Verein für
Association pour le
Associazione per il

medizinische Qualitätskontrolle
contrôle de qualité médical
controllo di qualità medico

Rapporto del controllo circolare

2015 - 3

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Campioni

Prima e durante la spedizione sono state controllate l'omogeneità e la stabilità di tutti i campioni e non sono state riscontrate irregolarità. I test sono stati condotti nei laboratori dell'ospedale universitario di Zurigo (http://www.uzl.usz.ch/).
I seguenti campioni sono stati prodotti appositamente per MQ in subappalto:
B1 Strep A Test, B2 Uricult, H4 Ematologia parassitaria, K14 Marker tumorale.

Determinazione dei valori assegnati

Per ogni valore assegnato è indicato il tipo di procedura secondo ISO17043:2010, B2.1 (colonna "tipo"):

a) Valori noti, derivati dalla formulazione del materiale
b) Valori di riferimento certificati per campioni particolari
c) Valori di riferimento, determinati da analisi
d) Valori di consenso da laboratori partecipanti esperti
e) Valori di consenso dai partecipanti

In gruppi con più di 9 partecipanti i valori assegnati vengono in genere determinati con il valore di consenso ("e"). Per la determinazione del valore bersaglio viene utilizzato il valore medio del collettivo di quel metodo. I valori con una deviazione rispetto al valore teorico superiore a 1.5 volte la tolleranza Qualab vengono considerati outlier ed eliminati dal calcolo del valore bersaglio. Come valore di partenza per l'eliminazione degli outlier si utilizzano i risultati degli esami di idoneità.
Per garantire a tutti i partecipanti valori assegnati rappresentativi, in gruppi più piccoli possono essere adottate anche altre procedure.

Incertezza dei valori assegnati

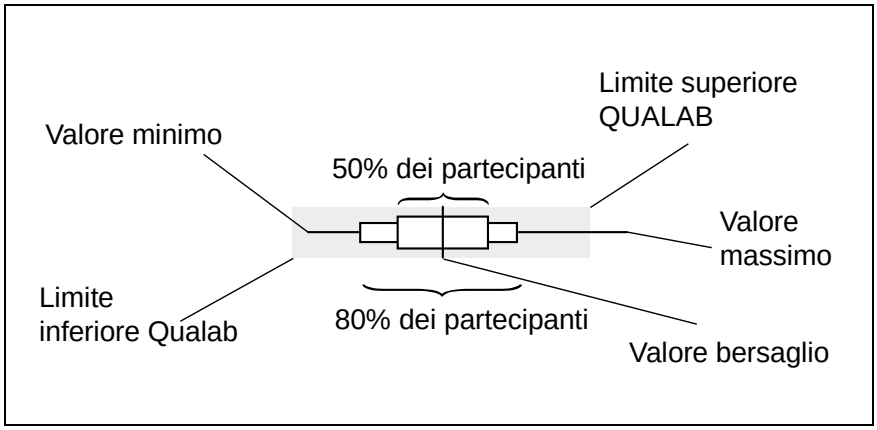
L'incertezza standard (ux) viene calcolata con la seguente formula (ISO13528):
ux = (valore assegnato/100) * 1.25/ radice quadrata del numero di partecipanti) * coeff. variazione (CV)%
ux ha la stessa unità di misura del valore assegnato
ux è paragonabile alla deviazione standard (SD) del collettivo dei partecipanti (SD: valore assegnato* CV%/100)
Se i numero dei partecipanti è superiore a 18, l'incertezza standard è molto inferiore alla variabilità' del collettivo e può essere ignorata

Tolleranze QUALAB e MQ

Per le analisi obbligatorie vengono utilizzate le tolleranze Qualab (www.qualab.ch, externe Qualitätskontrolle). Per le analisi non obbligatorie le tolleranze vengono definite dal direttore dei controlli circolari MQ.
Se l'incertezza calcolata ux del valore assegnato è superiore al 15% della tolleranza QUALAB o MQ, appare un asterisco accanto alla lettera che descrive la procedura di calcolo del valore assegnato (per esempio "e*"), per avvisare il partecipante che l'incertezza del valore assegnato può avere un'influenza sull'esito del controllo.

Rappresentazioni grafiche

I risultati sono rappresentati graficamente come segue:



Confronto degli strumenti

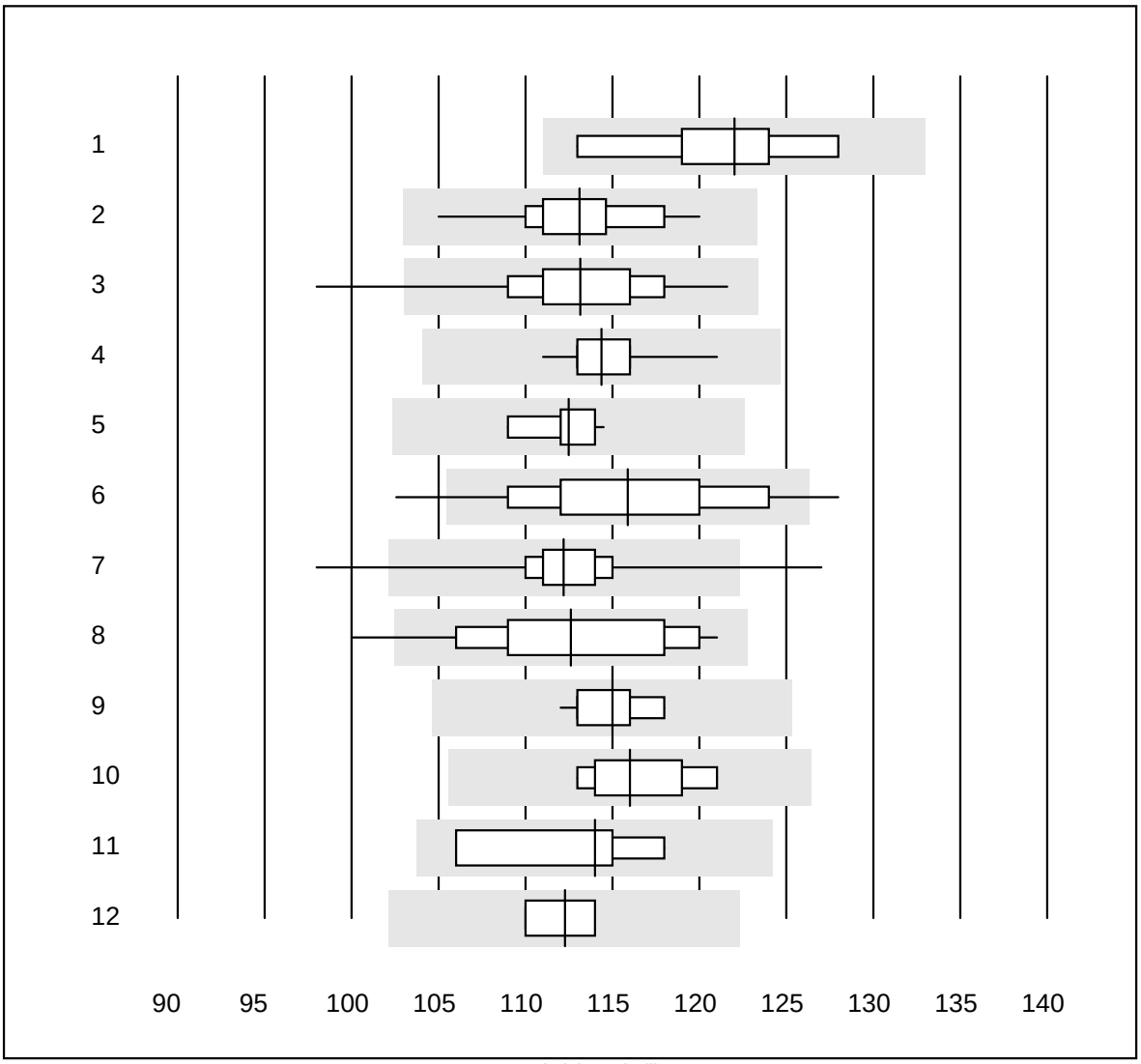
I dati in questa parte del rapporto consentono di paragonare l'efficienza dei vari strumenti. Non vanno però dimenticati i seguenti dettagli:

- Il campione di controllo K1 è un siero di controllo commerciale pronto per l'uso. Nonostante il campione sia di origine umana, non si può escludere l'insorgenza di effetti di matrice. Questi dipendono dallo strumento e portano a valori assegnati differenti.
- E' stato analizzato solo un campione. Poiché la distribuzione dei risultati dipende dalla natura del campione (effetto matrice) e dal valore stesso, i coefficienti di variazione determinati (in %) non hanno una validità generale.
- Gran parte dei valori anomali deriva da errori amministrativi (unità di misura sbagliata, scambio dei risultati) o da errori di manualità (campione sbagliato, non correttamente disciolto, non abbastanza mescolato) e non ha a che fare con lo strumento.

Zurigo, 28.9.2015

Dr. R. Fried
Direttore controlli circolari

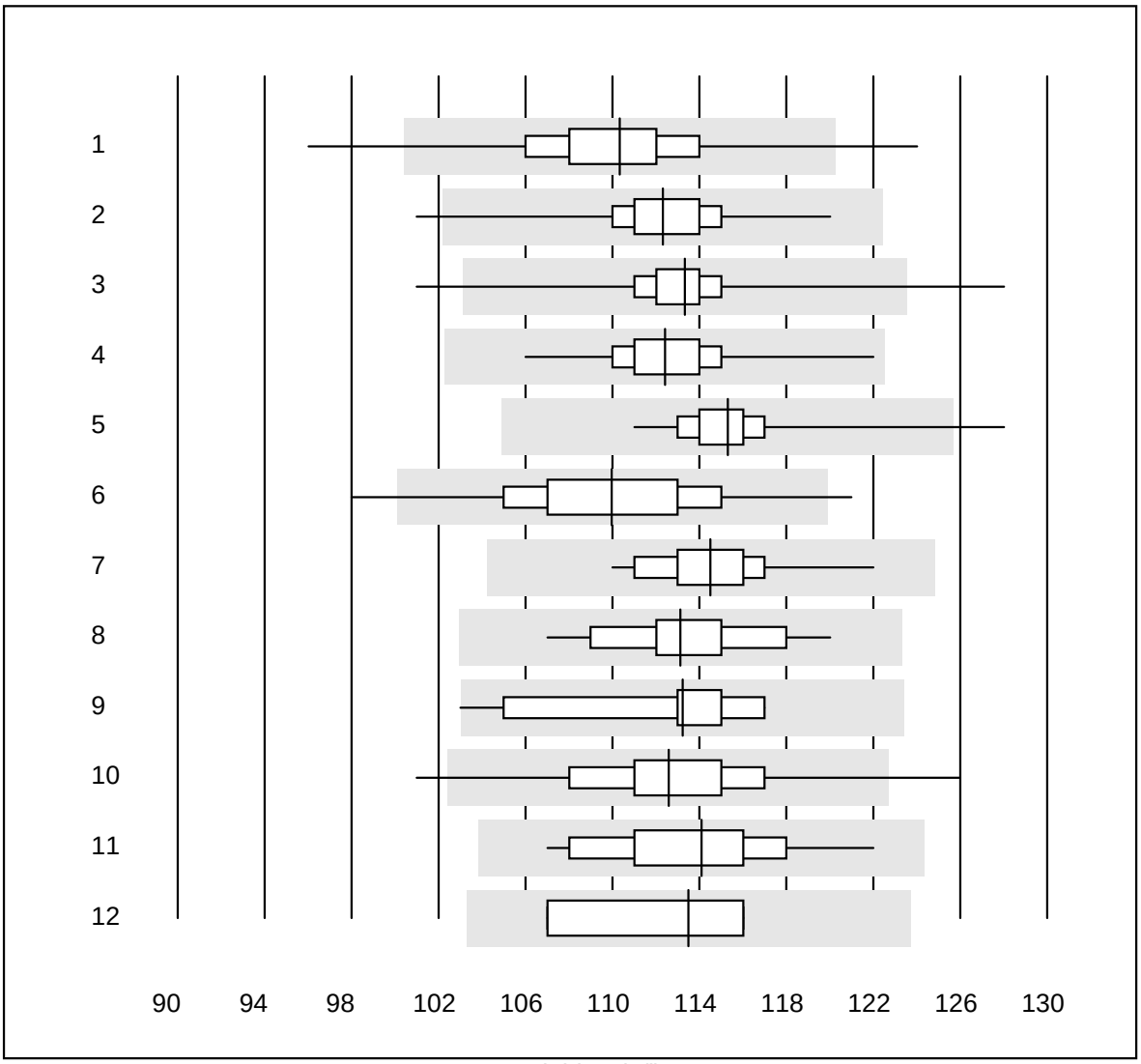
Emoglobina



Deviatione QUALAB : 9 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	DiaSpect	5	100.0	0.0	0.0	122.0	4.7	e*
2	Automatico	44	100.0	0.0	0.0	113.1	2.7	e
3	Cianometemoglobina	50	96.0	2.0	2.0	113.2	3.5	e
4	Sysmex XT/XE/XS	39	100.0	0.0	0.0	114.4	1.6	e
5	ABX Pentra	11	90.9	0.0	9.1	112.5	1.5	e
6	Reflotron	72	87.5	5.6	6.9	115.9	4.7	e
7	Hemocue	345	94.2	1.7	4.1	112.2	2.5	e
8	Dr. Lange	23	87.0	8.7	4.3	112.6	5.2	e*
9	Hemocontrol	12	100.0	0.0	0.0	115.0	1.6	e
10	Eurolyser	5	100.0	0.0	0.0	116.0	2.9	e*
11	MS4	4	100.0	0.0	0.0	114.0	4.5	e*
12	Beckman	4	100.0	0.0	0.0	112.3	1.8	a

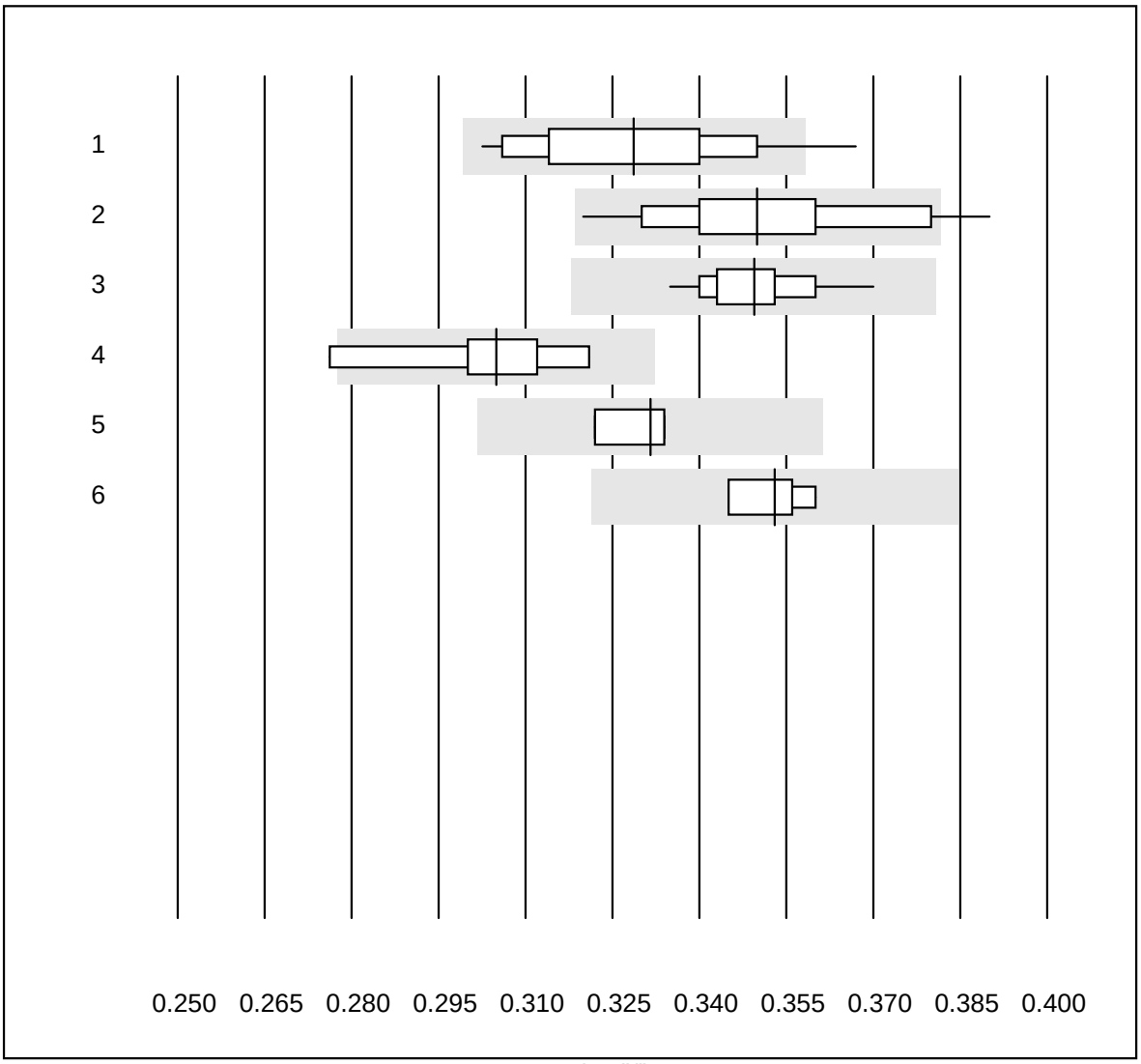
Emoglobina



Deviatione QUALAB : 9 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Abx Micros	835	96.8	0.7	2.5	110.3	3.1	e
2	Microsemi	262	97.3	0.4	2.3	112.3	2.0	e
3	Sysmex KX21	428	95.3	1.2	3.5	113.3	2.0	e
4	Sysmex PocH - 100i	213	100.0	0.0	0.0	112.4	2.1	e
5	Sysmex XP 300	192	96.9	0.5	2.6	115.3	1.8	e
6	Mythic	247	96.0	2.4	1.6	110.0	3.7	e
7	Swelab	70	98.6	0.0	1.4	114.5	2.0	e
8	Abacus Junior	13	100.0	0.0	0.0	113.1	3.2	e
9	Medonic	17	94.1	5.9	0.0	113.2	3.3	e
10	Nihon Kohden Celltac	39	89.8	5.1	5.1	112.6	3.9	e
11	Samsung HC10	44	97.7	0.0	2.3	114.1	3.2	e
12	Norma Icon 3	4	100.0	0.0	0.0	113.5	3.9	e*

