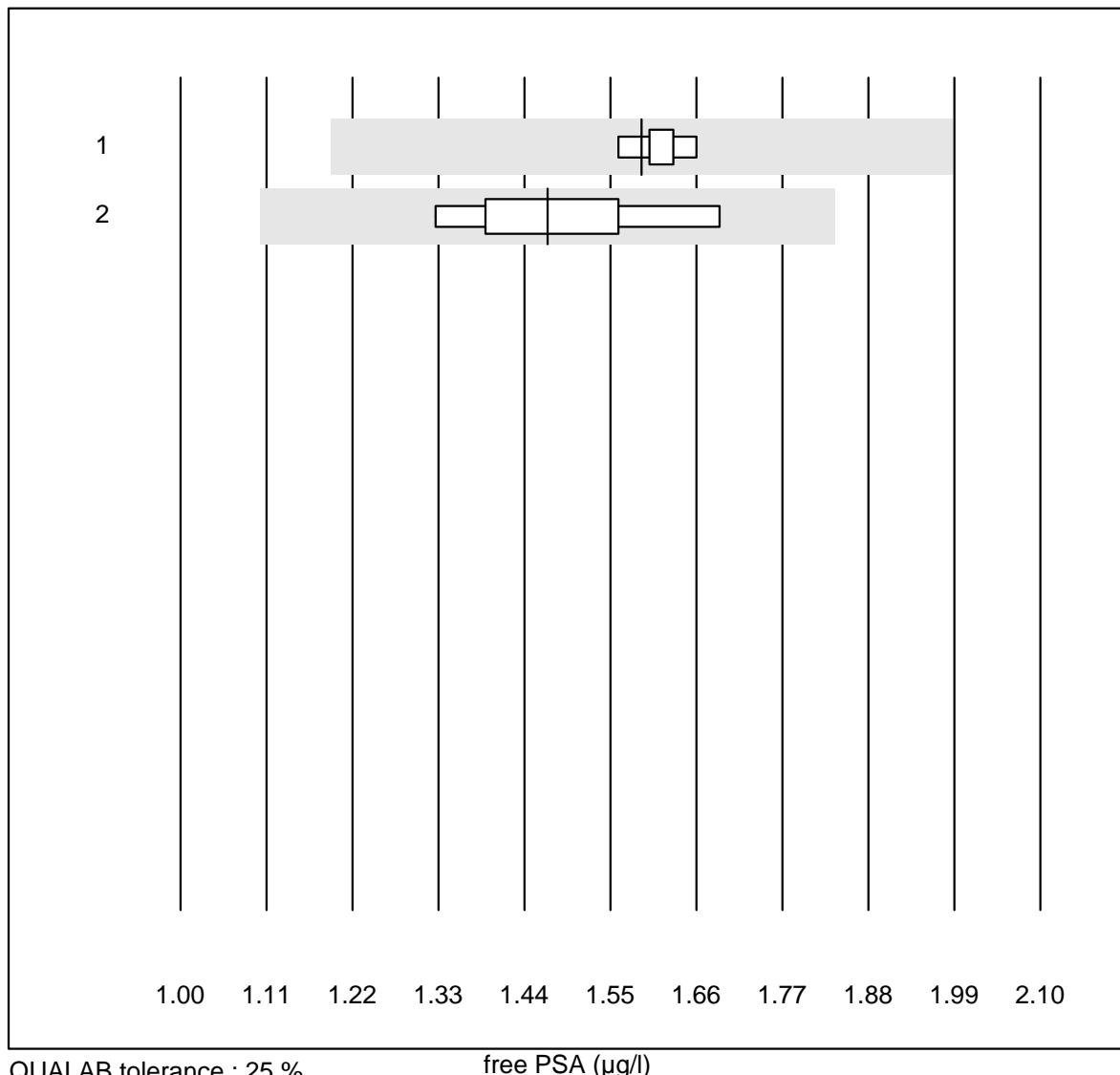
K14 Tumor Markers

PSA



K14 Tumor Markers

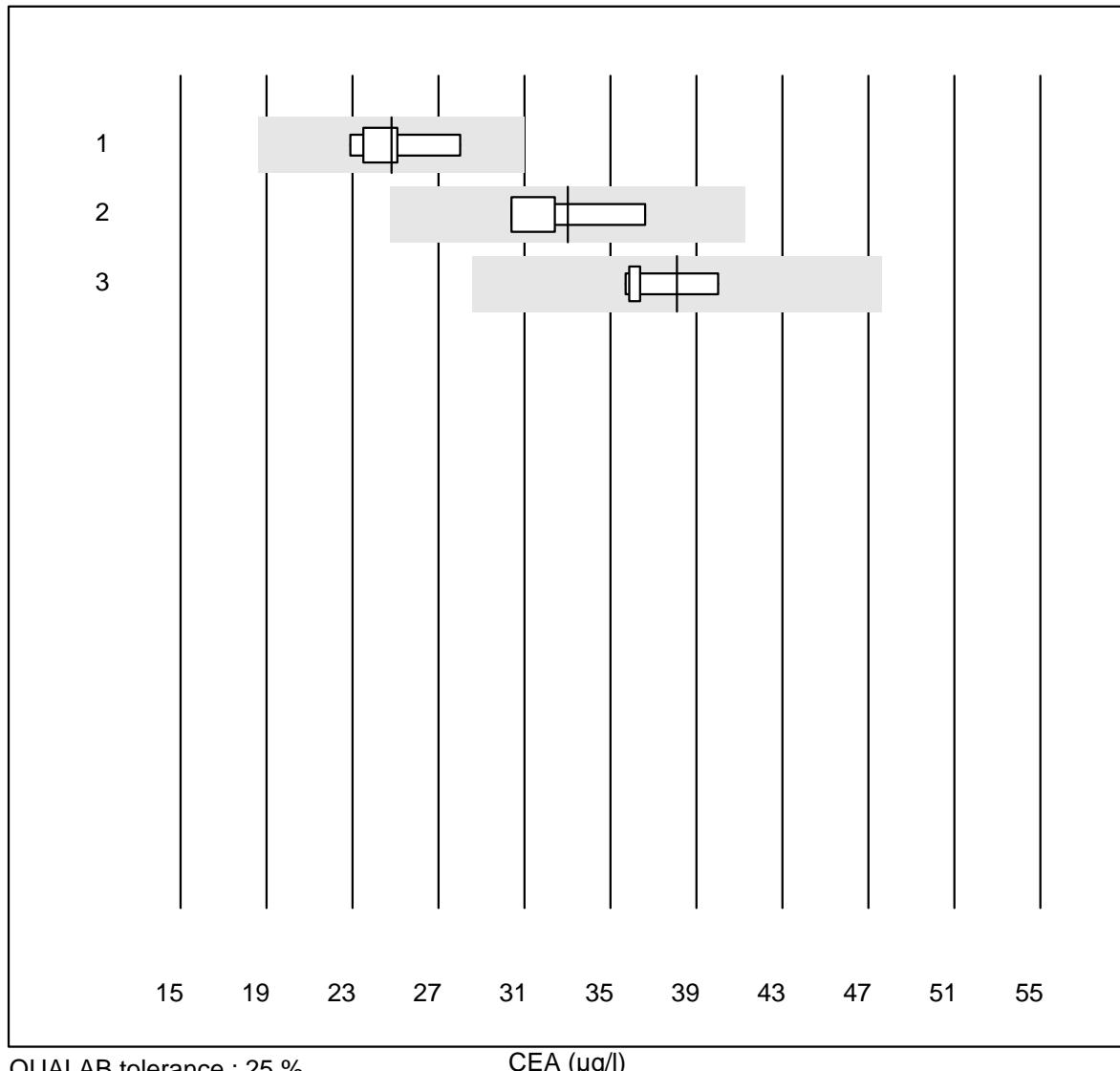
free PSA



| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|-------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Cobas E / Elecsys | 5 | 100.0 | 0.0 | 0.0 | 1.59 | 2.3 | a |
| 2 | Architect | 5 | 100.0 | 0.0 | 0.0 | 1.47 | 9.8 | a |

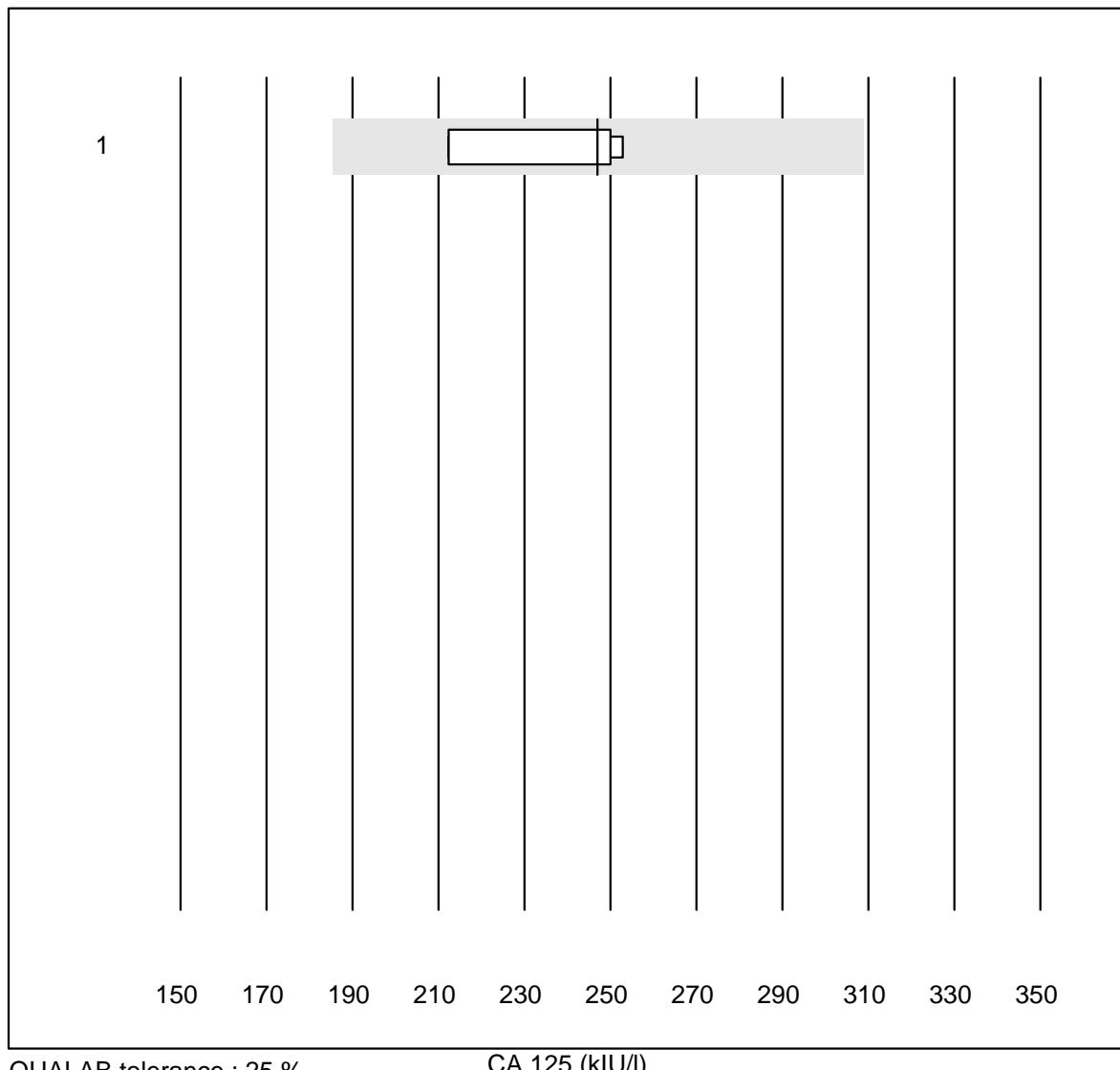
K14 Tumor Markers

CEA

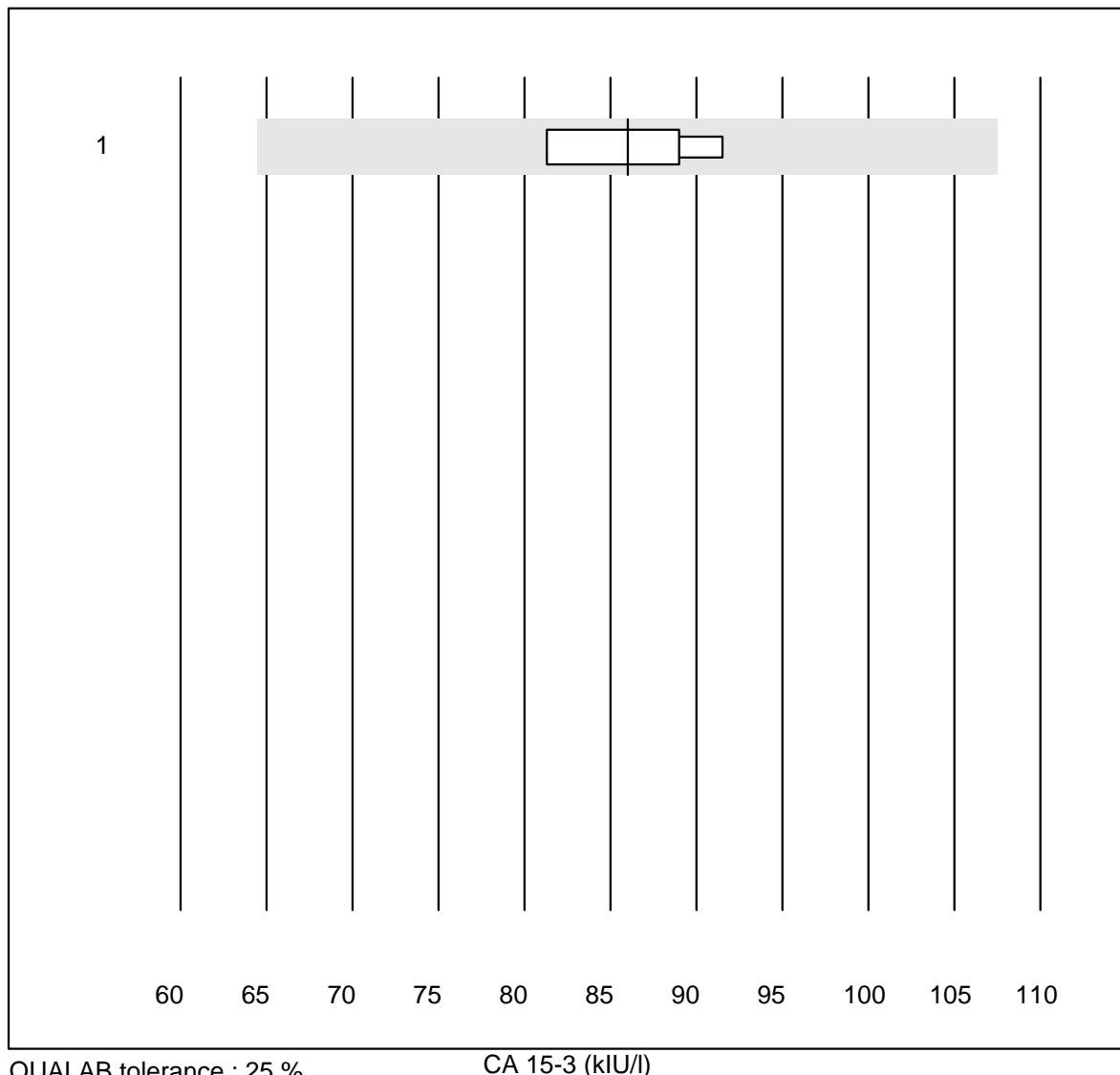


K14 Tumor Markers

CA 125



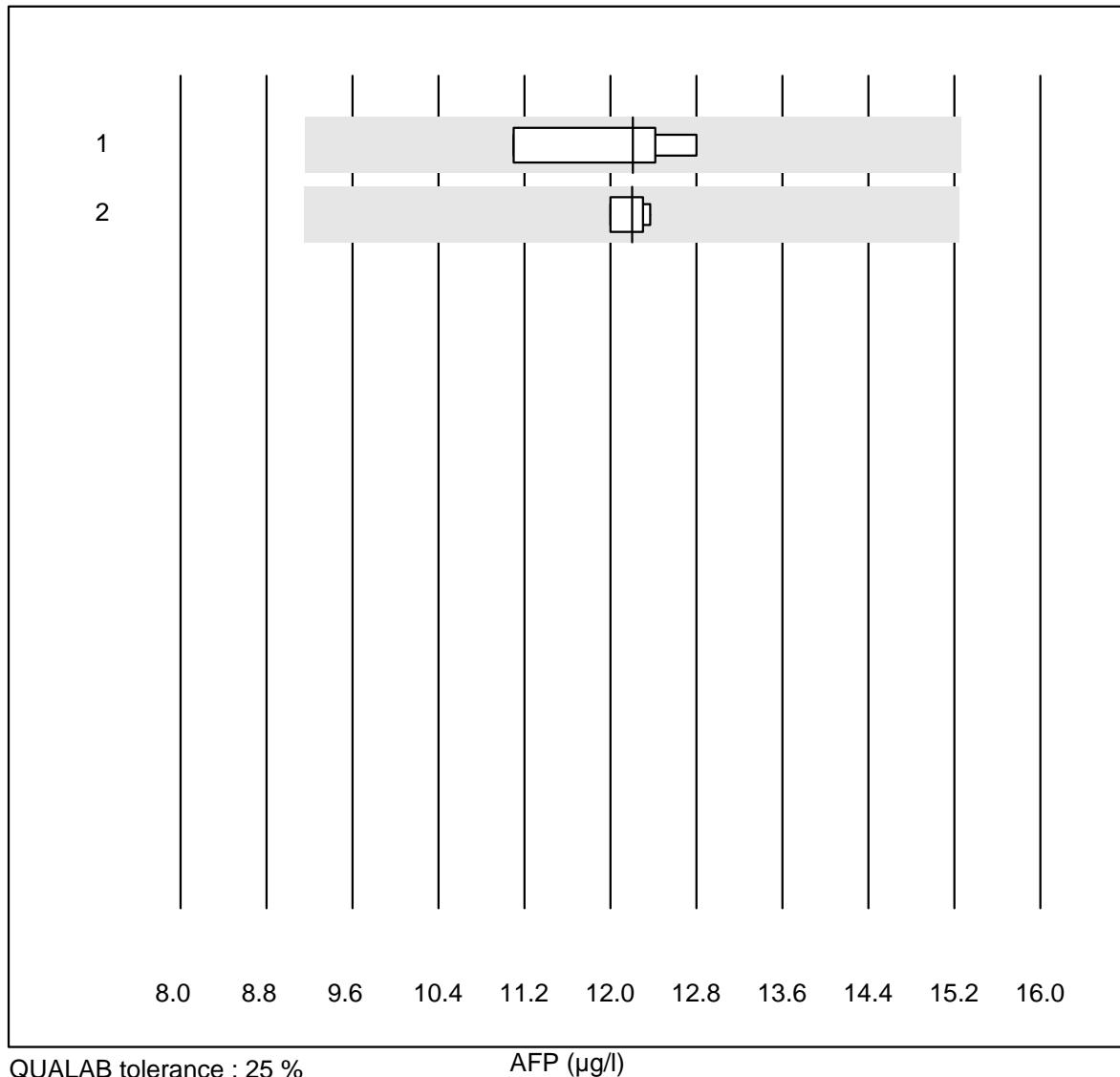
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Architect | 4 | 100.0 | 0.0 | 0.0 | 247.0 | 7.7 | a |

CA 15-3

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Architect | 4 | 100.0 | 0.0 | 0.0 | 86.0 | 5.6 | a |

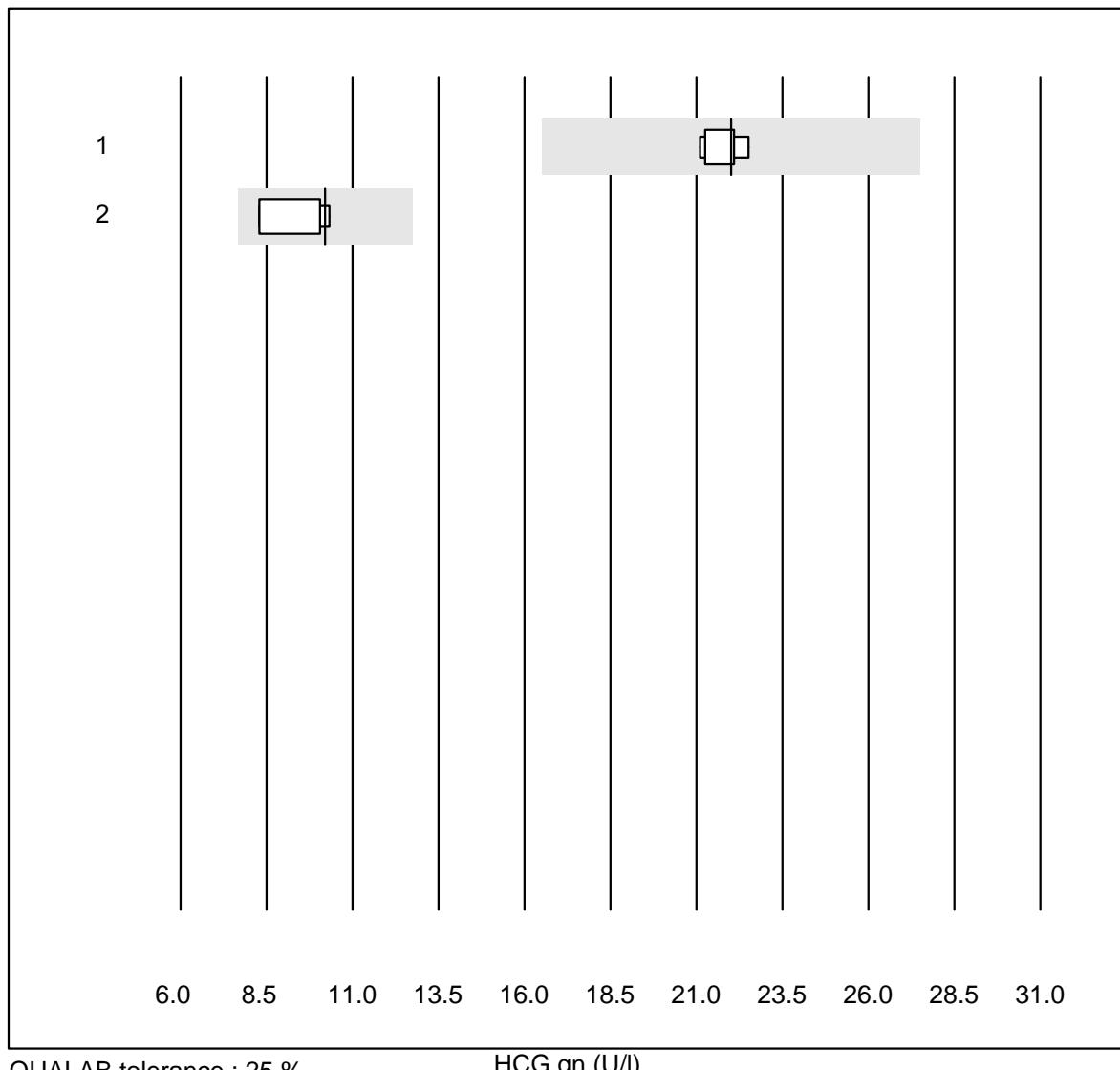
K14 Tumor Markers

AFP

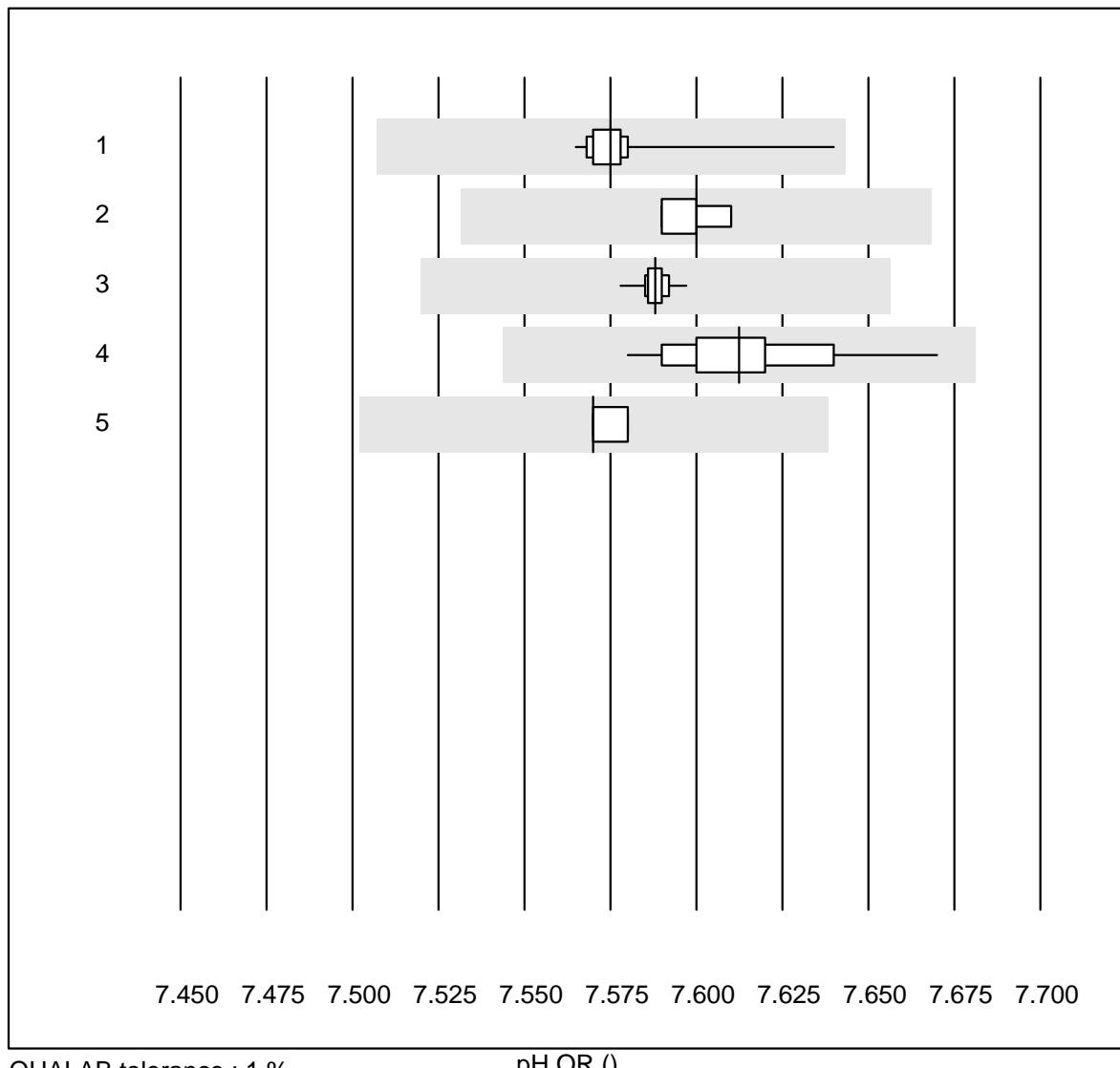


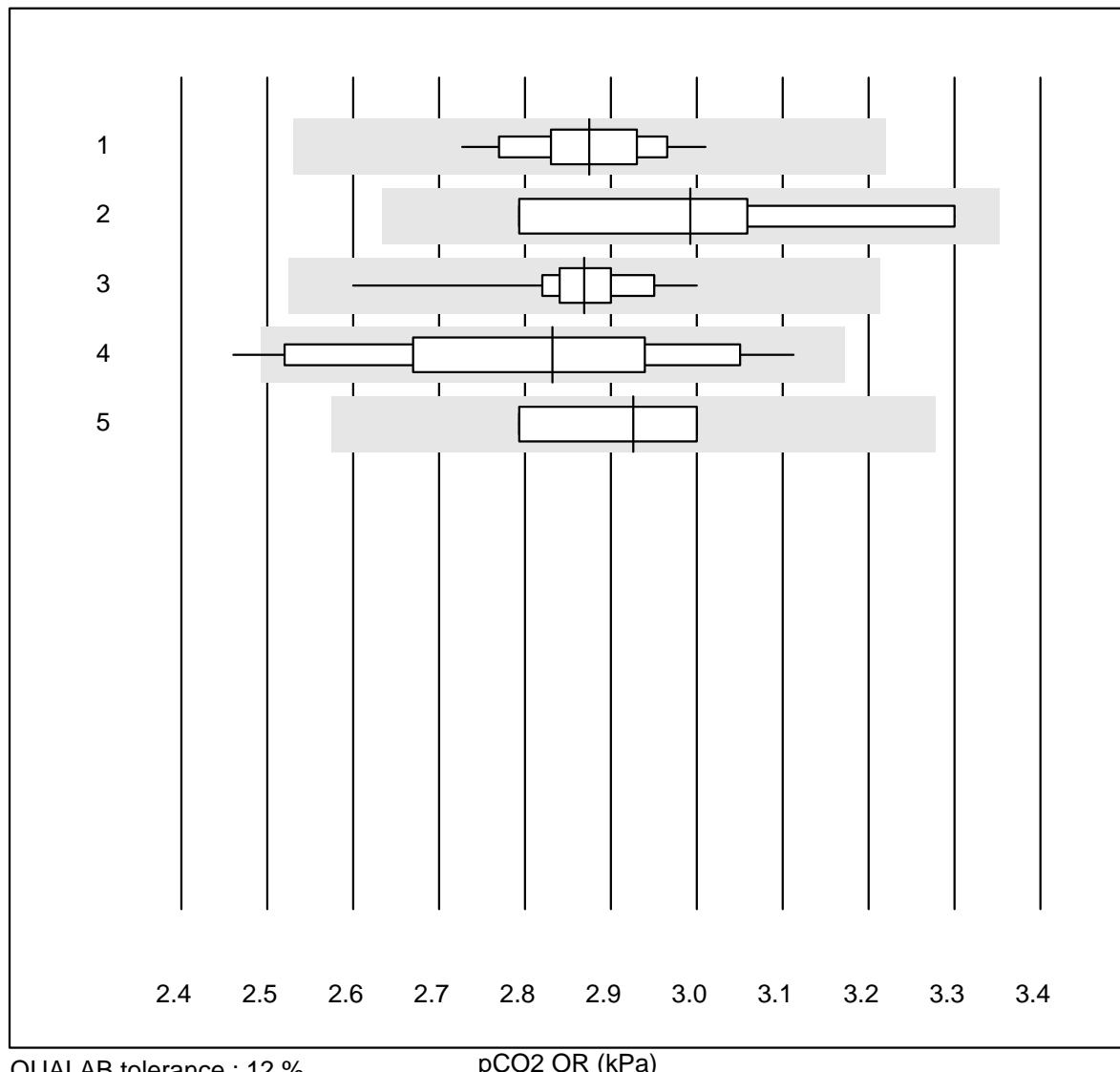
K14 Tumor Markers

HCG qn



| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|-------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Cobas E / Elecsys | 5 | 100.0 | 0.0 | 0.0 | 22 | 2.7 | a |
| 2 | Vidas | 4 | 100.0 | 0.0 | 0.0 | 10 | 9.5 | a |

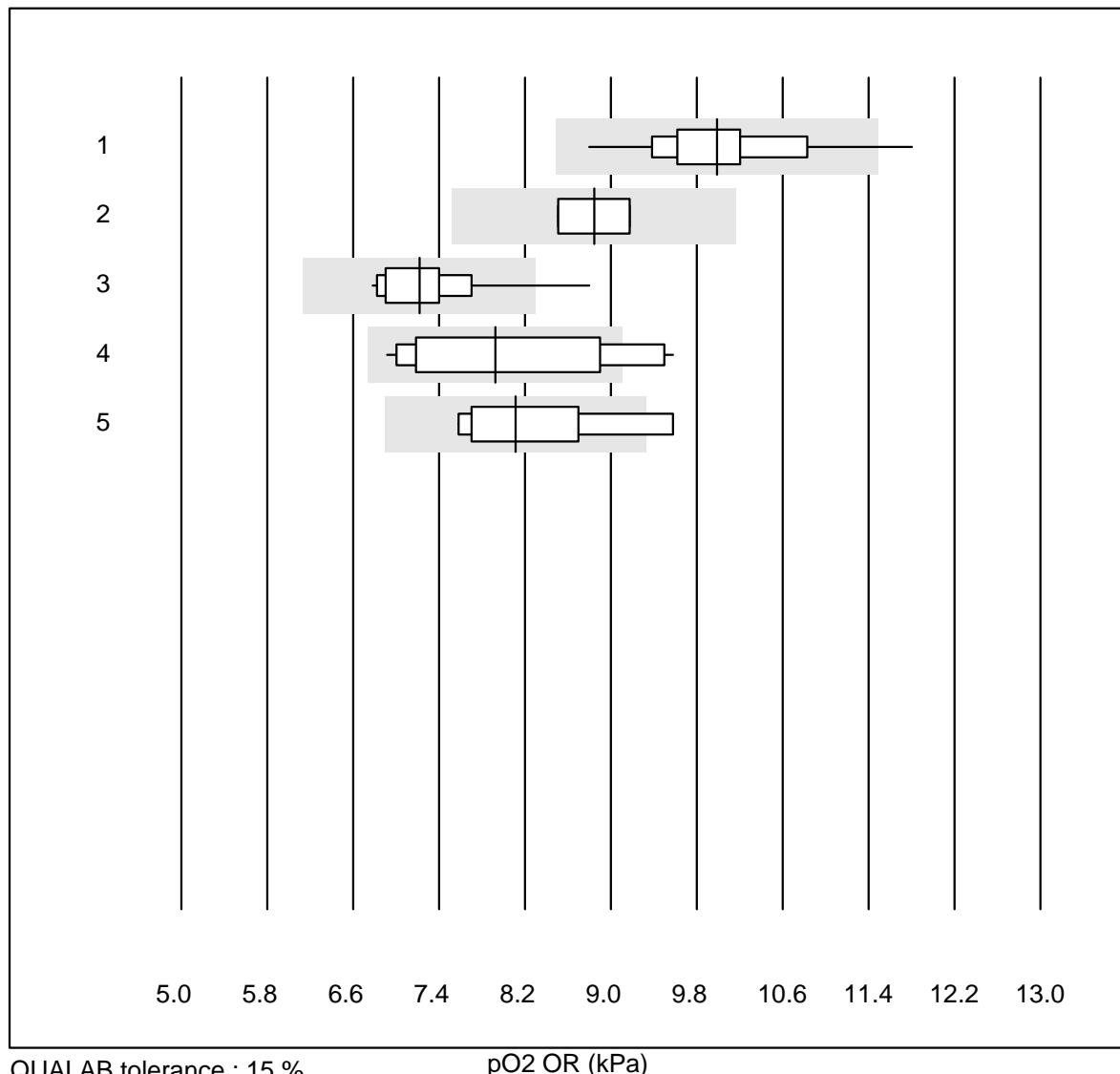
pH OR

pCO₂ OR

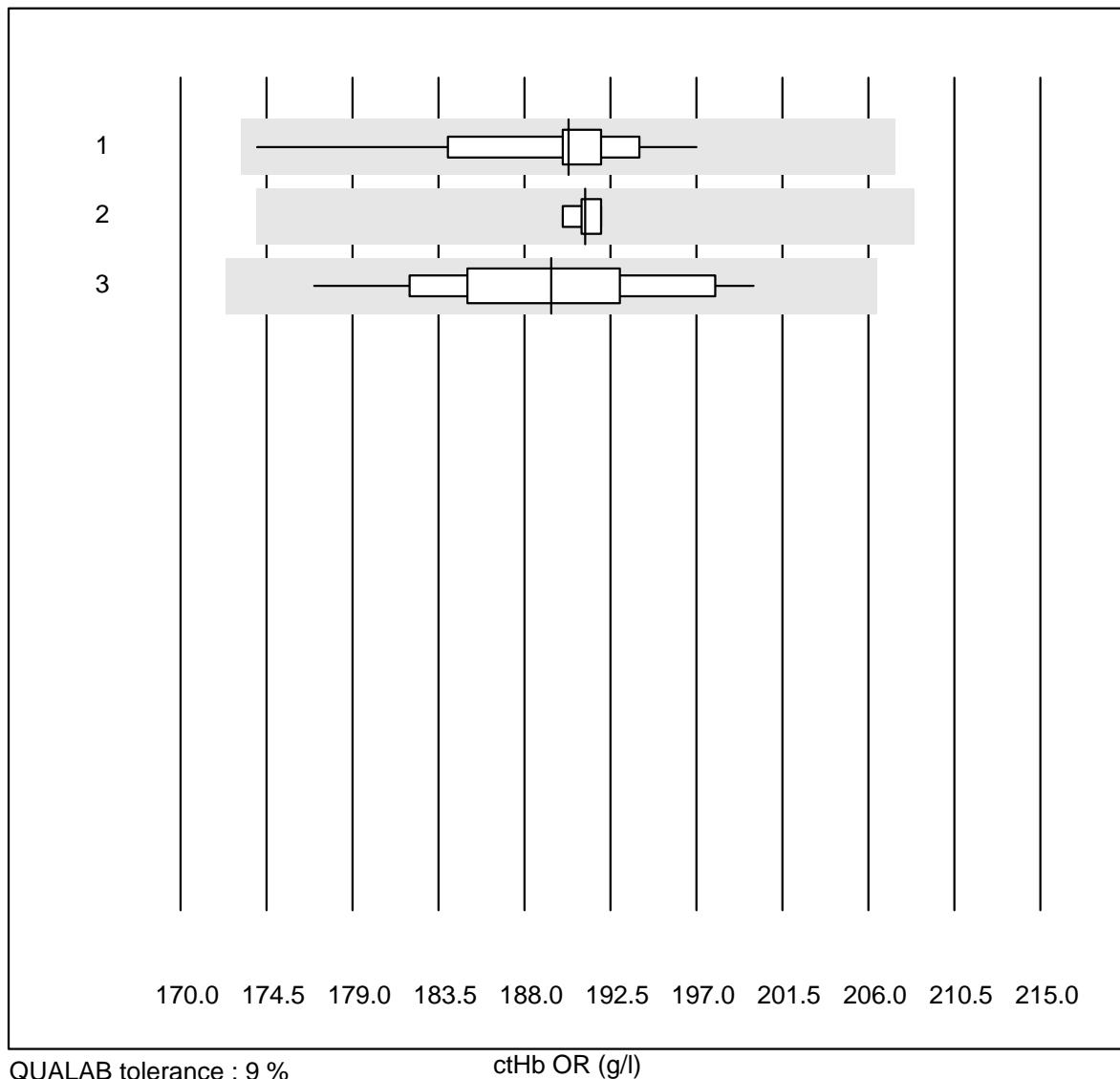
QUALAB tolerance : 12 %

pCO₂ OR (kPa)