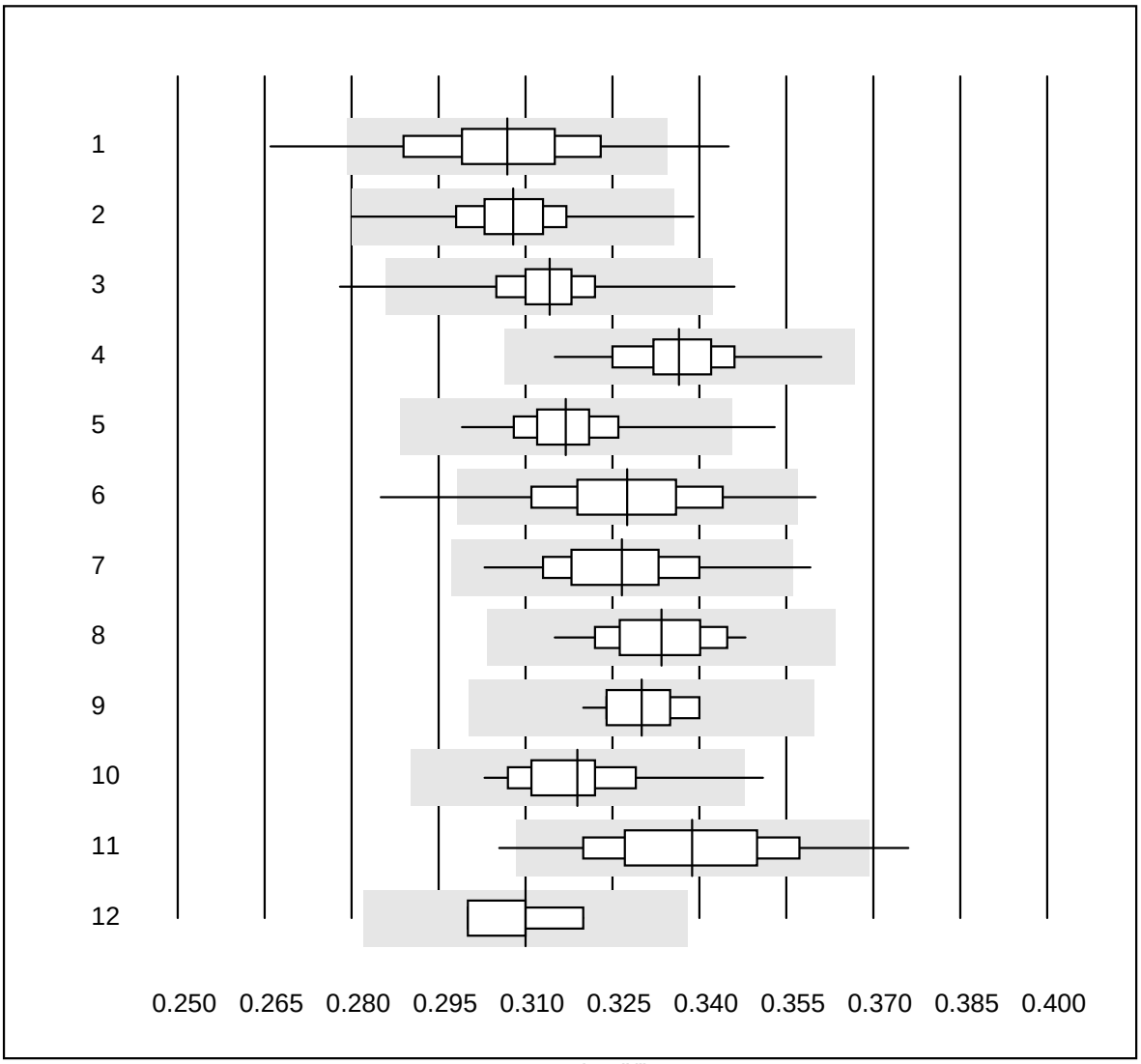
Ematocrito



Deviatione QUALAB : 9 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Automatico	37	83.8	5.4	10.8	0.33	5.2	e
2	Centrifuga	13	84.6	7.7	7.7	0.35	5.7	e*
3	Sysmex XT/XE/XS	38	100.0	0.0	0.0	0.35	2.3	e
4	ABX Pentra	10	80.0	10.0	10.0	0.30	4.4	e*
5	MS4	4	100.0	0.0	0.0	0.33	1.7	e
6	Beckman	4	100.0	0.0	0.0	0.35	1.8	a

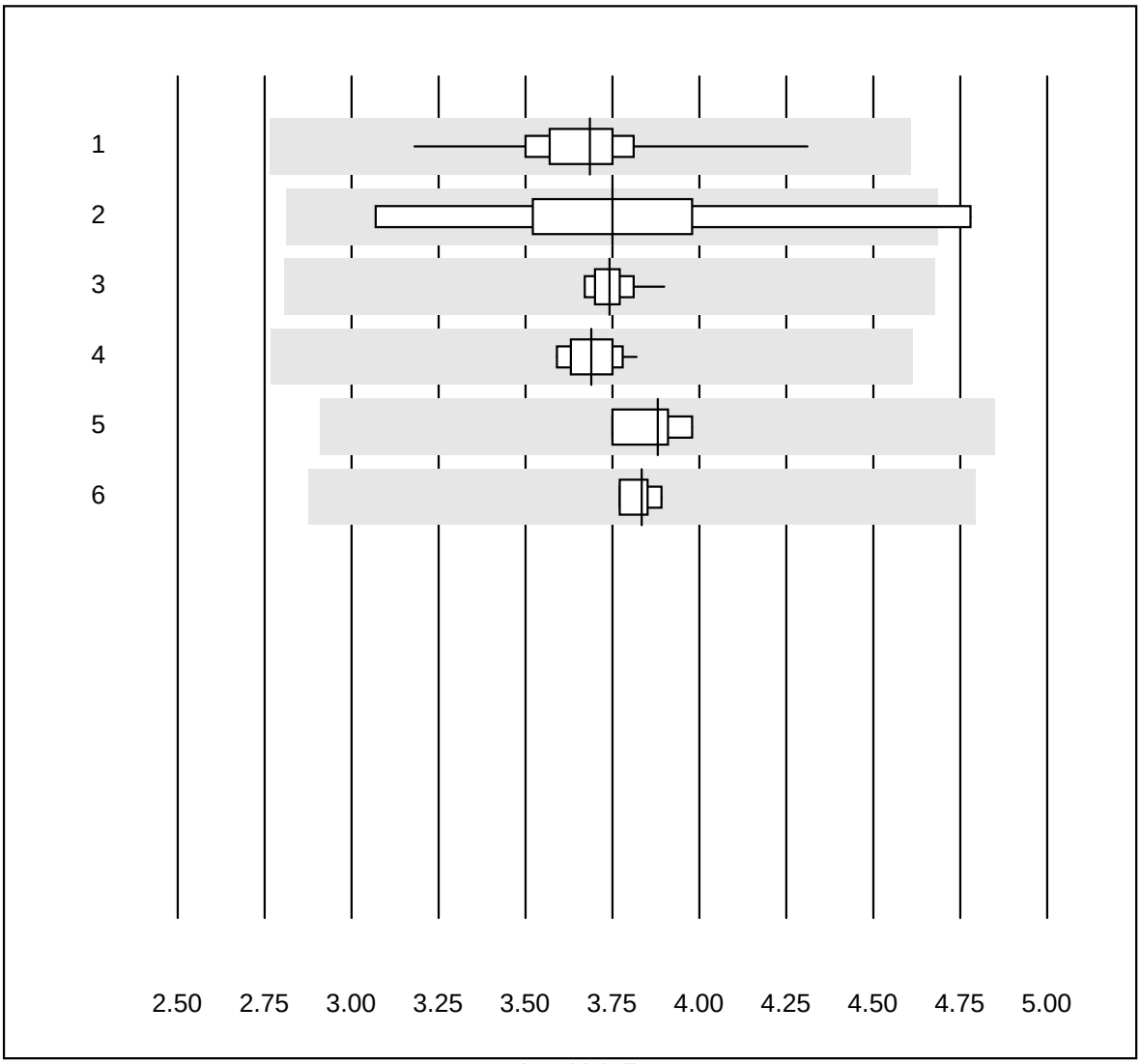
Ematocrito



Deviatione QUALAB : 9 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Abx Micros	834	91.2	4.8	4.0	0.31	4.3	e
2	Microsemi	262	98.4	0.8	0.8	0.31	2.6	e
3	Sysmex KX21	428	97.0	0.7	2.3	0.31	2.4	e
4	Sysmex PocH - 100i	213	100.0	0.0	0.0	0.34	2.4	e
5	Sysmex XP 300	188	96.8	0.5	2.7	0.32	2.3	e
6	Mythic	247	95.6	2.4	2.0	0.33	4.0	e
7	Swelab	70	97.2	1.4	1.4	0.33	3.4	e
8	Abacus Junior	13	100.0	0.0	0.0	0.33	2.9	e
9	Medonic	17	94.1	0.0	5.9	0.33	2.0	e
10	Nihon Kohden Celltac	39	92.3	2.6	5.1	0.32	3.0	e
11	Samsung HC10	44	90.9	6.8	2.3	0.34	4.7	e
12	Norma Icon 3	4	100.0	0.0	0.0	0.31	2.6	e*

Eritrociti

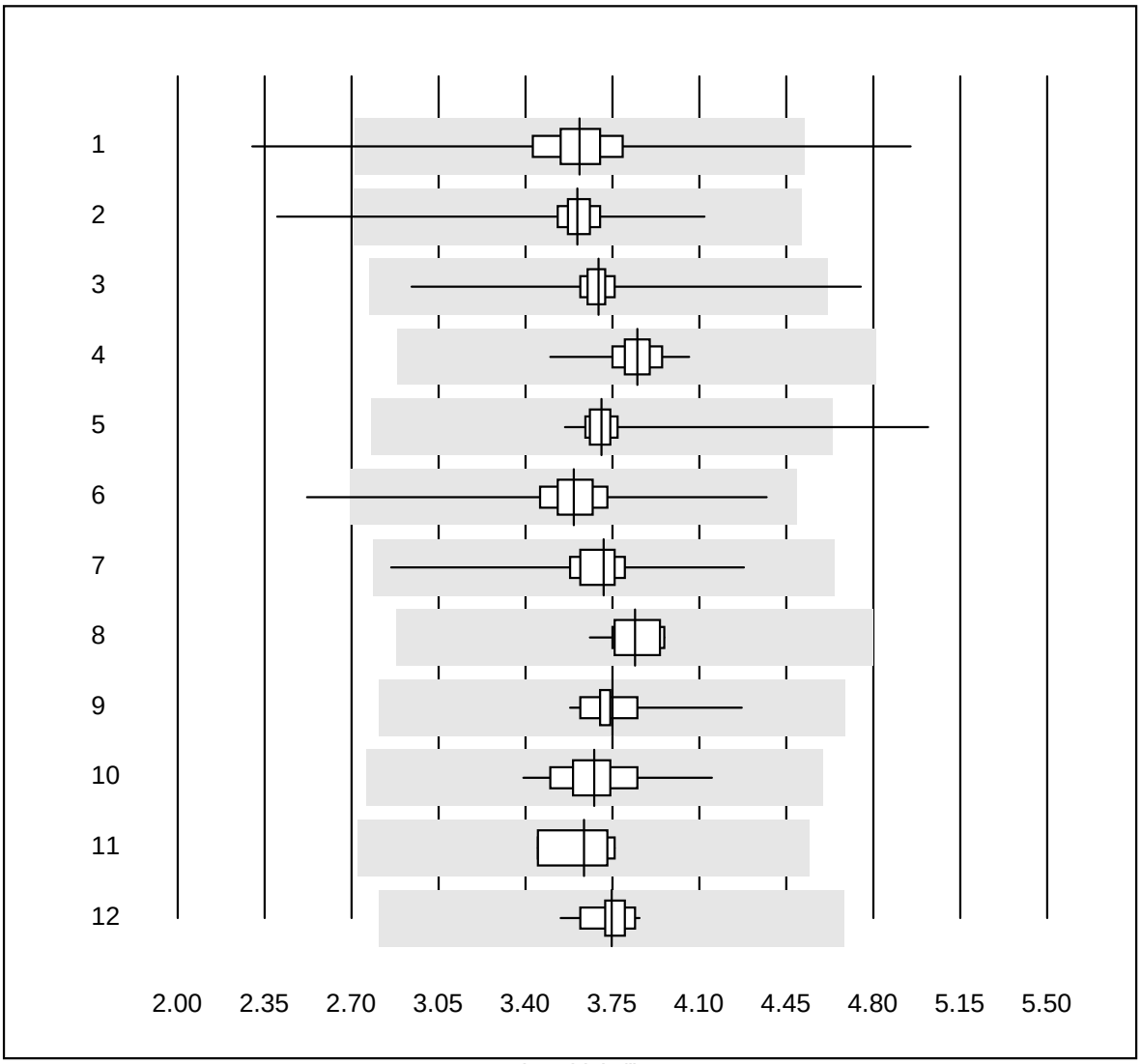


Deviatione QUALAB : 25 %

Eritrociti (T/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Automatico	35	100.0	0.0	0.0	3.69	4.8	e
2	Microscopio	9	88.9	11.1	0.0	3.75	13.5	e*
3	Sysmex XT/XE/XS	40	100.0	0.0	0.0	3.74	1.5	e
4	ABX Pentra	11	90.9	0.0	9.1	3.69	2.0	e
5	MS4	4	100.0	0.0	0.0	3.88	2.5	e
6	Beckman	4	100.0	0.0	0.0	3.84	1.3	a

Eritrociti

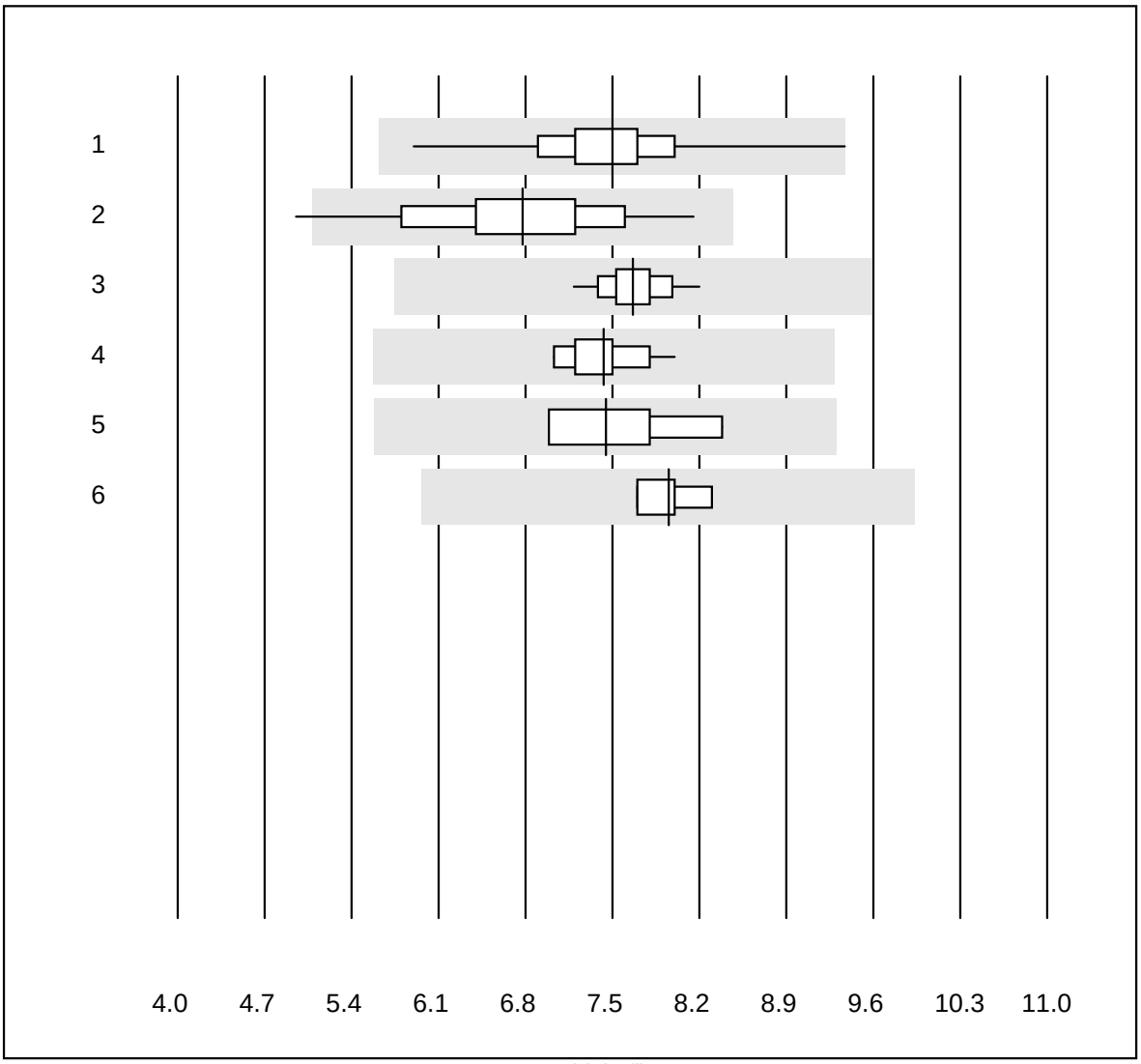


Deviatione QUALAB : 25 %

Eritrociti (T/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Abx Micros	833	97.7	1.2	1.1	3.62	5.6	e
2	Microsemi	263	99.6	0.4	0.0	3.61	3.6	e
3	Sysmex KX21	428	98.2	0.2	1.6	3.70	3.1	e
4	Sysmex PocH - 100i	213	100.0	0.0	0.0	3.85	2.1	e
5	Sysmex XP 300	190	97.4	0.5	2.1	3.71	3.1	e
6	Mythic	247	98.4	0.4	1.2	3.59	4.0	e
7	Swelab	70	100.0	0.0	0.0	3.72	4.2	e
8	Abacus Junior	13	100.0	0.0	0.0	3.84	2.6	e
9	Medonic	17	100.0	0.0	0.0	3.75	4.0	e
10	Samsung HC10	44	97.7	0.0	2.3	3.68	4.1	e
11	Norma Icon 3	4	100.0	0.0	0.0	3.64	4.1	e
12	Nihon Kohden Celltac	39	97.4	0.0	2.6	3.75	2.0	e

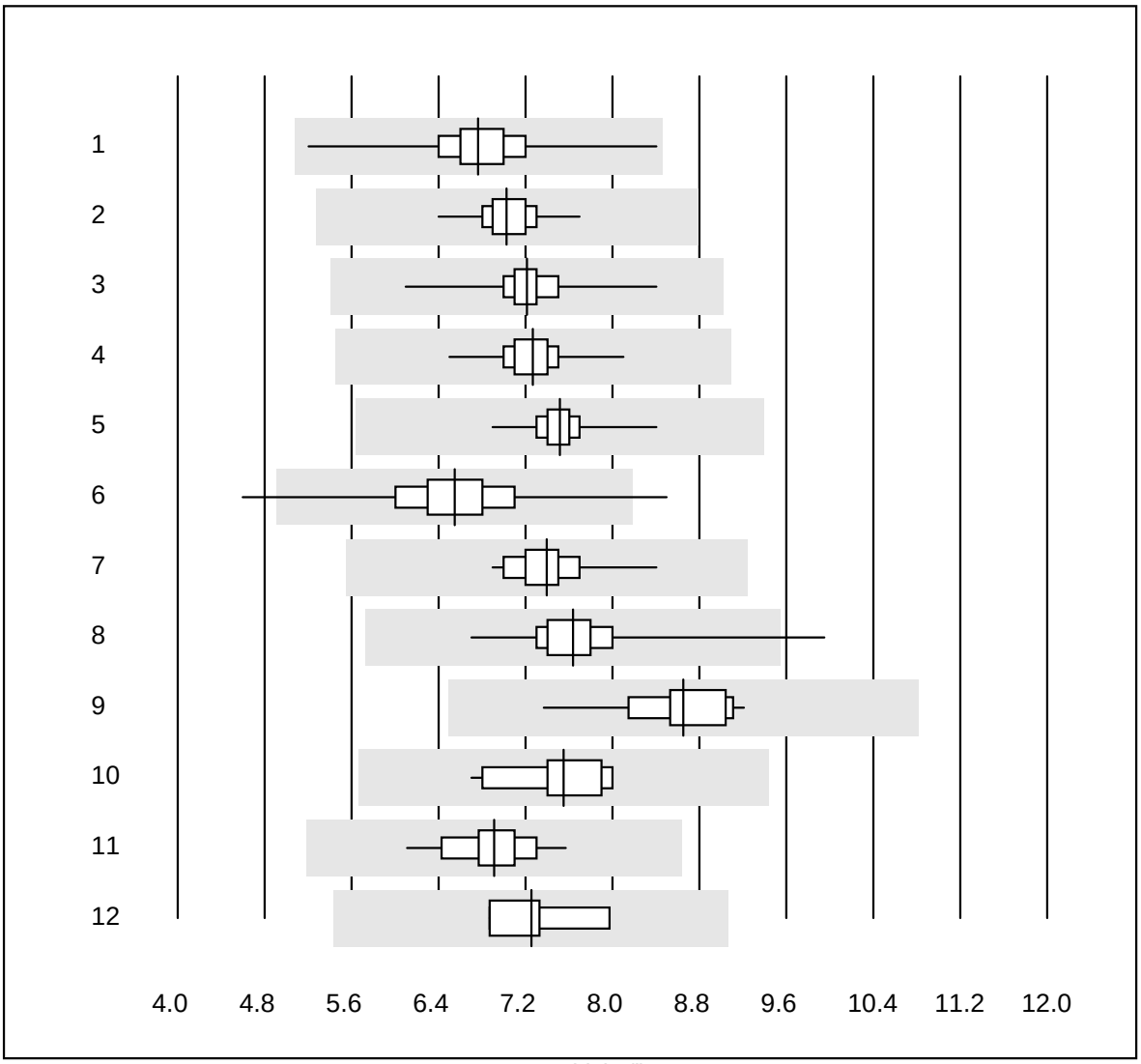
Leucociti



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Automatico	33	100.0	0.0	0.0	7.50	7.8	e
2	Microscopio	55	92.7	1.8	5.5	6.78	10.0	e
3	Sysmex XT/XE/XS	39	100.0	0.0	0.0	7.66	2.8	e
4	ABX Pentra	11	90.9	0.0	9.1	7.43	4.0	e
5	MS4	4	100.0	0.0	0.0	7.45	8.6	e*
6	Beckman	4	100.0	0.0	0.0	7.95	3.3	a

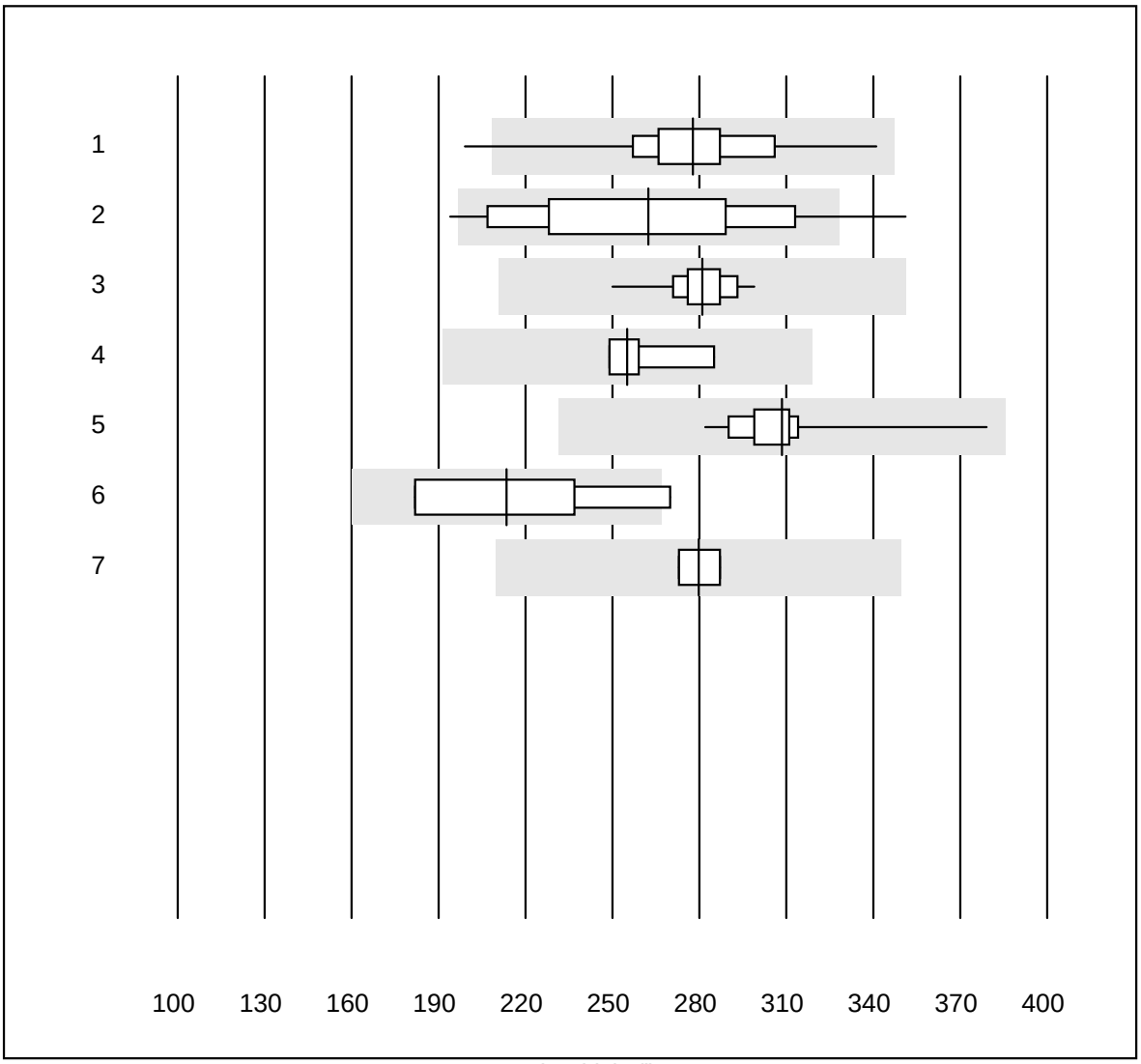
Leucociti



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Abx Micros	833	99.9	0.0	0.1	6.76	5.3	e
2	Microsemi	264	100.0	0.0	0.0	7.03	3.0	e
3	Sysmex KX21	428	99.5	0.0	0.5	7.22	3.2	e
4	Sysmex PocH - 100i	213	100.0	0.0	0.0	7.27	3.0	e
5	Sysmex XP 300	192	100.0	0.0	0.0	7.51	2.7	e
6	Mythic	246	98.0	1.6	0.4	6.55	7.2	e
7	Nihon Kohden Celltac	39	100.0	0.0	0.0	7.39	4.2	e
8	Swelab	70	98.6	1.4	0.0	7.63	5.8	e
9	Abacus Junior	13	100.0	0.0	0.0	8.66	5.8	e
10	Medonic	17	100.0	0.0	0.0	7.55	5.4	e
11	Samsung HC10	44	100.0	0.0	0.0	6.91	4.7	e
12	Norma Icon 3	4	100.0	0.0	0.0	7.26	6.3	e*

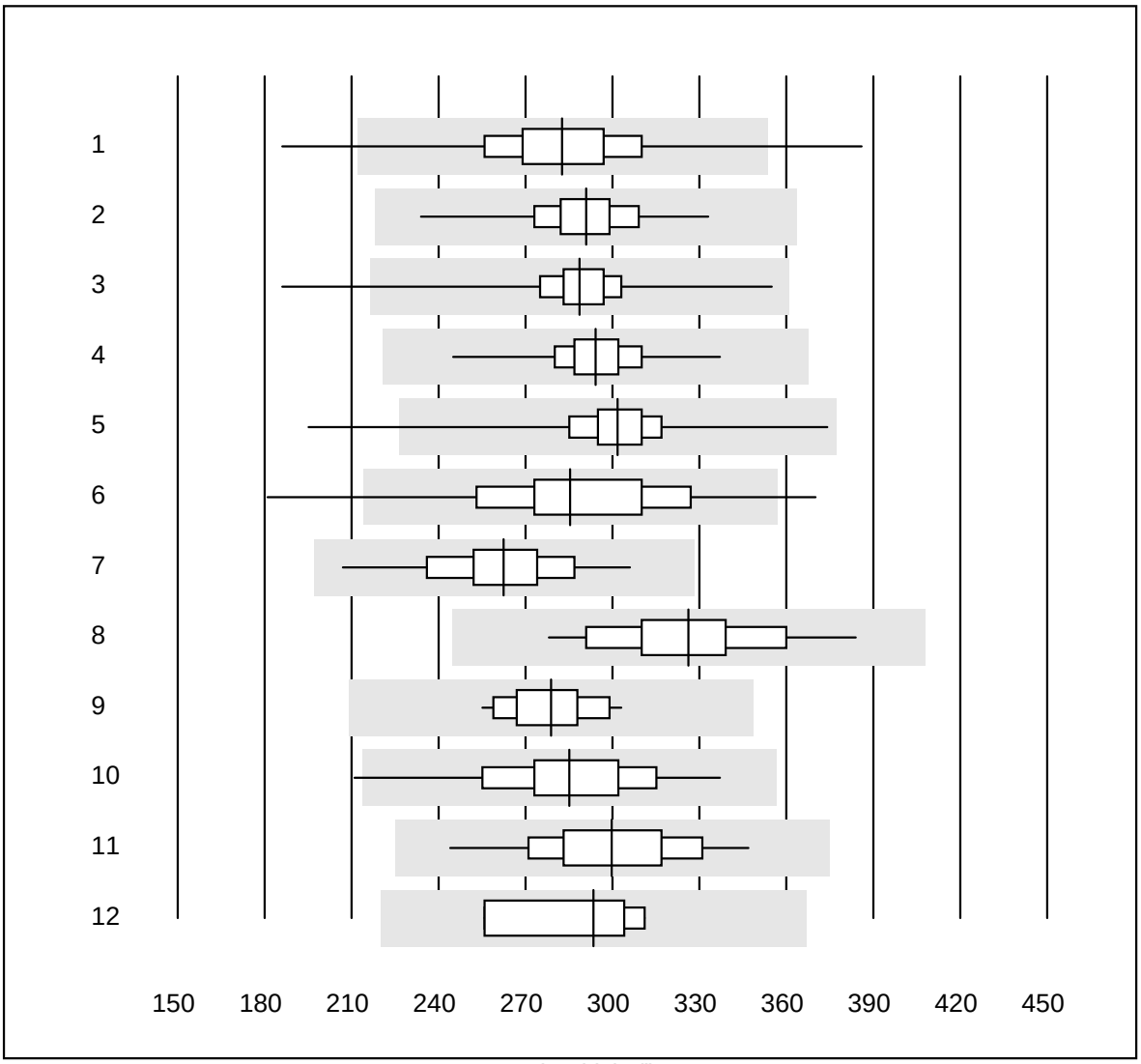
Trombociti



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Automatico	31	96.8	3.2	0.0	277.8	9.3	e
2	Microscopio	34	94.1	5.9	0.0	262.4	15.3	e
3	Sysmex XT/XE/XS	39	100.0	0.0	0.0	281.1	3.4	e
4	Advia 120	4	100.0	0.0	0.0	255.0	6.3	e*
5	ABX Pentra	11	100.0	0.0	0.0	308.5	8.1	e
6	MS4	4	75.0	25.0	0.0	213.5	18.8	e*
7	Beckman	4	75.0	0.0	25.0	279.7	2.5	a

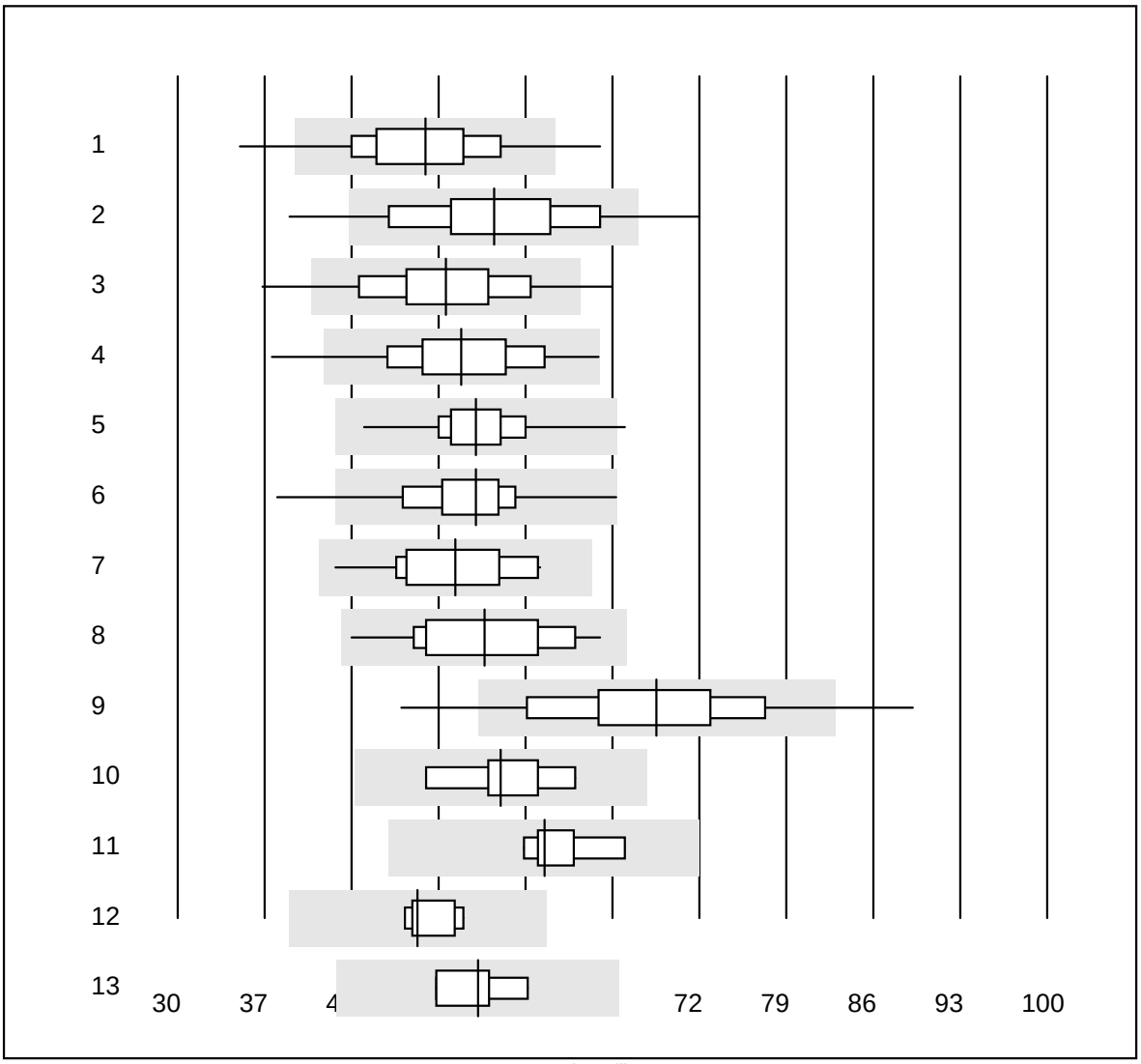
Trombociti



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Abx Micros	833	97.4	1.8	0.8	282.7	8.5	e
2	Microsemi	265	100.0	0.0	0.0	290.9	5.0	e
3	Sysmex KX21	428	99.3	0.7	0.0	288.5	5.1	e
4	Sysmex PocH - 100i	212	100.0	0.0	0.0	294.3	4.4	e
5	Sysmex XP 300	191	99.5	0.5	0.0	301.7	5.9	e
6	Mythic	247	97.6	2.4	0.0	285.5	10.2	e
7	Swelab	70	97.1	0.0	2.9	262.5	7.5	e
8	Abacus Junior	13	100.0	0.0	0.0	326.3	8.5	e
9	Medonic	17	100.0	0.0	0.0	278.7	5.5	e
10	Nihon Kohden Celltac	39	94.8	2.6	2.6	285.2	8.5	e
11	Samsung HC10	44	100.0	0.0	0.0	299.9	7.8	e
12	Norma Icon 3	4	100.0	0.0	0.0	293.5	8.6	e*