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 ABL700/800 | 82 | 100.0 | 0.0 | 0.0 | 2.87 | 2.5 | e |
| 2 Radiometer NPT-7 | 4 | 100.0 | 0.0 | 0.0 | 2.99 | 7.2 | e* |
| 3 ABL 90 | 33 | 100.0 | 0.0 | 0.0 | 2.87 | 2.3 | e |
| 4 ABL 80 / Coox | 22 | 86.4 | 4.5 | 9.1 | 2.83 | 6.6 | e |
| 5 ABL 5 | 5 | 100.0 | 0.0 | 0.0 | 2.93 | 3.6 | e* |

pO₂ OR

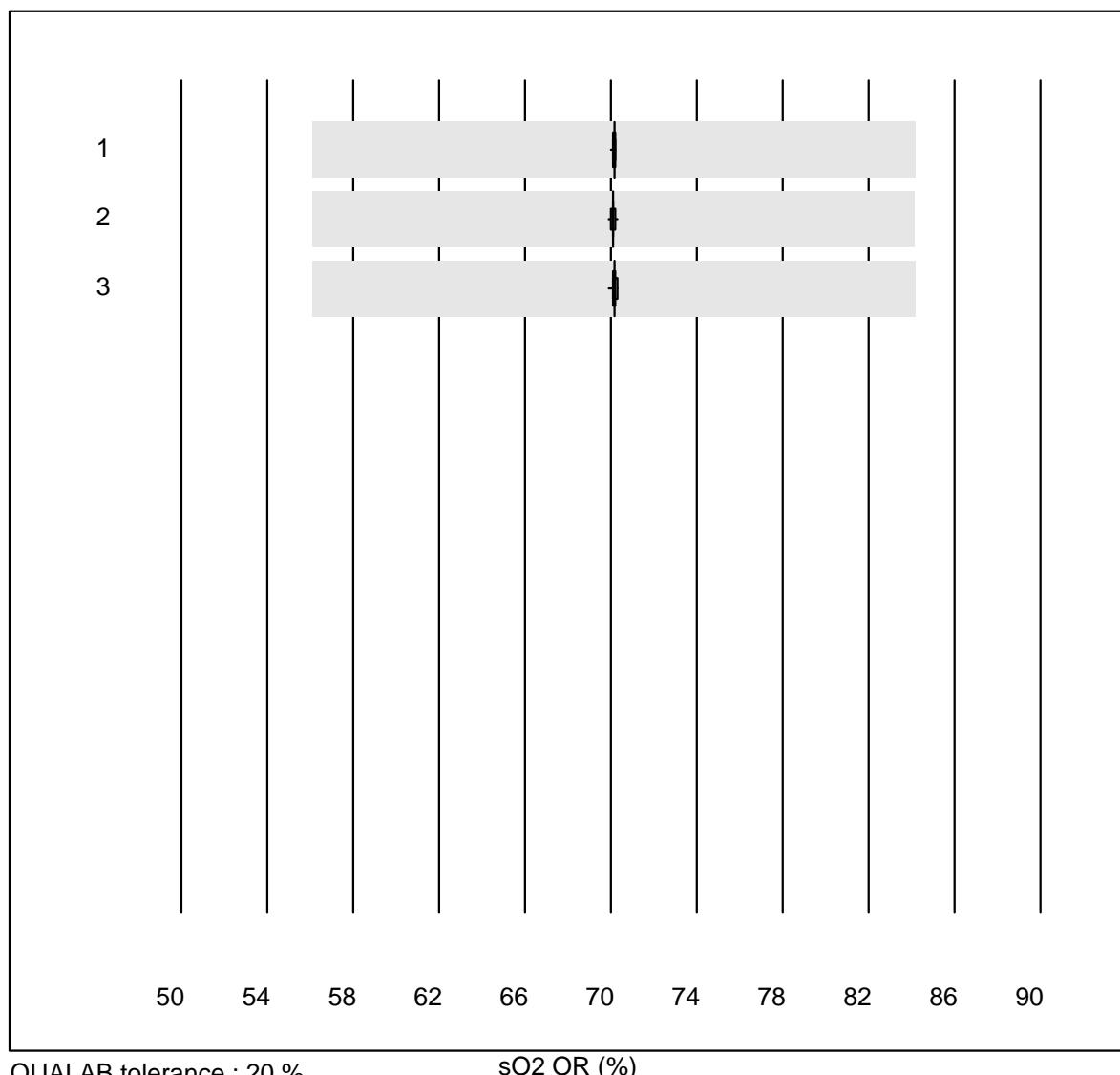
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 ABL700/800 | 82 | 96.4 | 2.4 | 1.2 | 9.99 | 5.8 | e |
| 2 Radiometer NPT-7 | 4 | 75.0 | 0.0 | 25.0 | 8.84 | 4.4 | e* |
| 3 ABL 90 | 33 | 75.8 | 3.0 | 21.2 | 7.22 | 6.3 | e |
| 4 ABL 80 / Coox | 22 | 68.2 | 13.6 | 18.2 | 7.92 | 12.1 | e* |
| 5 ABL 5 | 5 | 80.0 | 20.0 | 0.0 | 8.11 | 9.8 | e* |

ctHb OR

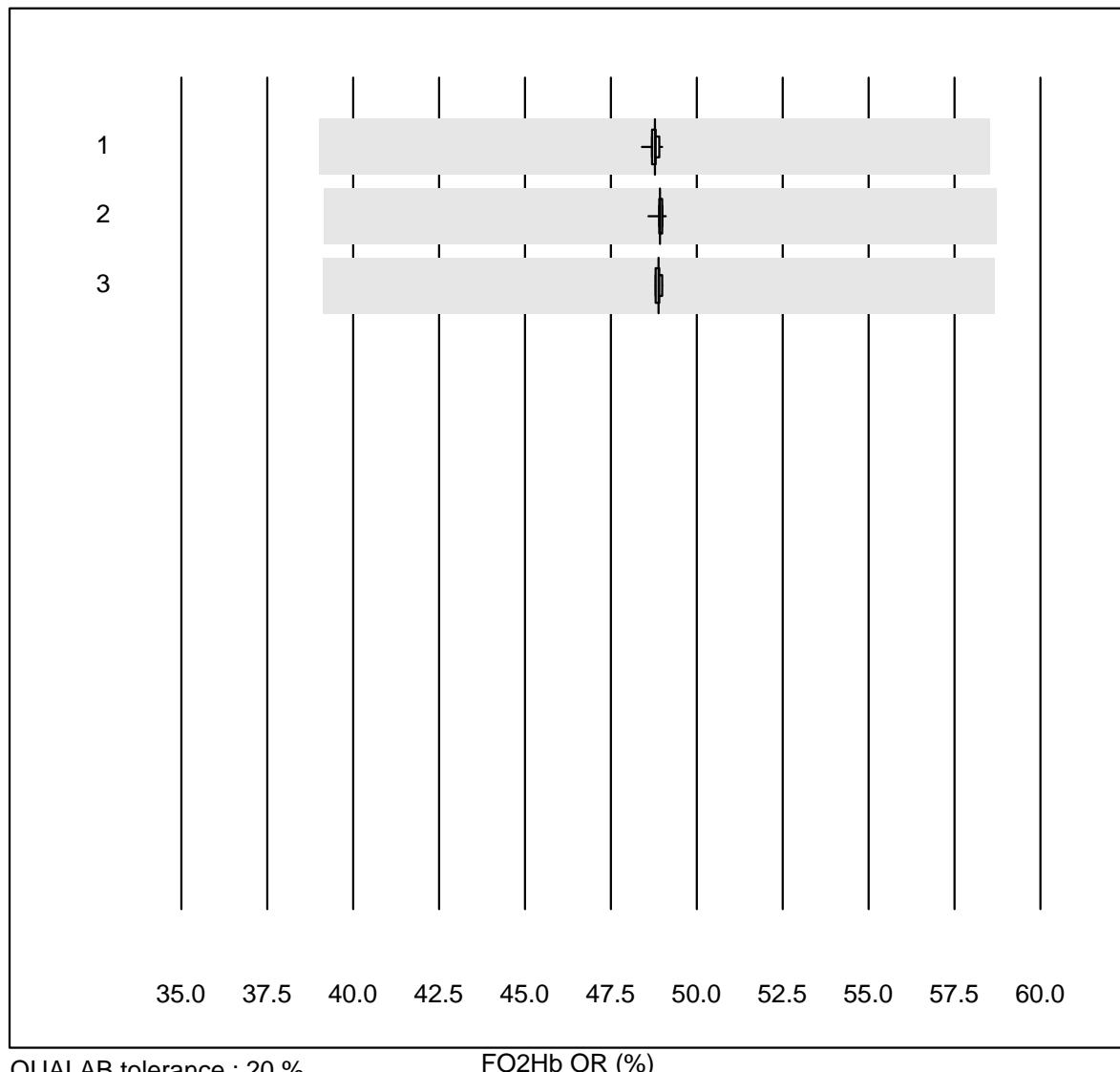
QUALAB tolerance : 9 %

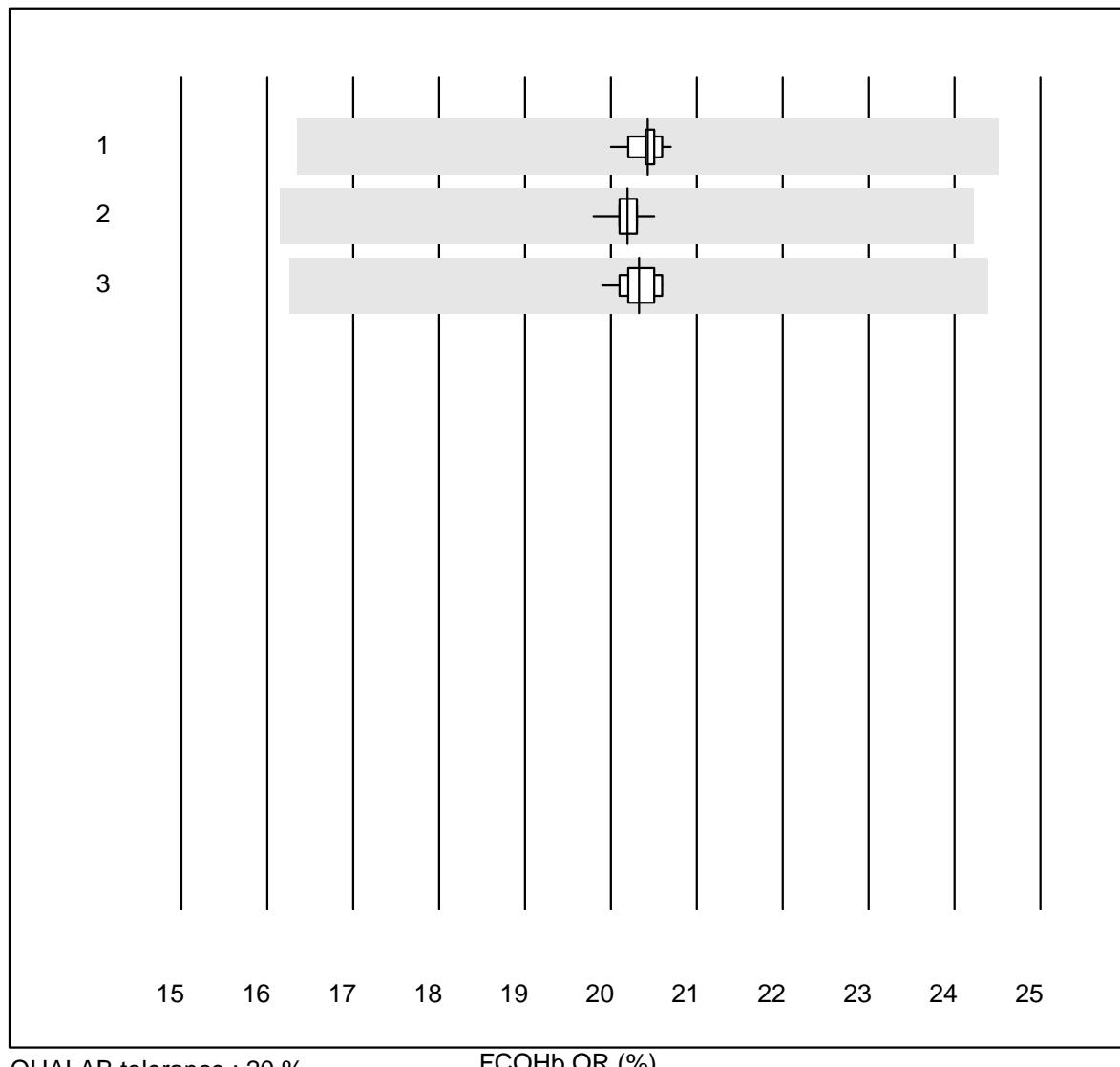
ctHb OR (g/l)

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 ABL700/800 | 71 | 100.0 | 0.0 | 0.0 | 190.3 | 2.3 | e |
| 2 ABL 90 | 33 | 93.9 | 0.0 | 6.1 | 191.2 | 0.3 | e |
| 3 ABL 80 / Coox | 15 | 100.0 | 0.0 | 0.0 | 189.4 | 3.2 | e |

sO₂ OR

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 ABL700/800 | 57 | 100.0 | 0.0 | 0.0 | 70.153 | 0.1 | e |
| 2 ABL 90 | 31 | 96.8 | 0.0 | 3.2 | 70.103 | 0.1 | e |
| 3 ABL 80 / Coox | 14 | 92.9 | 0.0 | 7.1 | 70.162 | 0.2 | e |

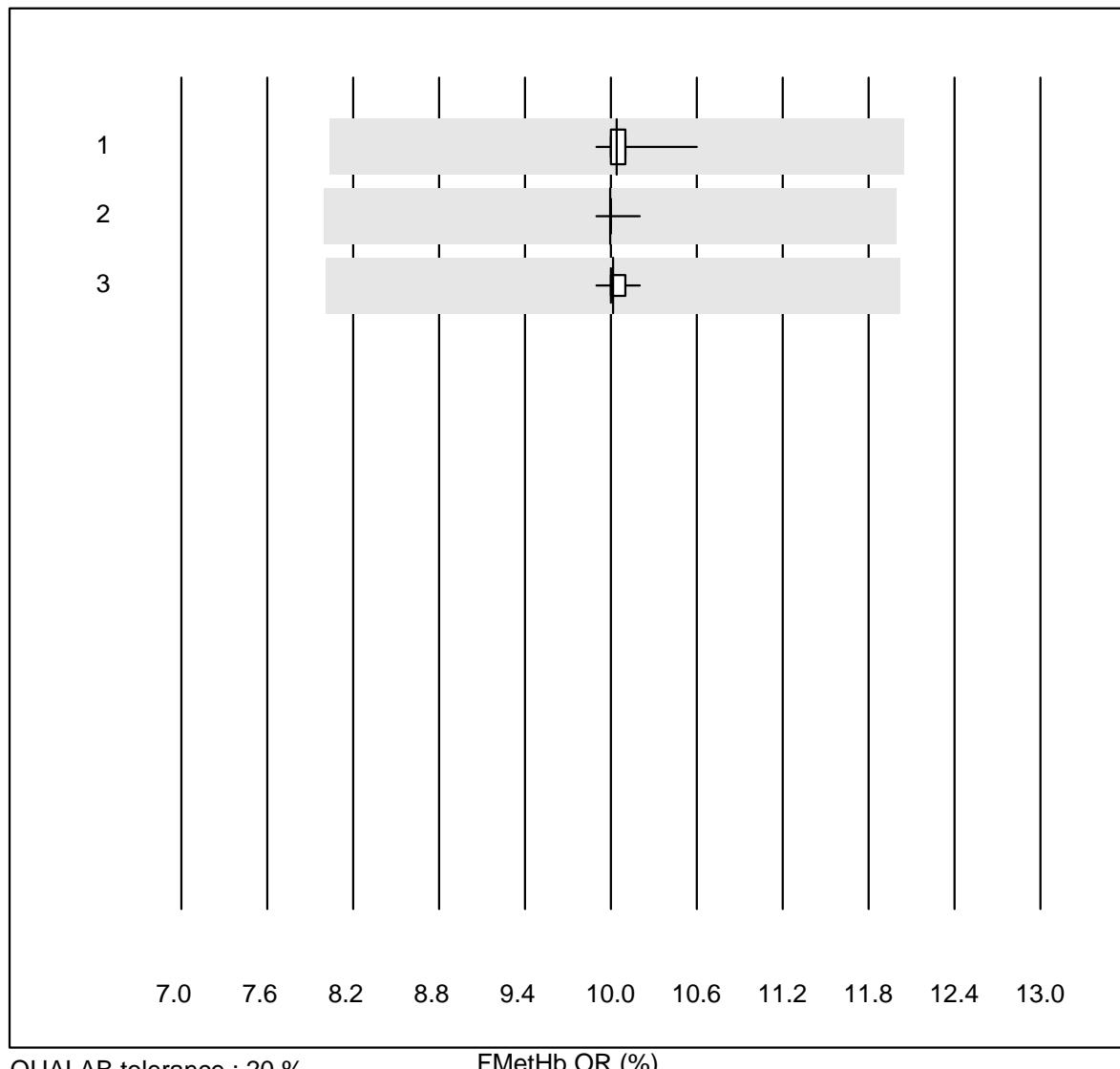
FO2Hb OR

FCOHb OR

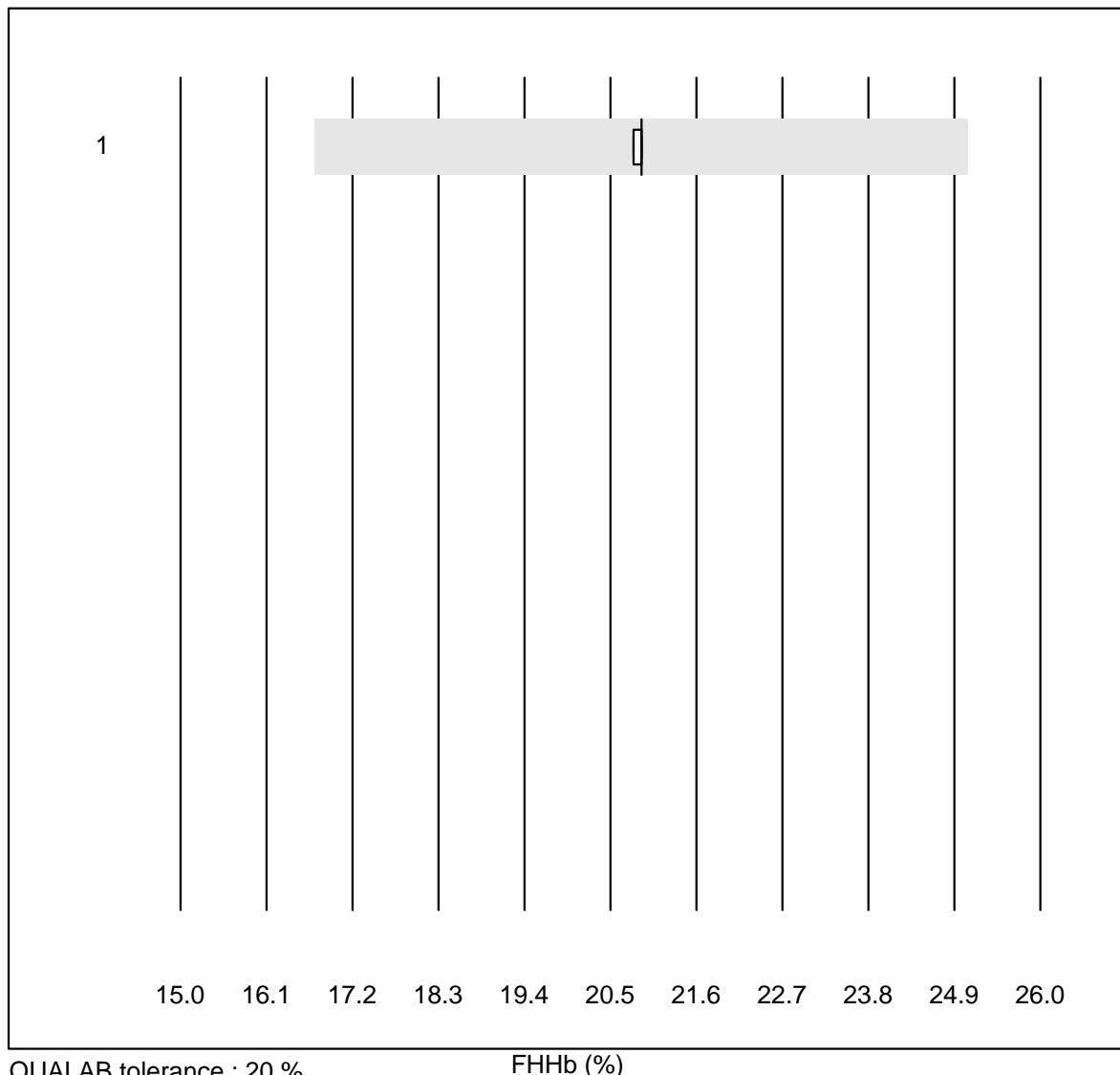
QUALAB tolerance : 20 %

FCOHb OR (%)

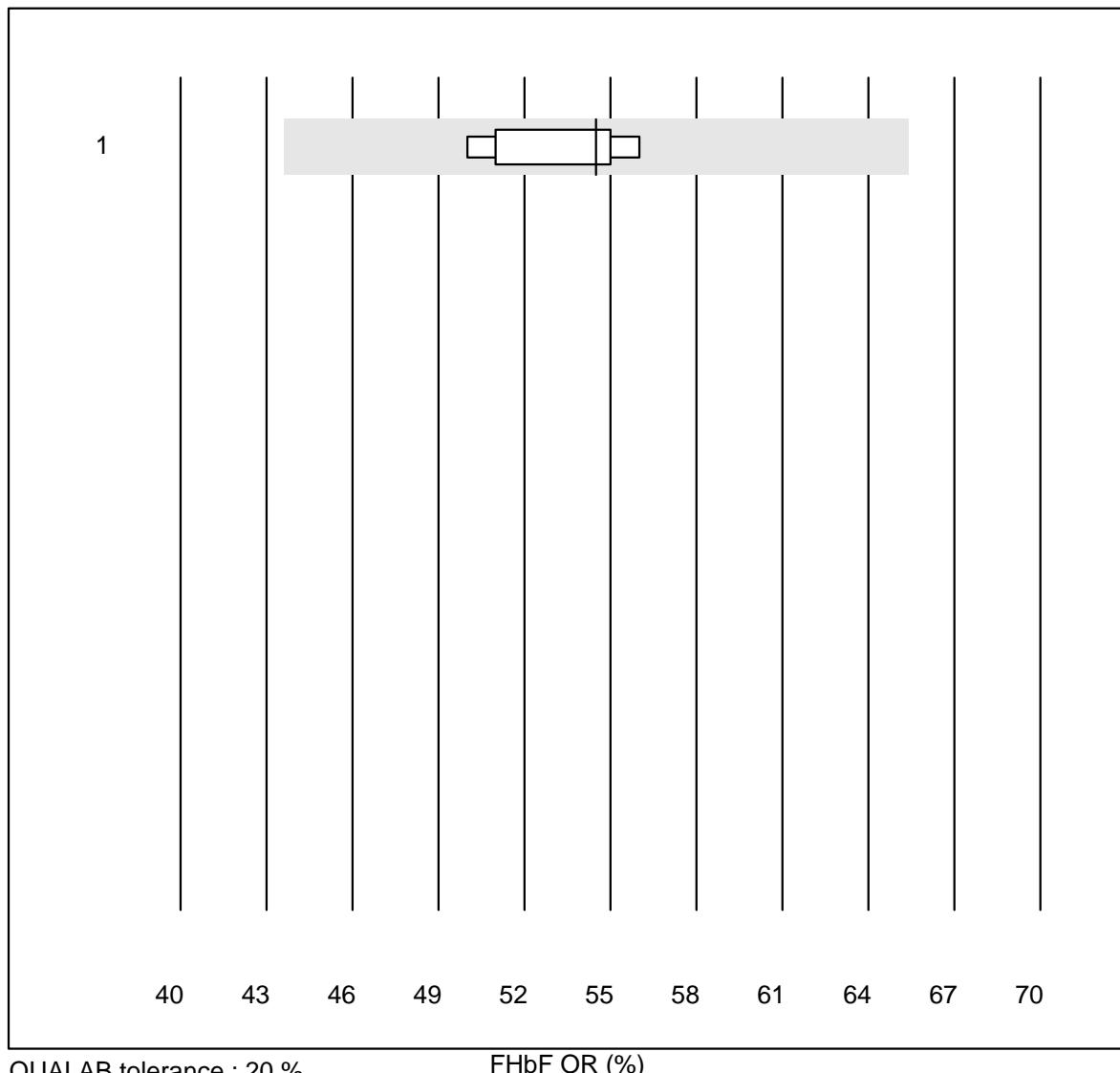
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 ABL700/800 | 52 | 100.0 | 0.0 | 0.0 | 20.427 | 0.7 | e |
| 2 ABL 90 | 31 | 100.0 | 0.0 | 0.0 | 20.190 | 0.6 | e |
| 3 ABL 80 / Coox | 15 | 100.0 | 0.0 | 0.0 | 20.327 | 1.0 | e |

FMetHb OR

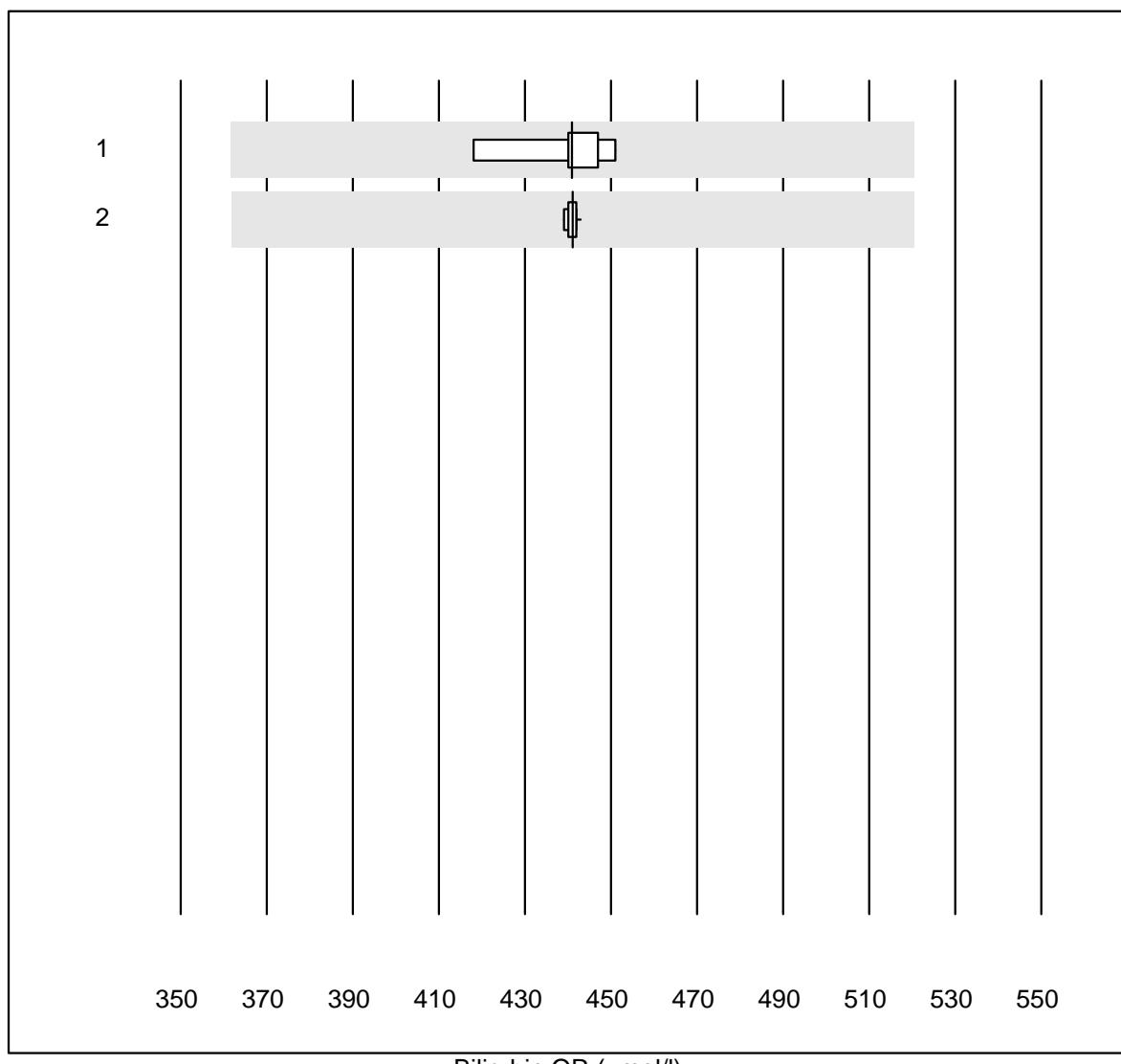
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 ABL700/800 | 54 | 100.0 | 0.0 | 0.0 | 10.041 | 1.2 | e |
| 2 ABL 90 | 31 | 100.0 | 0.0 | 0.0 | 9.997 | 0.5 | e |
| 3 ABL 80 / Coox | 15 | 100.0 | 0.0 | 0.0 | 10.013 | 0.6 | e |

FHHb

| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | ABL 80 / Coox | 5 | 100.0 | 0.0 | 0.0 | 20.900 | 0.3 | e |

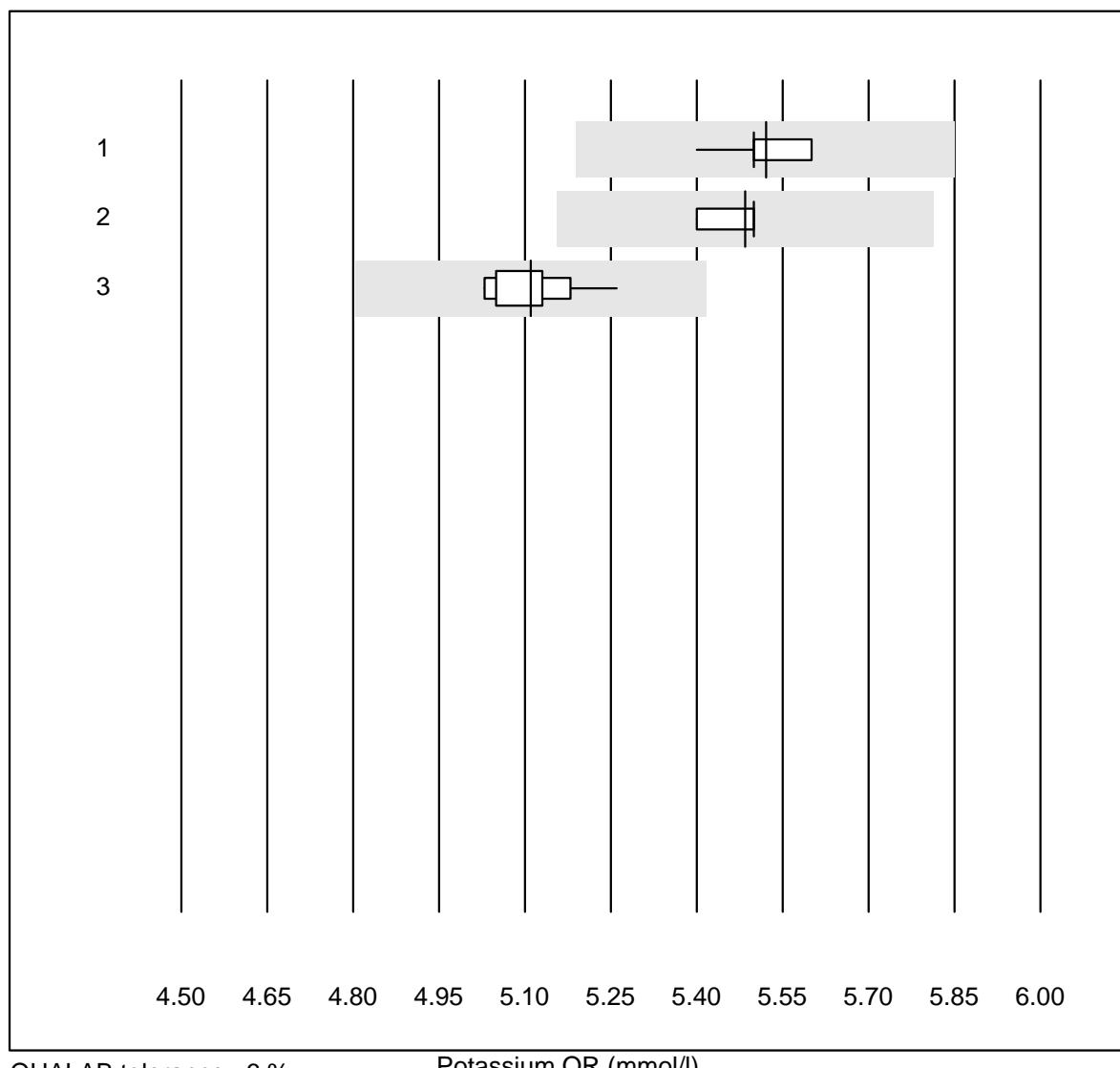
FHbF OR

| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | ABL 90 | 8 | 100.0 | 0.0 | 0.0 | 54.500 | 4.0 | e |

Bilirubin OR

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 ABL700/800 | 5 | 100.0 | 0.0 | 0.0 | 441.0 | 2.9 | e |
| 2 ABL 90 | 12 | 100.0 | 0.0 | 0.0 | 441.1 | 0.3 | e |