CRP

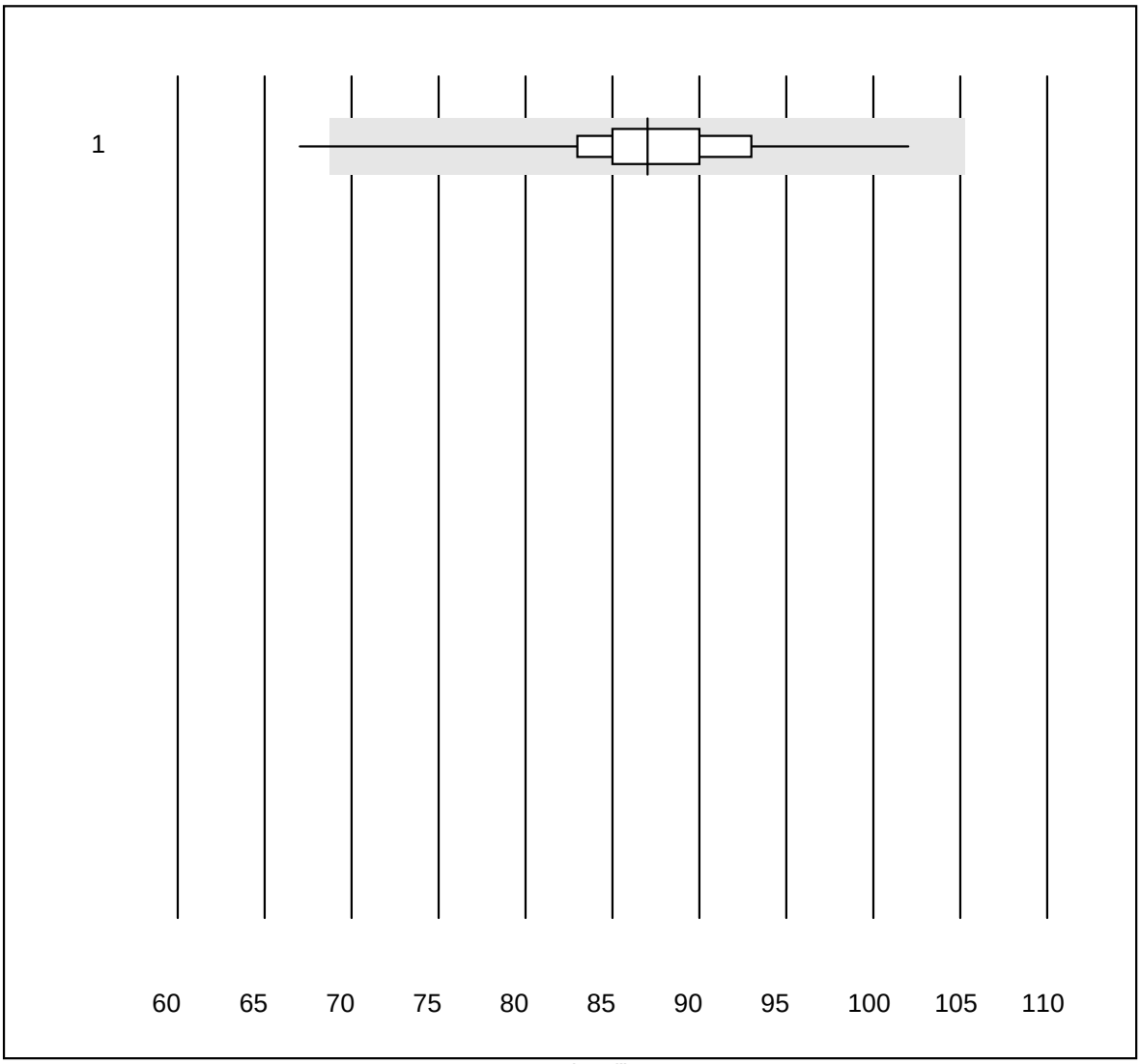


Deviatione QUALAB : 21 %

CRP (mg/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Afinion	1174	97.8	2.0	0.2	49.9	9.3	e
2	NycoCard SingleTest-	434	79.5	7.8	12.7	55.5	12.0	e
3	Abx Micros	120	90.8	5.0	4.2	51.6	10.4	e
4	ABX Micros CRP200	321	95.0	3.4	1.6	52.8	9.8	e
5	Quick Read go	100	98.0	1.0	1.0	54.0	6.0	e
6	Turbidimetrie	39	97.4	2.6	0.0	54.0	8.9	e
7	Cobas	11	100.0	0.0	0.0	52.4	9.6	e*
8	Fuji Dri-Chem	24	100.0	0.0	0.0	54.7	10.3	e
9	Eurolyser	131	74.8	9.9	15.3	68.5	12.0	e
10	AQT 90 FLEX	6	100.0	0.0	0.0	56.0	7.3	e*
11	Spotchem D-Concept	7	100.0	0.0	0.0	59.5	4.5	e
12	Spotchem SI-3510	5	100.0	0.0	0.0	49.3	4.1	e
13	altro	4	100.0	0.0	0.0	54.2	5.7	e*

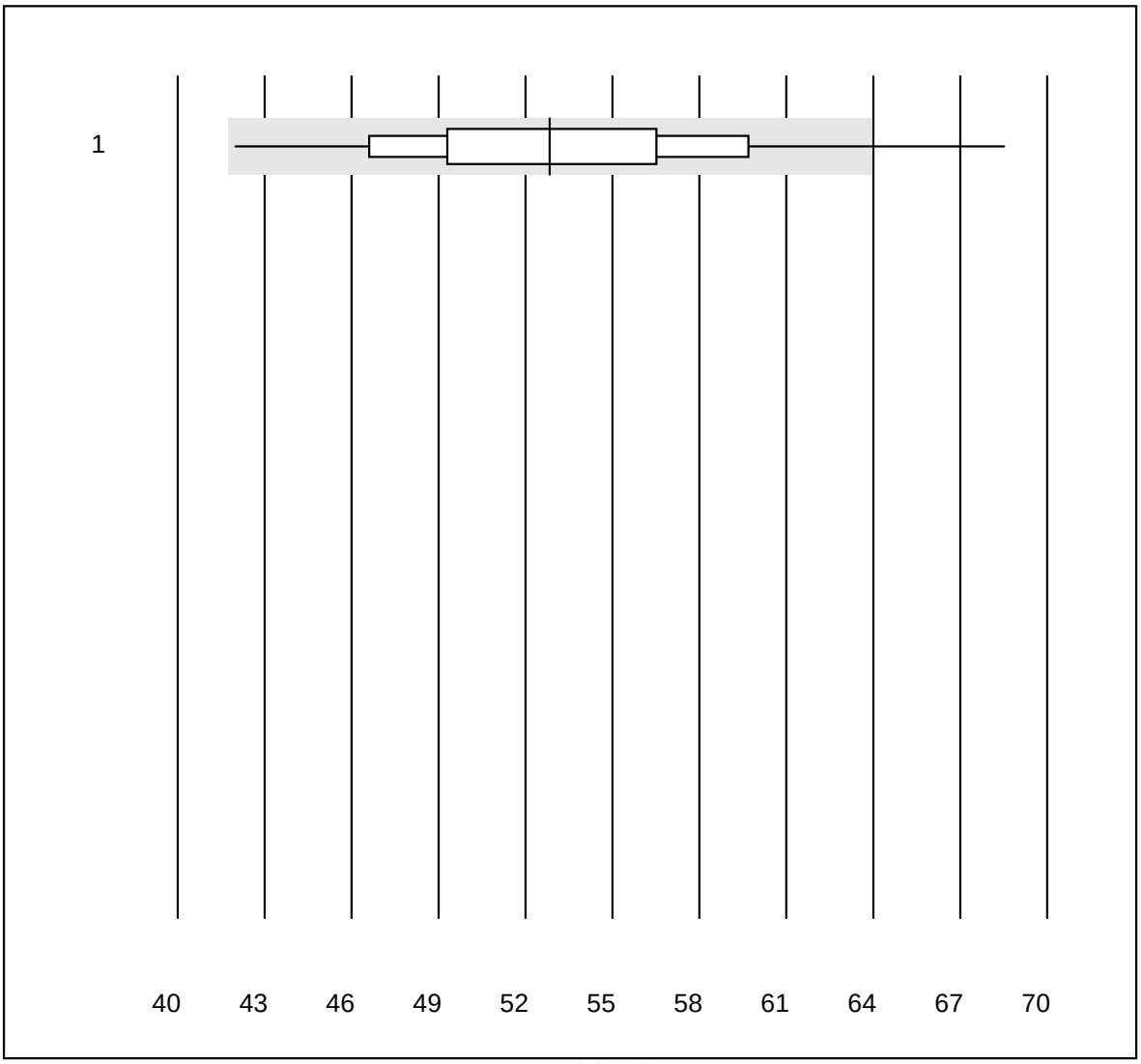
CRP



Deviatione QUALAB : 21 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	QuickRead (sangue)	175	96.5	0.6	2.9	87.0	5.4	e

CRP emi

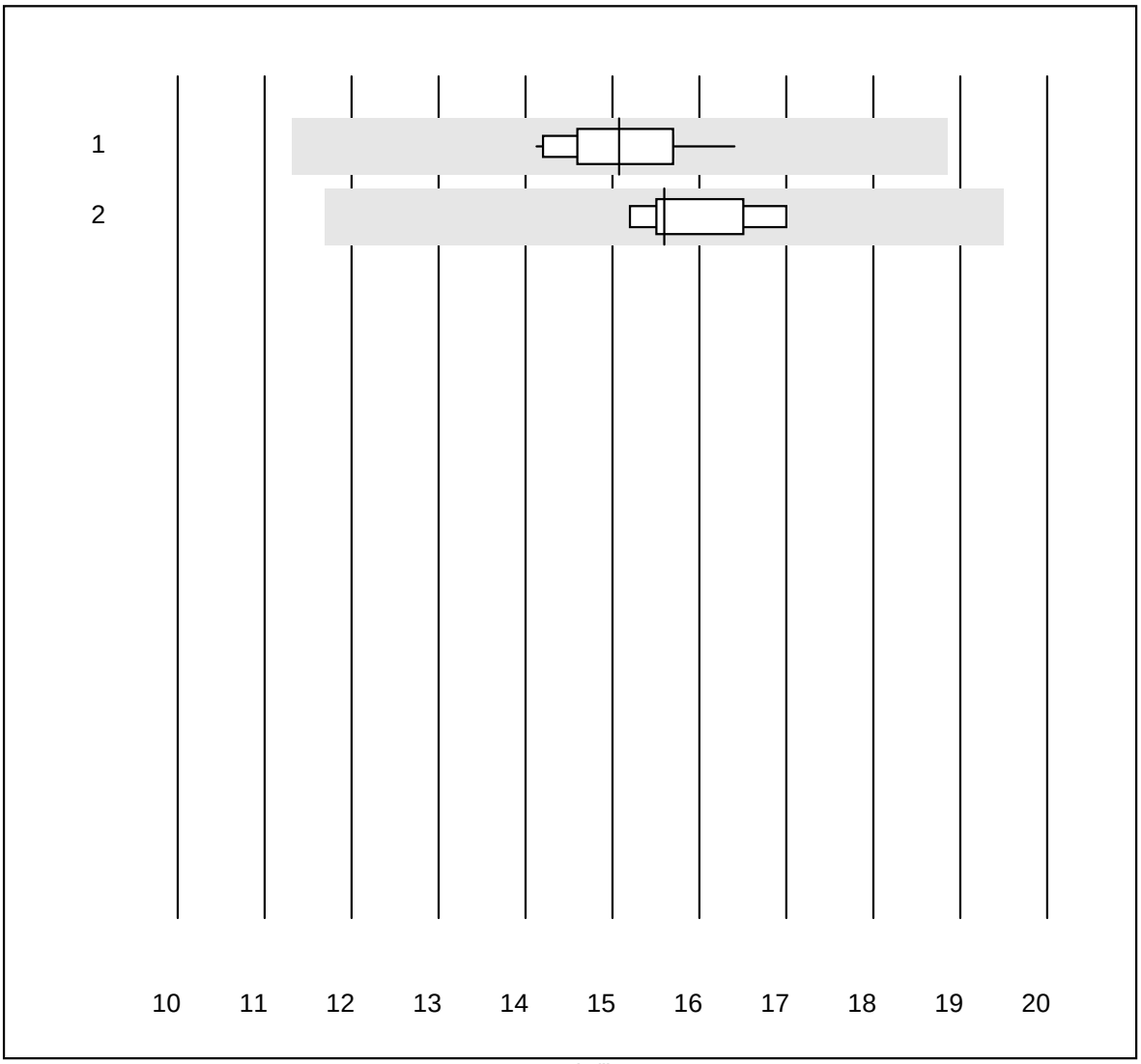


Deviatione QUALAB : 21 %

CRP emi (mg/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Microsemi	263	96.2	1.5	2.3	52.8	9.4	e

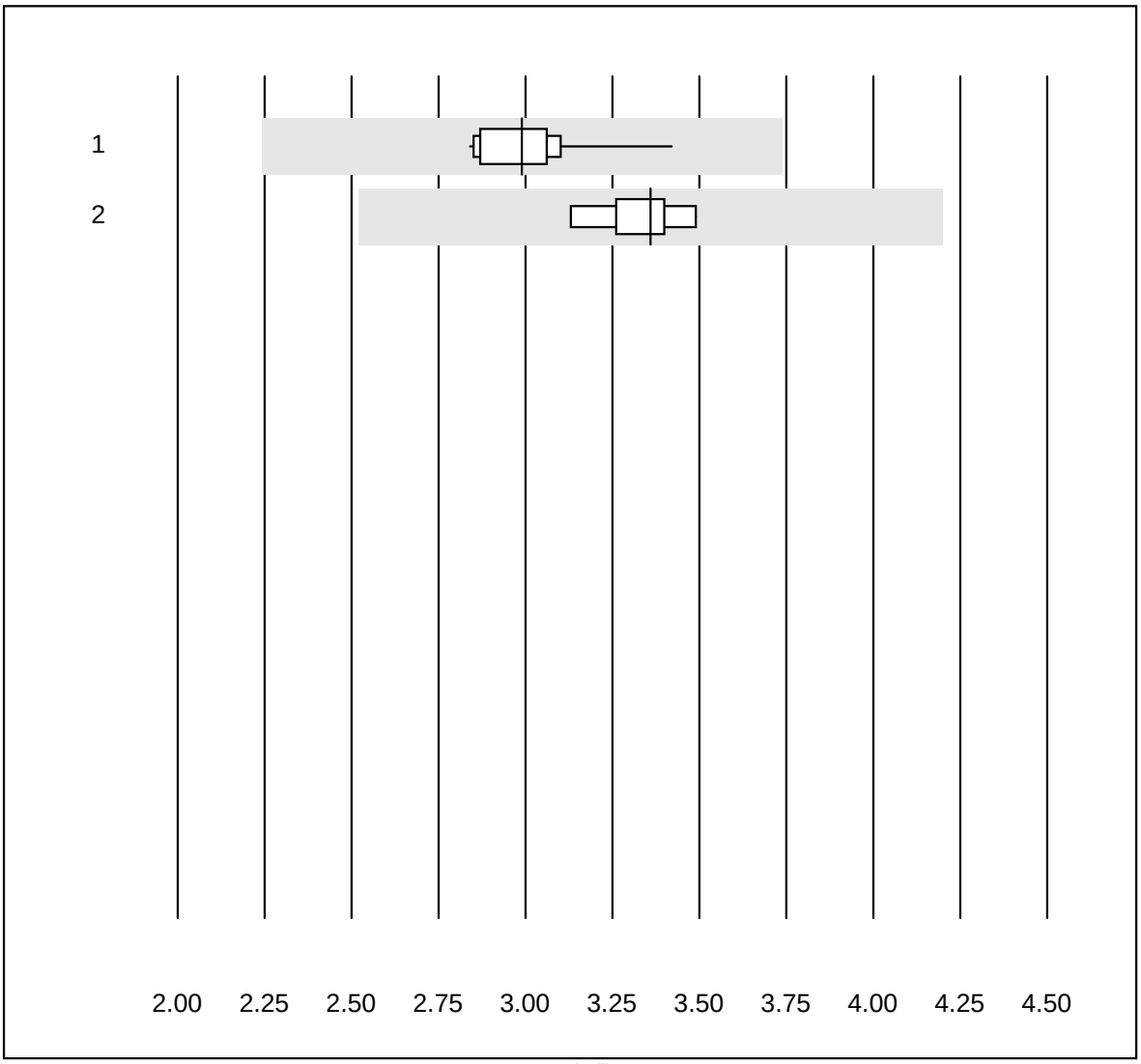
IgG



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Turbidimetrie	11	100.0	0.0	0.0	15.1	4.5	e
2	Nephelometrie	7	100.0	0.0	0.0	15.6	4.1	e

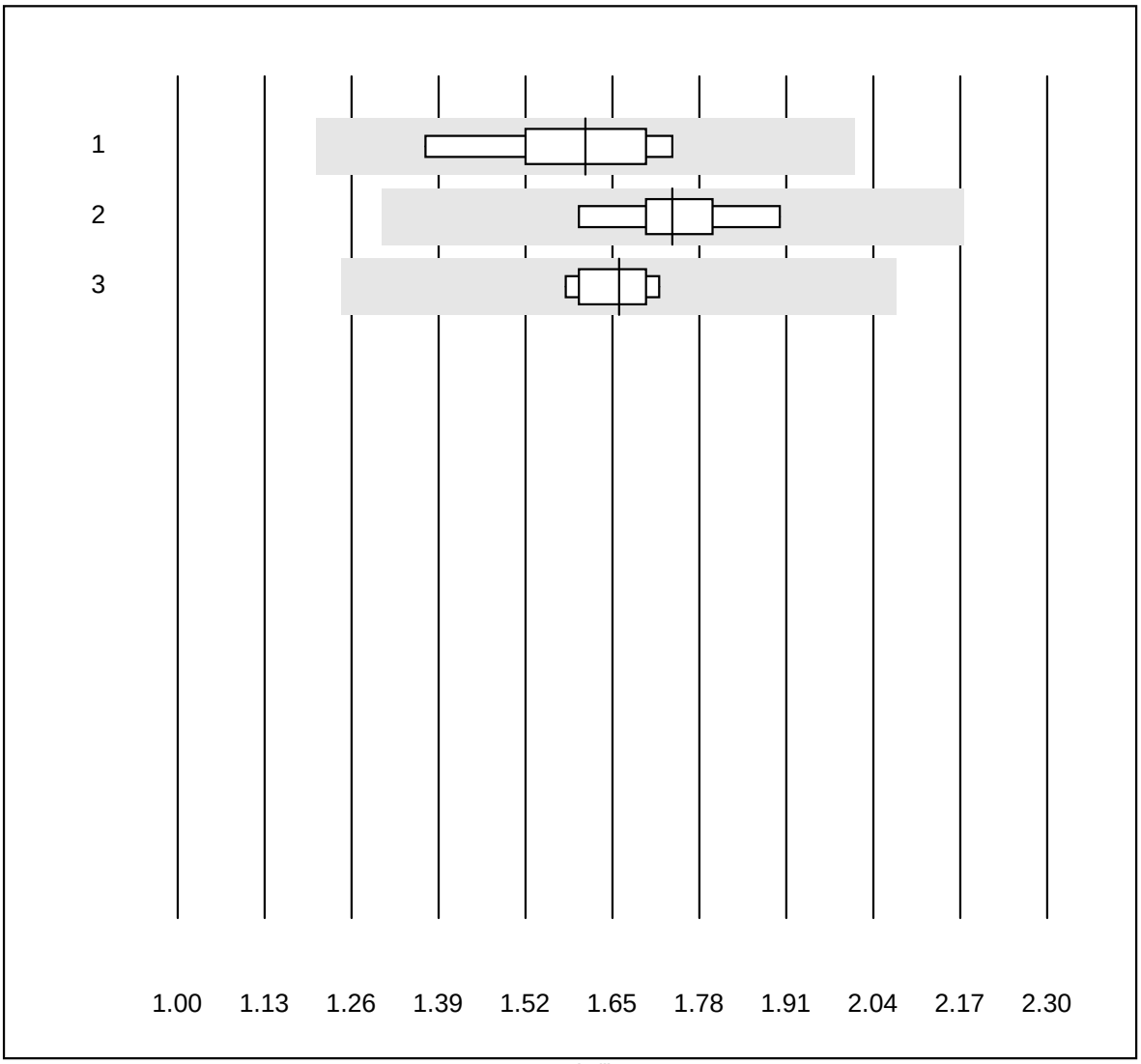
IgA



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Turbidimetrie	11	100.0	0.0	0.0	3.0	5.5	e
2	Nephelometrie	7	100.0	0.0	0.0	3.4	3.5	e

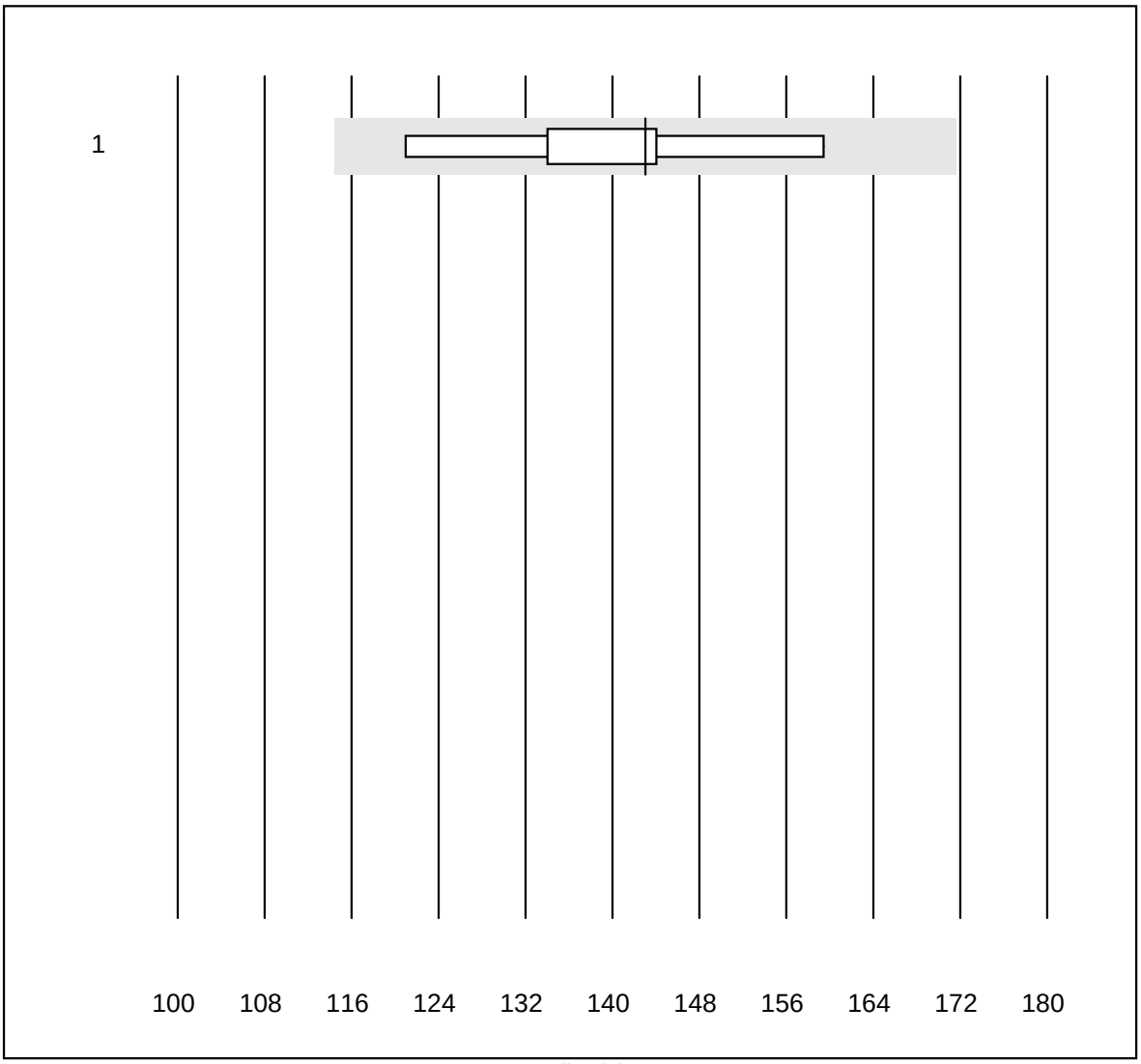
IgM



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Turbidimetrie	5	100.0	0.0	0.0	1.6	9.4	e*
2	Nephelometrie	7	100.0	0.0	0.0	1.7	5.4	e
3	Cobas Integra 800/40	6	100.0	0.0	0.0	1.7	3.4	e

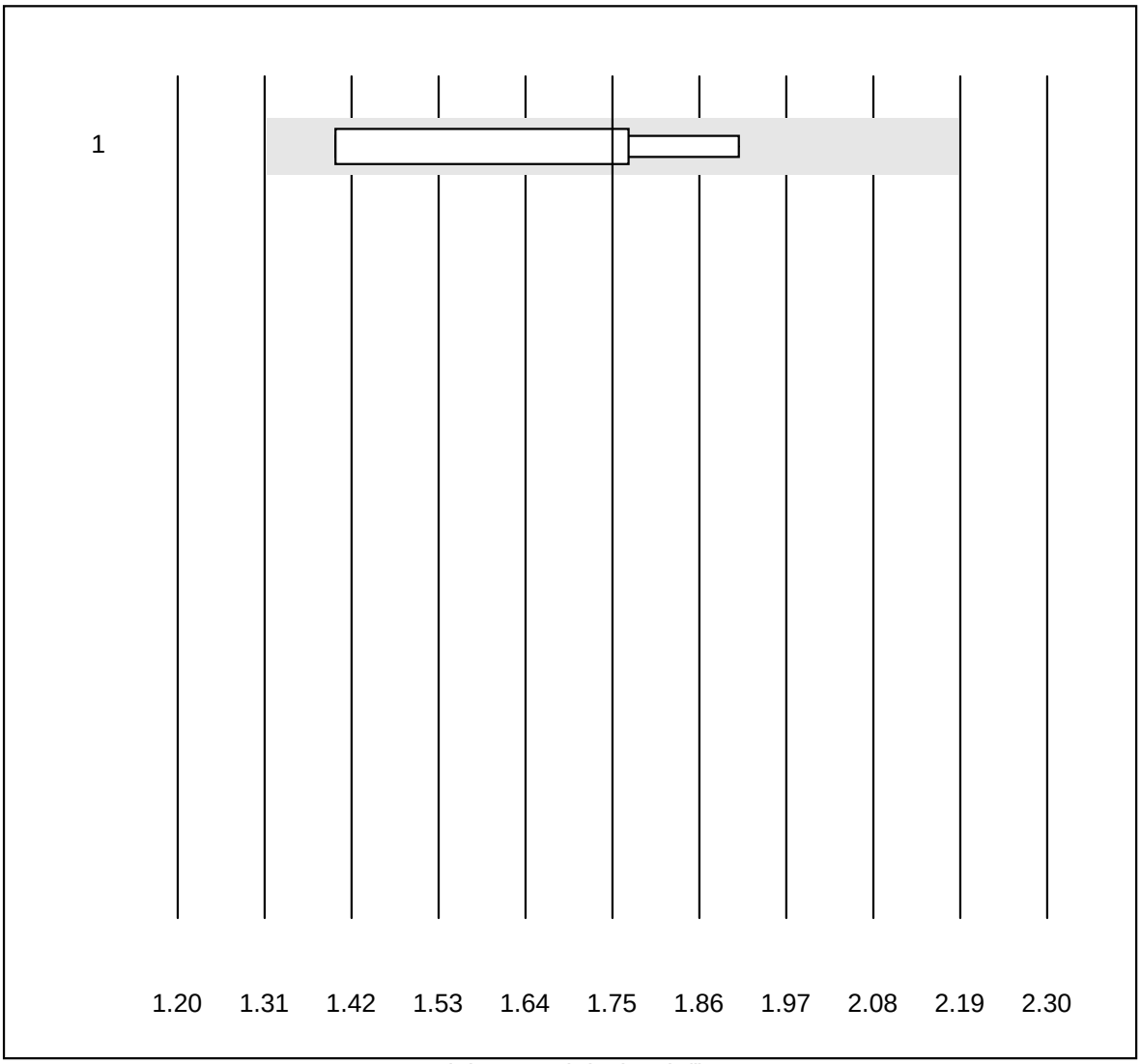
IgE



Deviatione QUALAB : 20 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	tutti	8	100.0	0.0	0.0	143	8.3	e*

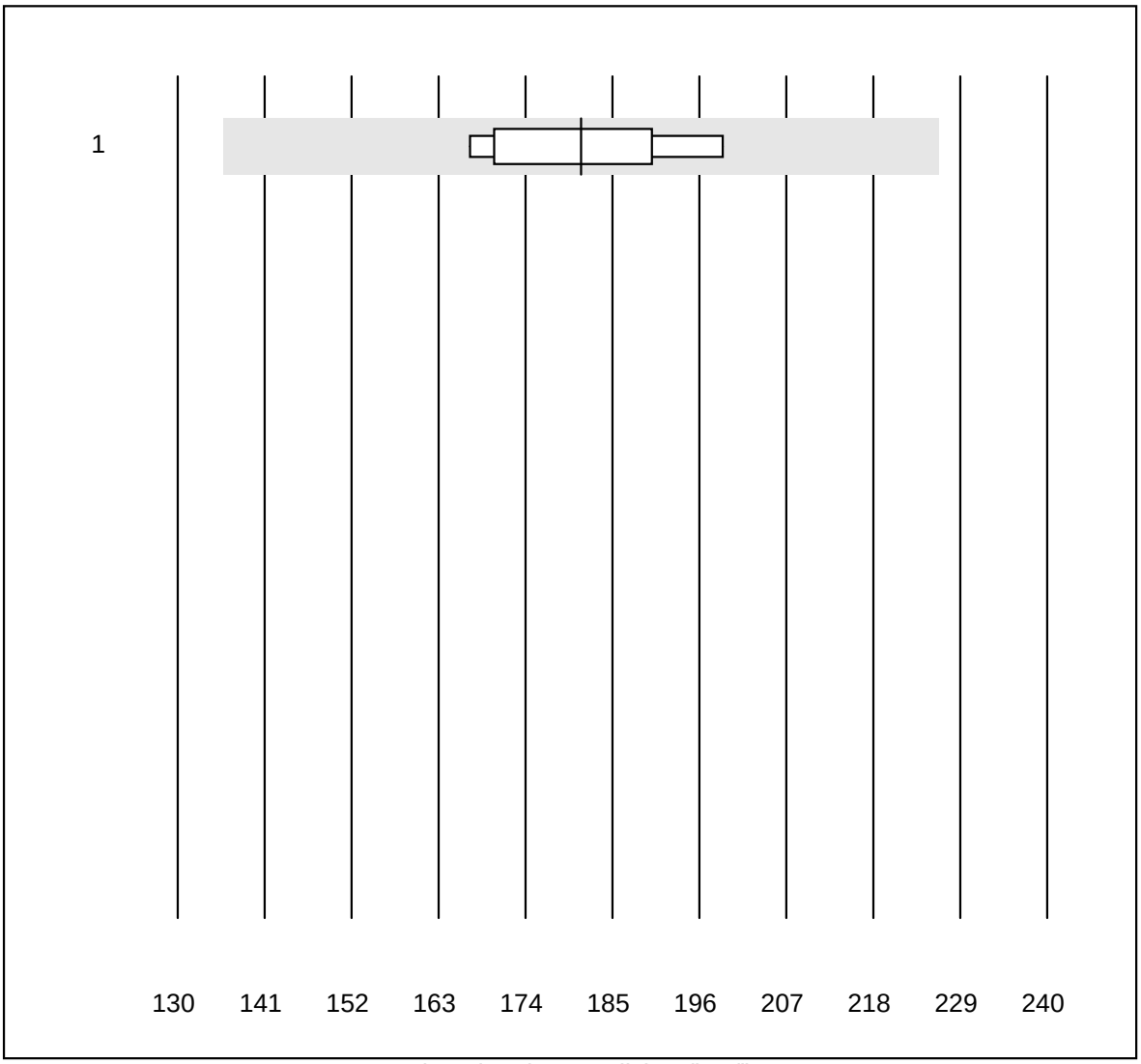
Alpha-1-Antitripsina



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Nephelometrie	4	100.0	0.0	0.0	1.75	12.7	e*

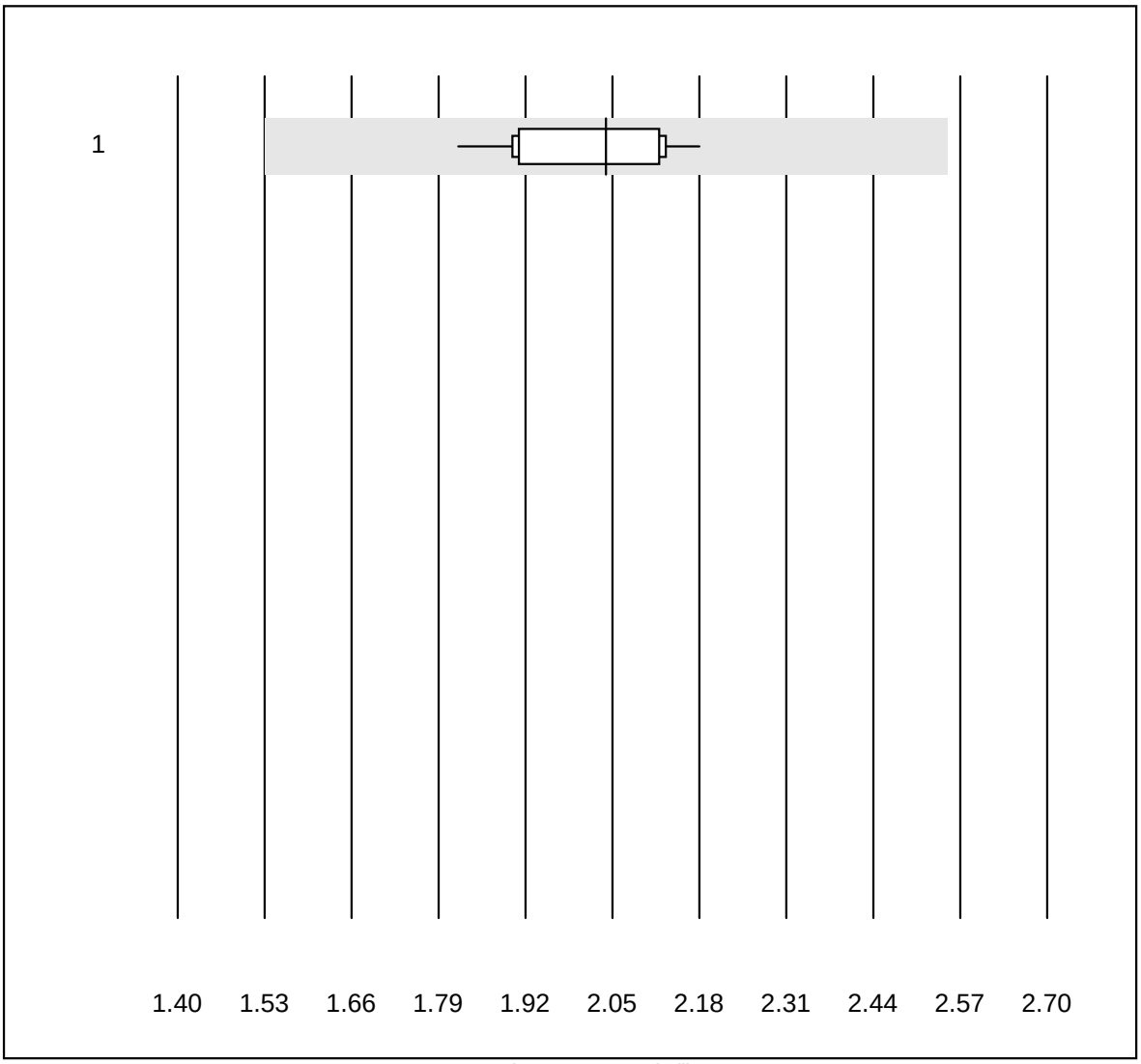
Anticorpi anti-streptolisina



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	tutti	10	90.0	0.0	10.0	181	6.4	a