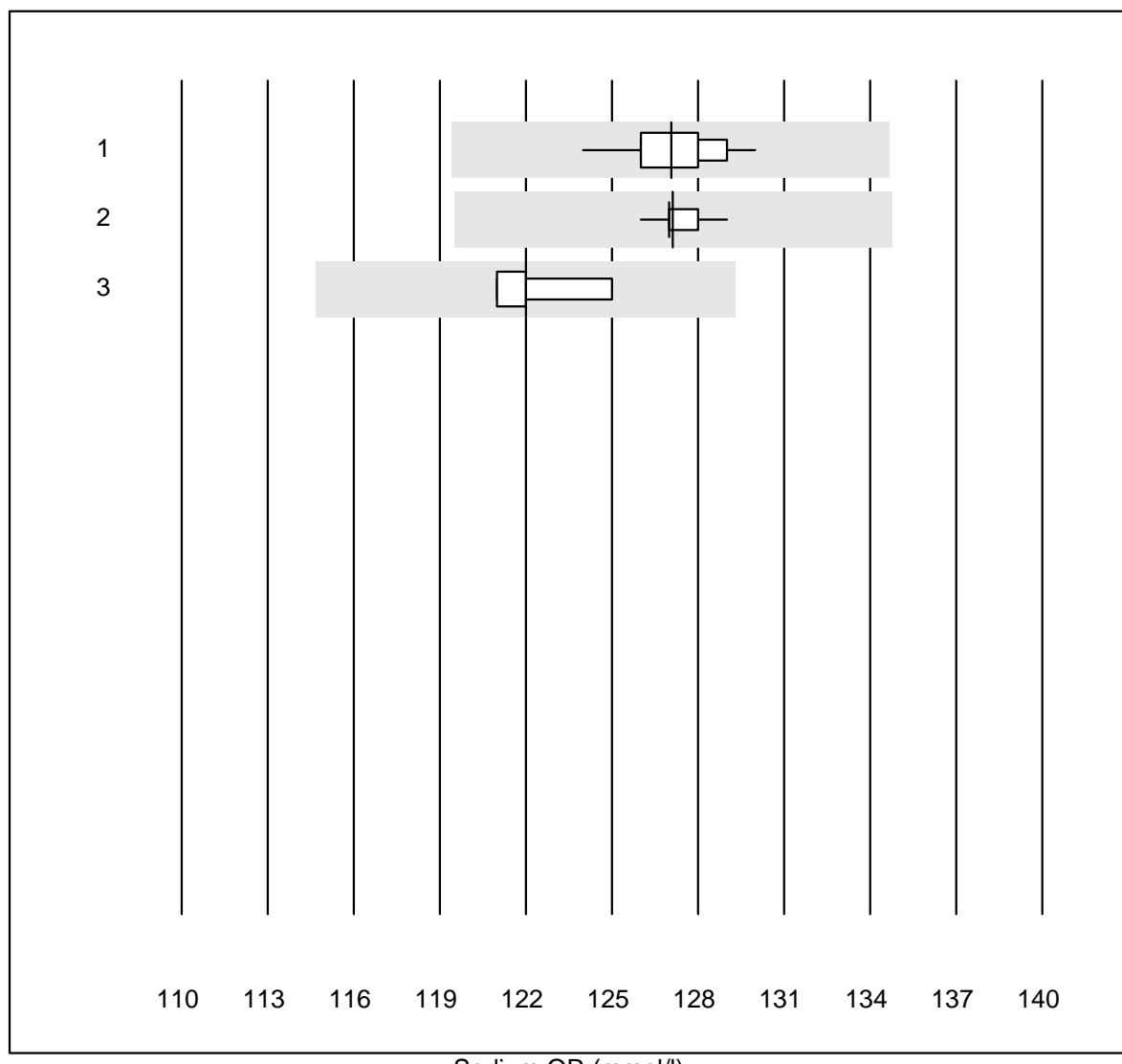
Potassium OR



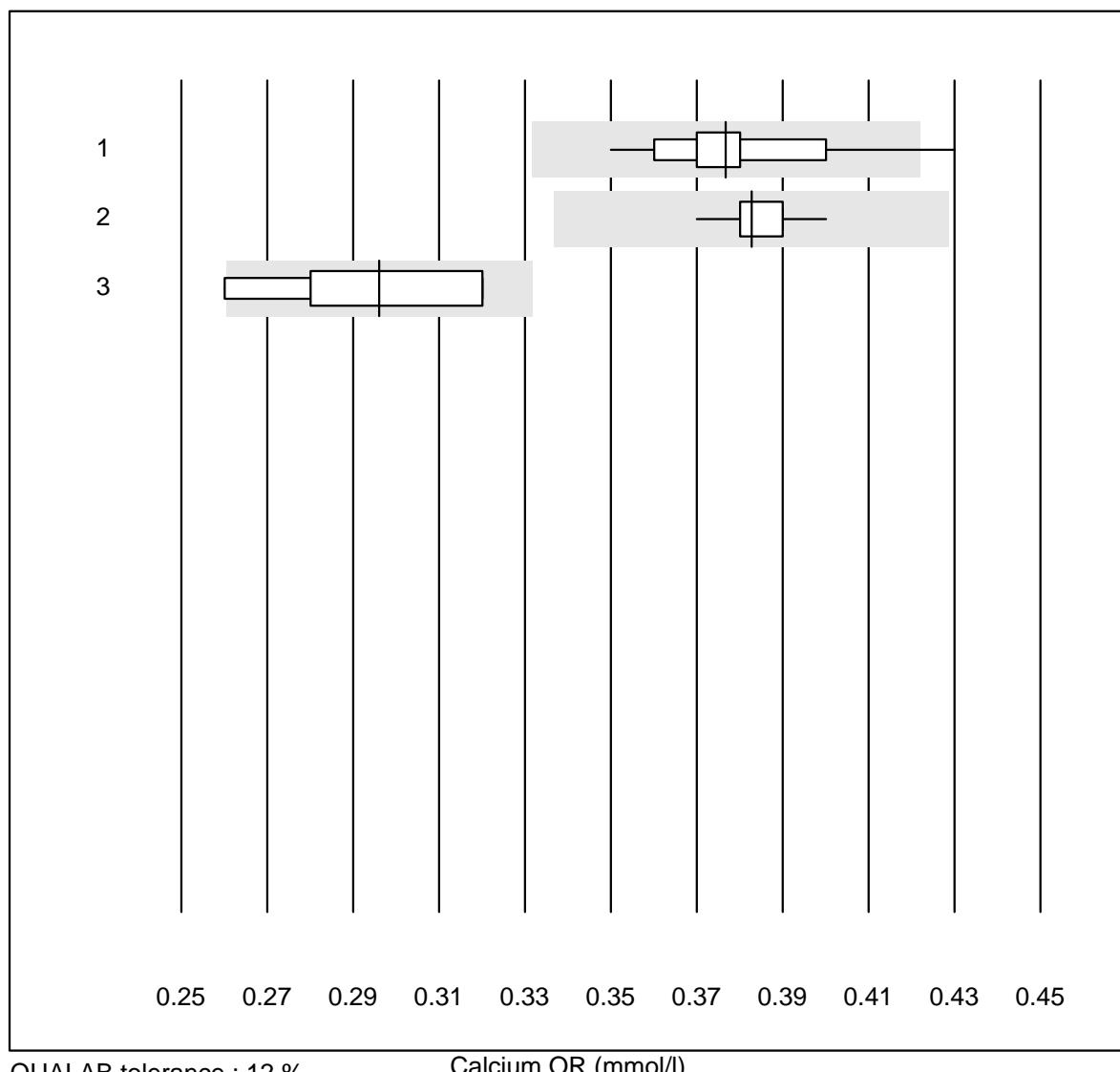
QUALAB tolerance : 6 %

Potassium OR (mmol/l)

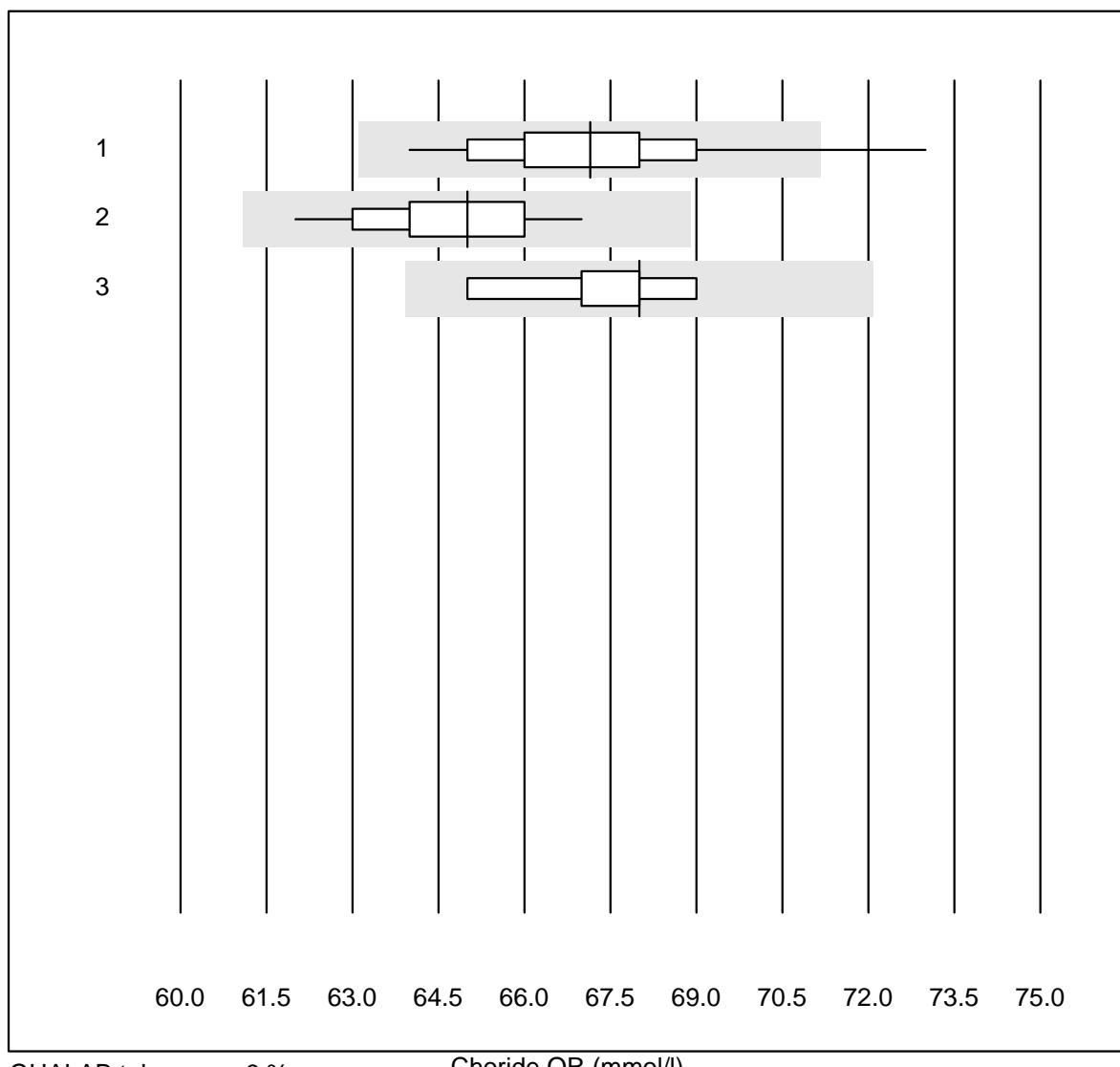
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 ABL700/800 | 68 | 100.0 | 0.0 | 0.0 | 5.5 | 0.9 | e |
| 2 ABL 90 | 33 | 100.0 | 0.0 | 0.0 | 5.5 | 0.7 | e |
| 3 ABL 80 / Coox | 10 | 100.0 | 0.0 | 0.0 | 5.1 | 1.4 | e |

Sodium OR

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 ABL700/800 | 65 | 100.0 | 0.0 | 0.0 | 127.1 | 0.9 | e |
| 2 ABL 90 | 33 | 100.0 | 0.0 | 0.0 | 127.1 | 0.5 | e |
| 3 ABL 80 / Coox | 8 | 100.0 | 0.0 | 0.0 | 122.0 | 1.2 | e |

Calcium OR

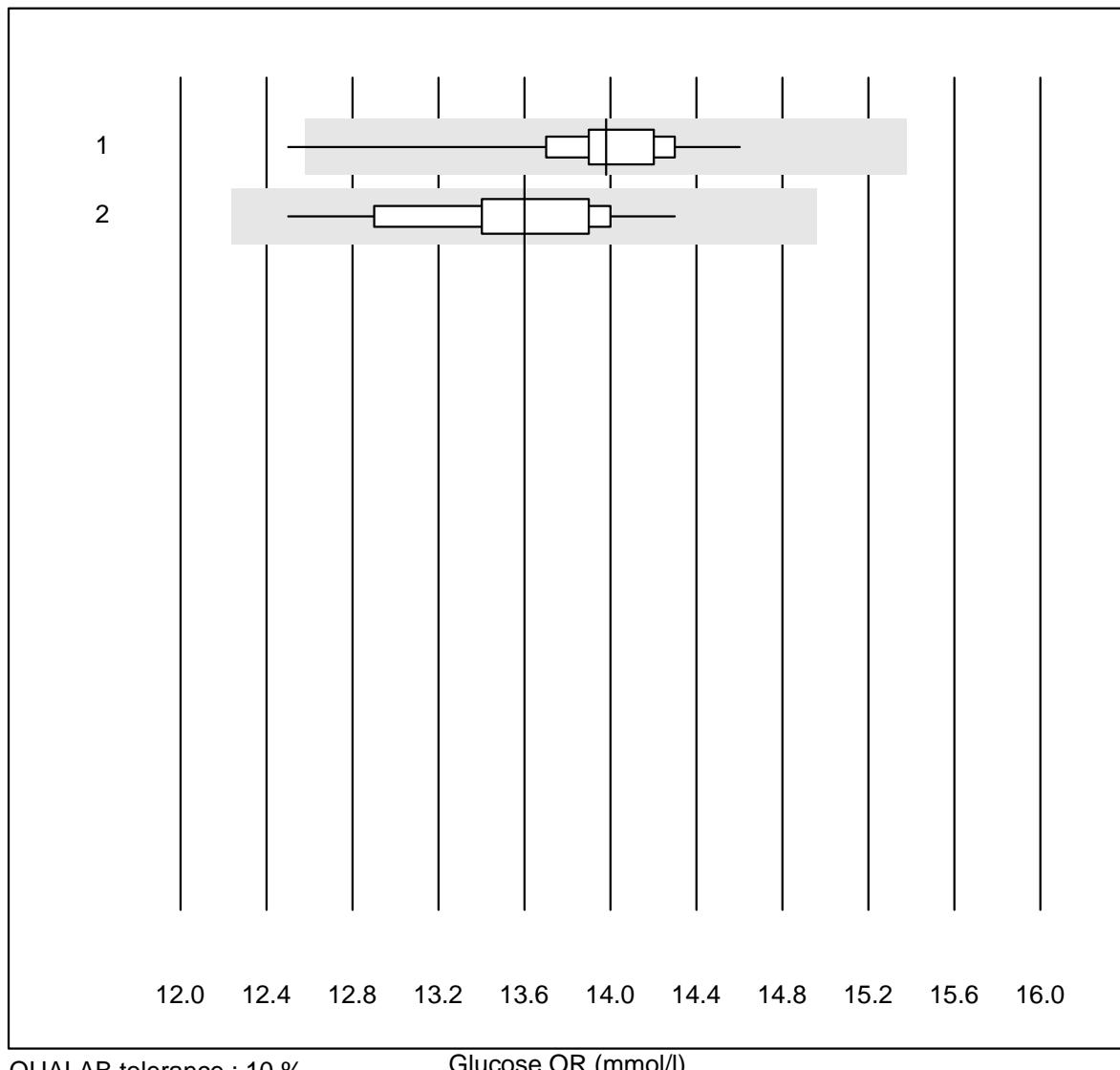
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 ABL700/800 | 66 | 98.5 | 1.5 | 0.0 | 0.38 | 4.1 | e |
| 2 ABL 90 | 33 | 100.0 | 0.0 | 0.0 | 0.38 | 1.8 | e |
| 3 ABL 80 / Coox | 8 | 62.5 | 12.5 | 25.0 | 0.30 | 8.2 | a |

Choride OR

QUALAB tolerance : 6 %

Choride OR (mmol/l)

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 ABL700/800 | 55 | 96.4 | 3.6 | 0.0 | 67.15 | 2.6 | e |
| 2 ABL 90 | 33 | 100.0 | 0.0 | 0.0 | 65.00 | 1.9 | e |
| 3 ABL 80 / Coox | 7 | 71.4 | 0.0 | 28.6 | 68.00 | 2.2 | e* |

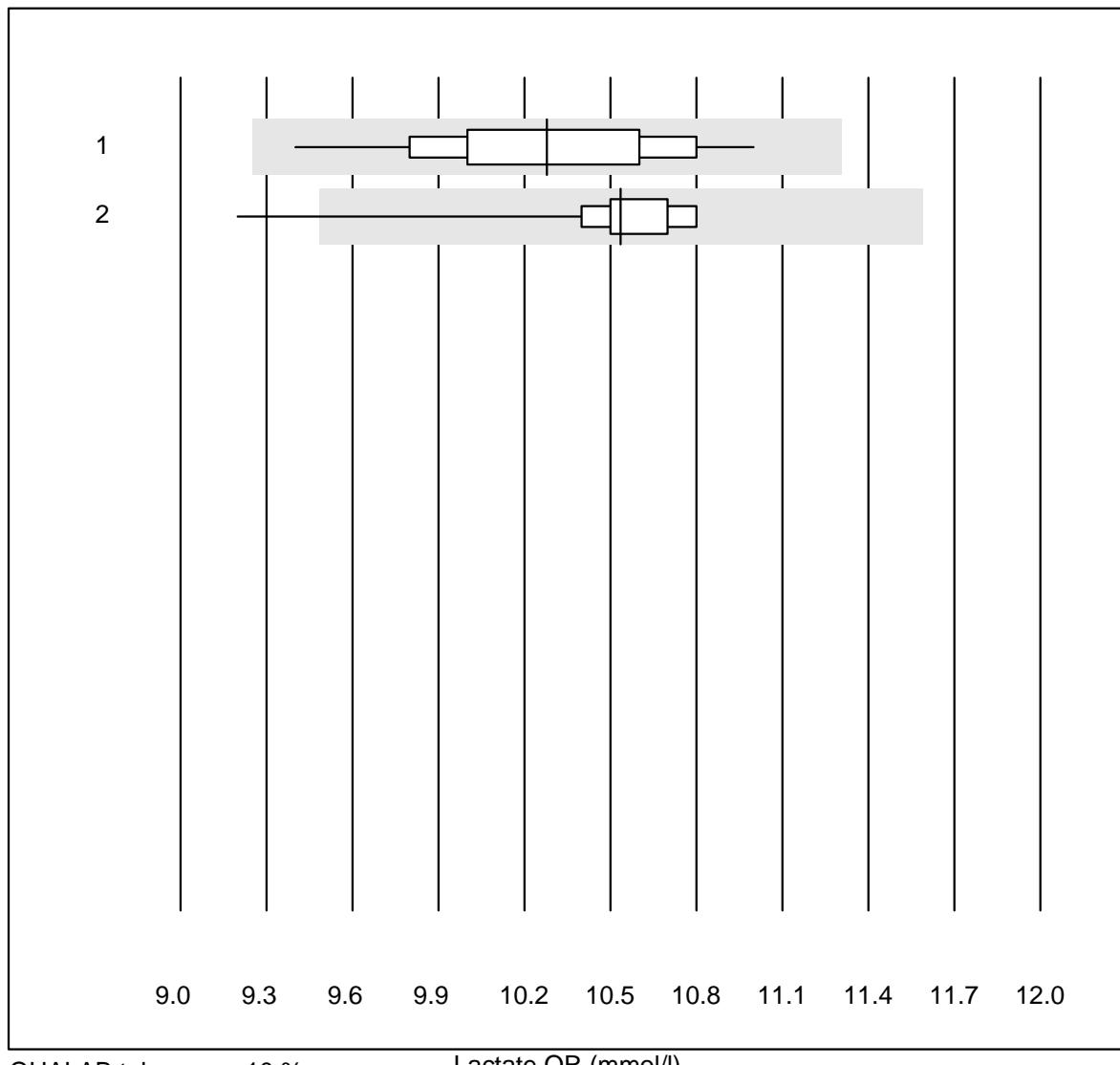
Glucose OR

QUALAB tolerance : 10 %

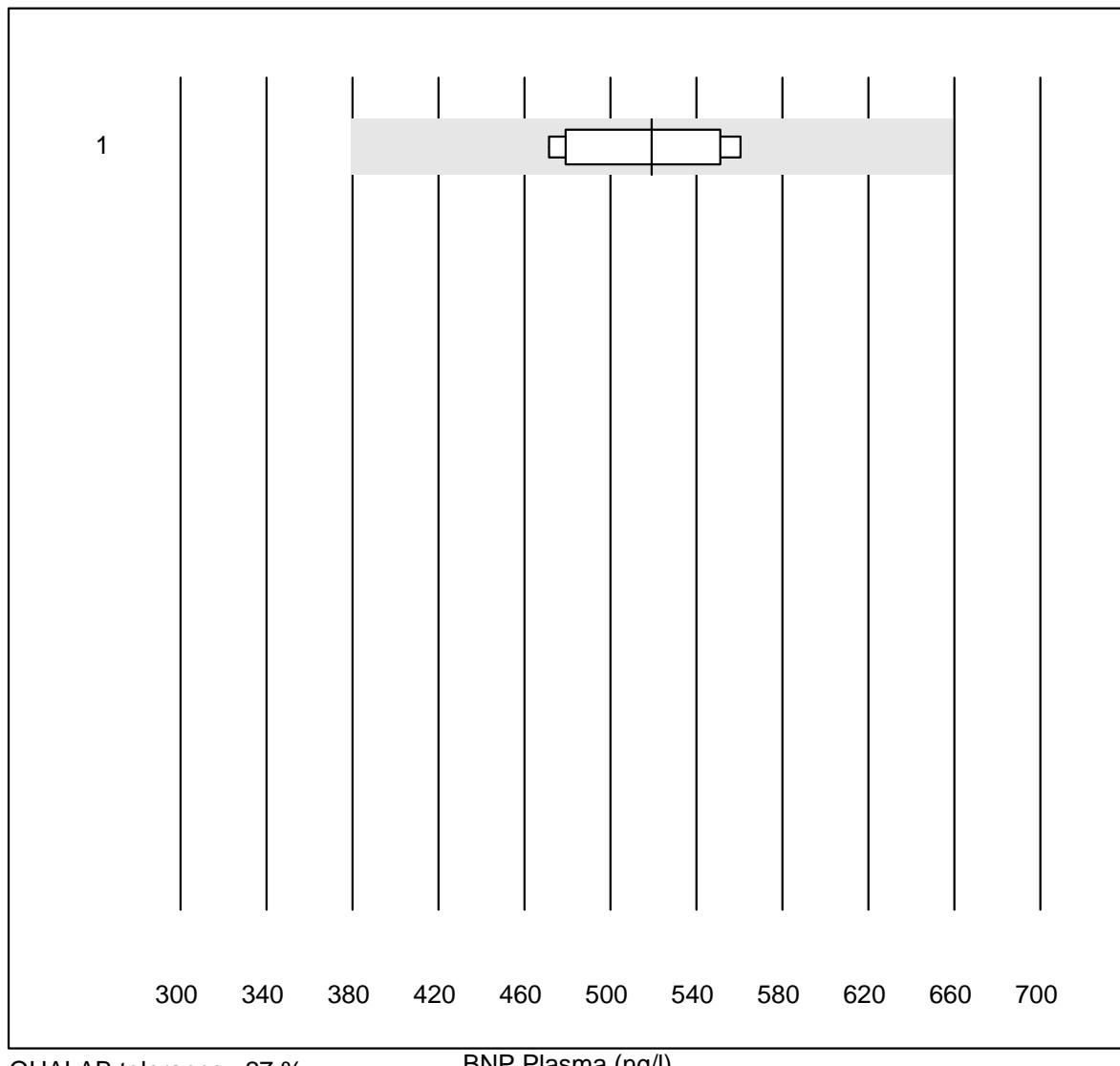
Glucose OR (mmol/l)

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 ABL700/800 | 67 | 98.5 | 1.5 | 0.0 | 14.0 | 2.3 | e |
| 2 ABL 90 | 33 | 100.0 | 0.0 | 0.0 | 13.6 | 3.3 | e |

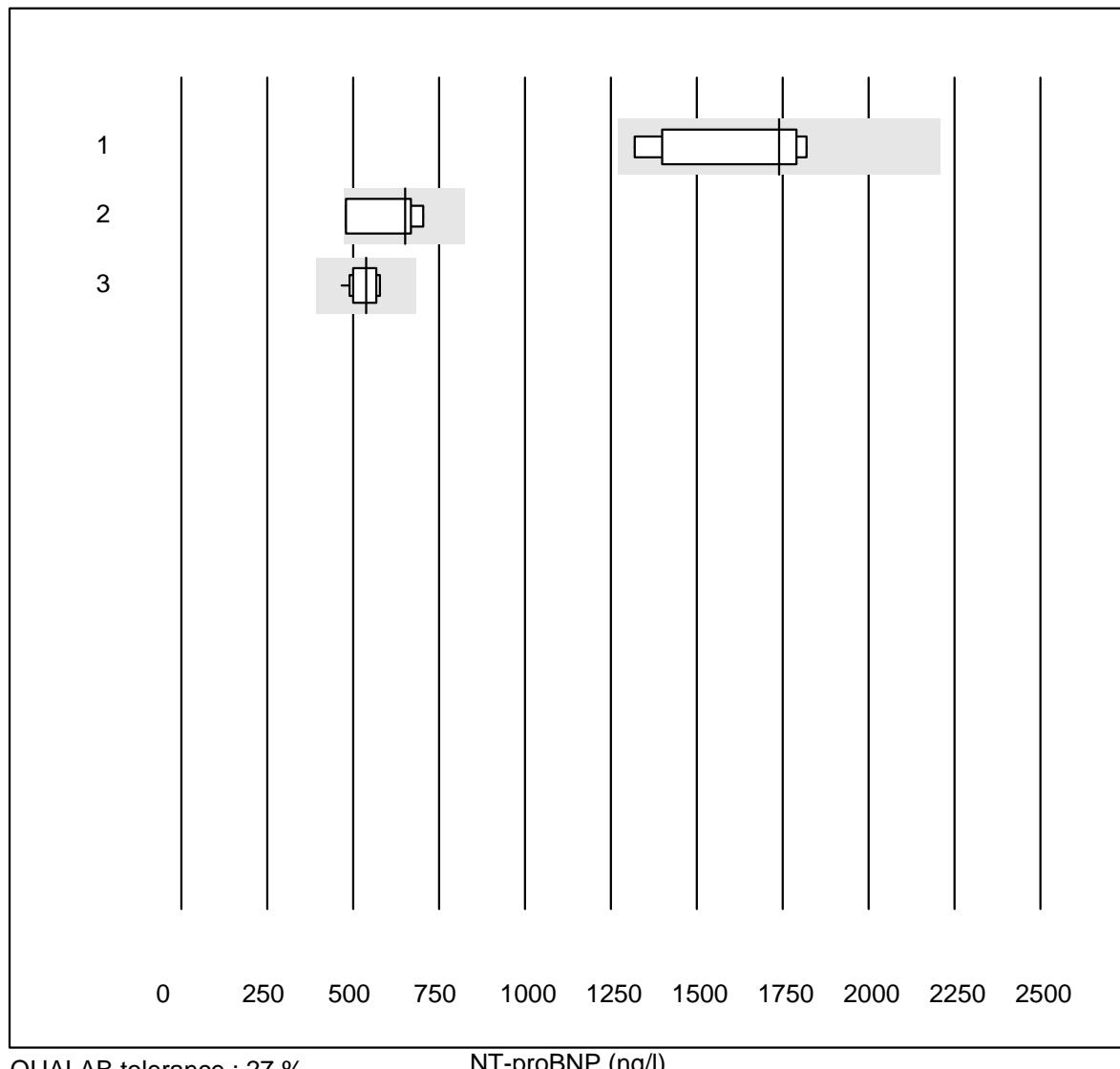
Lactate OR



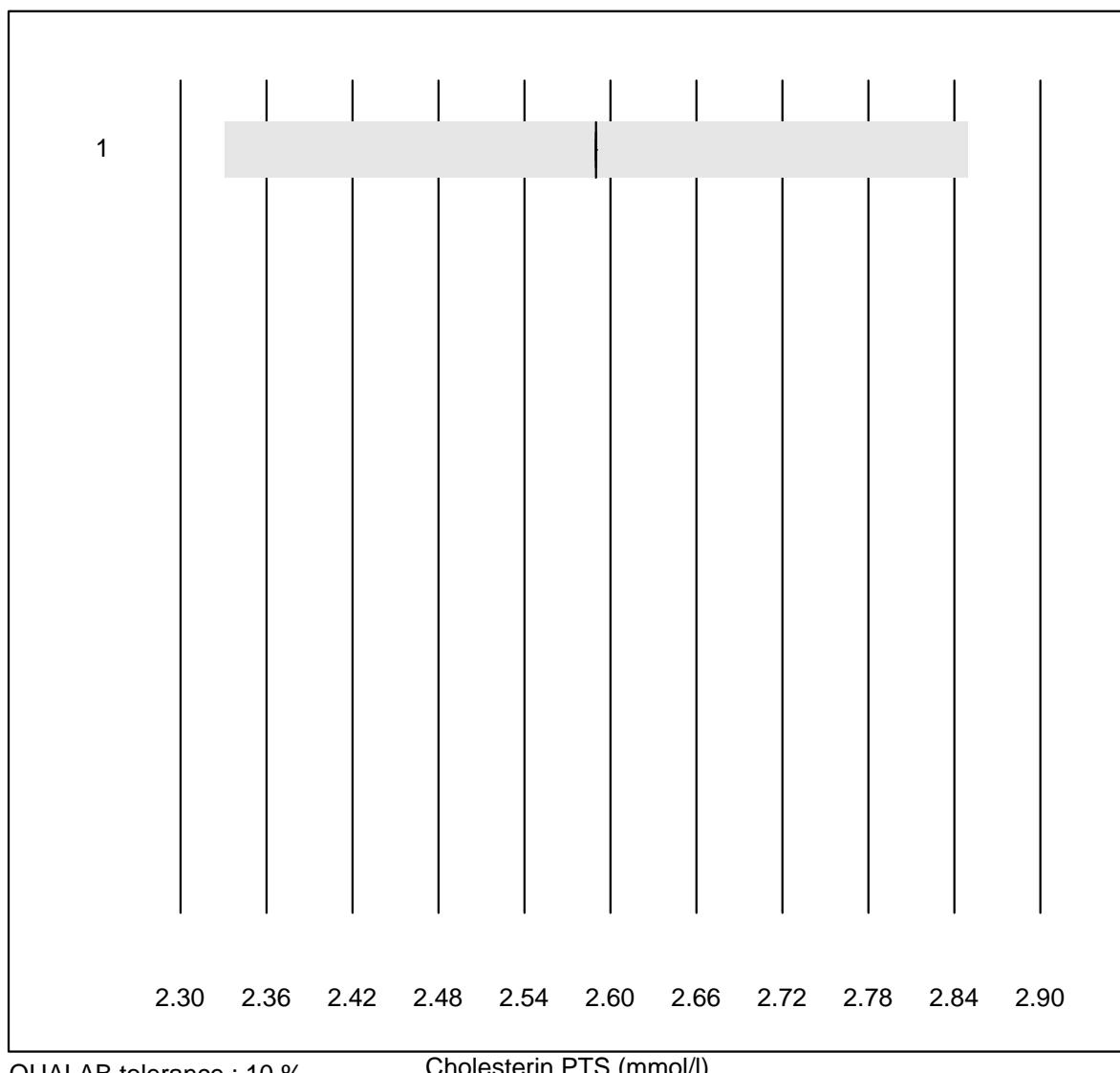
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 ABL700/800 | 71 | 100.0 | 0.0 | 0.0 | 10.28 | 4.0 | e |
| 2 ABL 90 | 33 | 97.0 | 3.0 | 0.0 | 10.54 | 2.8 | e |

BNP Plasma

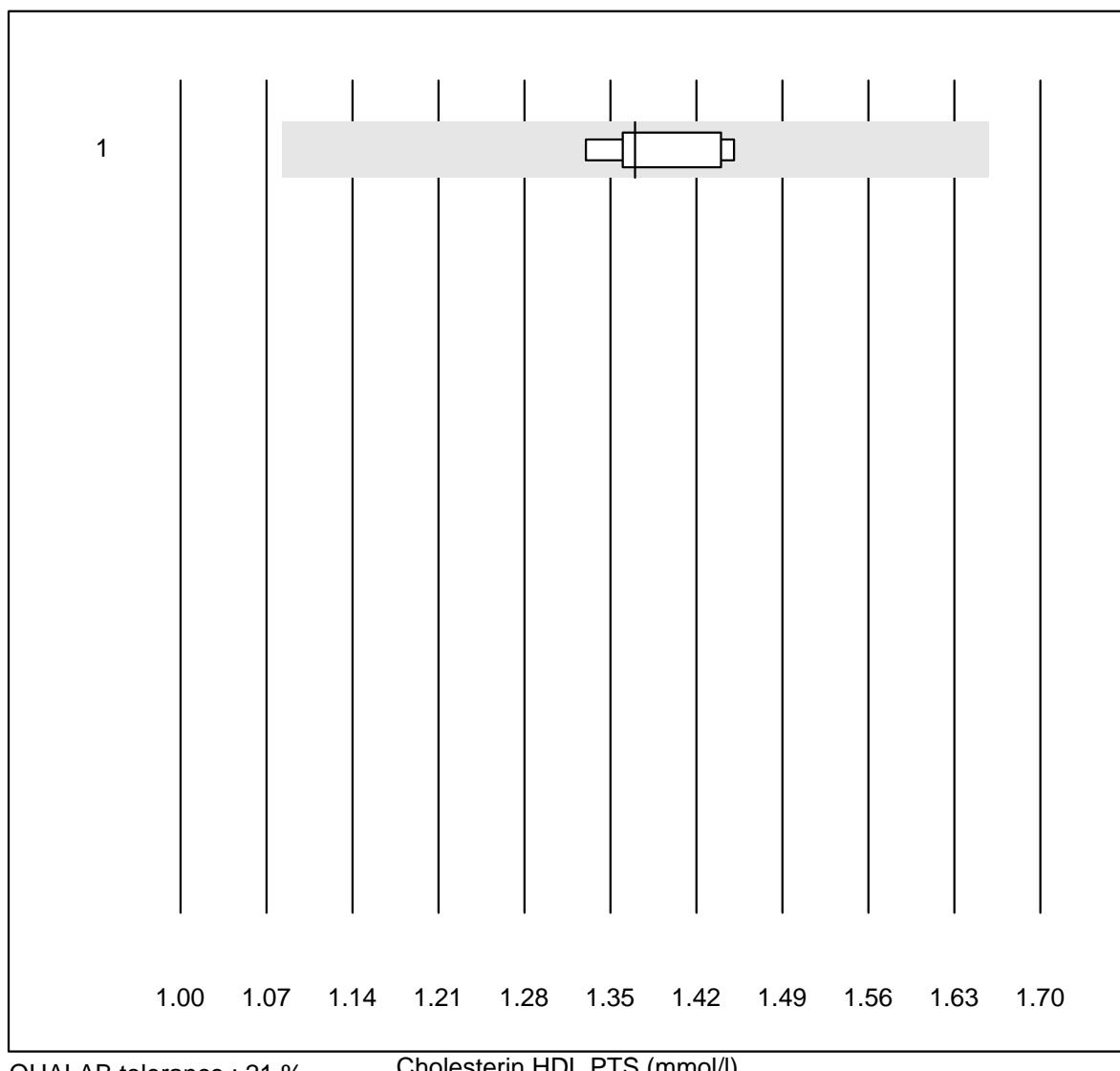
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | ADVIA Centaur XP/CP | 5 | 100.0 | 0.0 | 0.0 | 519.0 | 7.9 | e* |

NT-proBNP

Cholesterin PTS

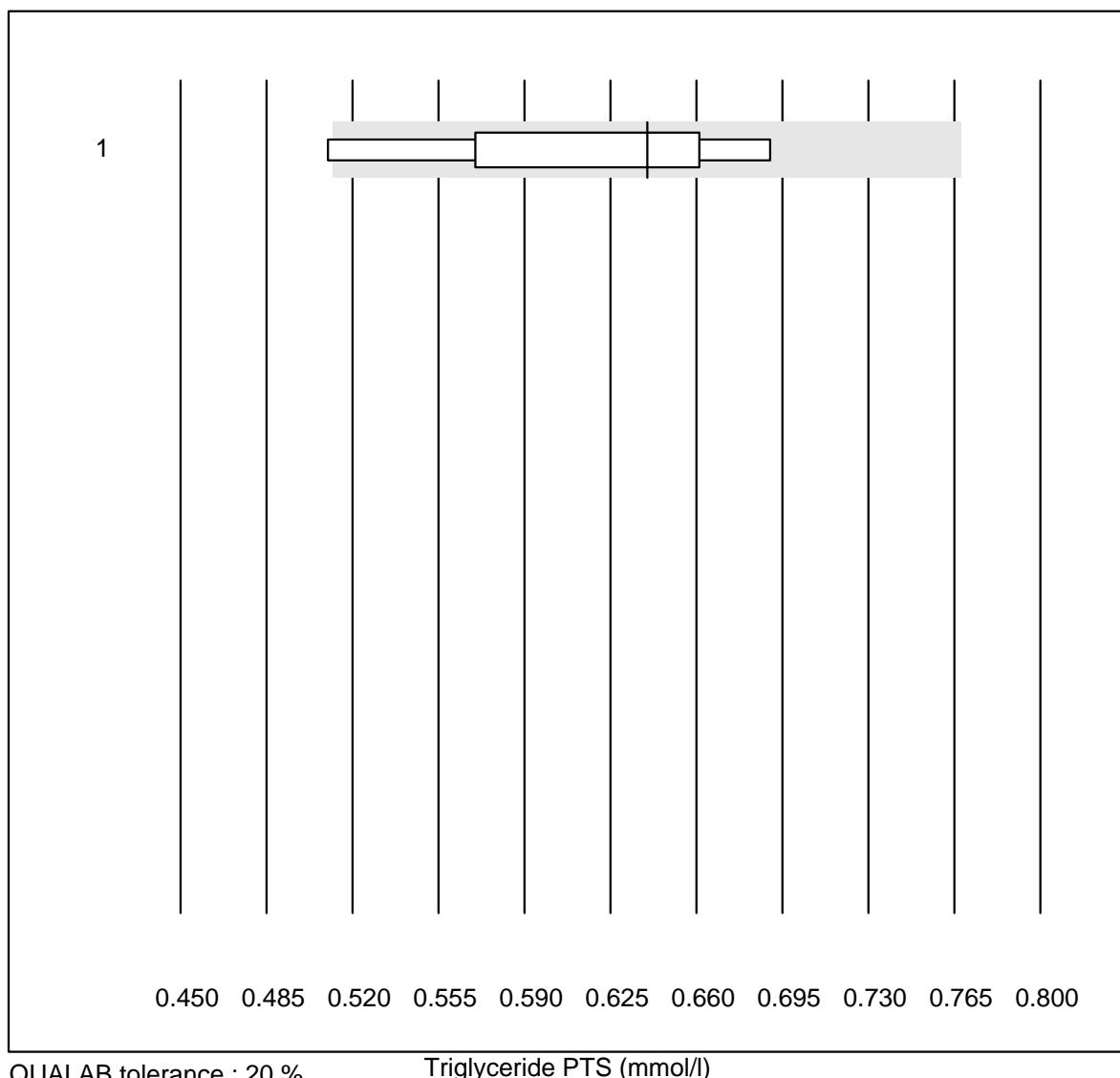


| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | CardioChek | 7 | 85.7 | 0.0 | 14.3 | 2.59 | 0.0 | e |