Complemento C3

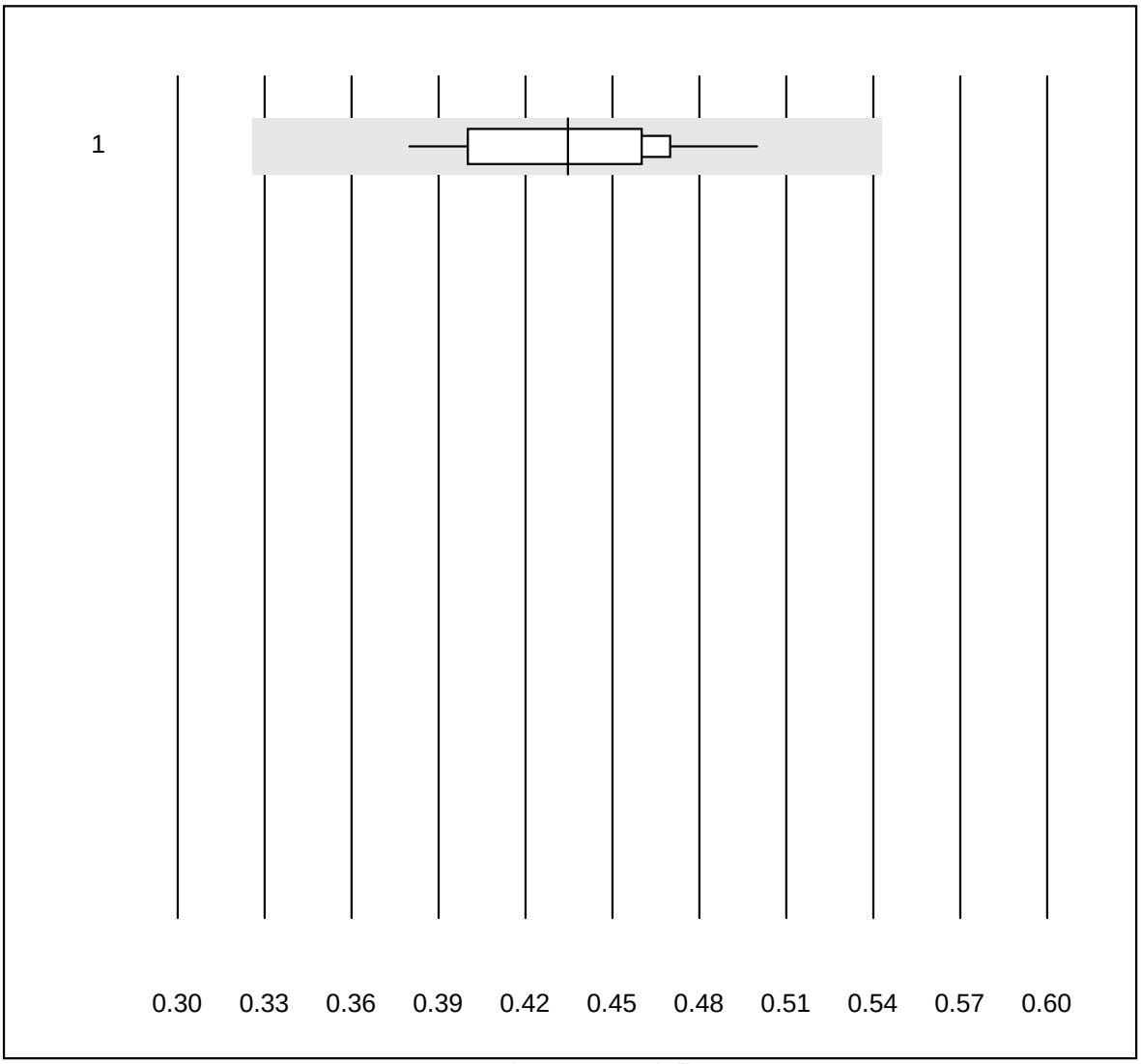


Deviatione QUALAB : 25 %

Complemento C3 (g/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	tutti	12	100.0	0.0	0.0	2.04	5.4	e

Complemento C4

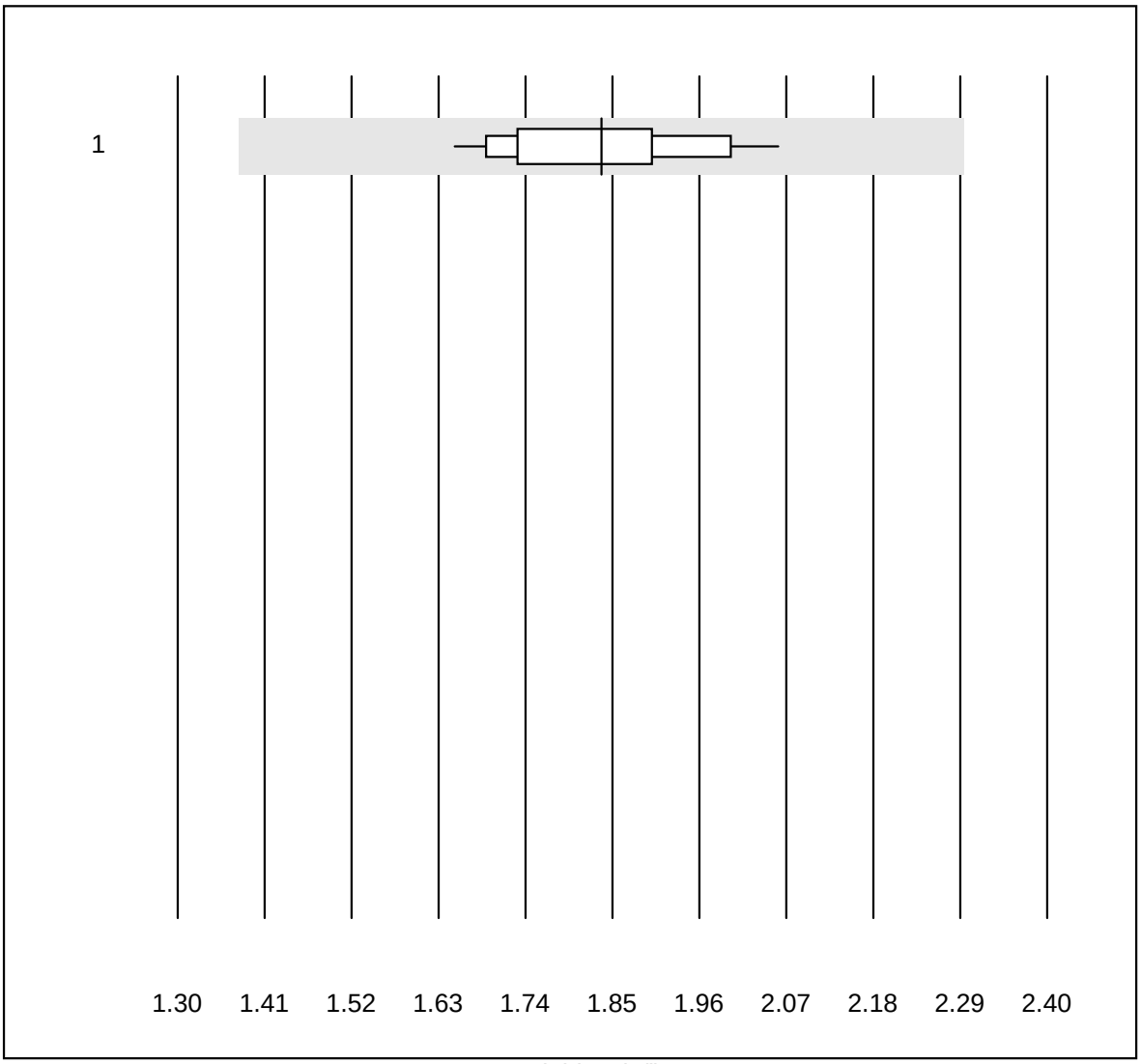


Deviatione QUALAB : 25 %

Complemento C4 (g/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	tutti	11	100.0	0.0	0.0	0.43	7.9	e

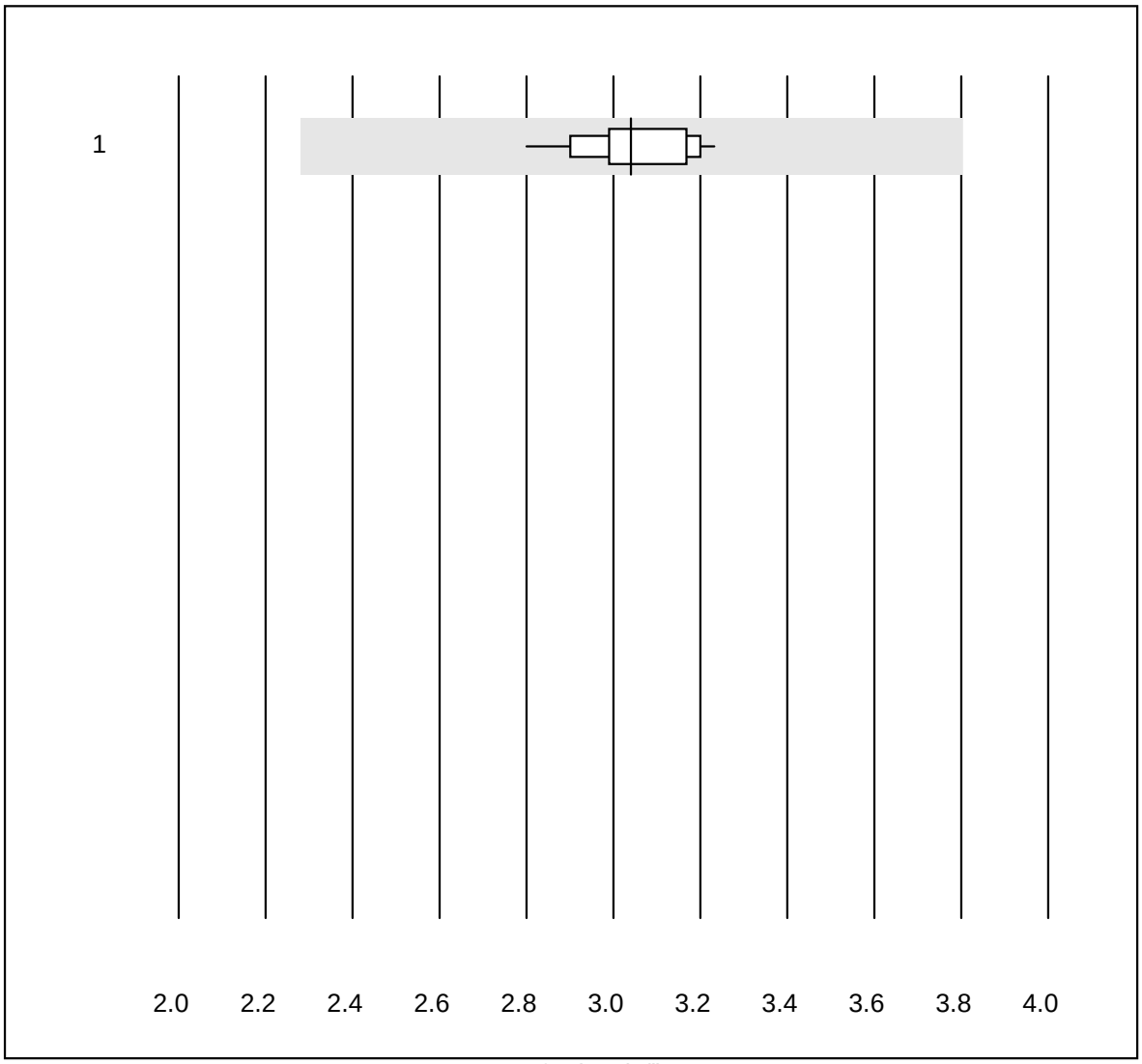
Aptoglobina



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	tutti	12	100.0	0.0	0.0	1.84	6.7	e

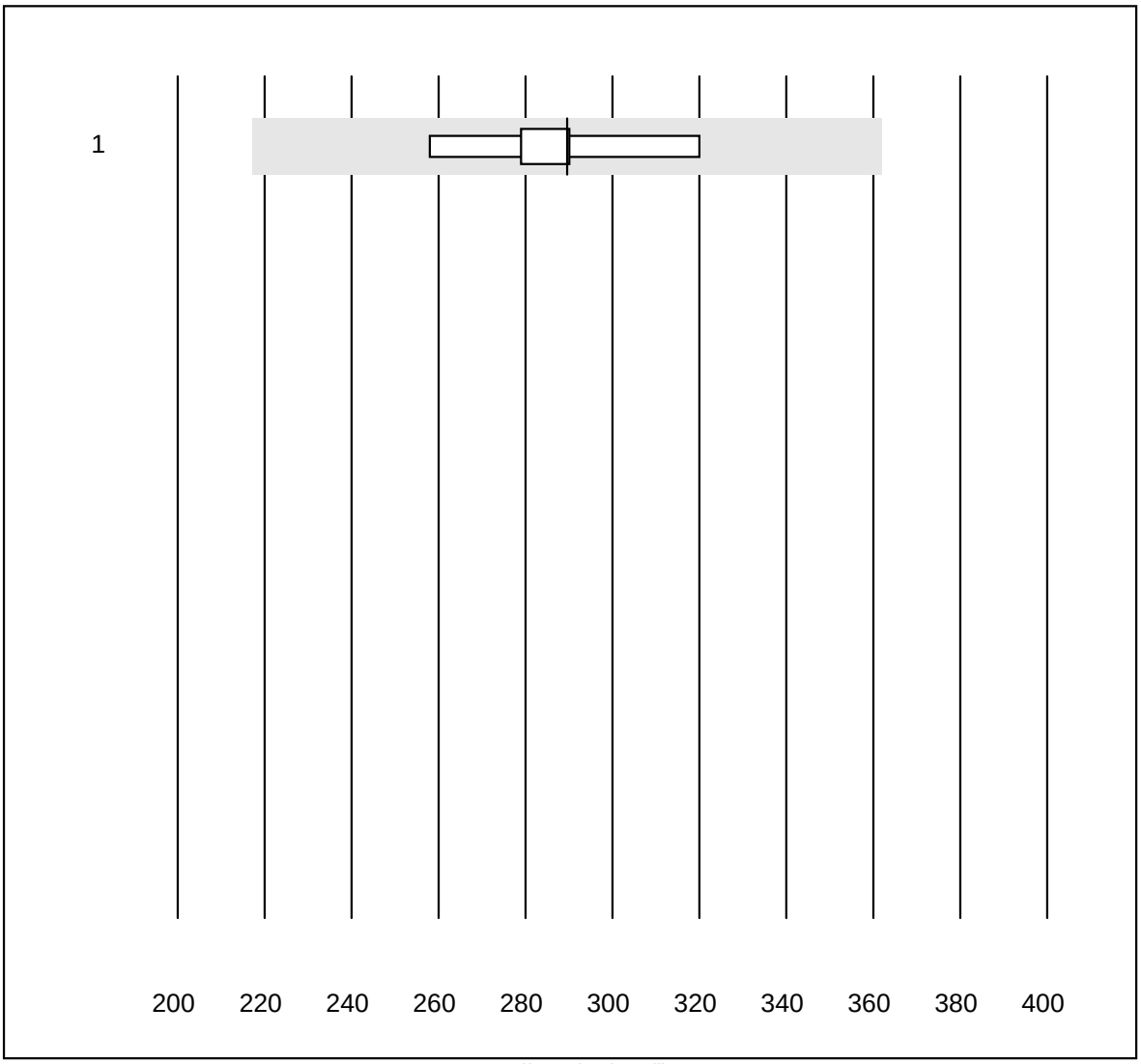
Transferrina



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	tutti	15	100.0	0.0	0.0	3.04	3.9	e

Präalbumin

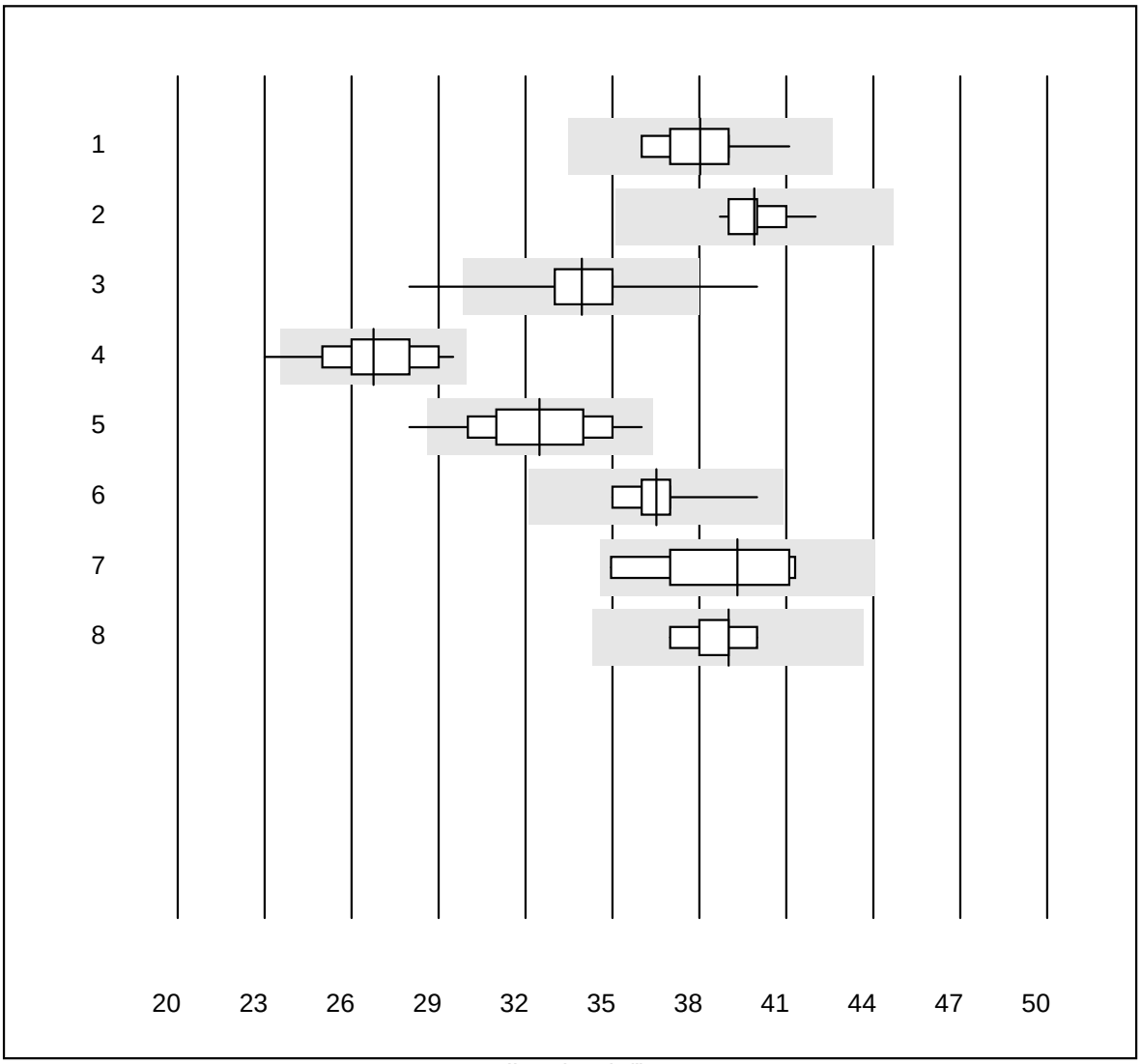


Deviatione QUALAB : 25 %

Präalbumin (mg/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	tutti	7	100.0	0.0	0.0	289.6	6.4	e