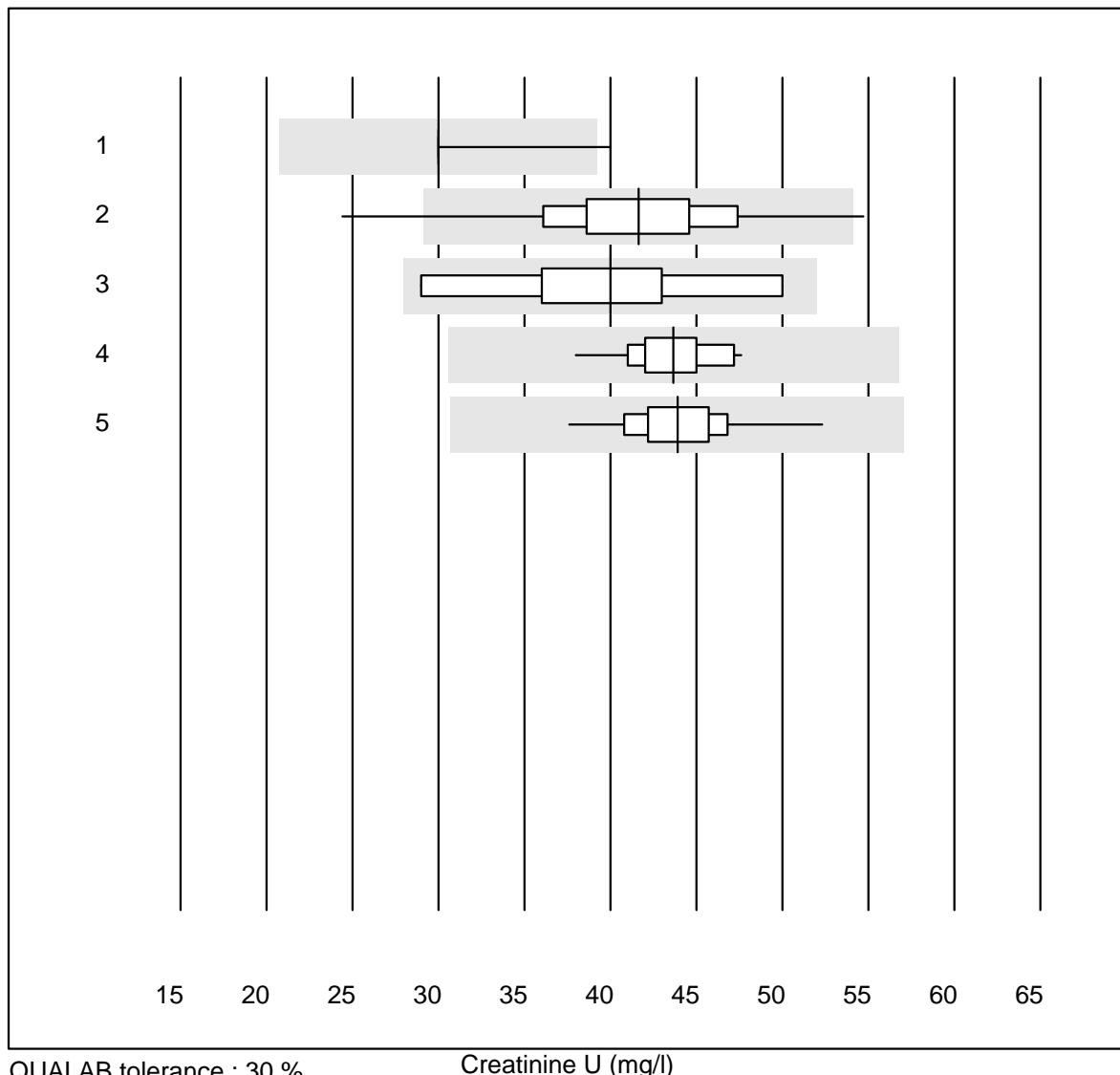
Cholesterin HDL PTS

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 CardioChek | 7 | 71.4 | 0.0 | 28.6 | 1.37 | 3.8 | e |

Triglyceride PTS



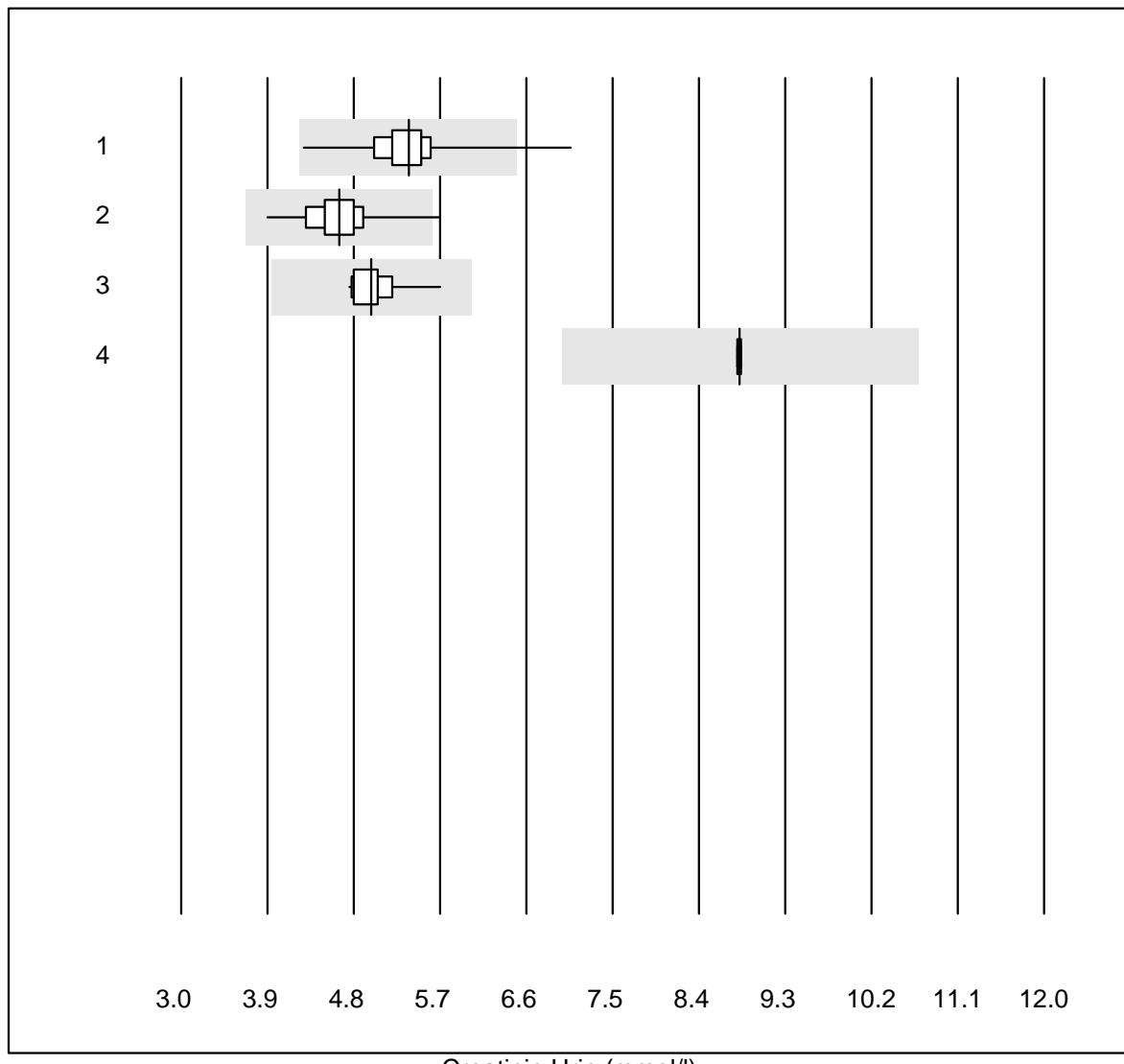
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 | CardioChek | 7 | 71.4 | 14.3 | 14.3 | 0.64 | 11.2 | e* |

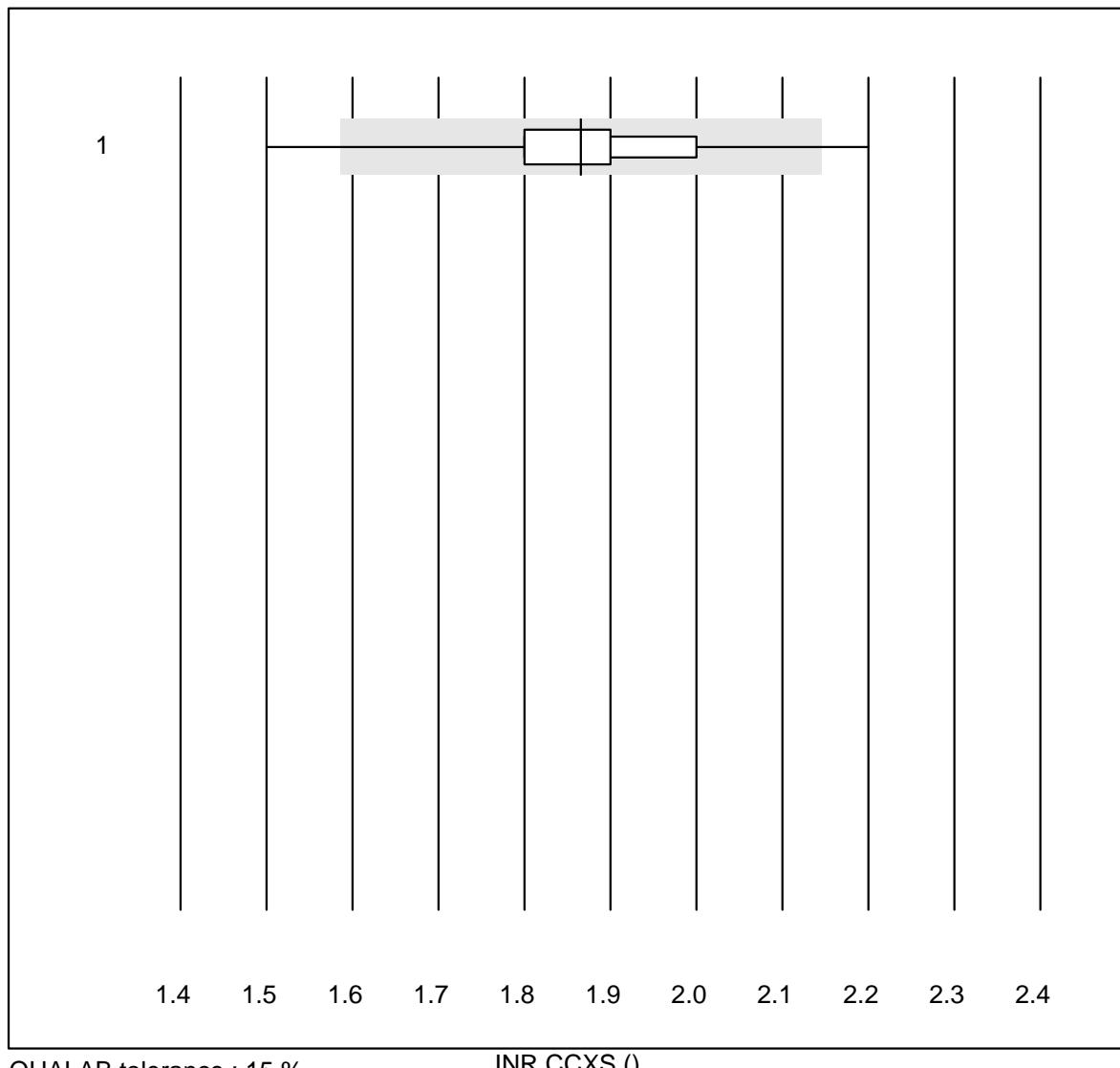
Creatinine U

QUALAB tolerance : 30 %

Creatinine U (mg/l)

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 Siemens Clinitek | 15 | 73.3 | 6.7 | 20.0 | 30.0 | 9.4 | a |
| 2 Afinion | 336 | 96.1 | 1.8 | 2.1 | 41.6 | 11.2 | e |
| 3 NycoCard | 10 | 90.0 | 0.0 | 10.0 | 40.0 | 17.8 | e* |
| 4 Turbidimetry | 17 | 100.0 | 0.0 | 0.0 | 43.7 | 5.8 | e |
| 5 DCA2000/Vantage | 124 | 97.6 | 0.0 | 2.4 | 43.9 | 6.0 | e |

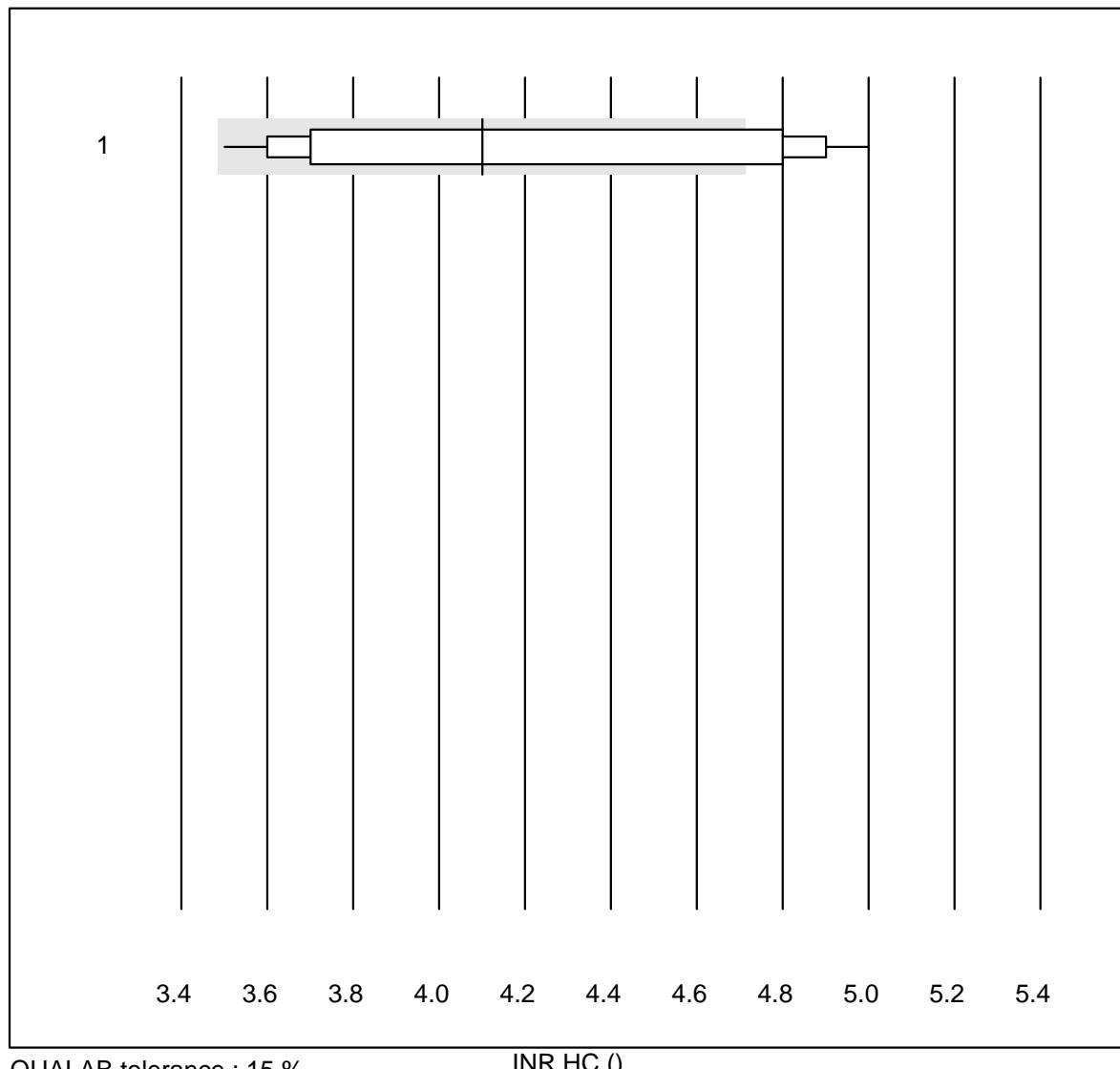
Creatinin Urin

INR CCXS

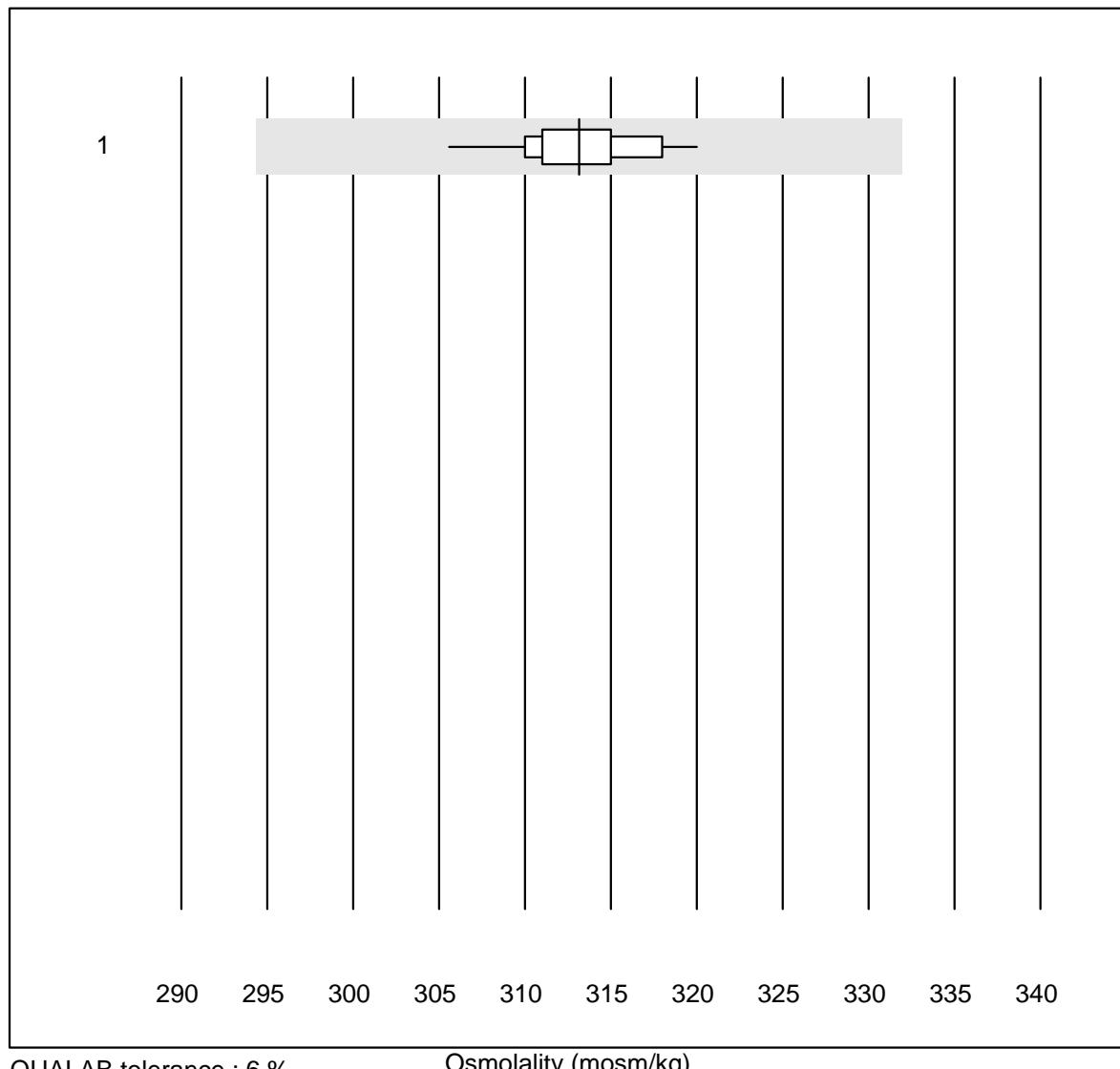
QUALAB tolerance : 15 %

INR CCXS ()

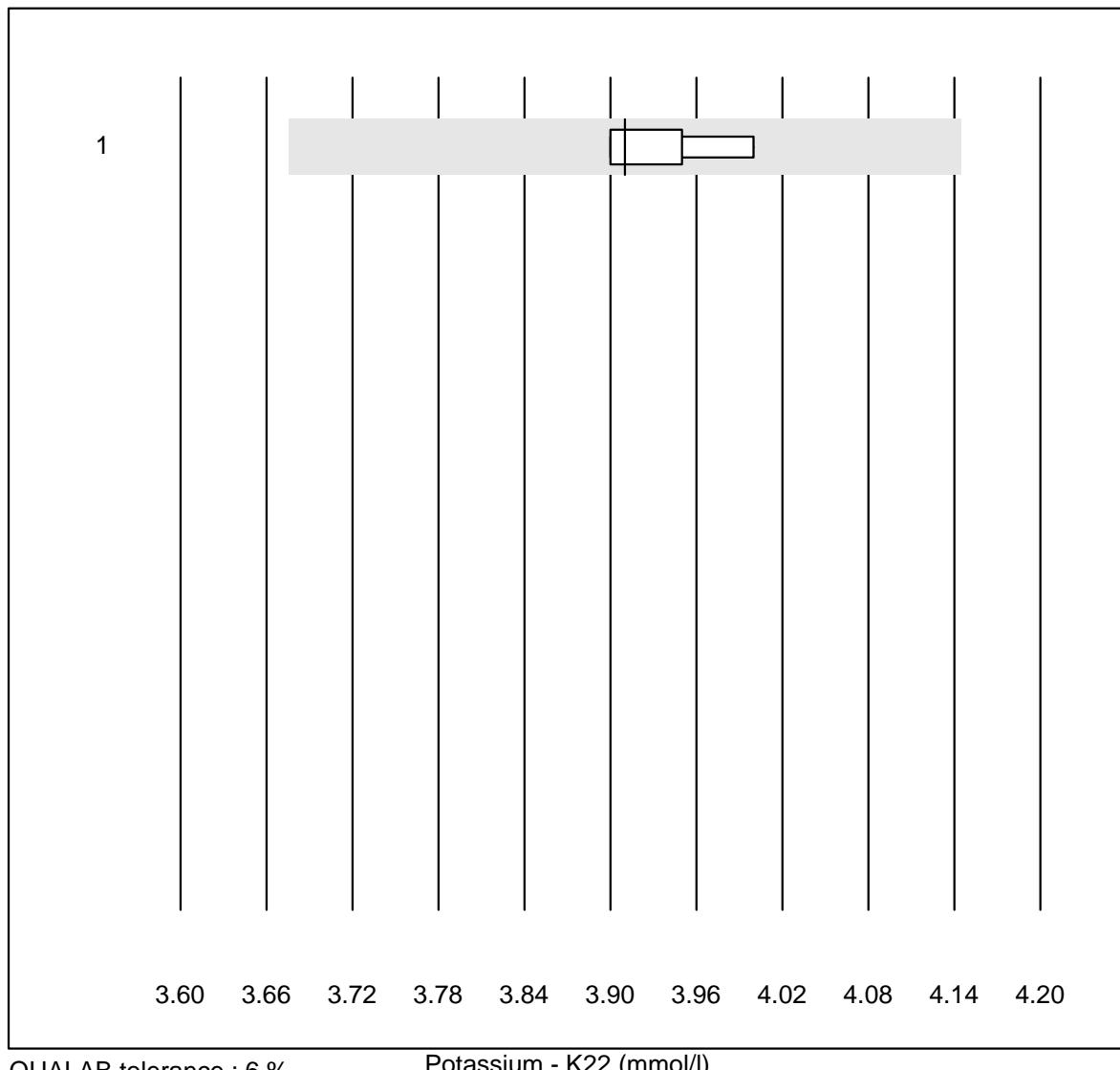
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 CoaguChek XS | 2325 | 99.4 | 0.3 | 0.3 | 1.9 | 4.4 | e |

INR HC

Osmolality



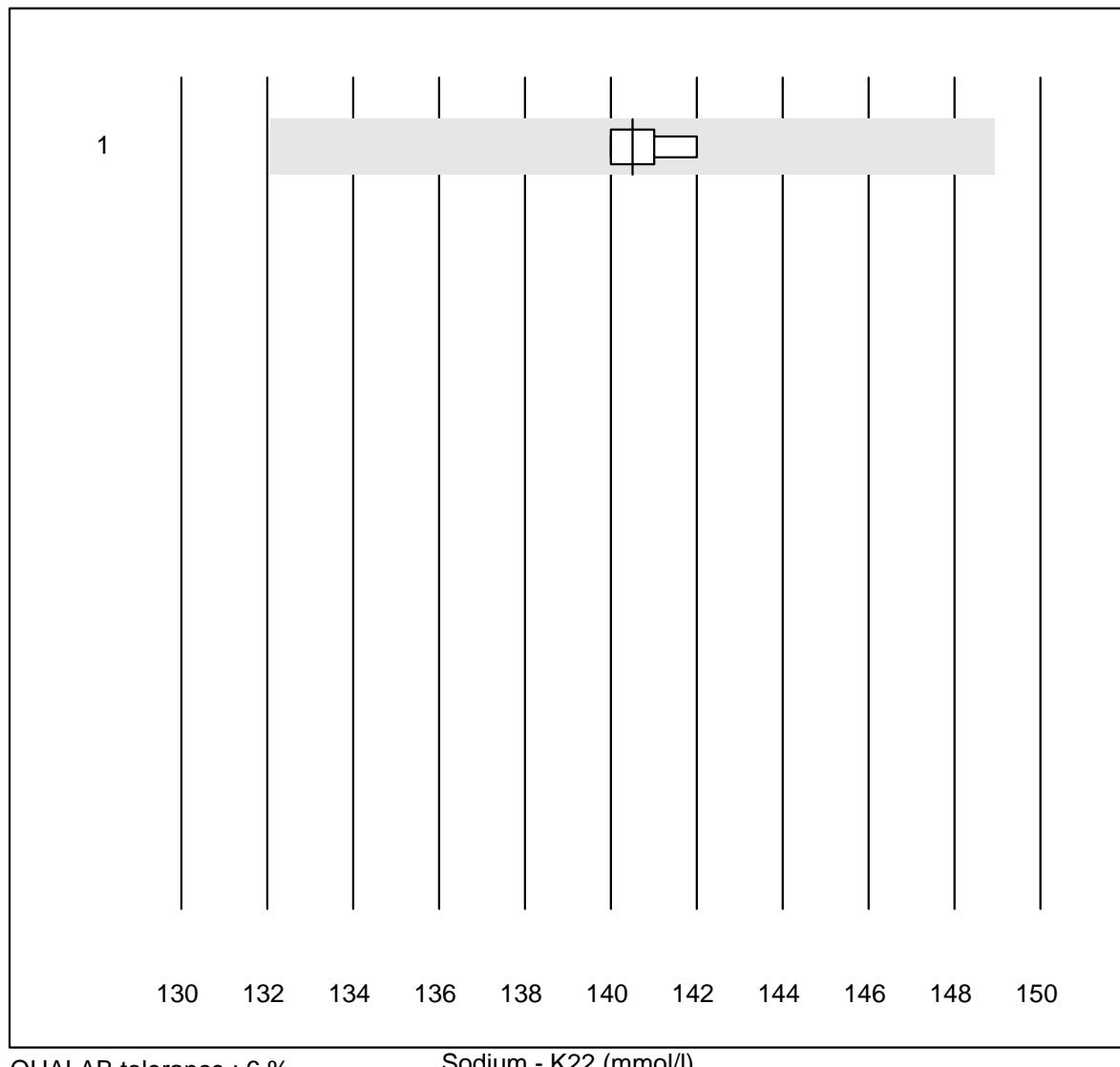
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|-----------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Cryoskopy | 13 | 100.0 | 0.0 | 0.0 | 313 | 1.2 | e |

Potassium - K22

| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | ISE | 9 | 100.0 | 0.0 | 0.0 | 3.9 | 1.0 | e |

K22 Osmolality

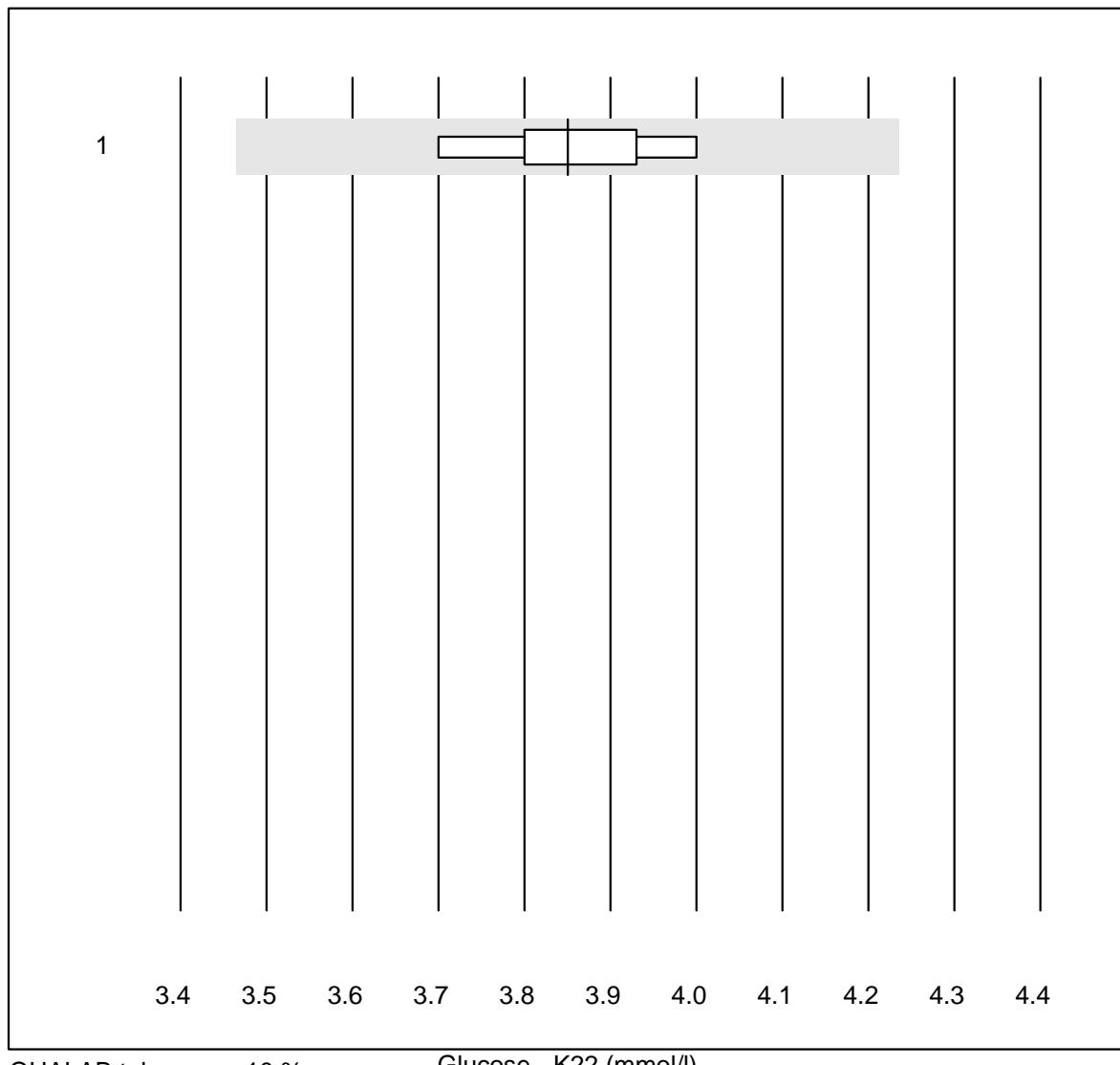
Sodium - K22



| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | ISE | 9 | 100.0 | 0.0 | 0.0 | 141 | 0.5 | e |

K22 Osmolality

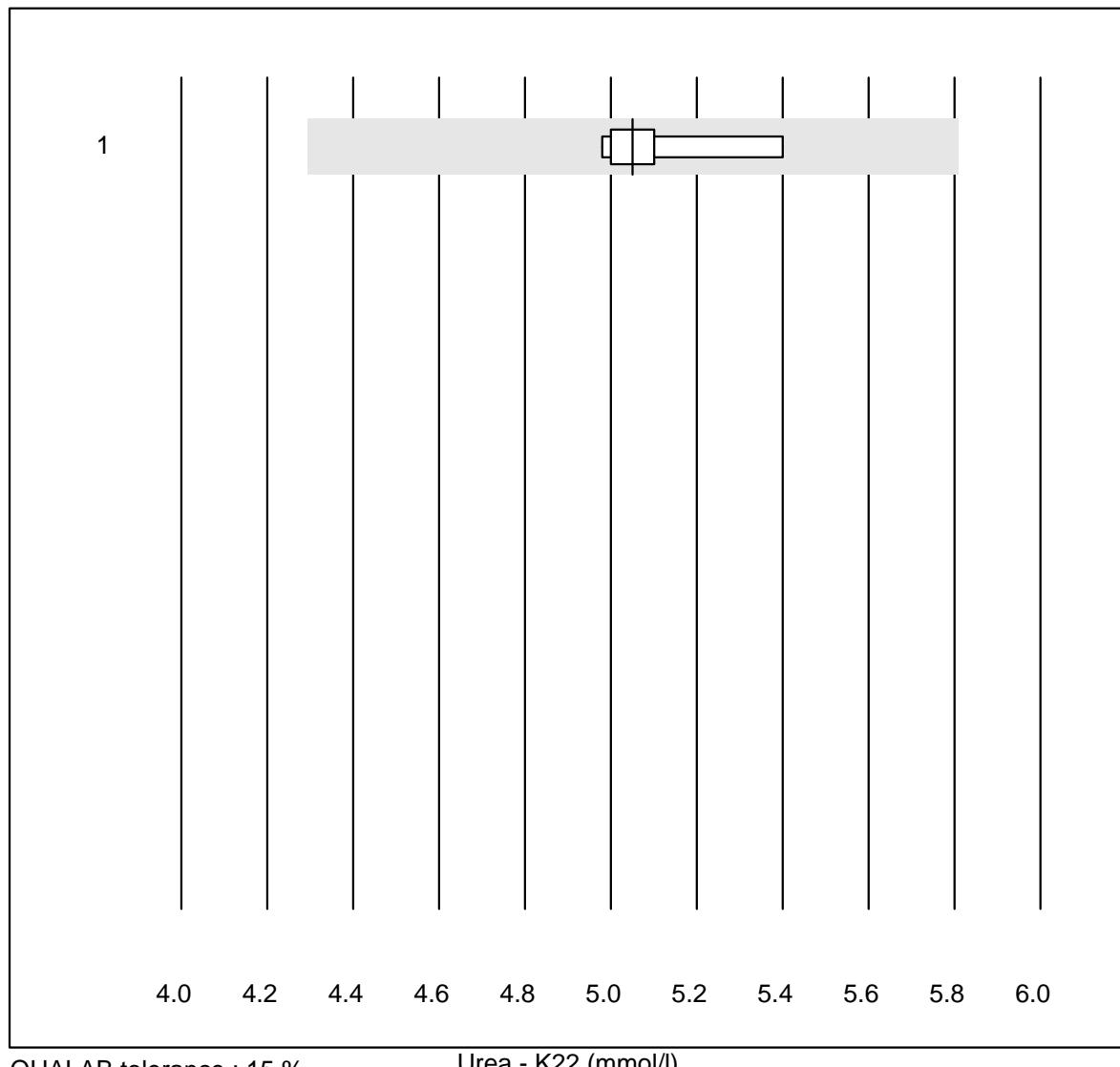
Glucose - K22



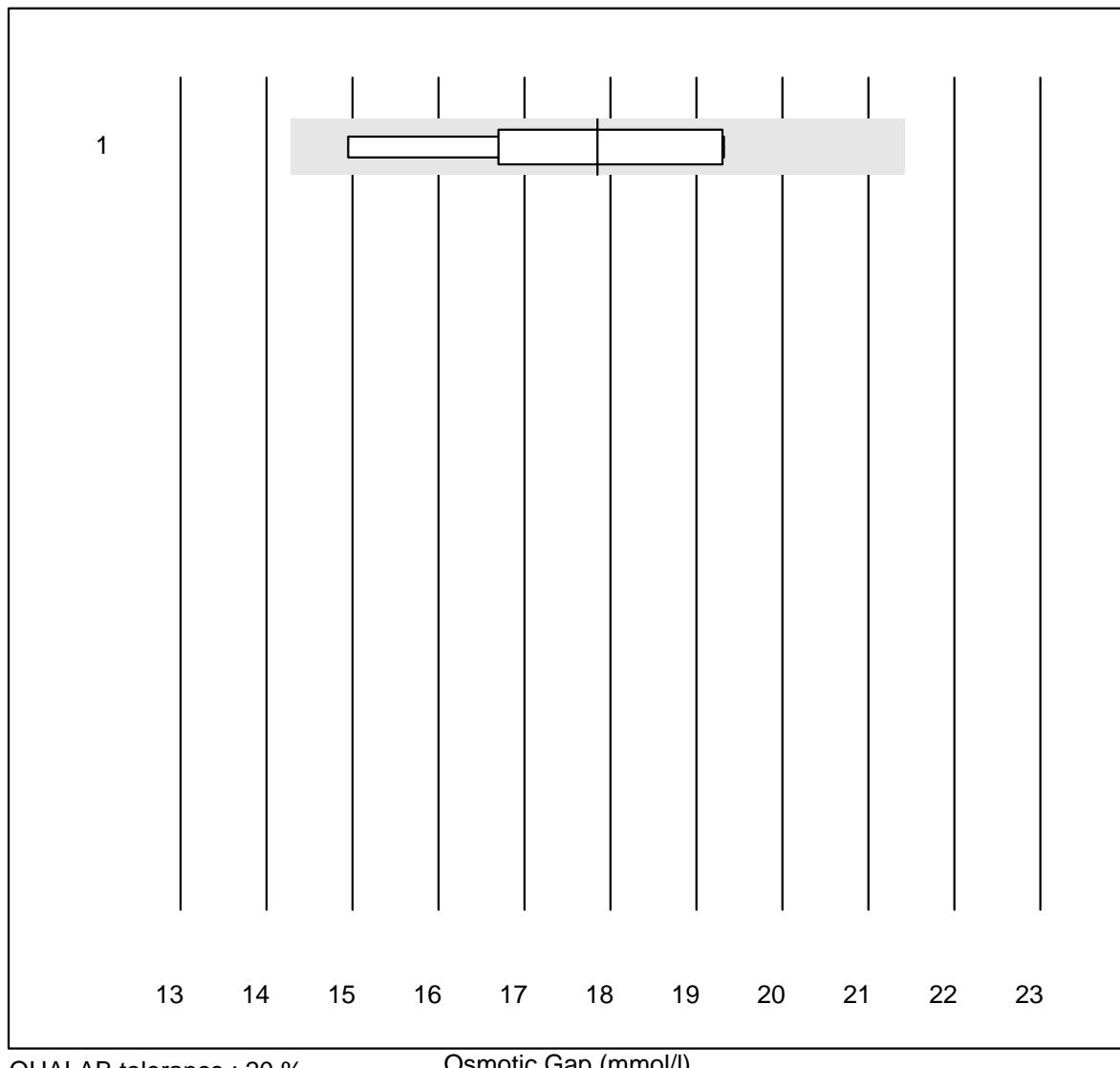
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Standard chemistry | 9 | 100.0 | 0.0 | 0.0 | 3.9 | 3.0 | e |