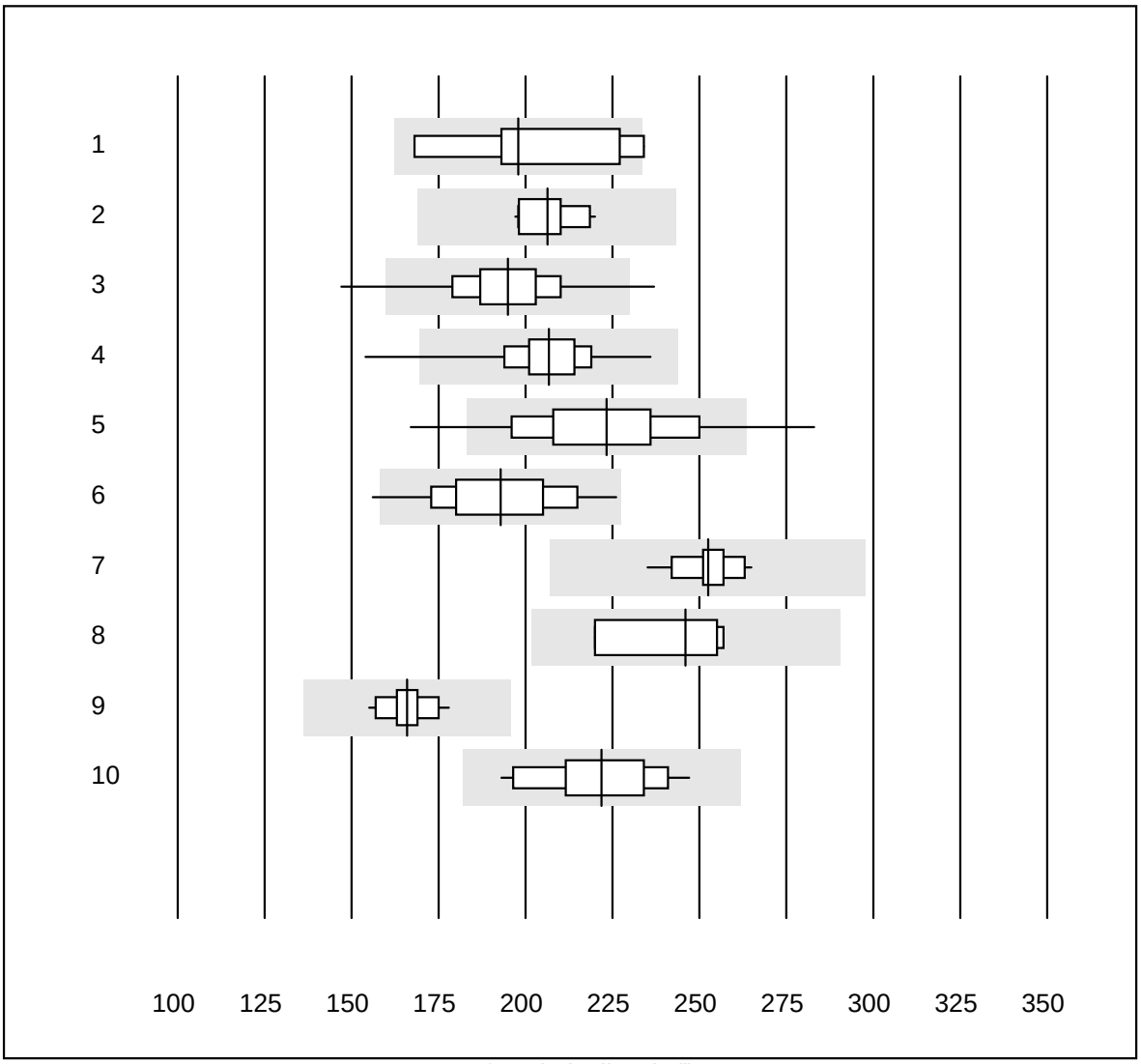
Albumina



Deviatione QUALAB : 12 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	20	100.0	0.0	0.0	38	3.5	e
2	Cobas	12	100.0	0.0	0.0	40	2.2	e
3	Fuji Dri-Chem	172	96.6	1.7	1.7	34	4.1	e
4	Spotchem/Ready	42	97.6	2.4	0.0	27	5.2	e
5	Spotchem D-Concept	70	97.2	1.4	1.4	32	5.4	e
6	Piccolo	30	100.0	0.0	0.0	37	2.7	e
7	Abx Mira	8	87.5	0.0	12.5	39	5.8	e*
8	Hitachi S40/M40	9	100.0	0.0	0.0	39	2.3	e

Fosfatasi alcalina

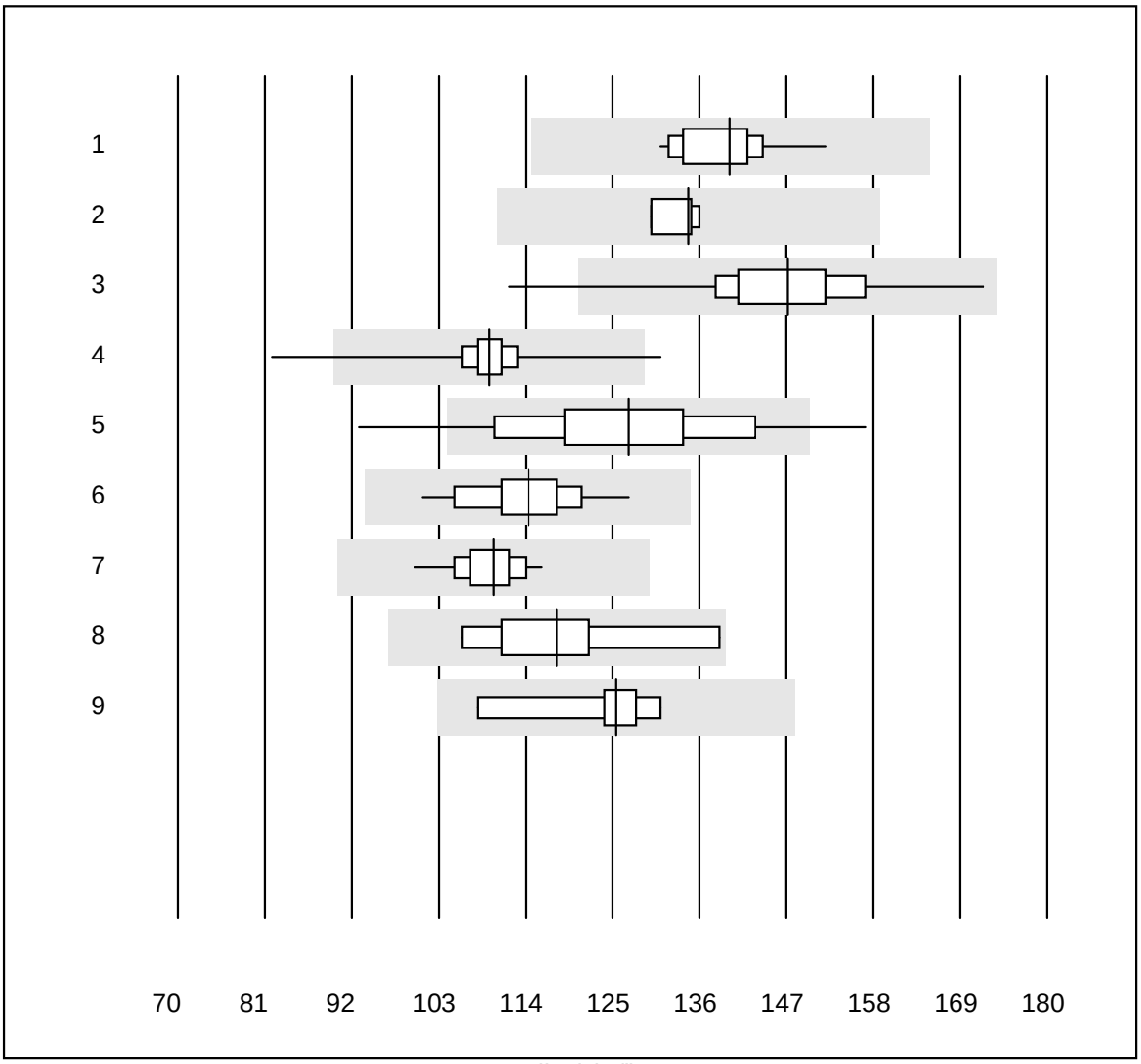


Deviatione QUALAB : 18 %

Fosfatasi alcalina (U/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	IFCC/SGKC/SFBC 37'C	11	72.7	9.1	18.2	198	11.1	e*
2	Cobas	17	100.0	0.0	0.0	206	3.8	e
3	Reflotron	676	96.5	1.6	1.9	195	6.7	e
4	Fuji Dri-Chem	666	98.6	0.6	0.8	207	4.9	e
5	Spotchem/Ready	116	90.5	7.8	1.7	223	9.8	e
6	Spotchem D-Concept	131	99.2	0.8	0.0	193	8.2	e
7	Hitachi S40/M40	14	100.0	0.0	0.0	253	3.2	e
8	Olympus	7	100.0	0.0	0.0	246	7.1	e*
9	Piccolo	29	100.0	0.0	0.0	166	3.6	e
10	Abx Mira	20	100.0	0.0	0.0	222	6.8	e

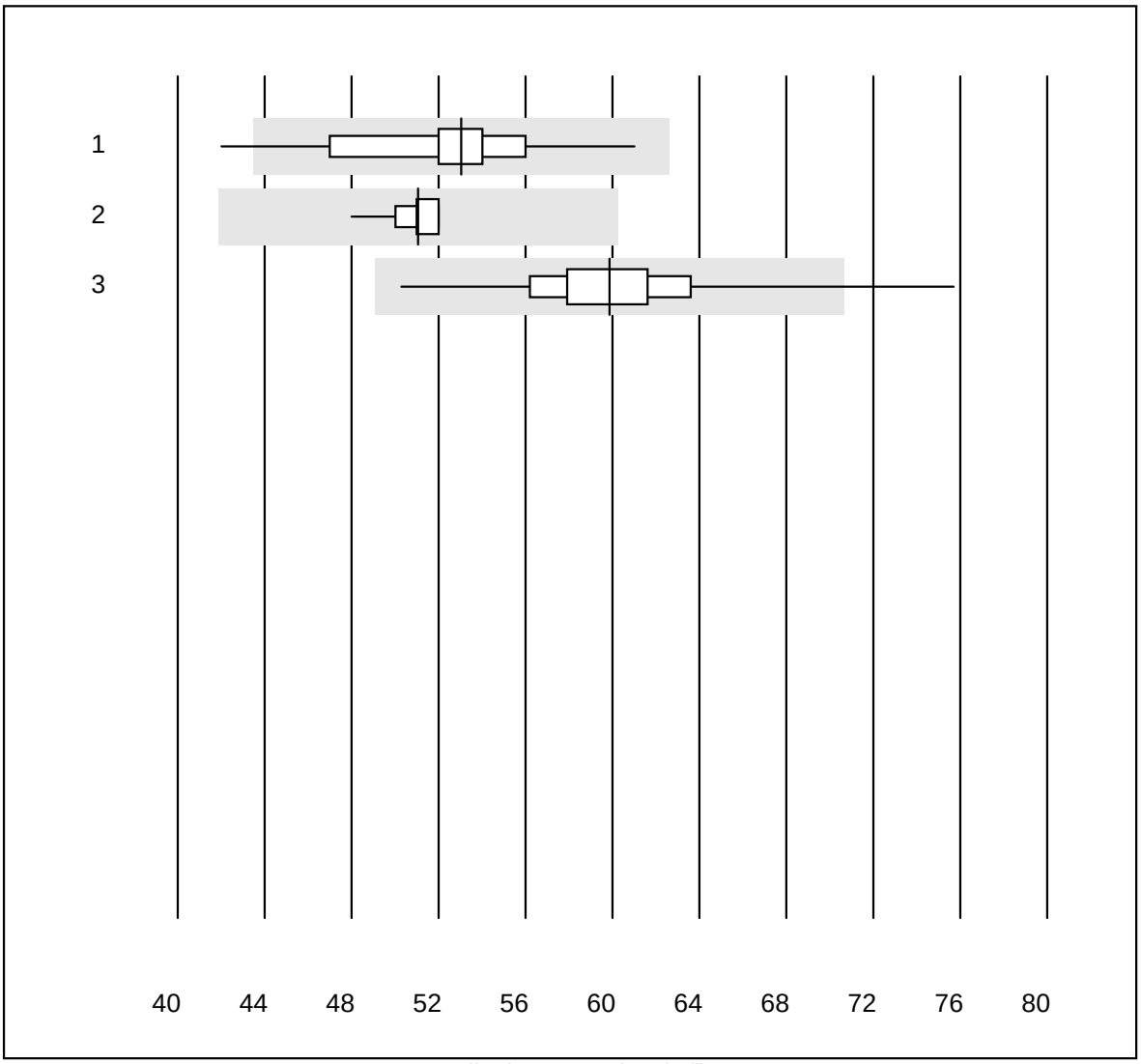
Amilasi



Deviatione QUALAB : 18 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	IFCC EPS liquid 37°C	12	100.0	0.0	0.0	140	4.2	e
2	Cobas	5	100.0	0.0	0.0	135	2.2	e
3	Reflotron	178	97.2	0.6	2.2	147	5.8	e
4	Fuji Dri-Chem	493	99.0	0.6	0.4	109	3.5	e
5	Spotchem/Ready	77	90.9	6.5	2.6	127	9.6	e
6	Spotchem D-Concept	98	100.0	0.0	0.0	114	5.2	e
7	Piccolo	27	100.0	0.0	0.0	110	3.2	e
8	Abx Mira	9	88.9	0.0	11.1	118	8.4	e*
9	Hitachi S40/M40	7	100.0	0.0	0.0	126	6.0	e*

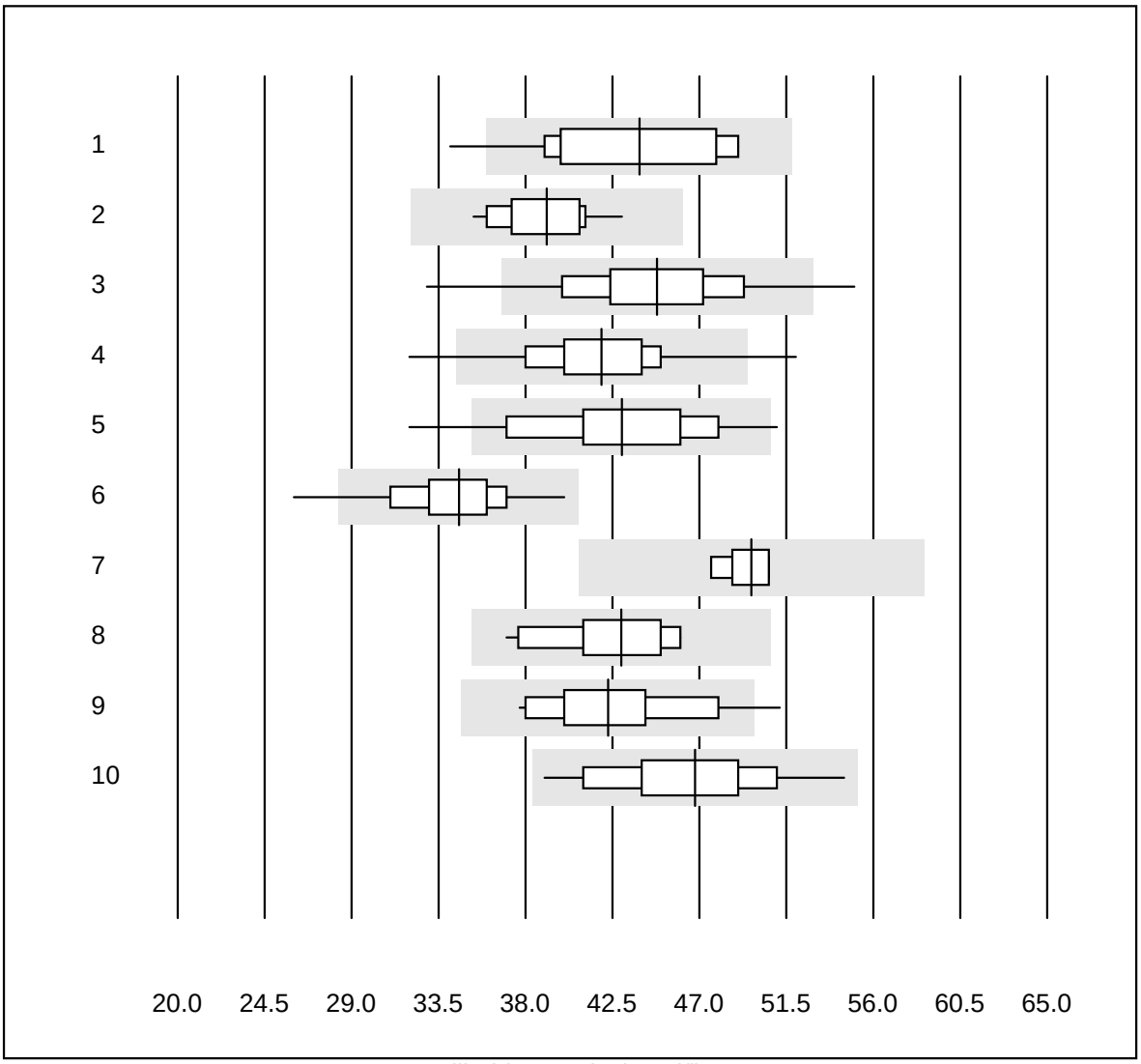
Amilasi pancreatica



Deviatione QUALAB : 18 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	IFCC EPS liquid 37°C	19	89.4	5.3	5.3	53	7.3	e
2	Cobas	13	100.0	0.0	0.0	51	2.2	e
3	Reflotron	441	97.7	0.9	1.4	60	5.4	e