K22 Osmolality

Urea - K22

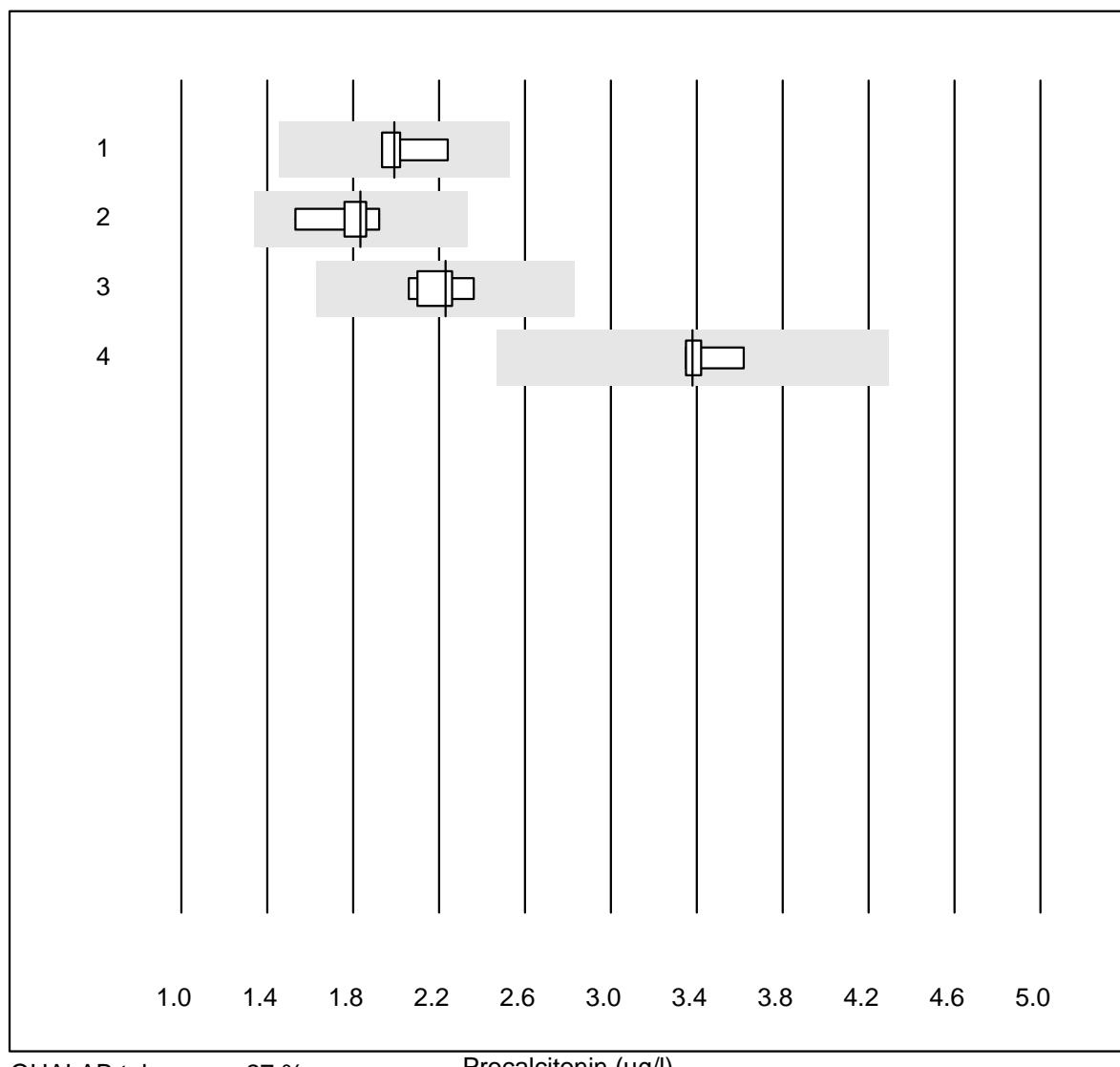


| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Standard chemistry | 9 | 100.0 | 0.0 | 0.0 | 5.1 | 2.7 | e |

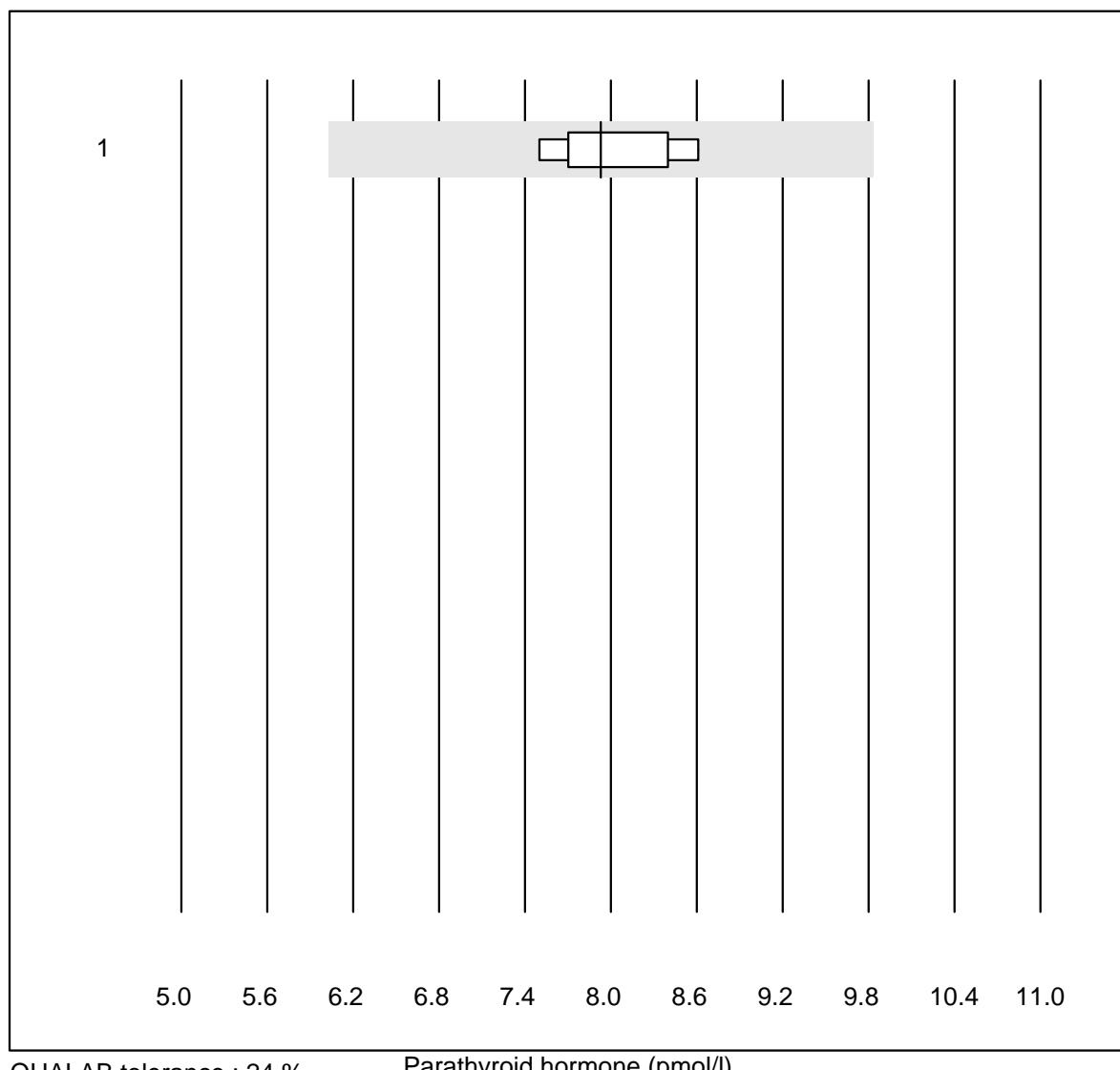
Osmotic Gap

| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|-----------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Formel 1 (2Na+K+Glu+) | 6 | 100.0 | 0.0 | 0.0 | 17.9 | 9.9 | e* |

Procalcitonin

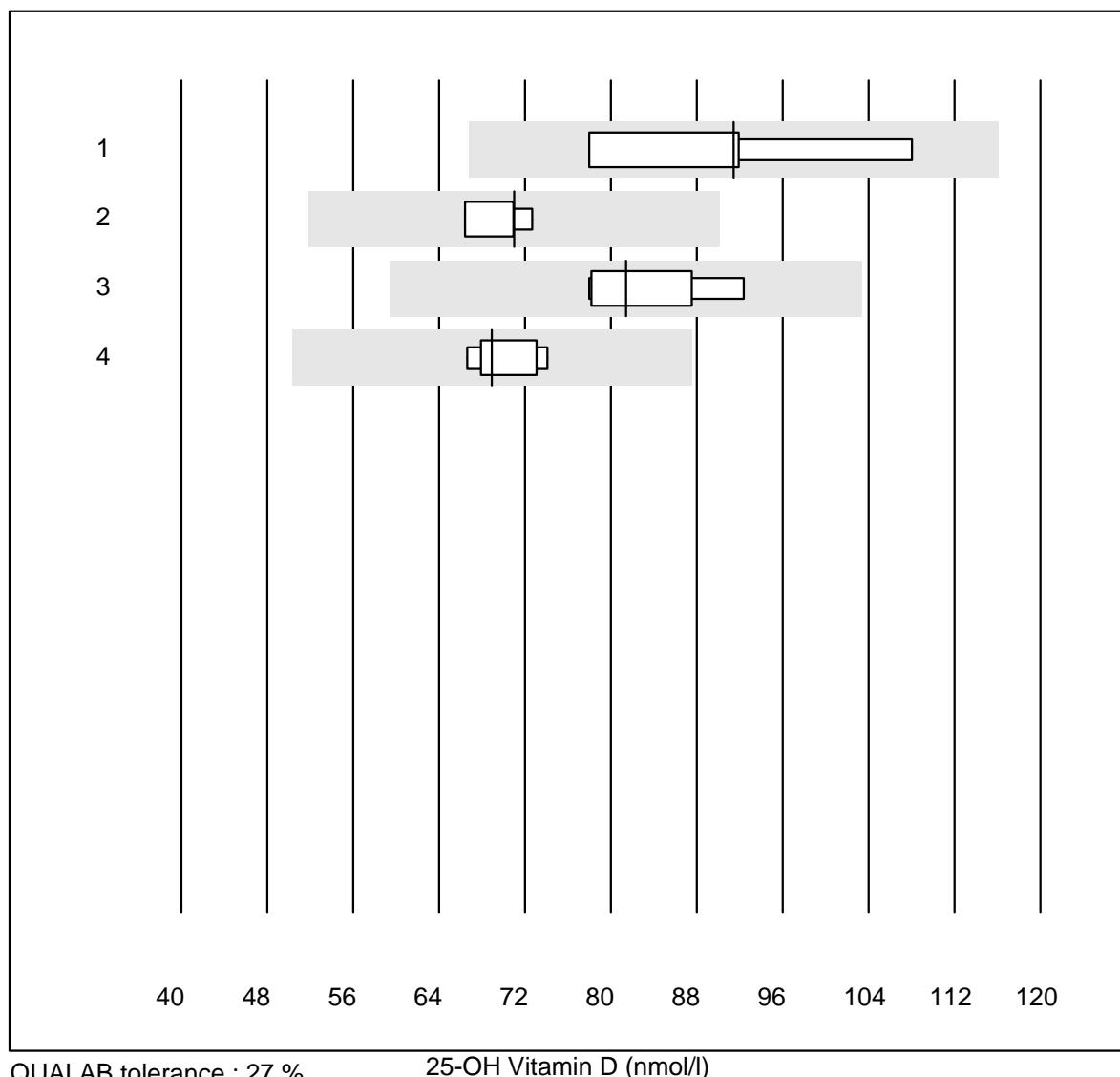


Parathyroid hormone

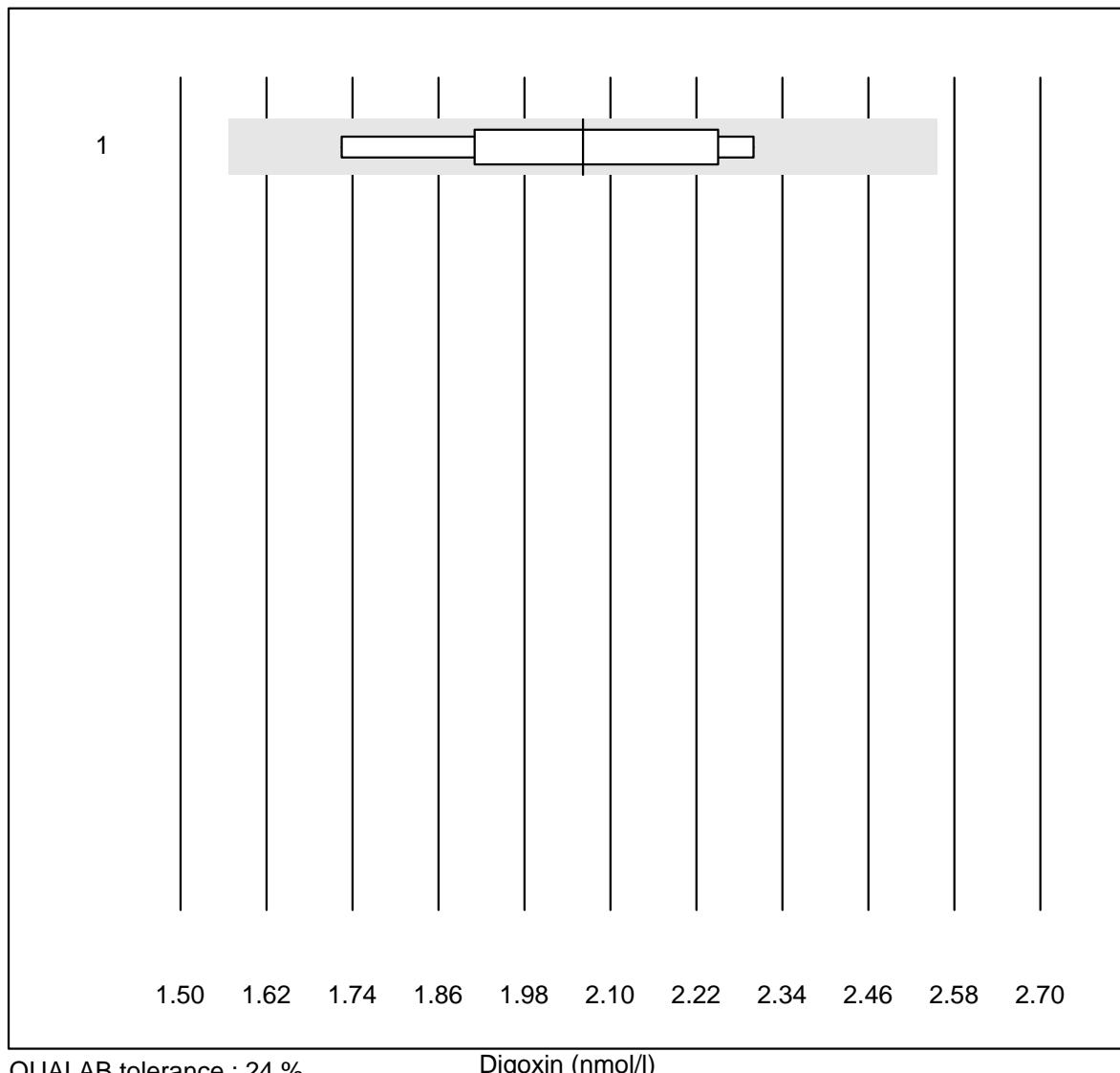


| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|----------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Cobas PTH STAT | 5 | 100.0 | 0.0 | 0.0 | 7.9 | 5.8 | e |

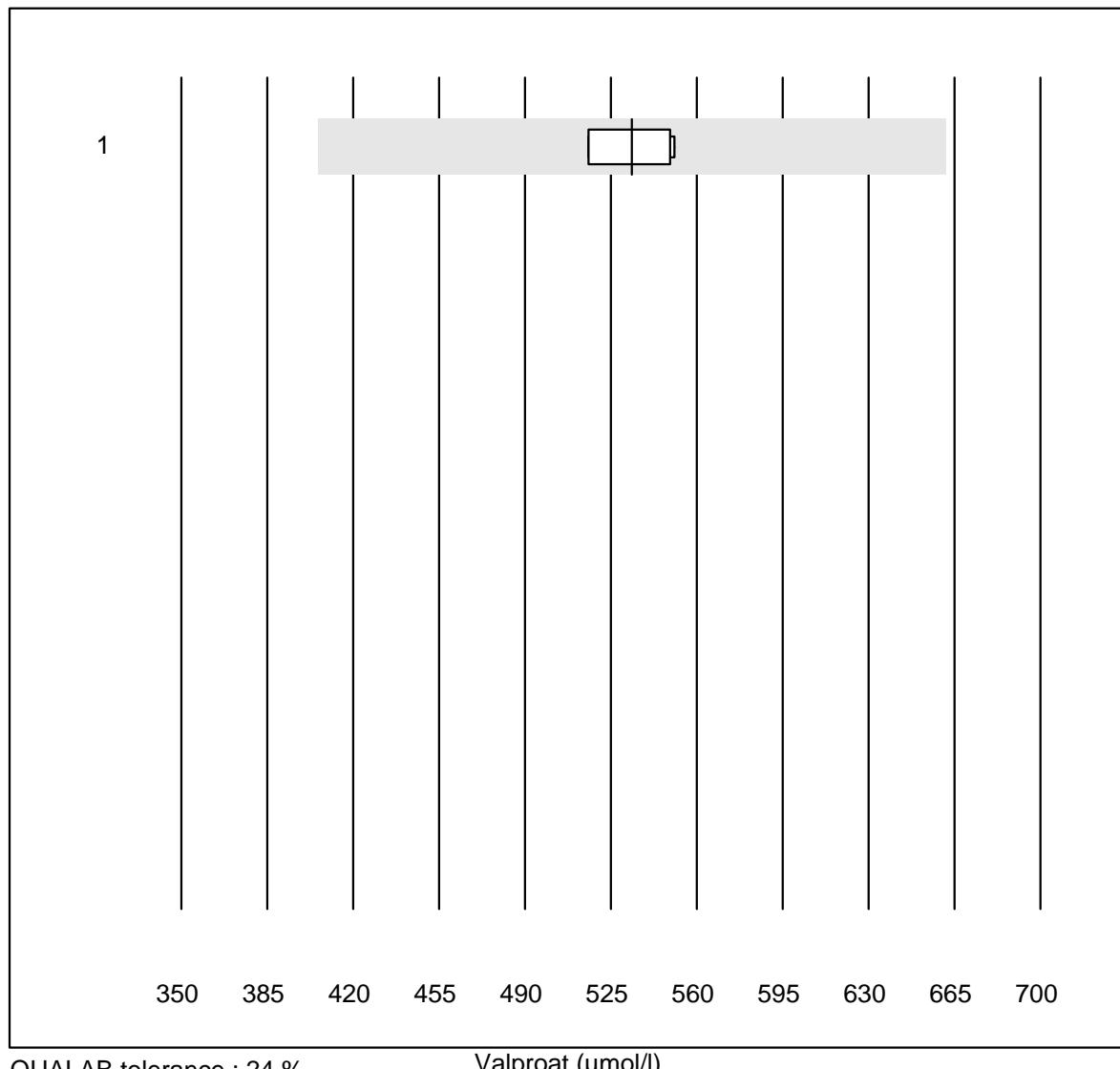
25-OH Vitamin D



Digoxin

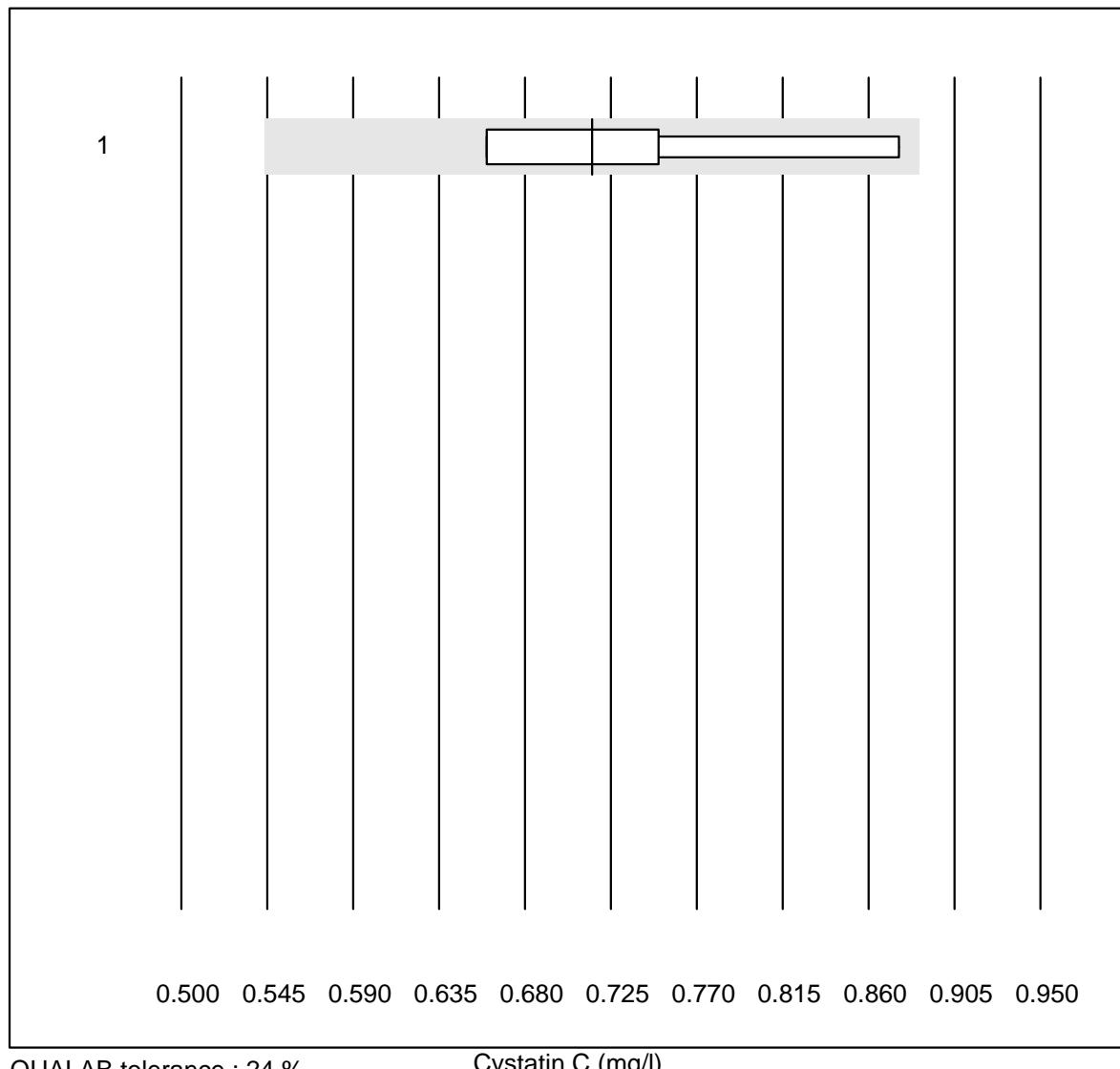


| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 | Other methods | 10 | 100.0 | 0.0 | 0.0 | 2.06 | 10.7 | e* |

Valproat

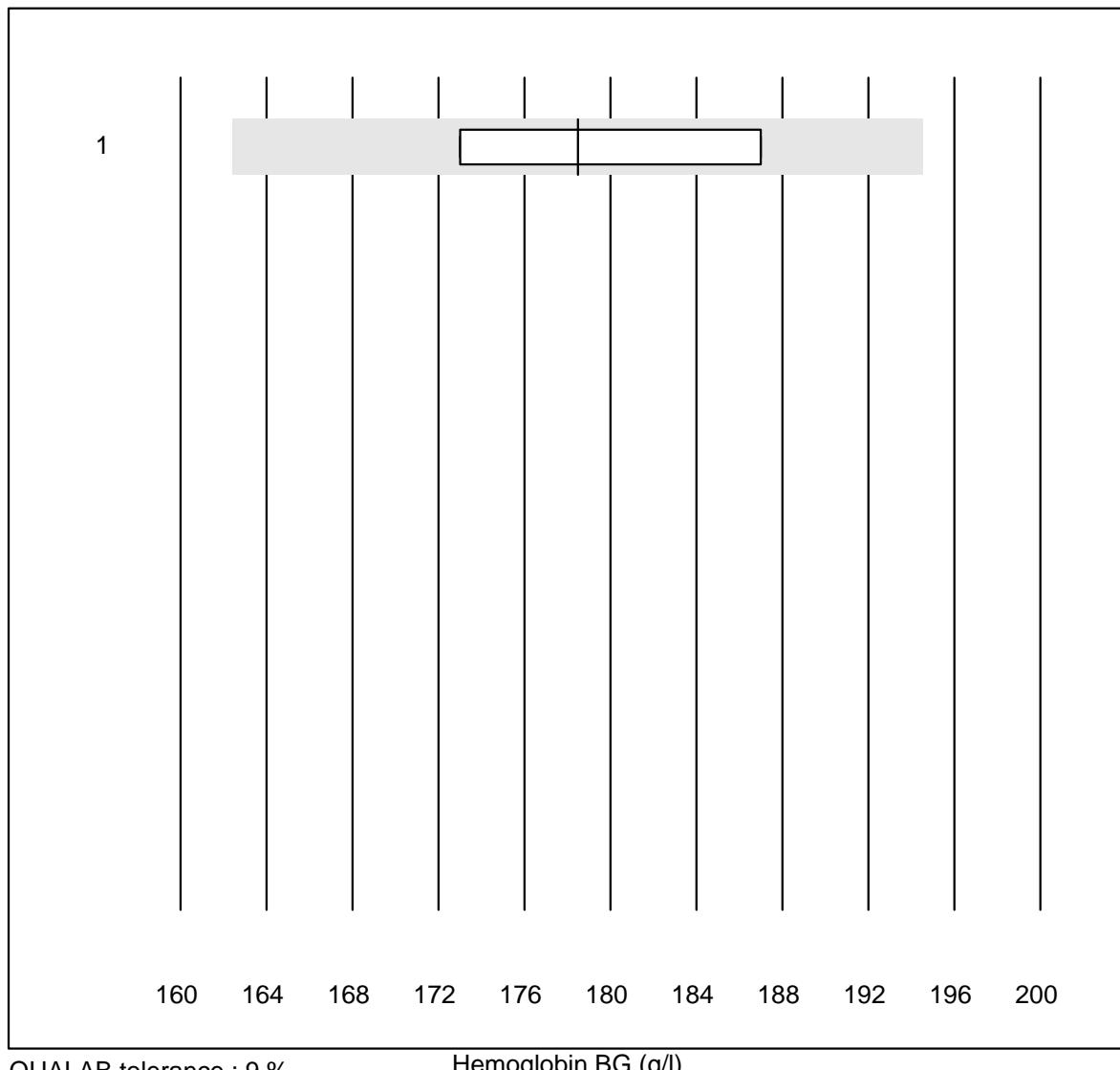
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 all Participants | 4 | 100.0 | 0.0 | 0.0 | 533.5 | 3.6 | e |

Cystatin C



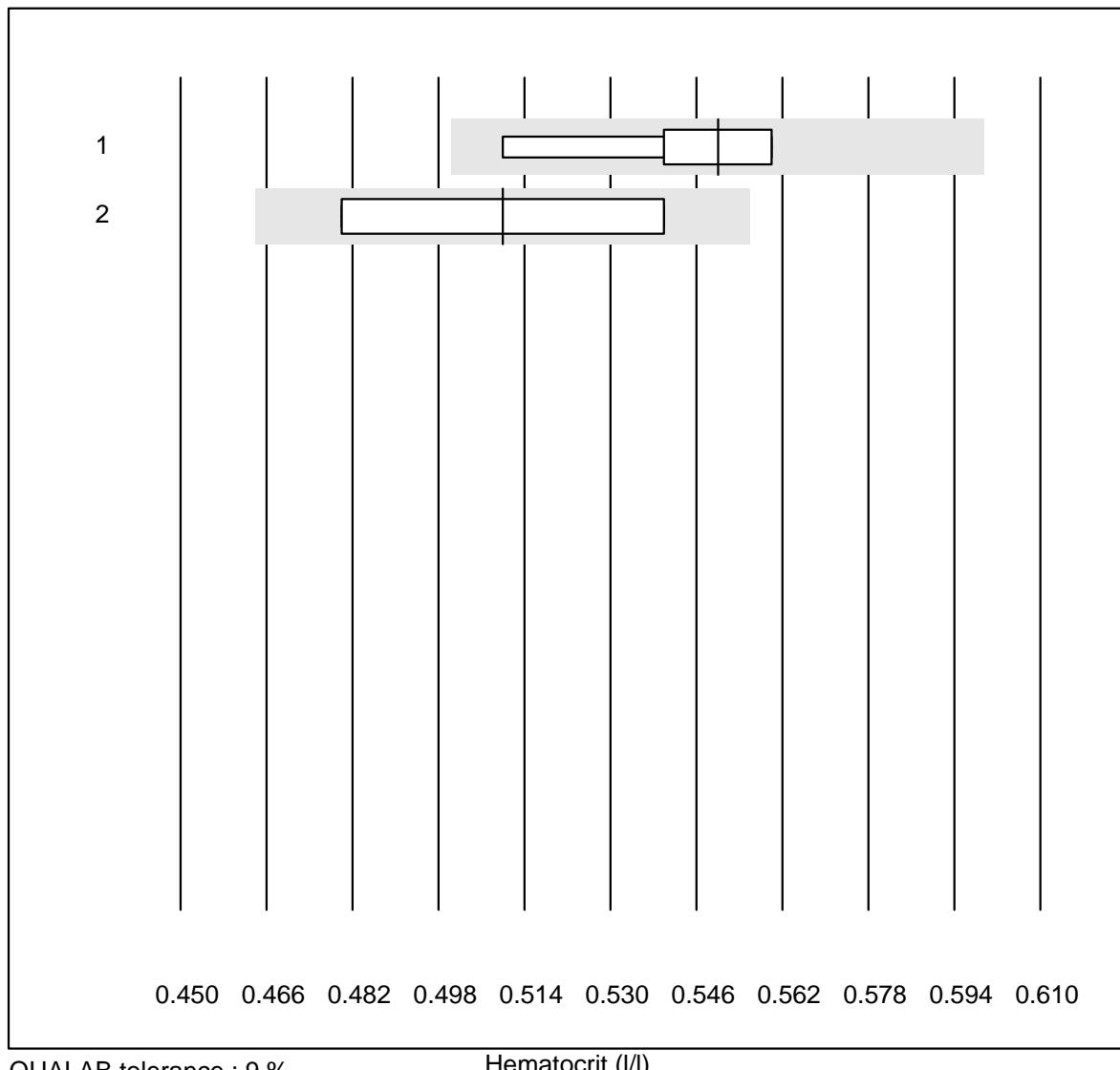
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 all Participants | 4 | 100.0 | 0.0 | 0.0 | 0.7 | 13.2 | e* |

Hemoglobin BG



| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | iStat | 4 | 75.0 | 0.0 | 25.0 | 178.5 | 4.1 | e* |

Hematocrit

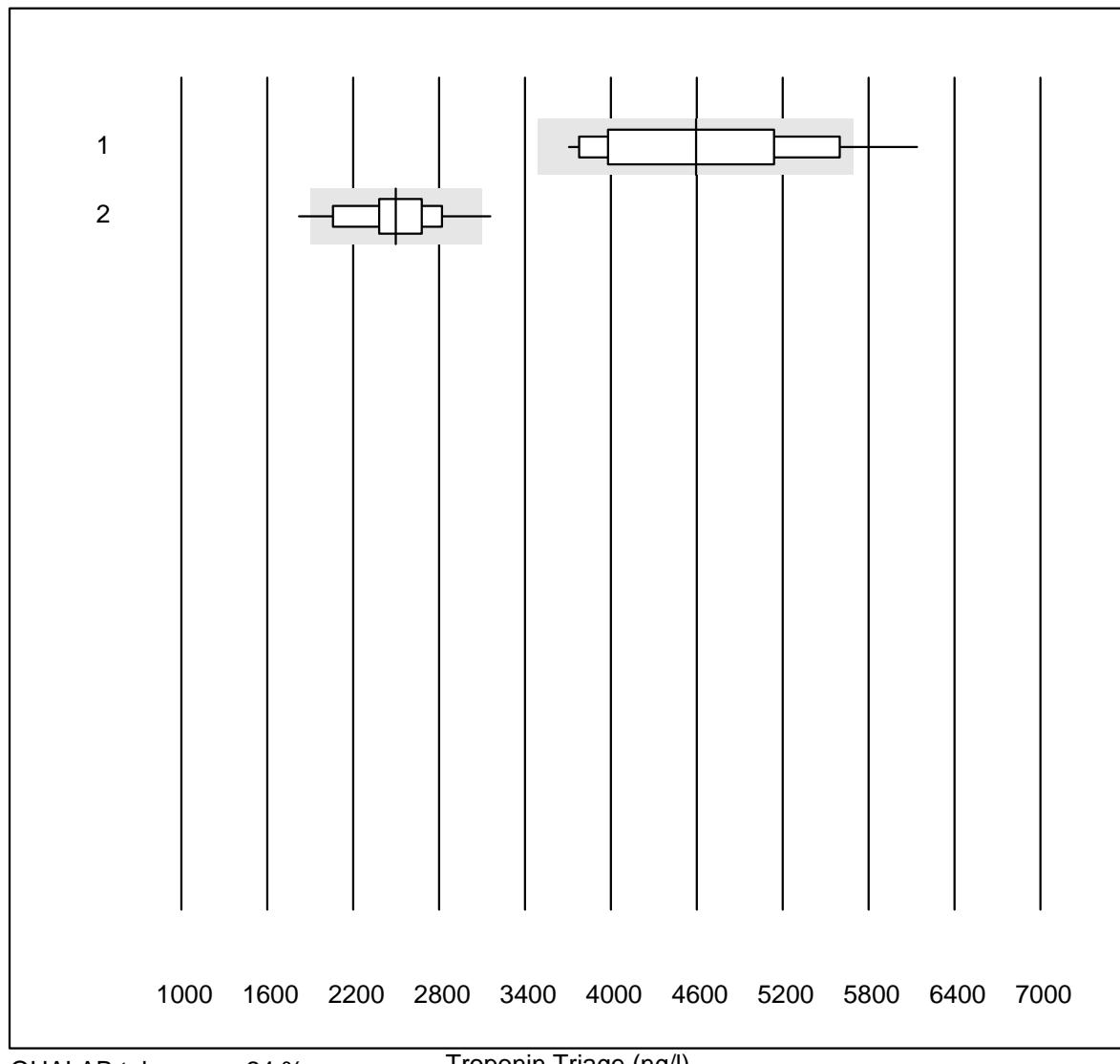


QUALAB tolerance : 9 %

Hematocrit (l/l)

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 iStat | 5 | 100.0 | 0.0 | 0.0 | 0.55 | 3.8 | e* |
| 2 EPOC | 5 | 80.0 | 0.0 | 20.0 | 0.51 | 6.8 | a |

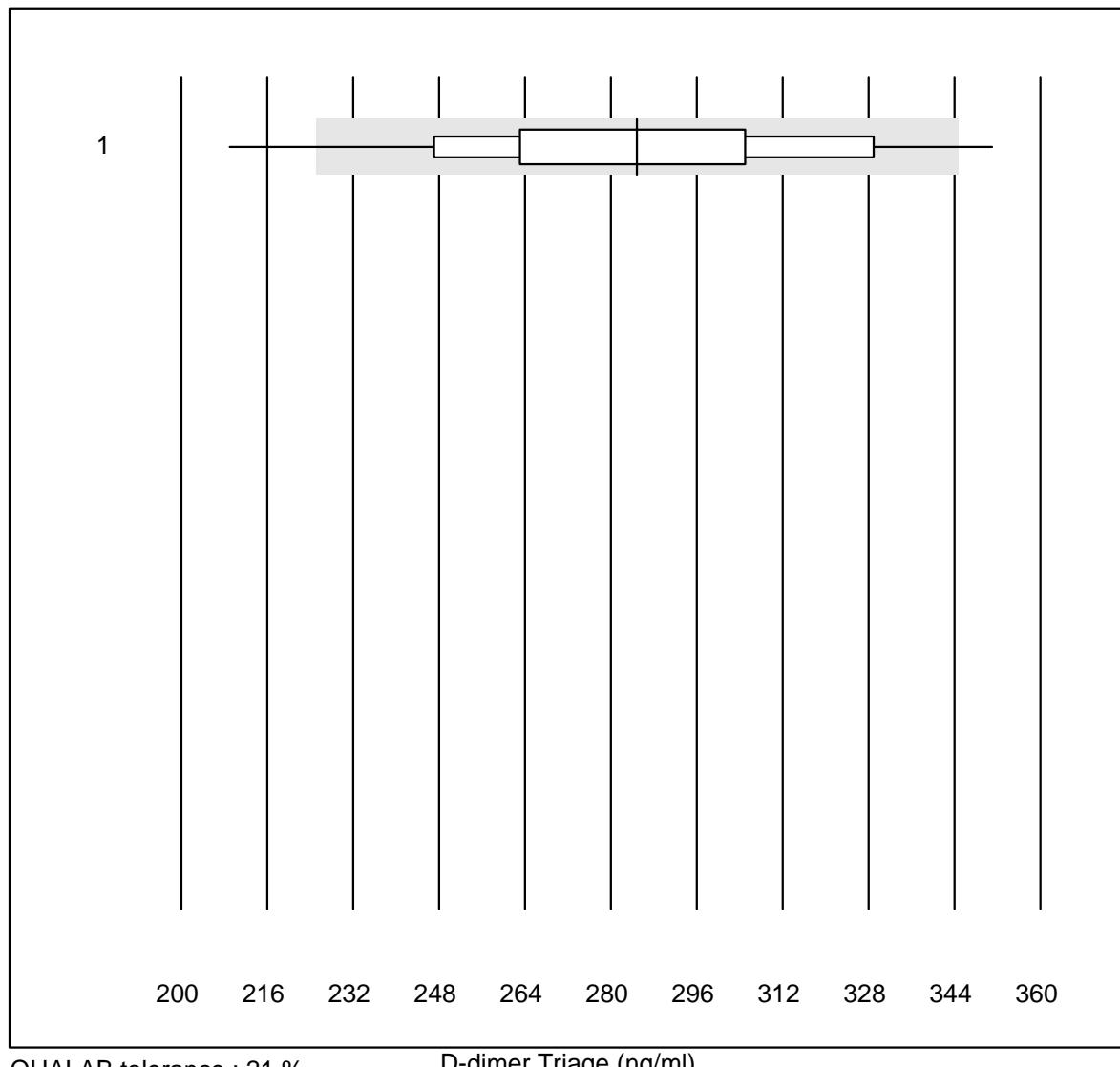
Troponin Triage



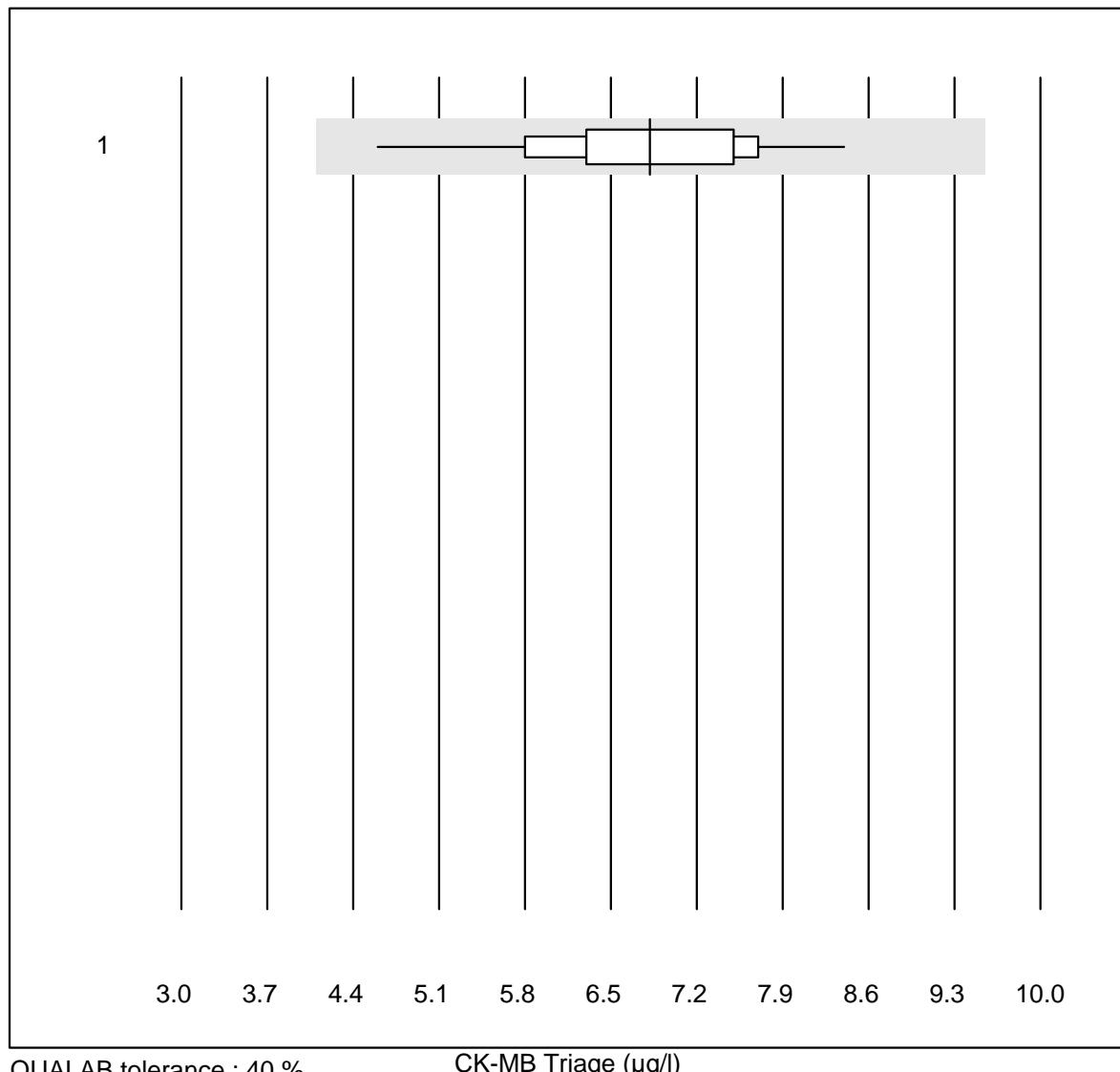
QUALAB tolerance : 24 %

Troponin Triage (ng/l)

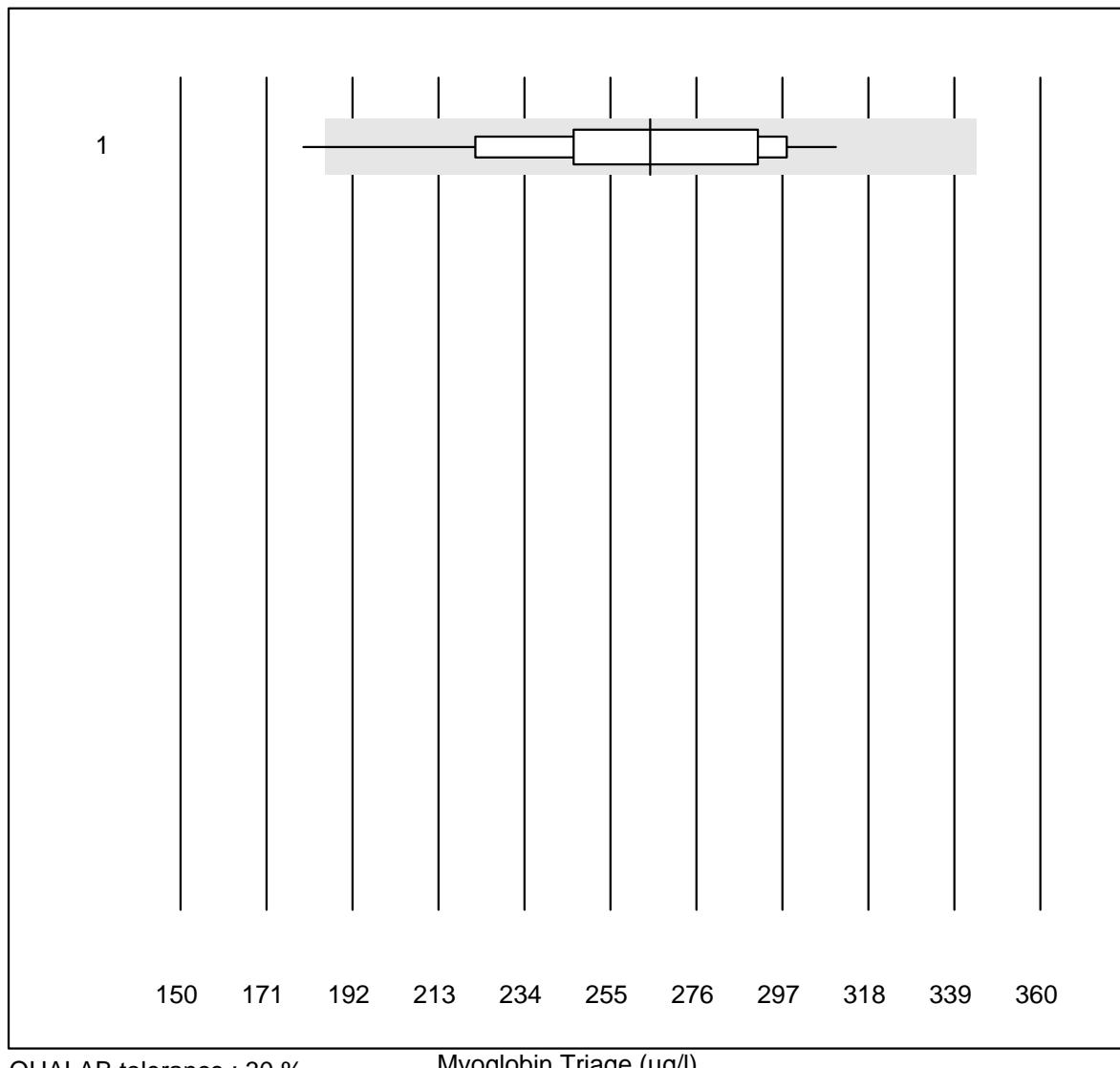
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|----------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 Triage Next Gen | 30 | 93.4 | 3.3 | 3.3 | 4592.90 | 14.9 | e |
| 2 Triage SOB/Cardiac | 23 | 78.3 | 13.0 | 8.7 | 2497.62 | 14.0 | e* |

D-dimer Triage

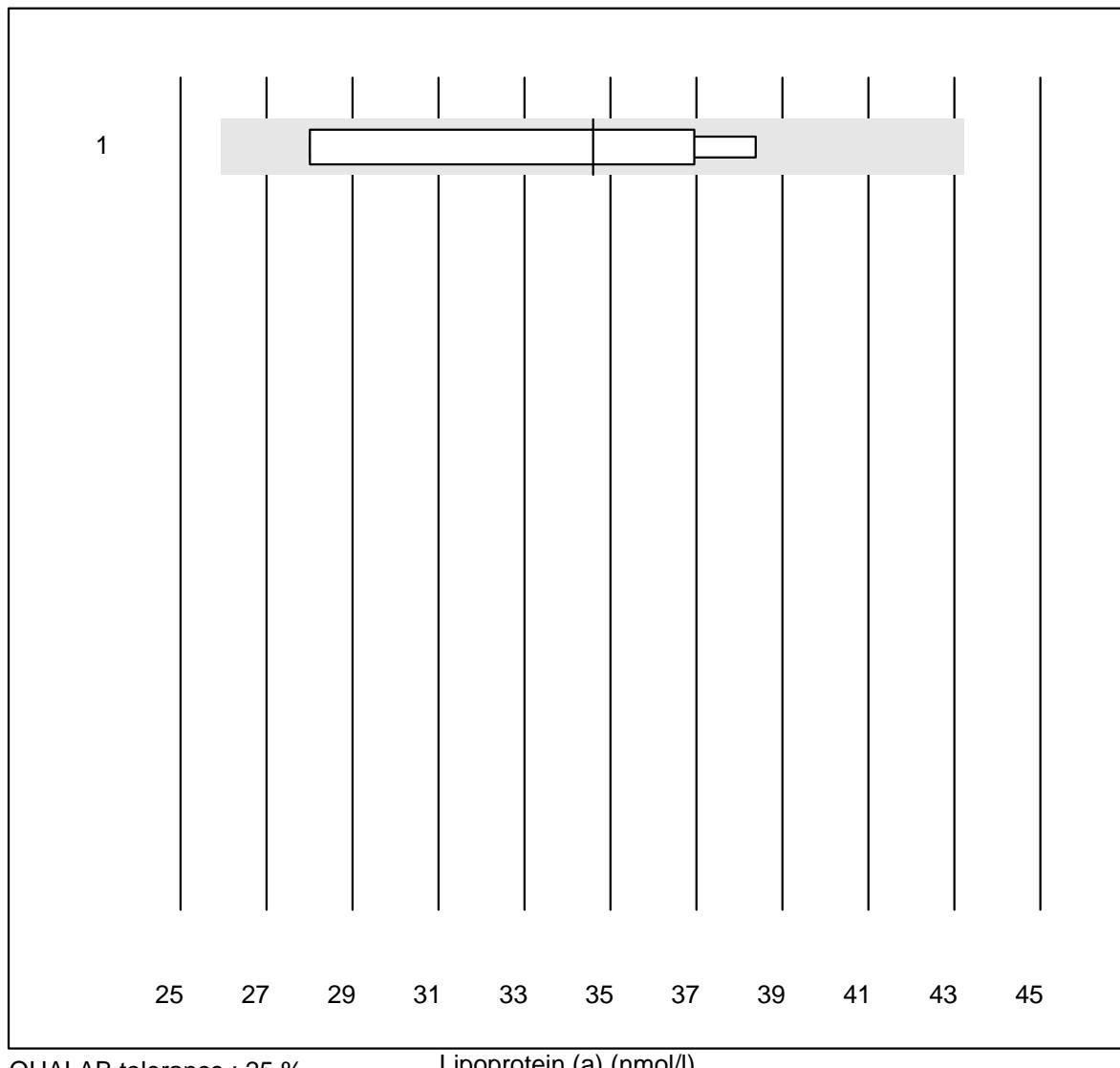
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 | Triage | 49 | 91.8 | 4.1 | 4.1 | 284.89 | 10.9 | e |

CK-MB Triage

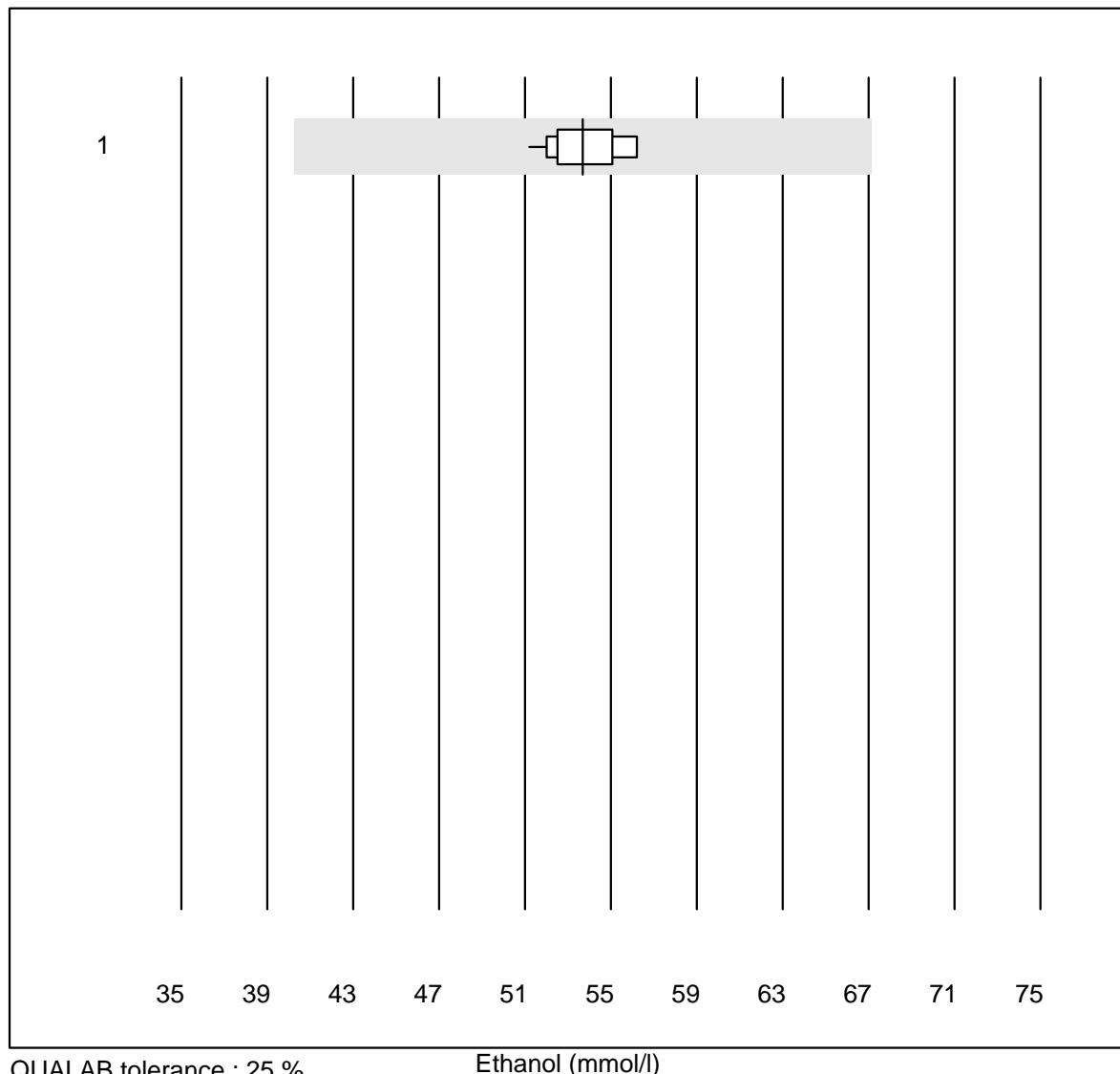
Myoglobin Triage



| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 Triage | 19 | 94.7 | 5.3 | 0.0 | 264.8 | 11.7 | e |

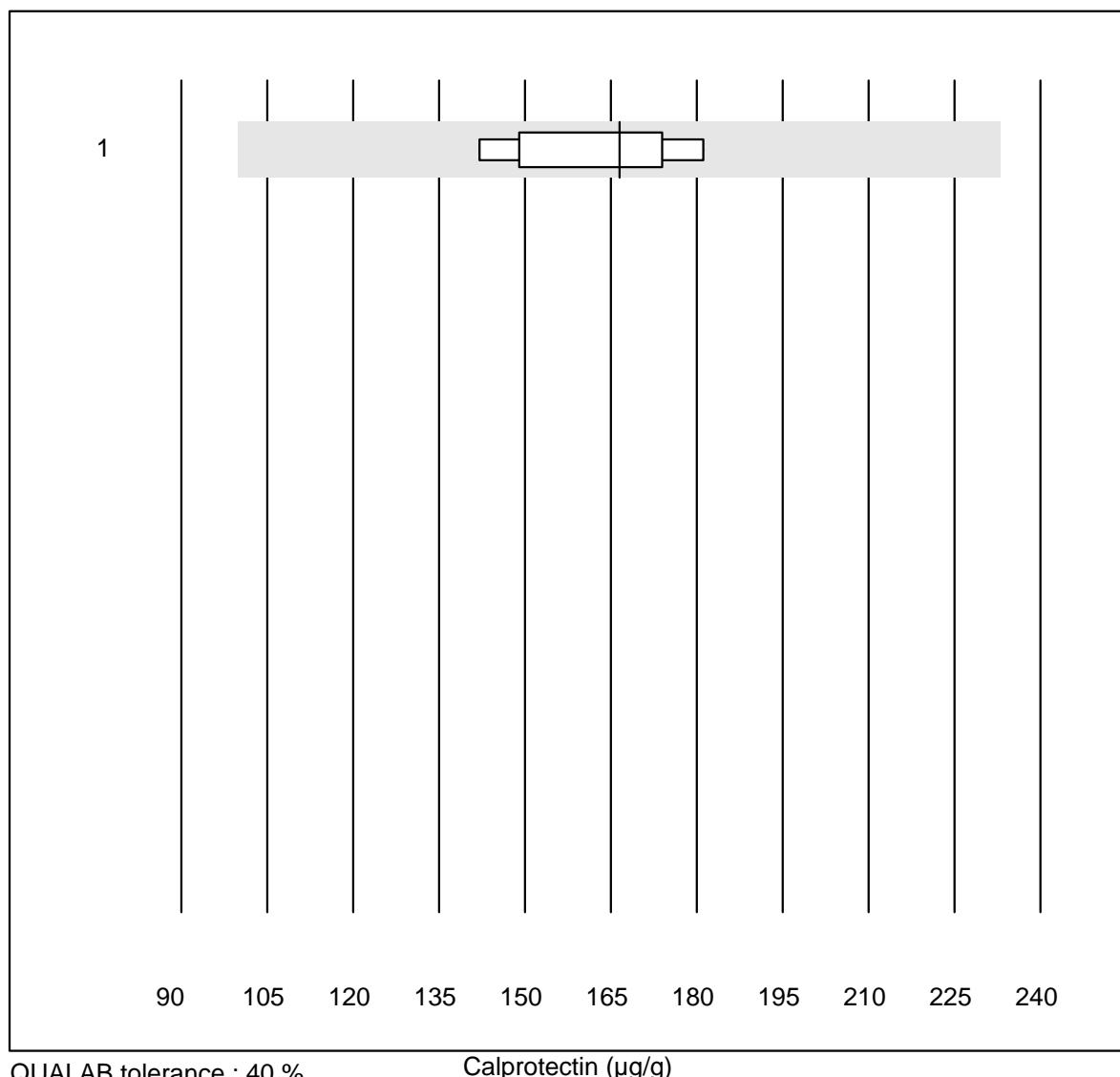
Lipoprotein (a)

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 all Participants | 4 | 100.0 | 0.0 | 0.0 | 35 | 13.3 | a |

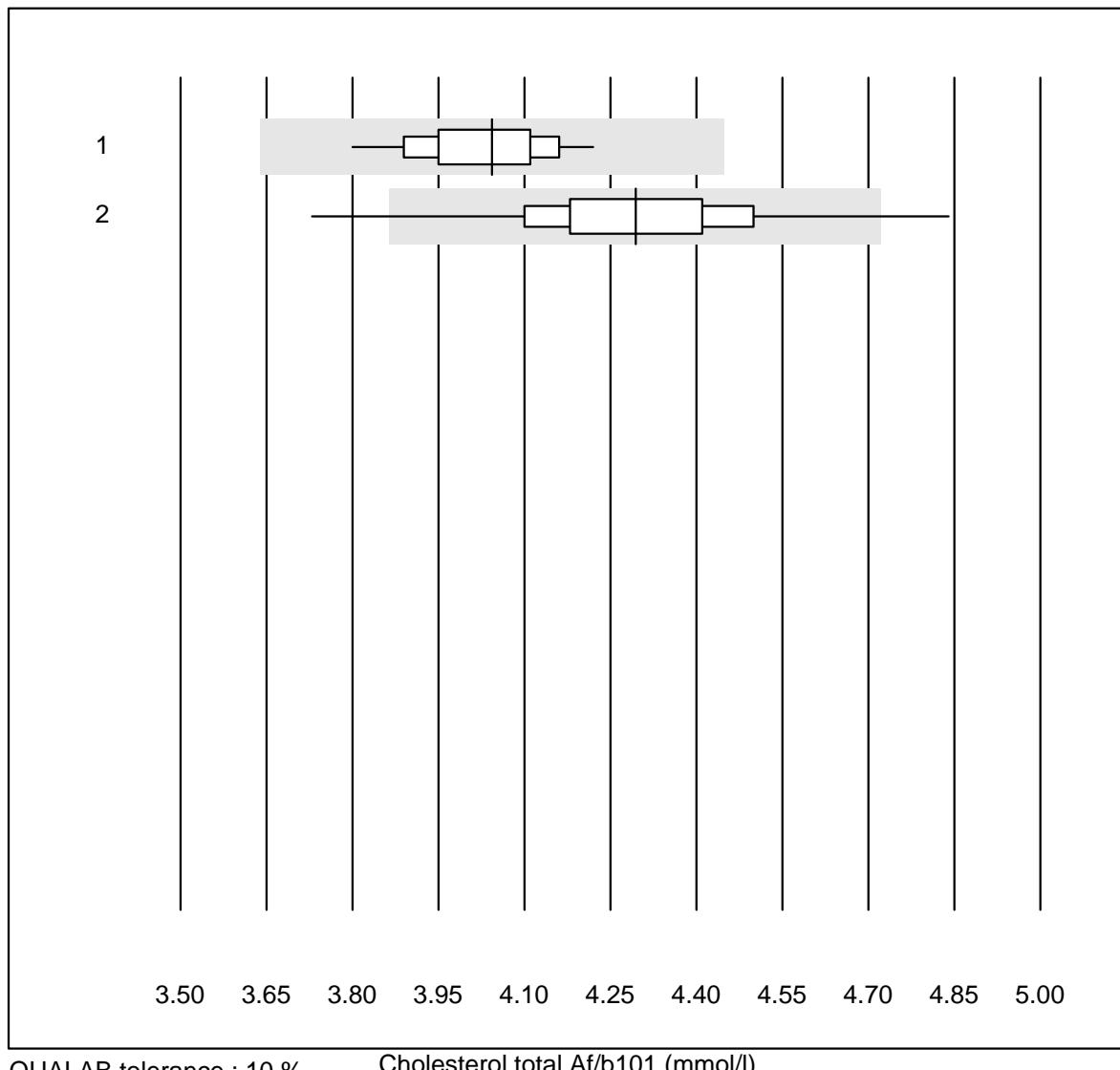
Ethanol

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 all Participants | 13 | 100.0 | 0.0 | 0.0 | 53.7 | 3.1 | e |

Calprotectin



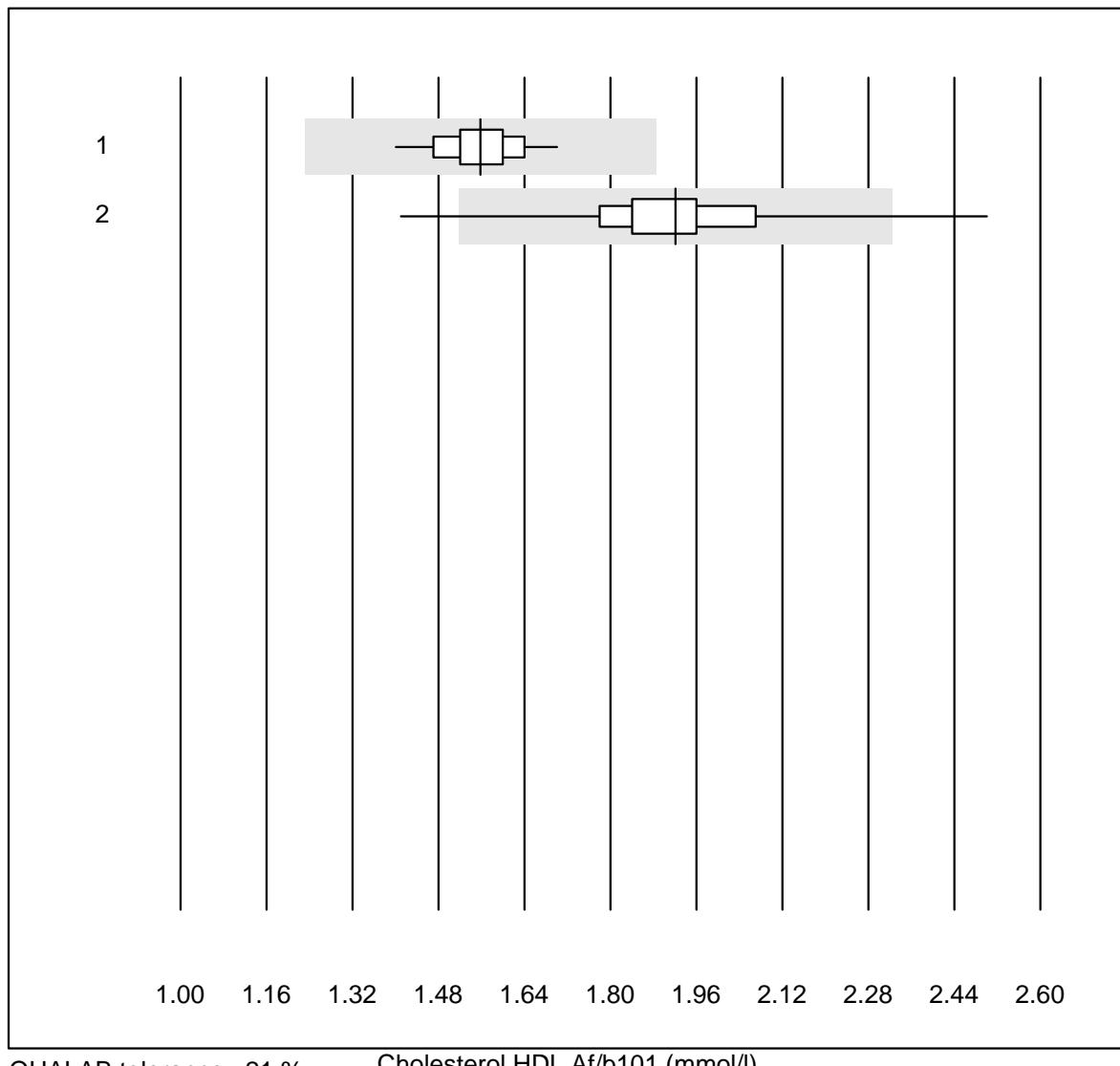
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|----------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Bühlmann | 7 | 100.0 | 0.0 | 0.0 | 167 | 9.0 | e |

Cholesterol total Af/b101

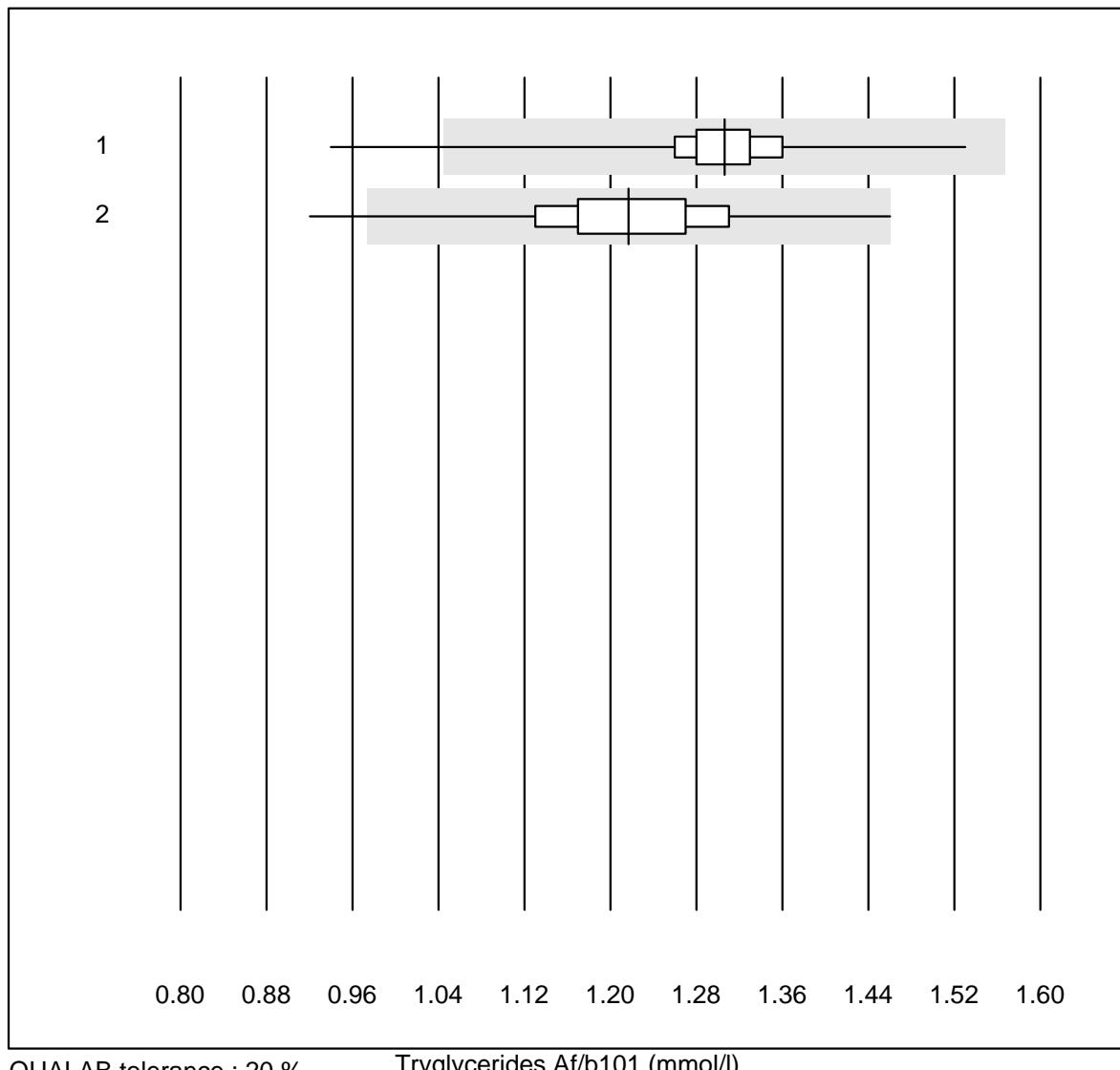
QUALAB tolerance : 10 %

Cholesterol total Af/b101 (mmol/l)

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Cobas b101 | 44 | 100.0 | 0.0 | 0.0 | 4.04 | 2.5 | e |
| 2 Afinion | 275 | 98.9 | 0.7 | 0.4 | 4.29 | 3.9 | e |

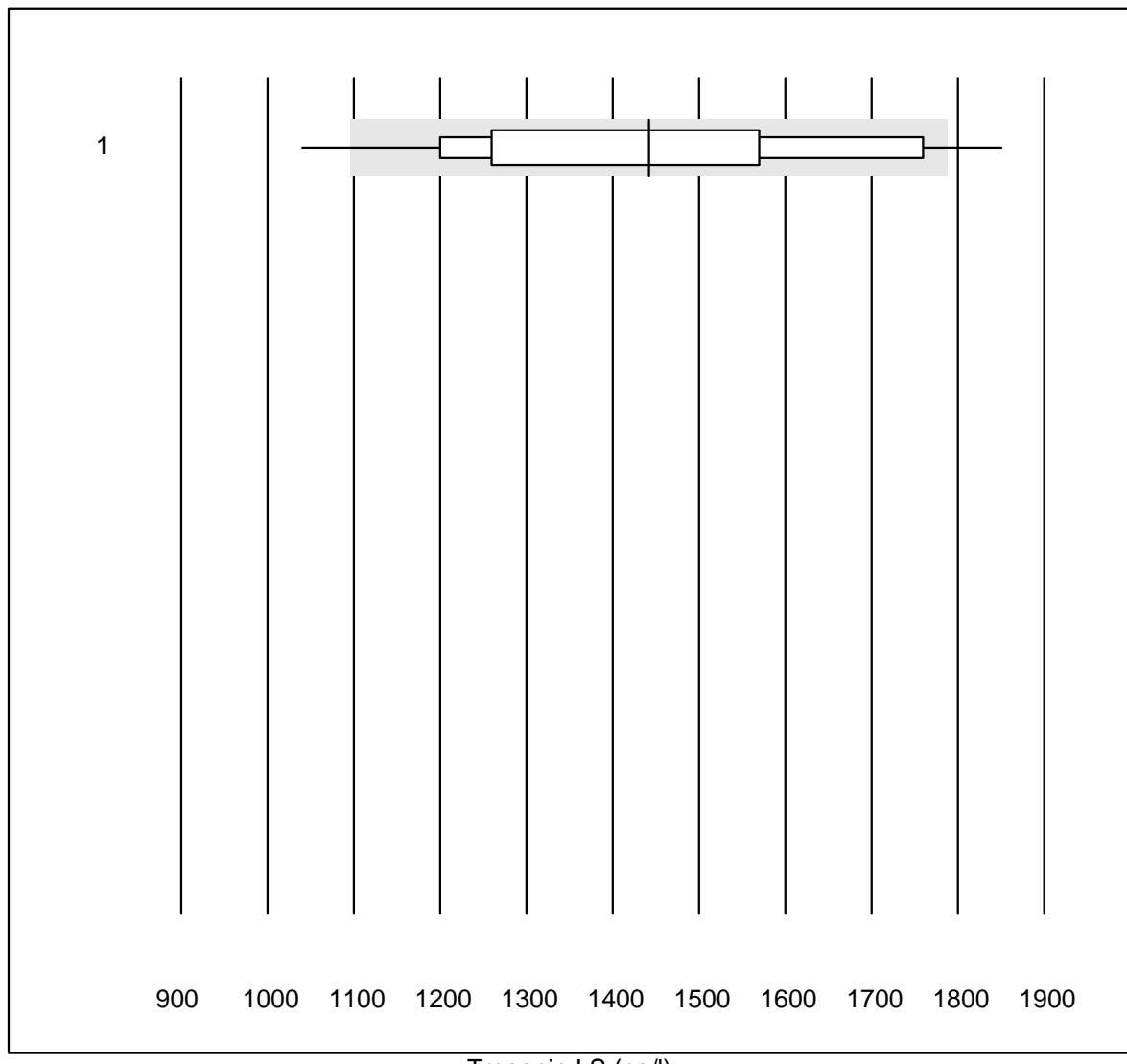
Cholesterol HDL Af/b101

Tryglycerides Af/b101

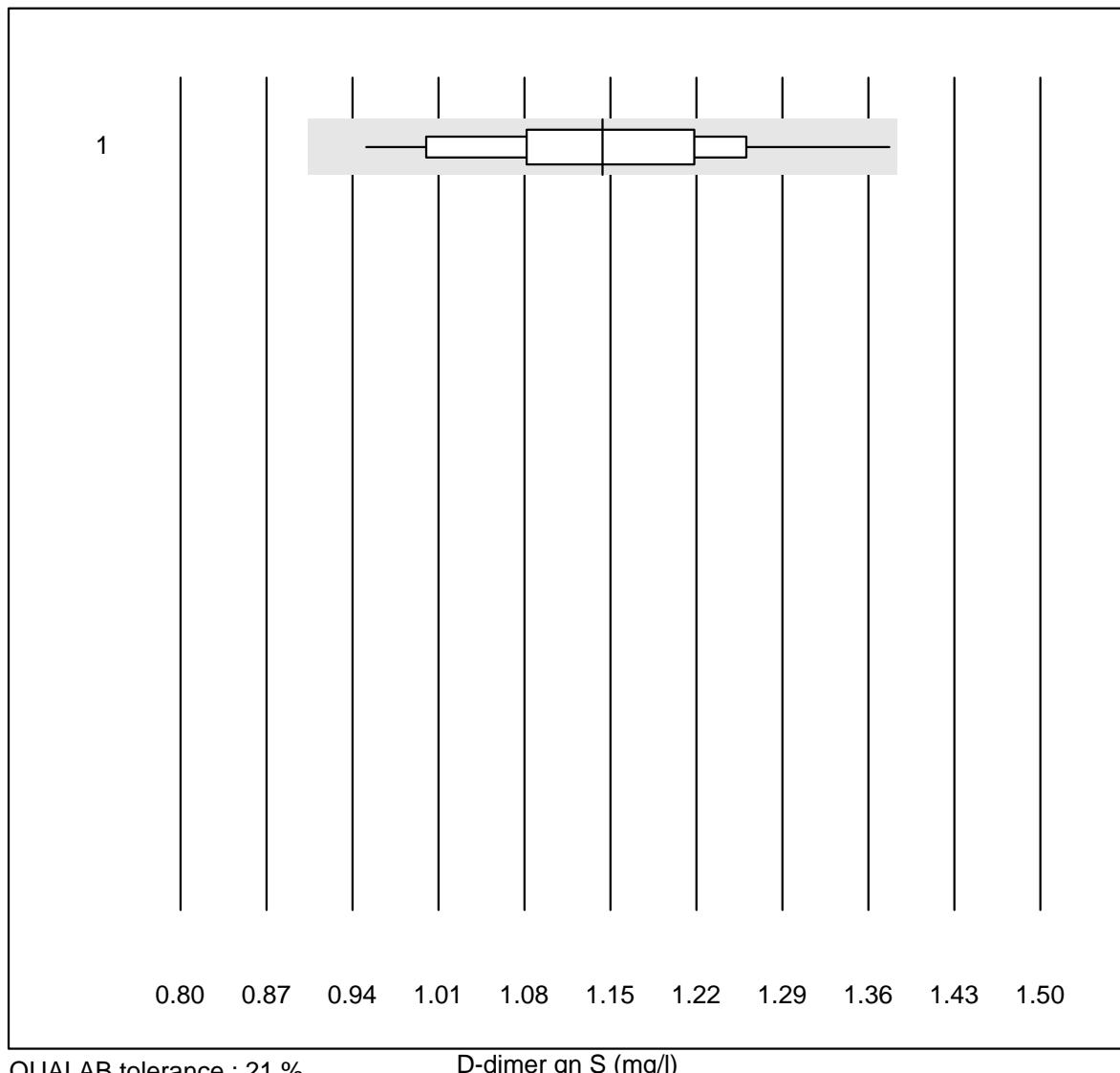


| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Cobas b101 | 44 | 84.1 | 2.3 | 13.6 | 1.31 | 6.3 | e |
| 2 Afinion | 272 | 88.6 | 1.1 | 10.3 | 1.22 | 6.7 | e |

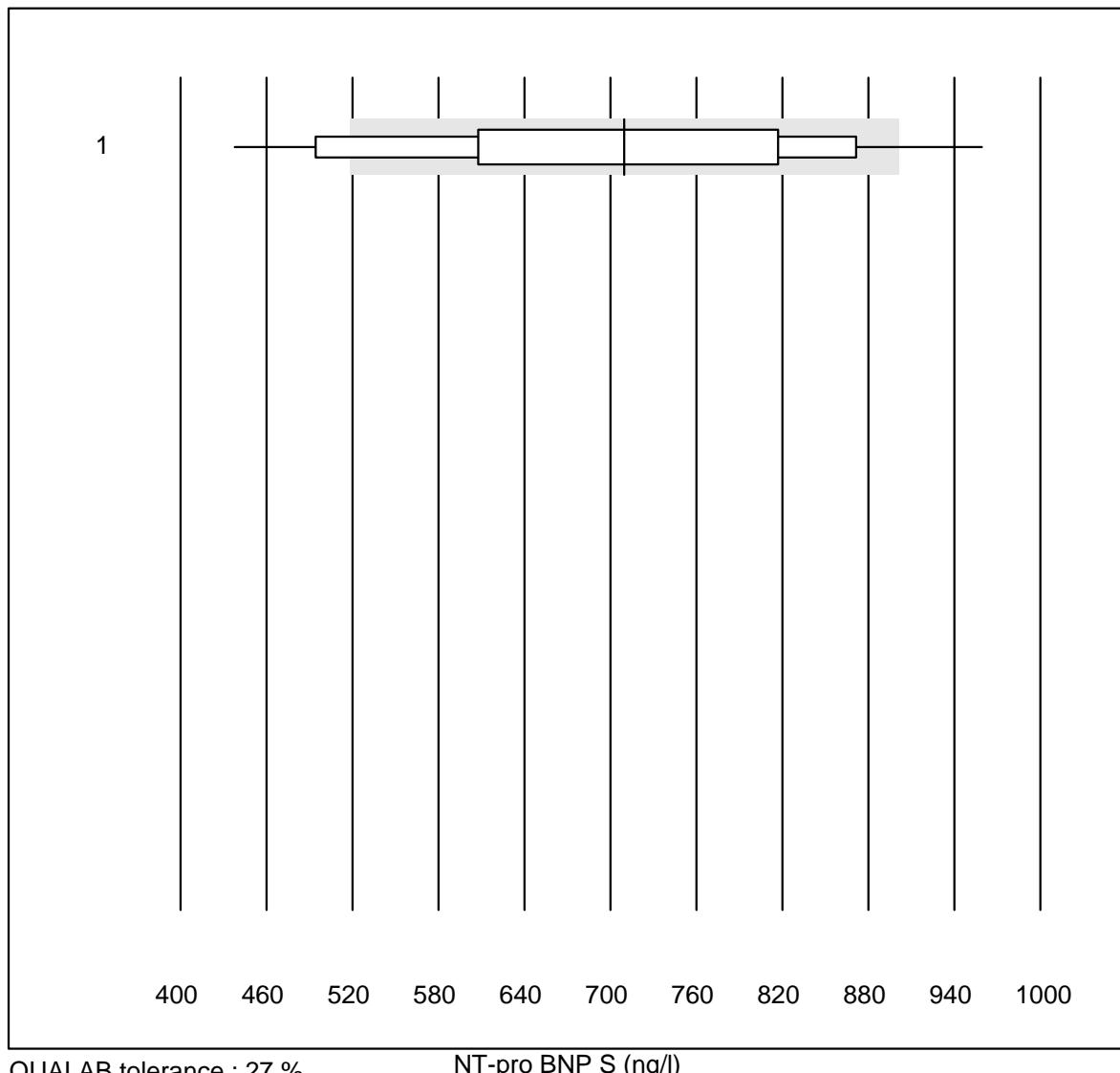
Troponin I S



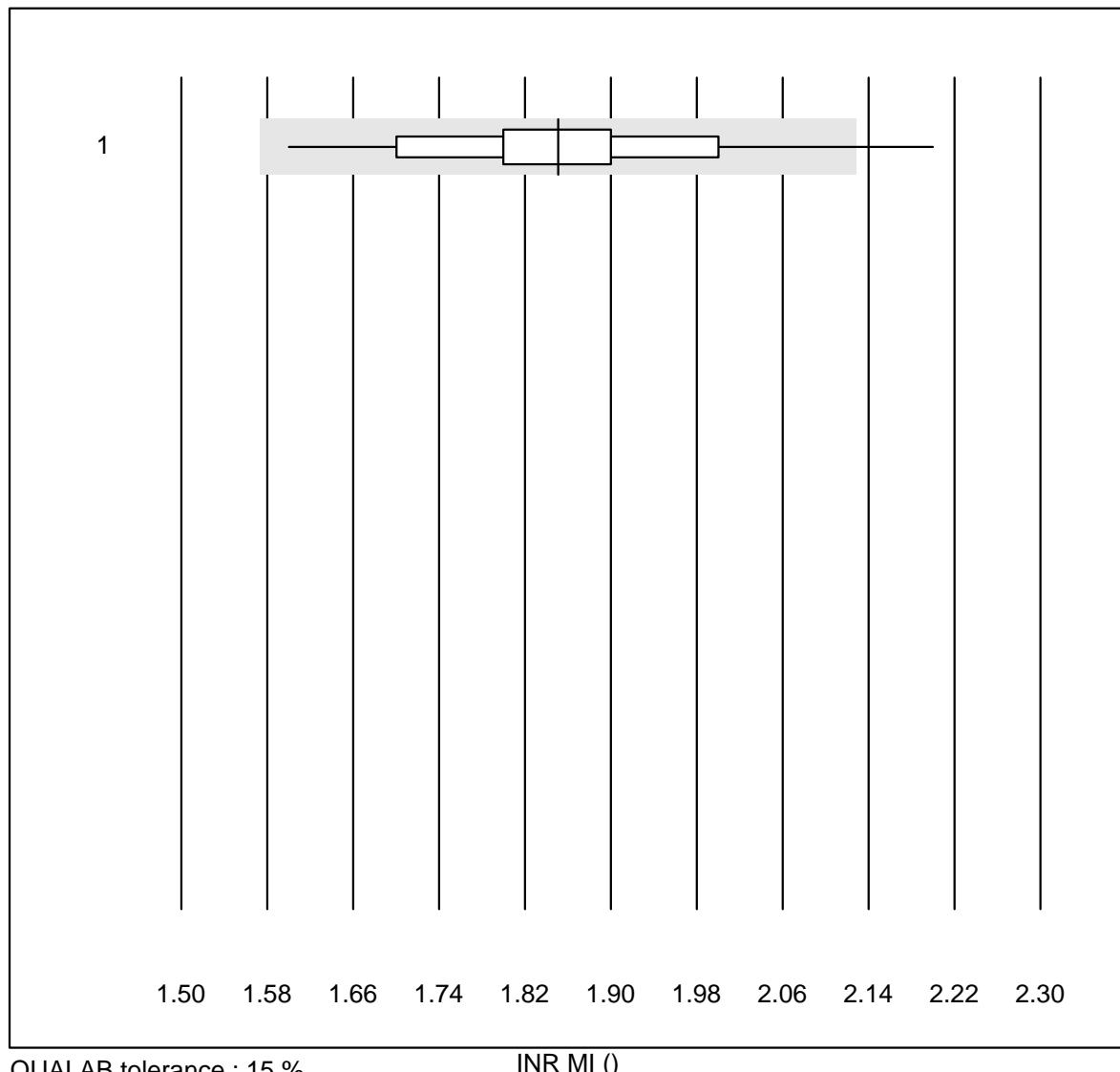
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 | Samsung LABGEO IB10 | 67 | 88.1 | 10.4 | 1.5 | 1441.89 | 14.9 | e |

D-dimer qn S

| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Samsung LABGEO IB10 | 81 | 98.8 | 0.0 | 1.2 | 1.14 | 8.9 | e |

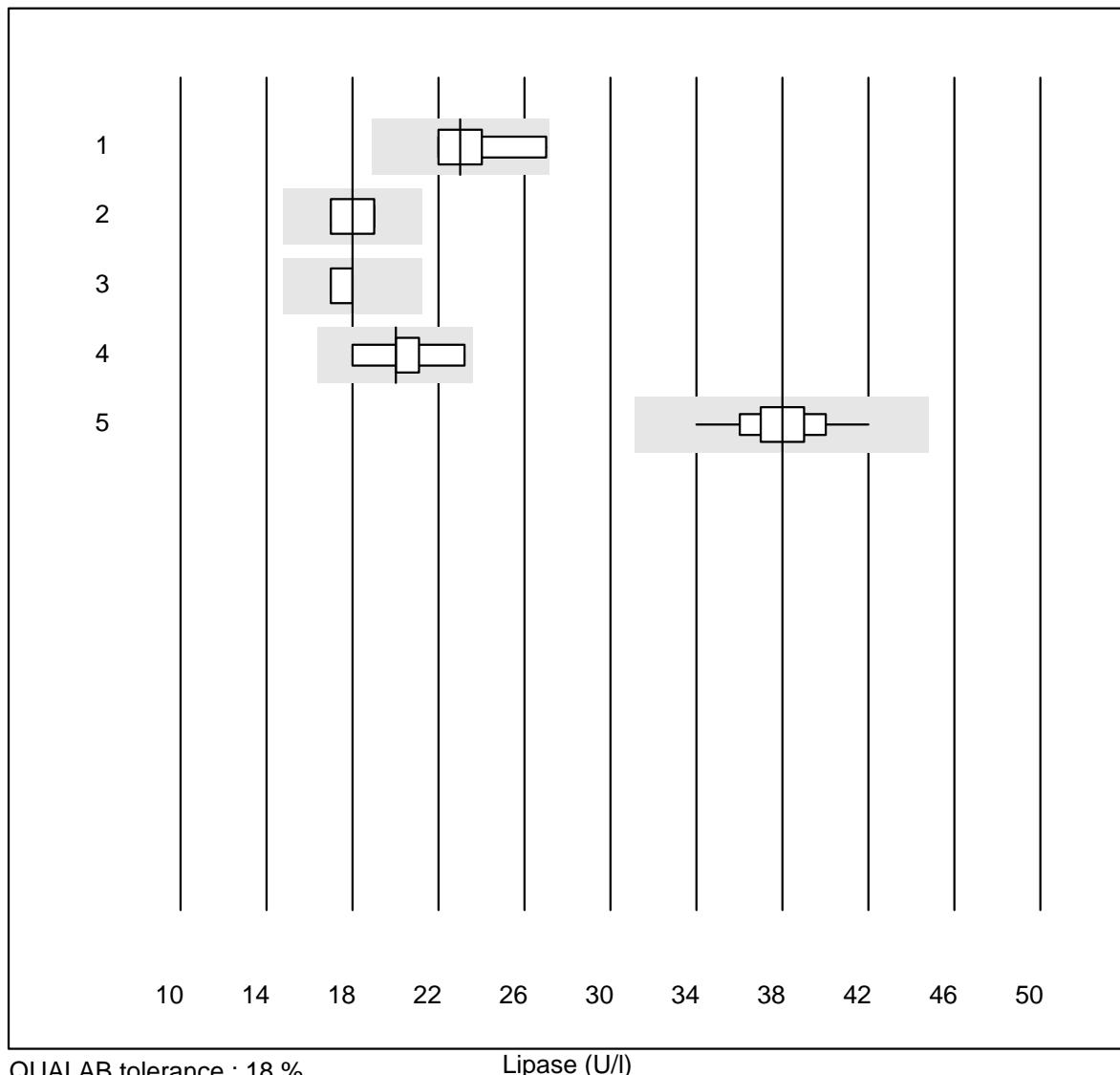
NT-pro BNP S

| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 | Samsung LABGEO IB10 | 53 | 71.7 | 20.8 | 7.5 | 709.7 | 19.7 | e |

INR MI

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 MicroINR | 72 | 88.9 | 1.4 | 9.7 | 1.9 | 7.5 | e |

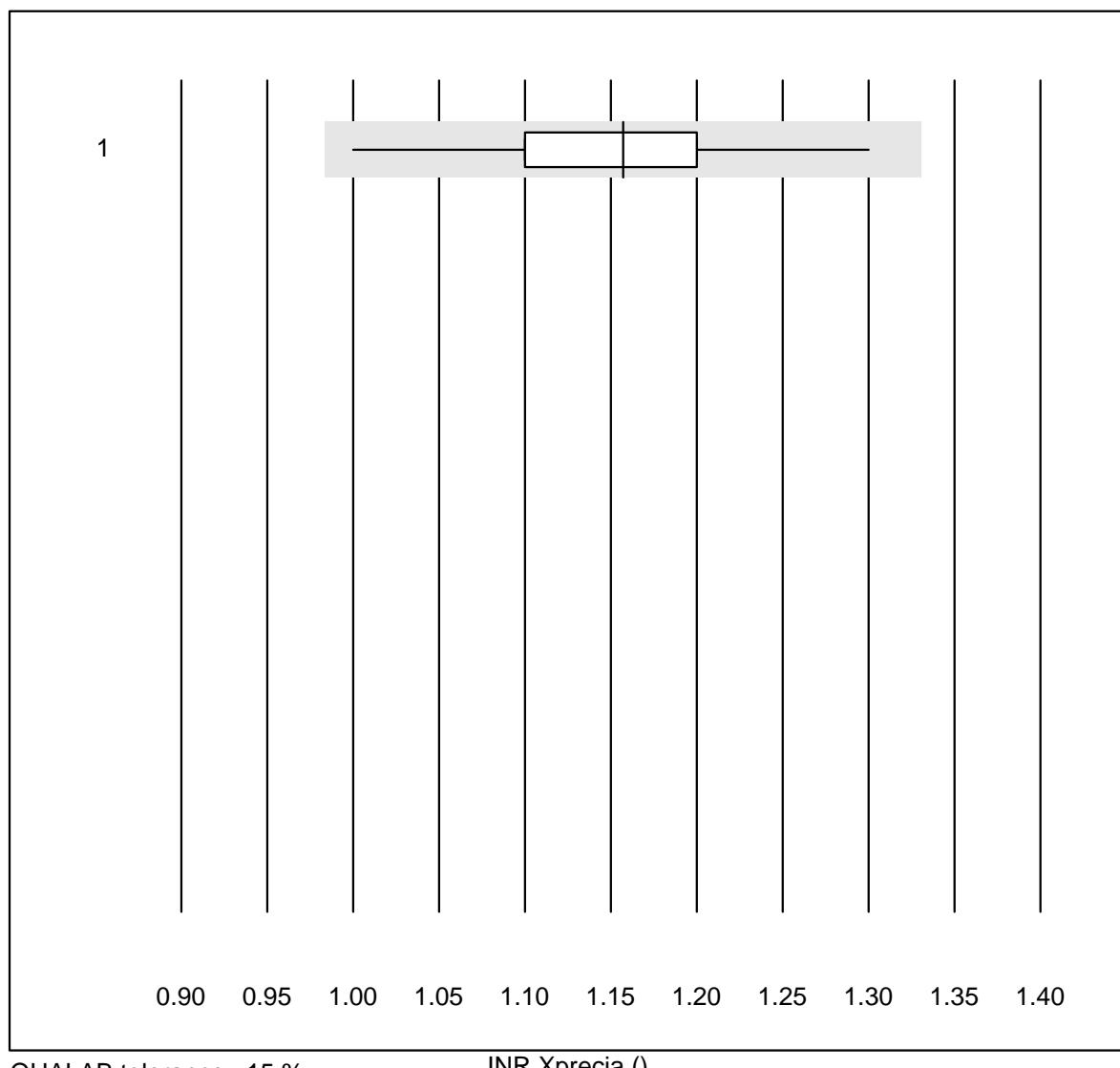
Lipase



QUALAB tolerance : 18 %

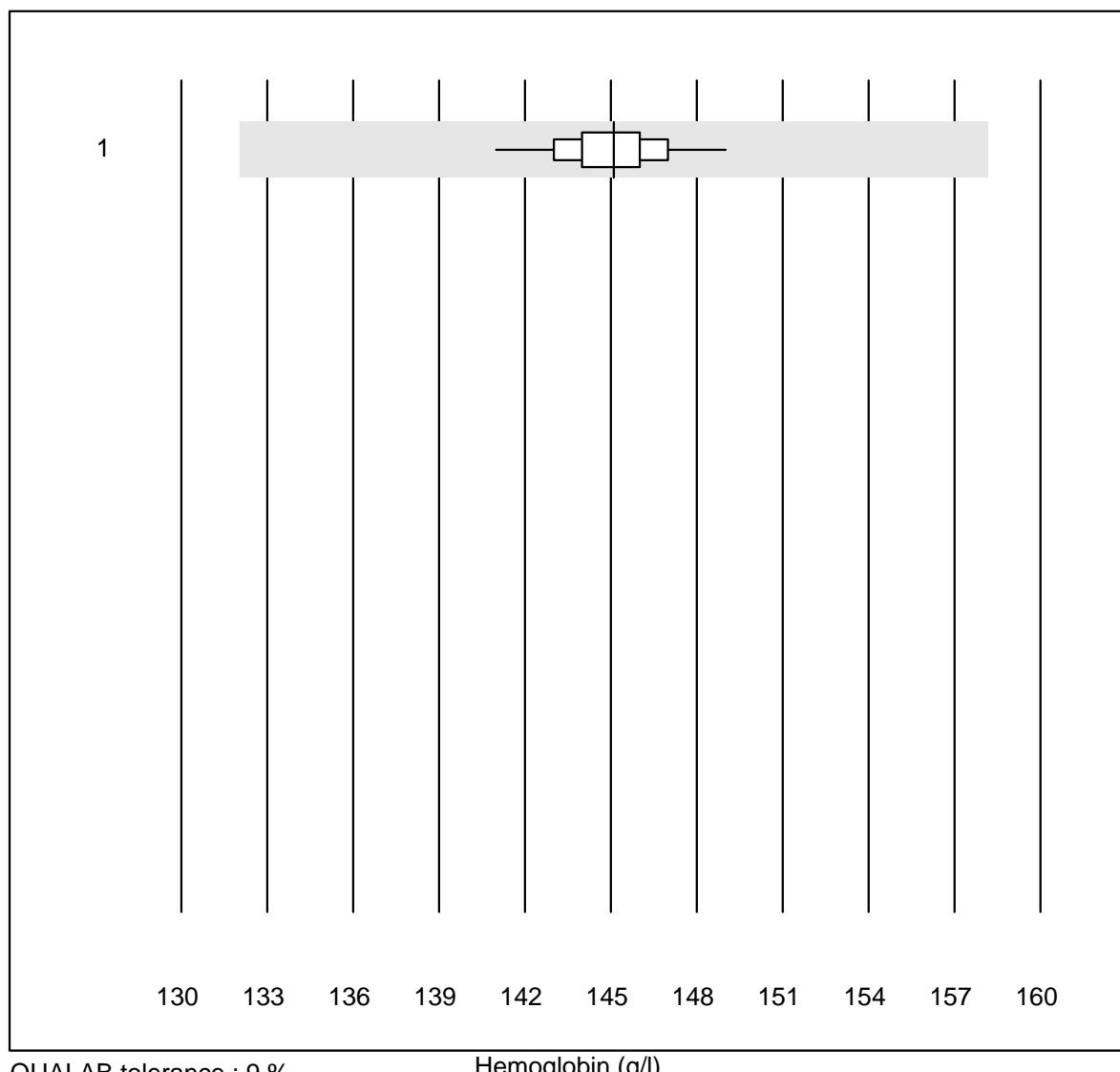
Lipase (U/l)

| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|--------------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Autolyser/DiaSys | 4 | 100.0 | 0.0 | 0.0 | 23.0 | 9.9 | e* |
| 2 Architect | 4 | 75.0 | 0.0 | 25.0 | 18.0 | 6.5 | e* |
| 3 Beckman | 5 | 80.0 | 0.0 | 20.0 | 18.0 | 2.8 | e |
| 4 Cobas | 9 | 100.0 | 0.0 | 0.0 | 20.0 | 6.9 | e* |
| 5 Fuji Dri-Chem | 88 | 98.9 | 0.0 | 1.1 | 38.0 | 4.4 | e |

INR Xprecia

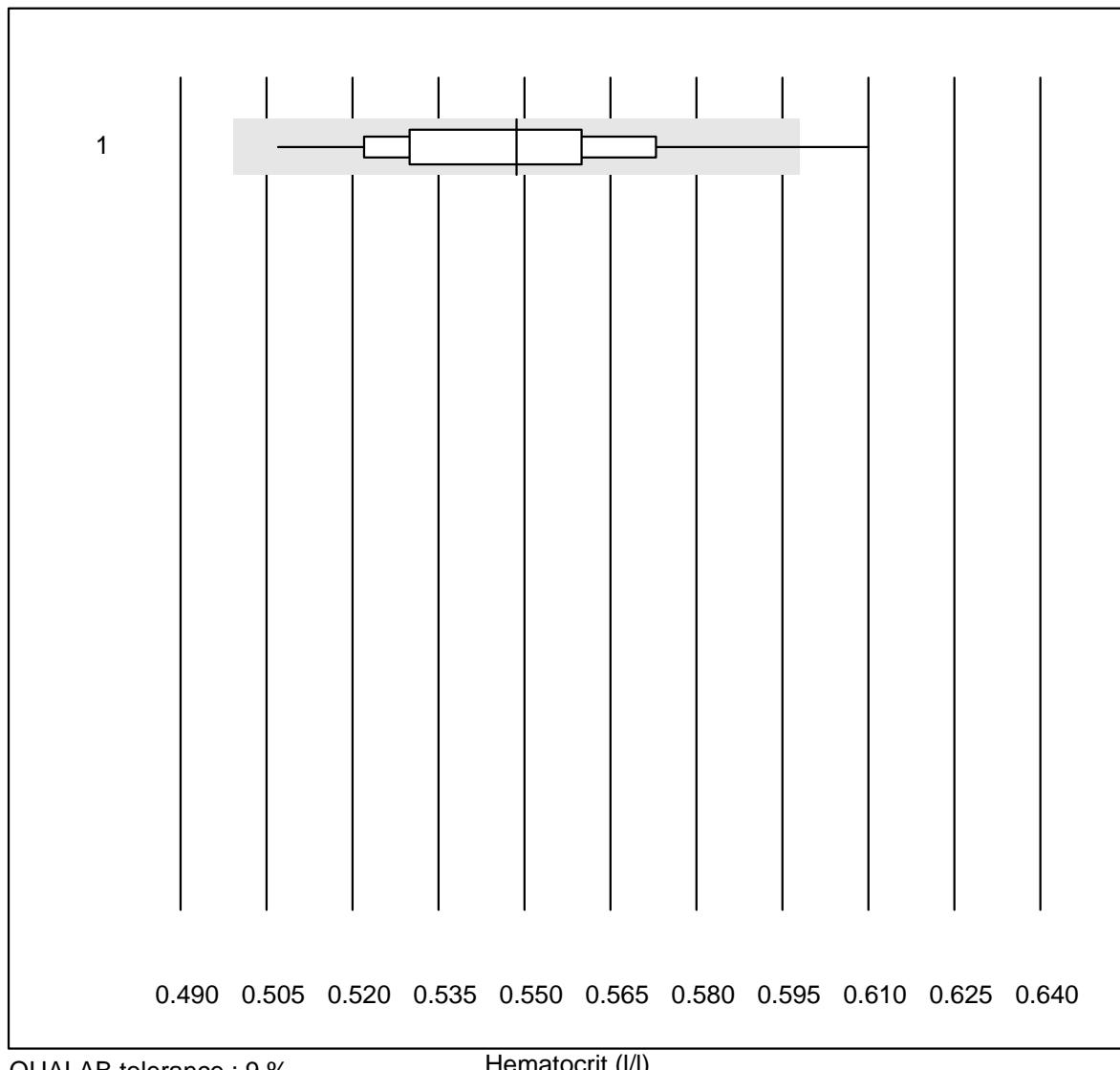
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Xprecia | 21 | 100.0 | 0.0 | 0.0 | 1.2 | 5.8 | e |

Hemoglobin



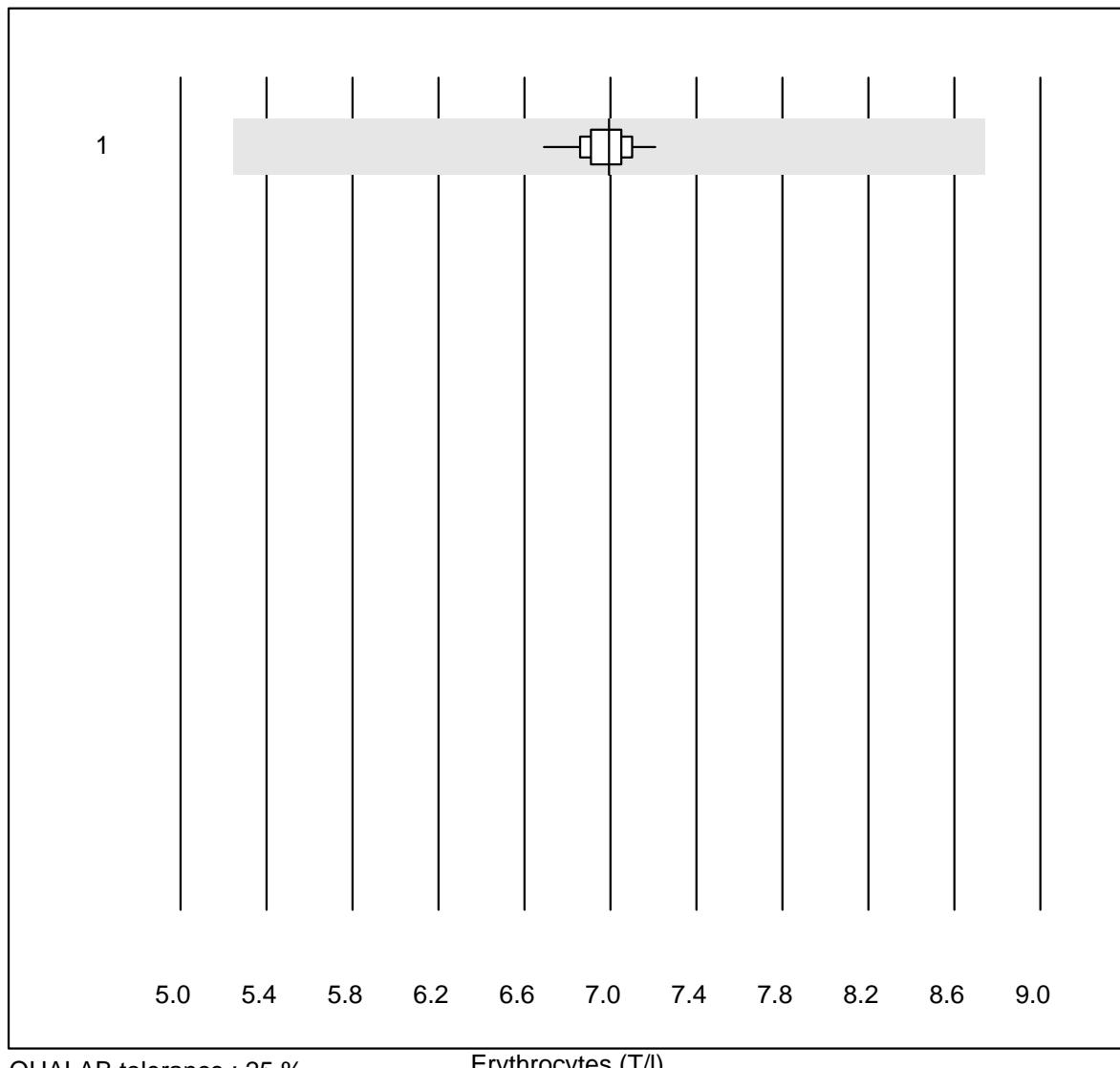
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Sysmex | 32 | 100.0 | 0.0 | 0.0 | 145.1 | 1.1 | e |

Hematocrit



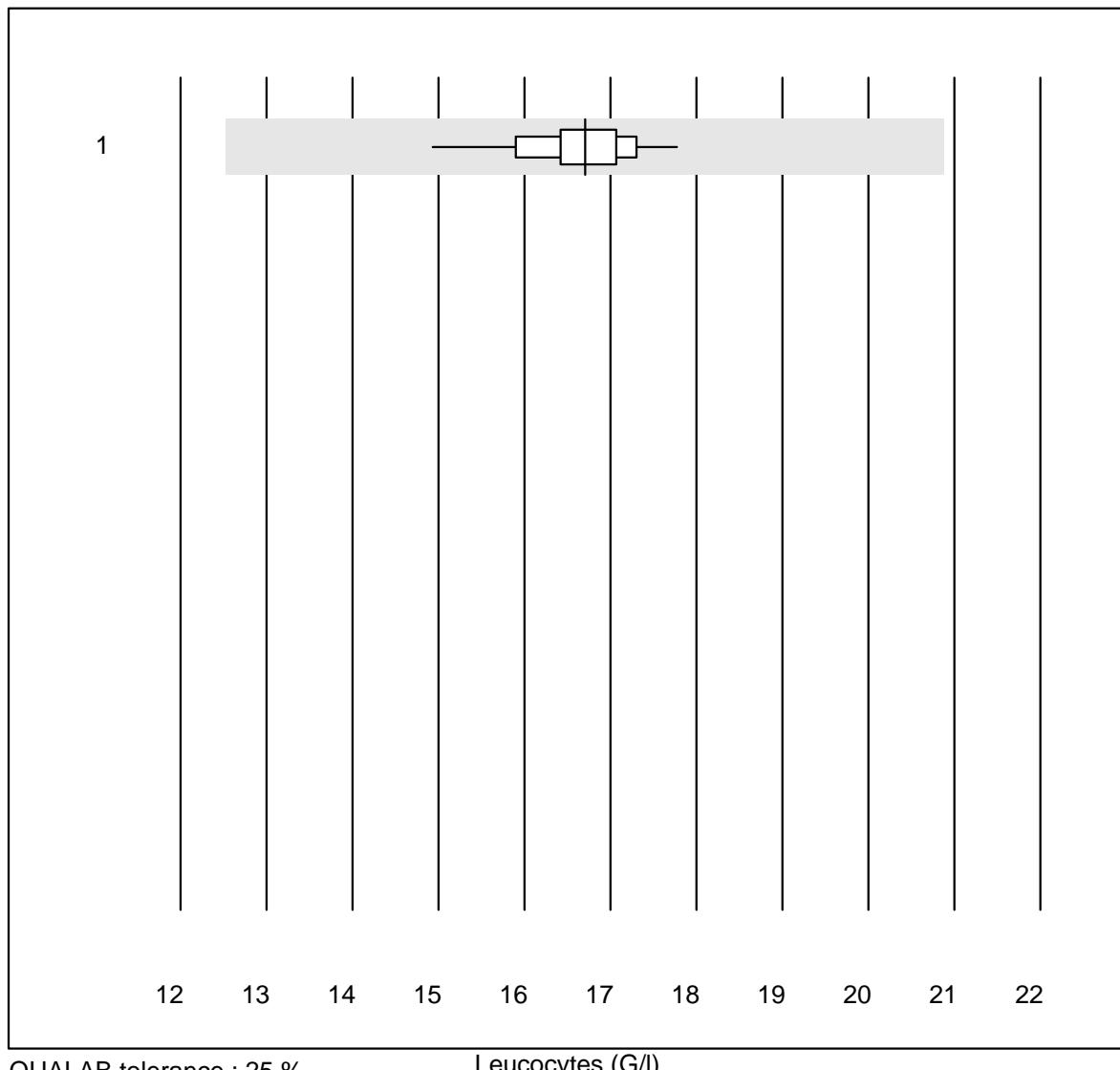
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Sysmex | 32 | 96.9 | 3.1 | 0.0 | 0.55 | 4.1 | e |

Erythrocytes



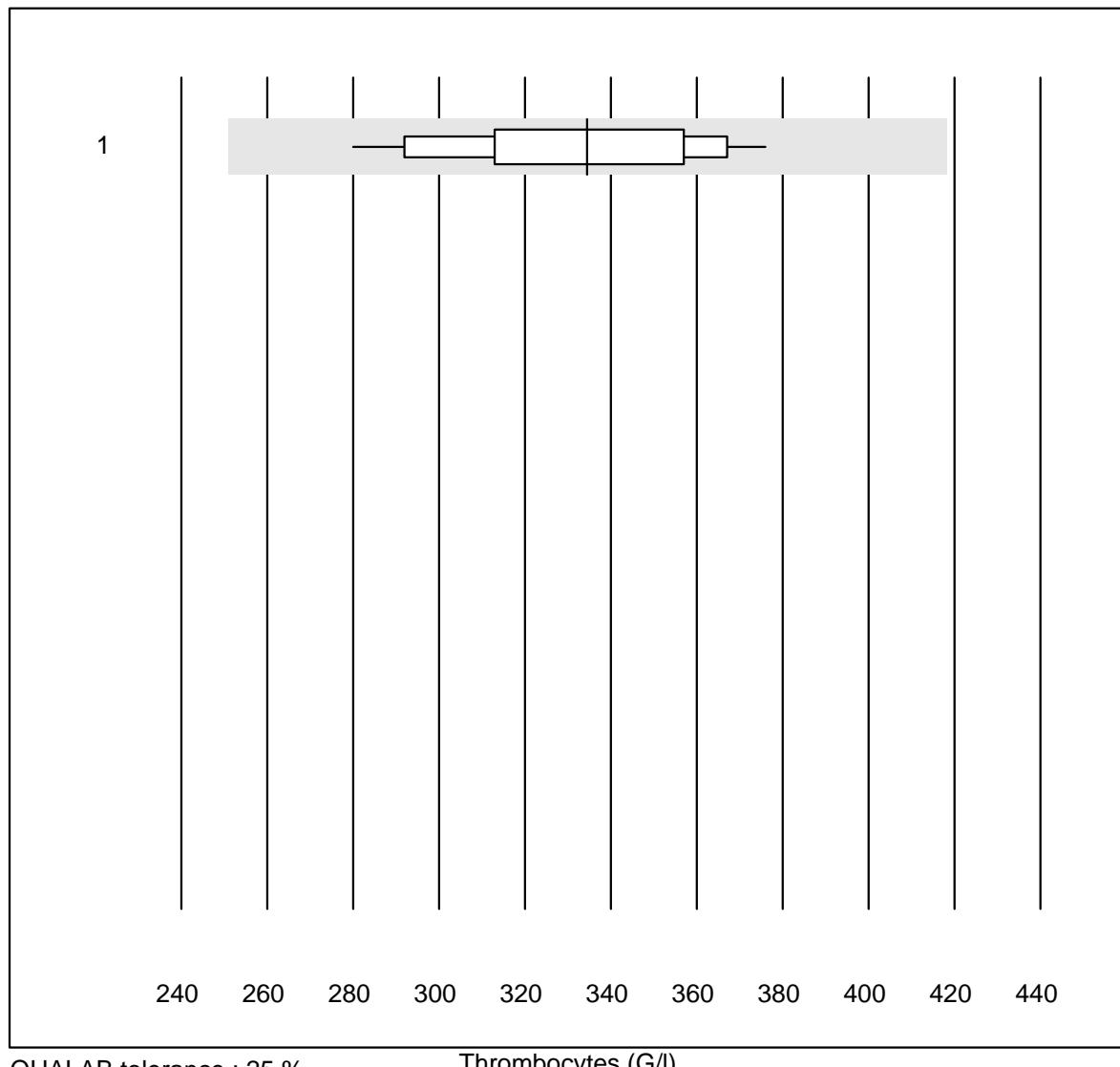
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Sysmex | 32 | 100.0 | 0.0 | 0.0 | 6.99 | 1.5 | e |

Leucocytes



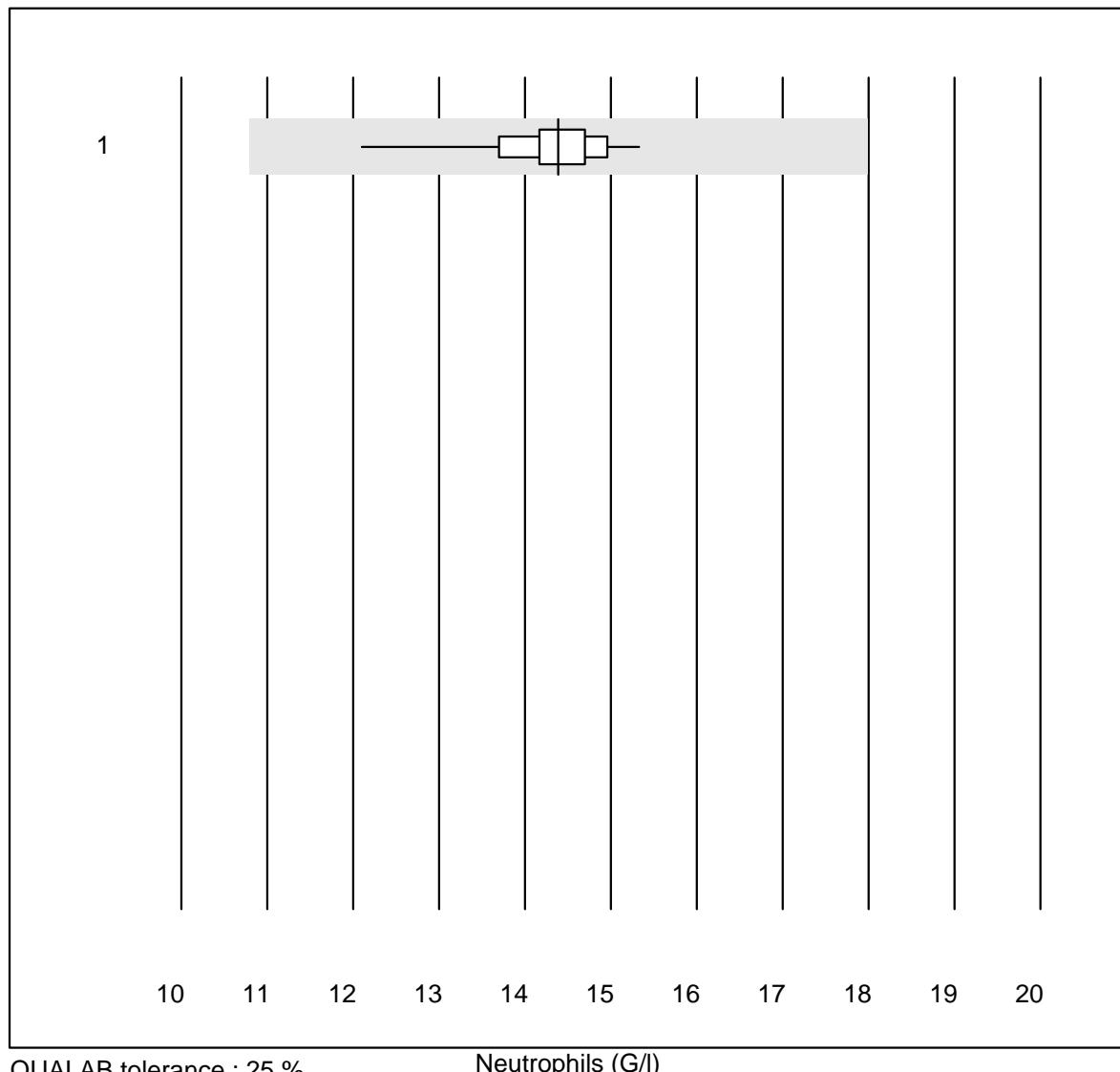
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Sysmex | 32 | 100.0 | 0.0 | 0.0 | 16.71 | 3.5 | e |

Thrombocytes



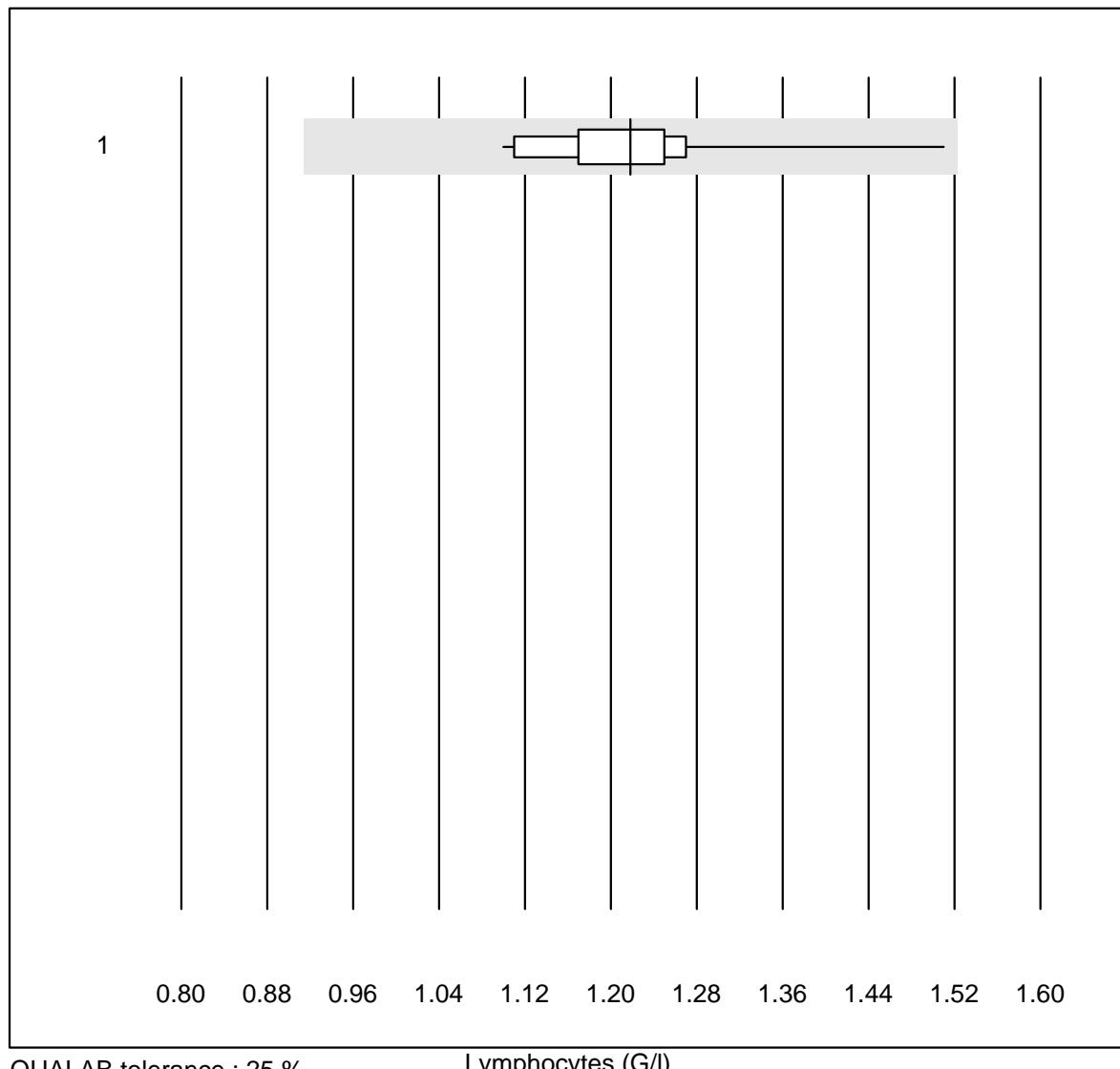
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Sysmex | 32 | 100.0 | 0.0 | 0.0 | 334.5 | 8.1 | e |

Neutrophils



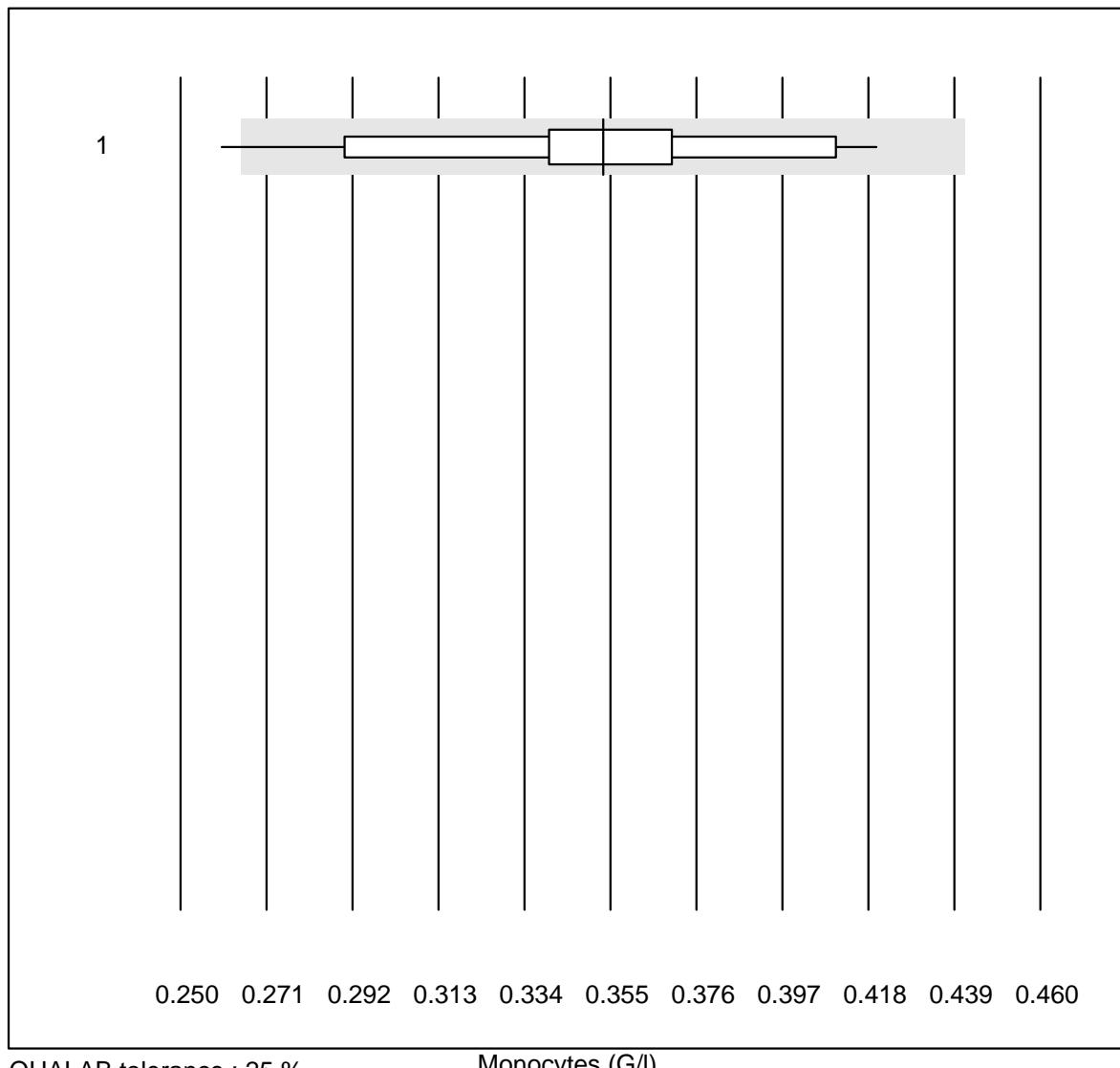
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Sysmex | 32 | 100.0 | 0.0 | 0.0 | 14.38 | 4.2 | e |

Lymphocytes



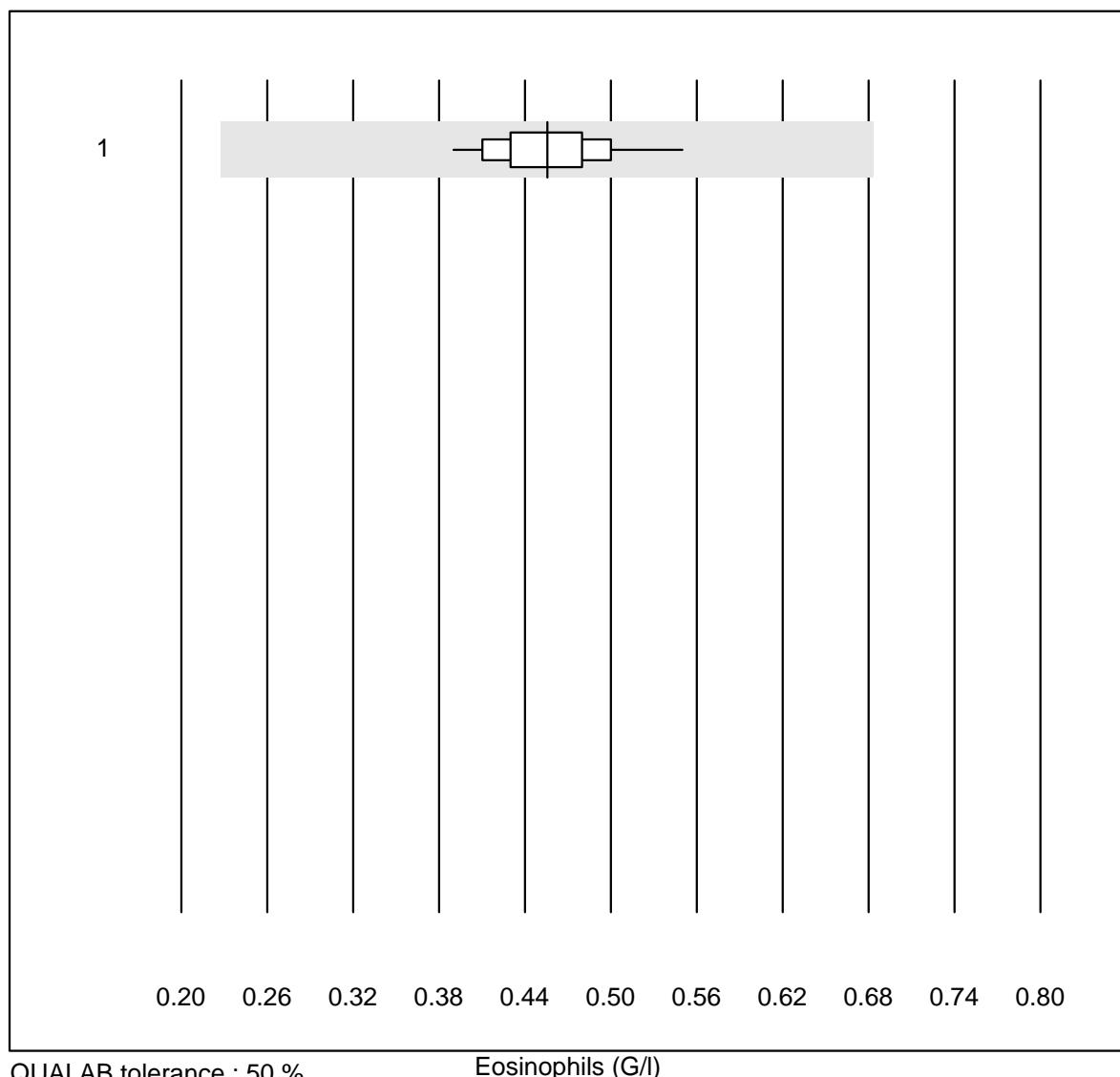
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Sysmex | 32 | 100.0 | 0.0 | 0.0 | 1.22 | 6.5 | e |

Monocytes



| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 | Sysmex | 32 | 81.3 | 3.1 | 15.6 | 0.35 | 11.6 | e |

Eosinophils

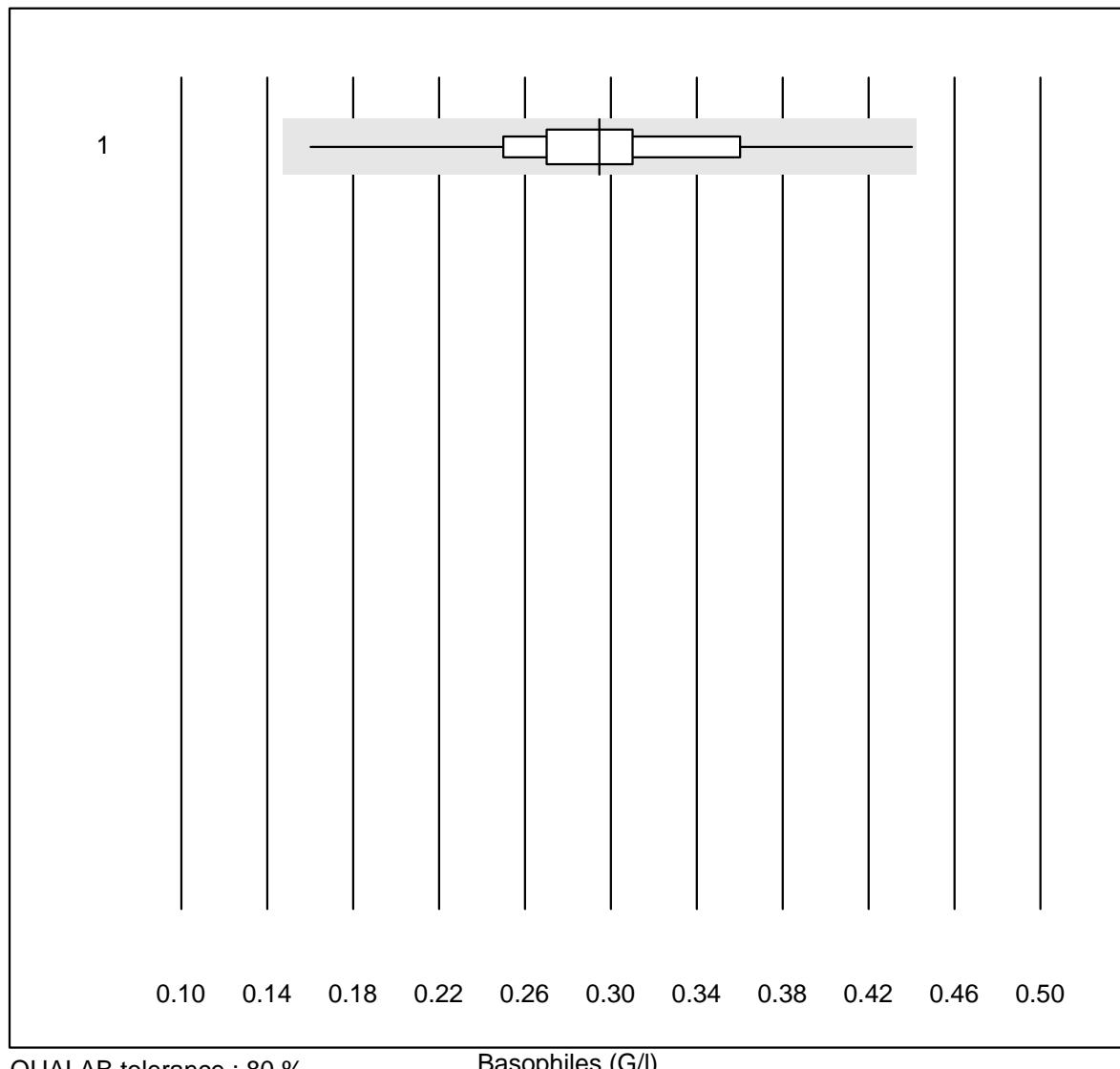


QUALAB tolerance : 50 %

Eosinophils (G/l)

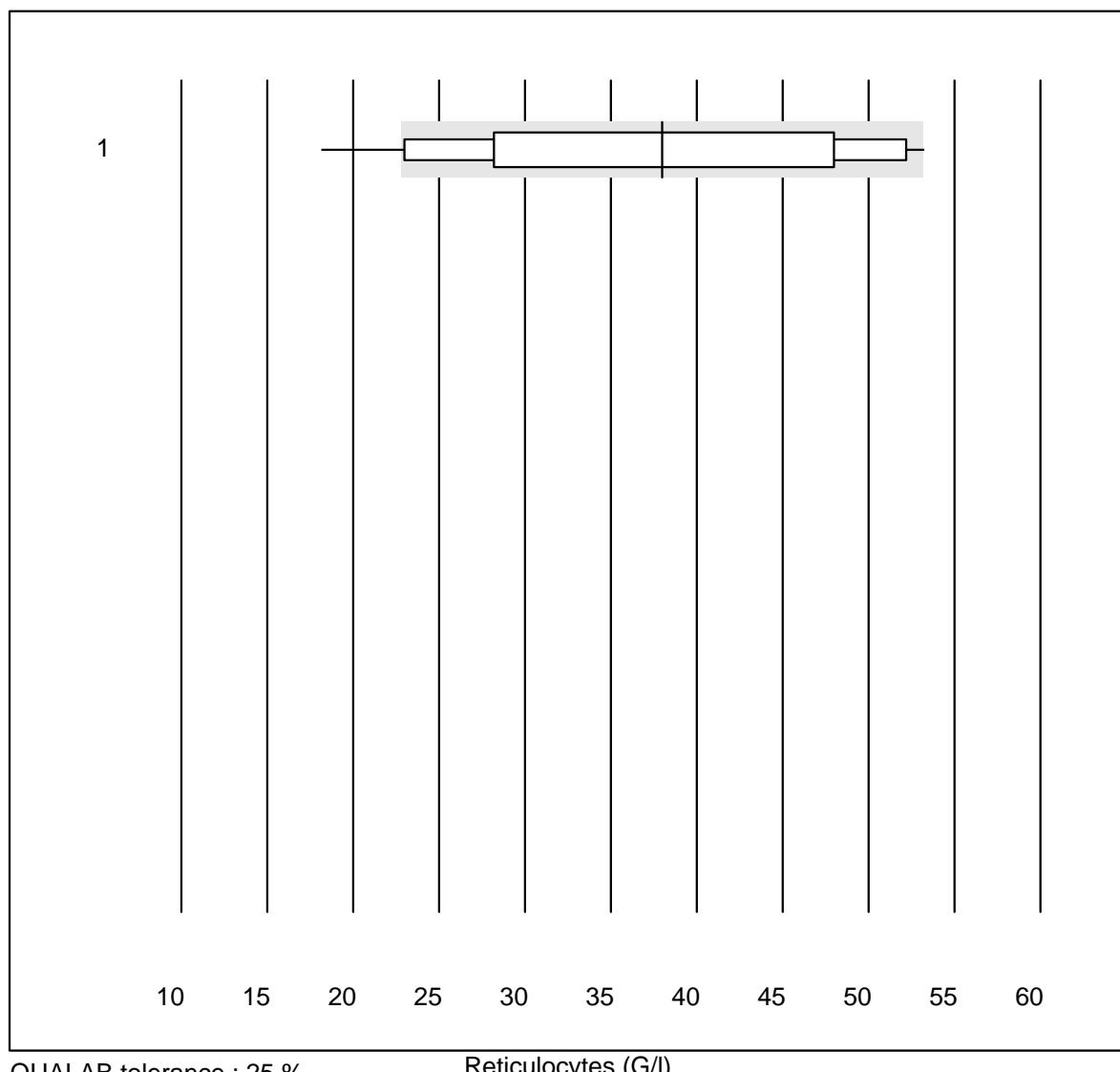
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Sysmex | 32 | 100.0 | 0.0 | 0.0 | 0.46 | 8.0 | e |

Basophiles



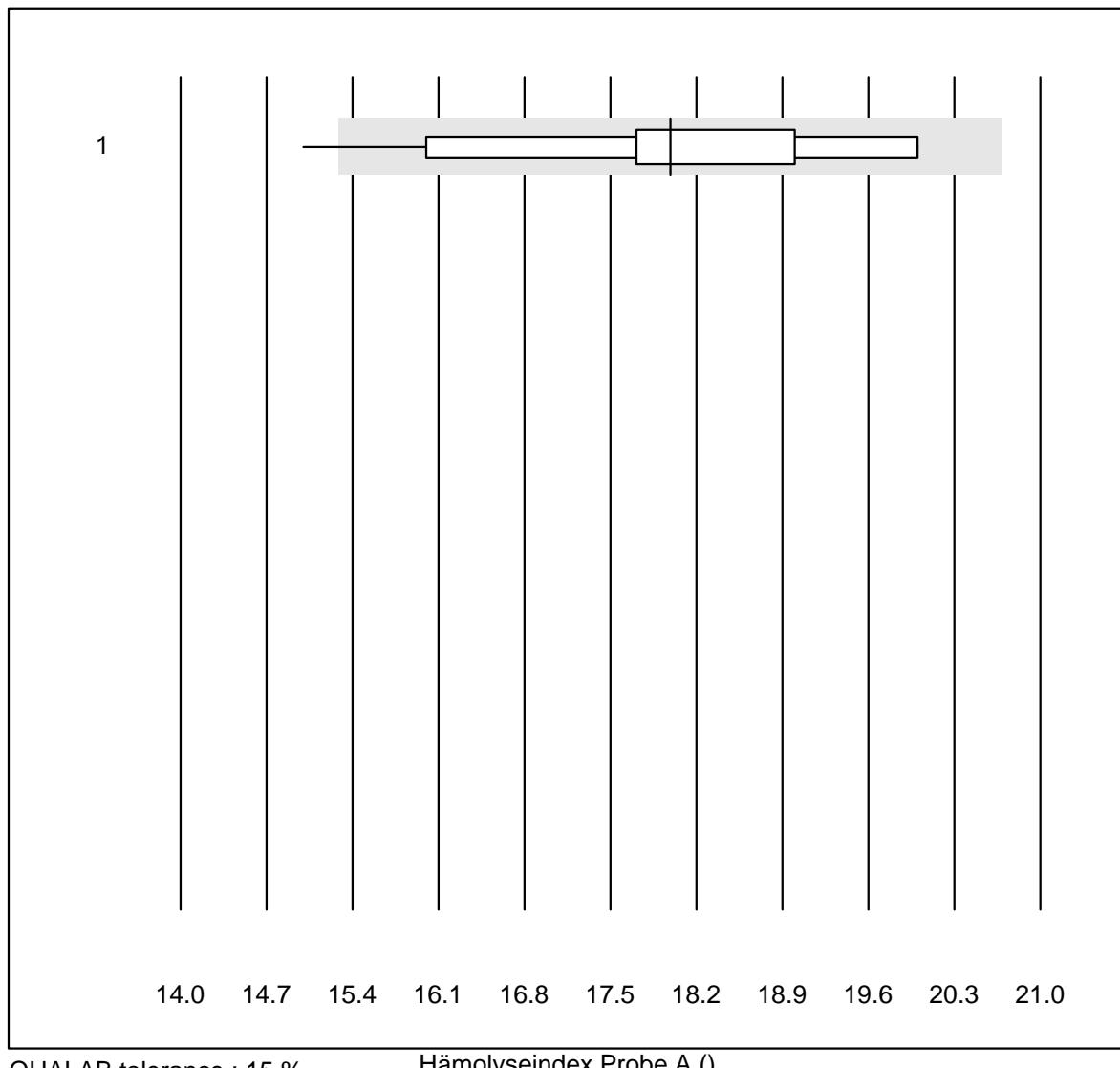
| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 | Sysmex | 32 | 100.0 | 0.0 | 0.0 | 0.29 | 18.6 | a |

Reticulocytes



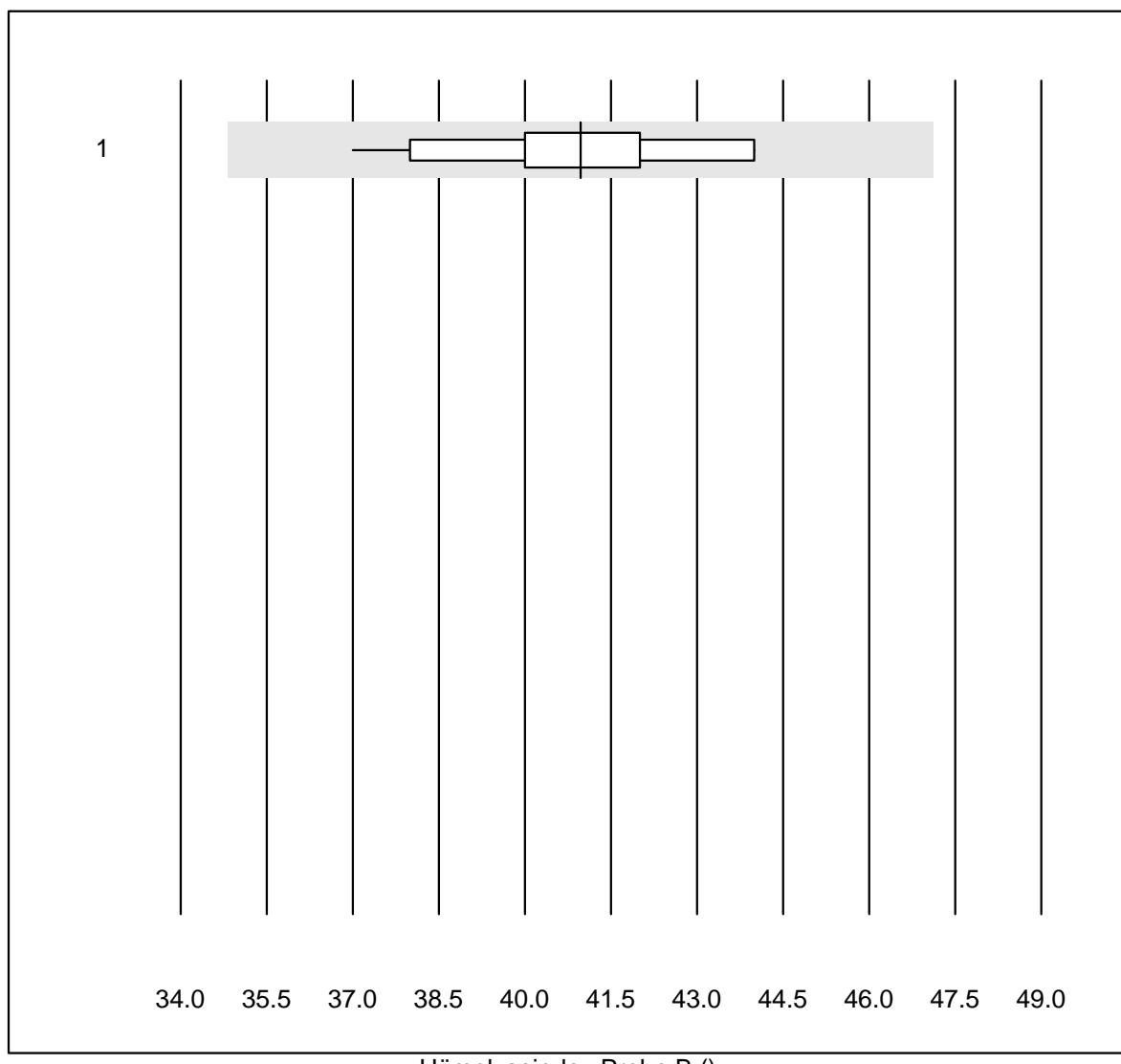
| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-------------|-------|--------|-----------|-----------|--------------|------|-----|
| 1 Sysmex | 16 | 81.2 | 12.5 | 6.3 | 38.0 | 29.8 | a |

Hämolyseindex Probe A



| No. | Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-----|---------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 | Cobas | 14 | 92.9 | 7.1 | 0.0 | 18.0 | 8.1 | e* |

Hämolyseindex Probe B



| No. Methode | Total | % good | % insuff. | % outlier | Target value | CV% | Typ |
|-------------|-------|--------|-----------|-----------|--------------|-----|-----|
| 1 Cobas | 14 | 100.0 | 0.0 | 0.0 | 41.0 | 5.3 | e |