Bilirubina totale

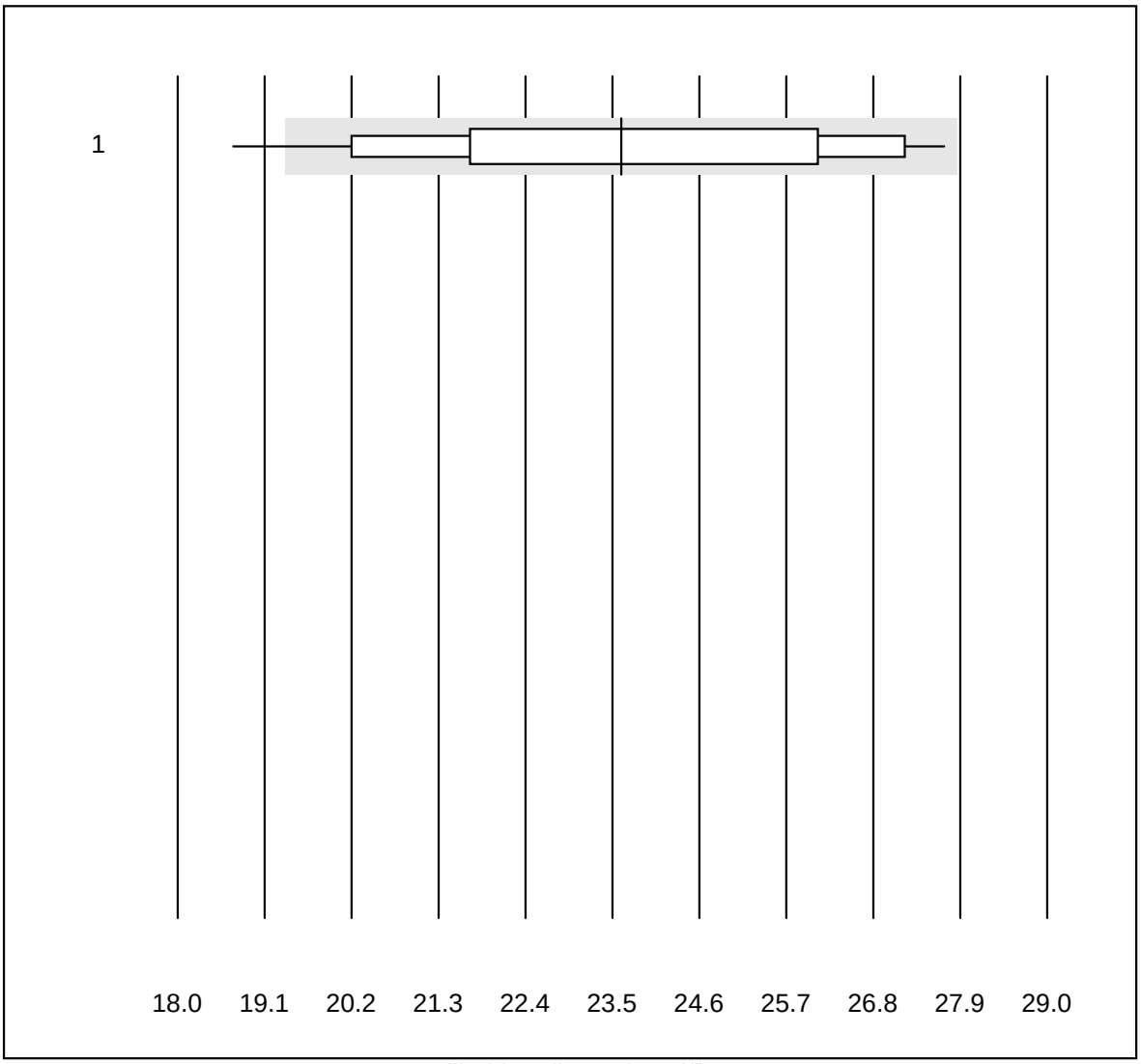


Deviatione QUALAB : 18 %

Bilirubina totale (µmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	15	93.3	6.7	0.0	43.9	10.3	e*
2	Cobas	16	100.0	0.0	0.0	39.1	5.3	e
3	Reflotron	491	93.3	2.6	4.1	44.8	8.1	e
4	Fuji Dri-Chem	496	97.4	1.6	1.0	41.9	6.8	e
5	Spotchem/Ready	94	91.5	5.3	3.2	43.0	8.9	e
6	Spotchem D-Concept	106	96.3	2.8	0.9	34.5	7.3	e
7	Beckman/Olympus	7	100.0	0.0	0.0	49.7	2.2	e
8	Piccolo	28	96.4	0.0	3.6	42.9	6.6	e
9	Abx Mira	20	90.0	5.0	5.0	42.3	8.2	e
10	Hitachi S40/M40	12	100.0	0.0	0.0	46.8	9.0	e*

Bilirubina diretto

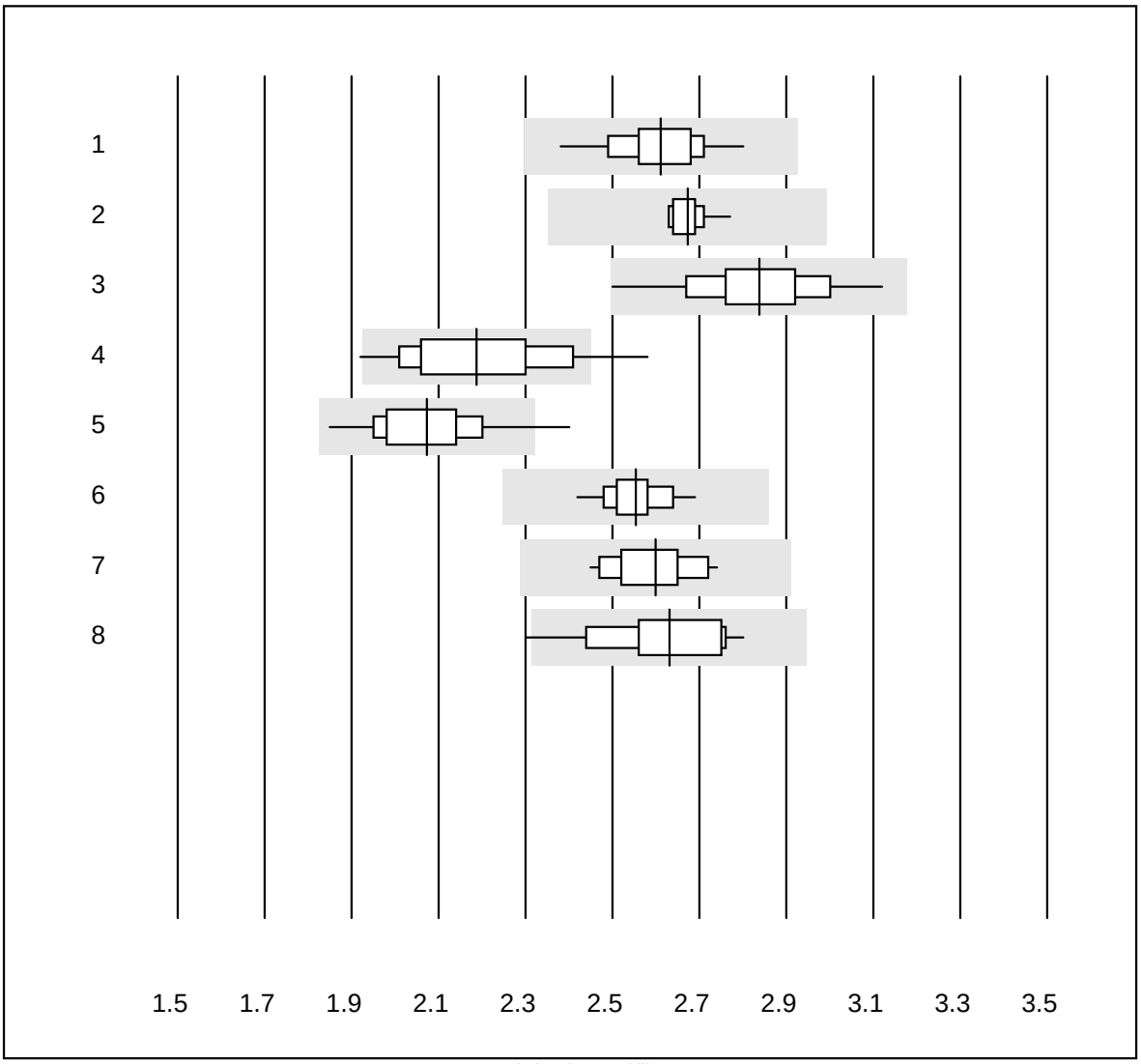


Deviatione QUALAB : 18 %

Bilirubina diretto (µmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Fuji Dri-Chem	32	87.5	3.1	9.4	23.6	10.8	e

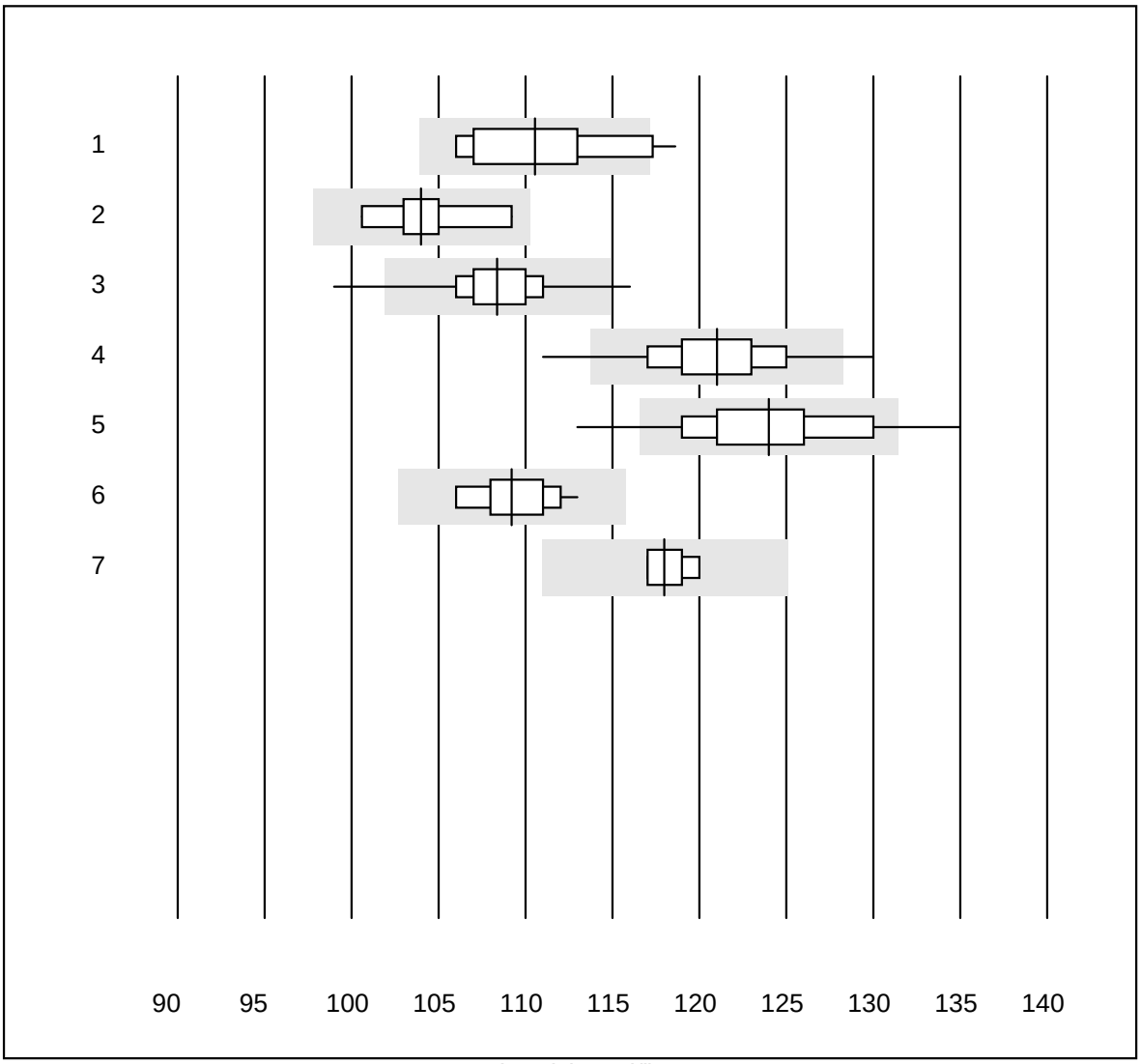
Calcio



Deviatione QUALAB : 12 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	23	95.7	0.0	4.3	2.61	3.6	e
2	Cobas	12	100.0	0.0	0.0	2.67	1.6	e
3	Fuji Dri-Chem	339	99.1	0.0	0.9	2.84	4.3	e
4	Spotchem/Ready	44	93.2	6.8	0.0	2.19	7.3	e
5	Spotchem D-Concept	66	95.5	3.0	1.5	2.07	5.5	e
6	Piccolo	29	100.0	0.0	0.0	2.55	2.4	e
7	Abx Mira	14	100.0	0.0	0.0	2.60	3.5	e
8	Hitachi S40/M40	11	90.9	9.1	0.0	2.63	5.7	e*

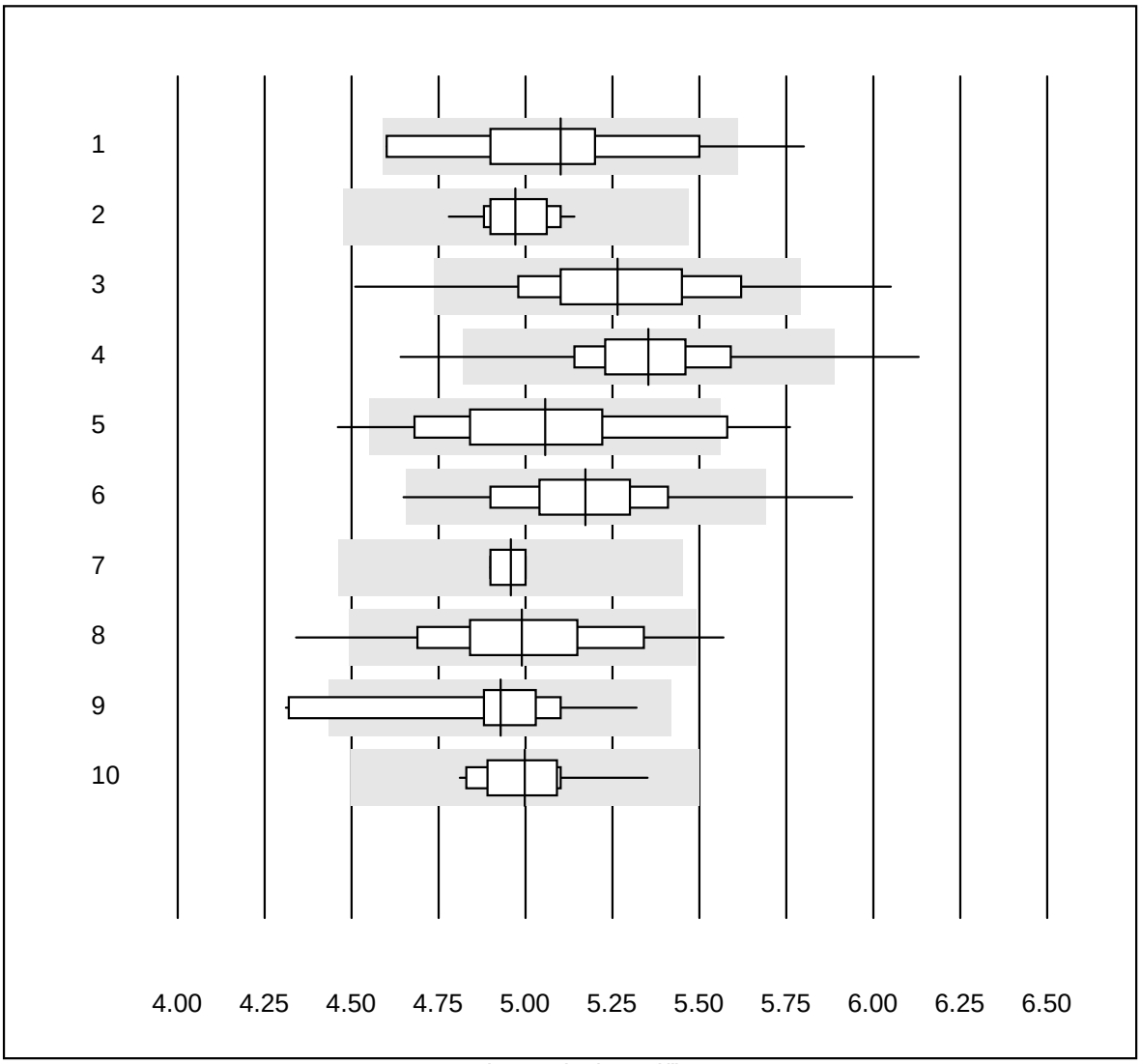
Cloruri



Deviatione QUALAB : 6 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ISE	16	81.2	12.5	6.3	111	3.8	e*
2	Cobas	9	100.0	0.0	0.0	104	2.6	e*
3	Fuji Dri-Chem	585	98.1	1.7	0.2	108	2.0	e
4	Spotchem D-Concept	120	93.4	3.3	3.3	121	2.7	e
5	Spotchem EL-SE 1520	110	85.5	12.7	1.8	124	3.6	e
6	Piccolo	18	100.0	0.0	0.0	109	1.9	e
7	iStat Chem8	4	100.0	0.0	0.0	118	1.3	e

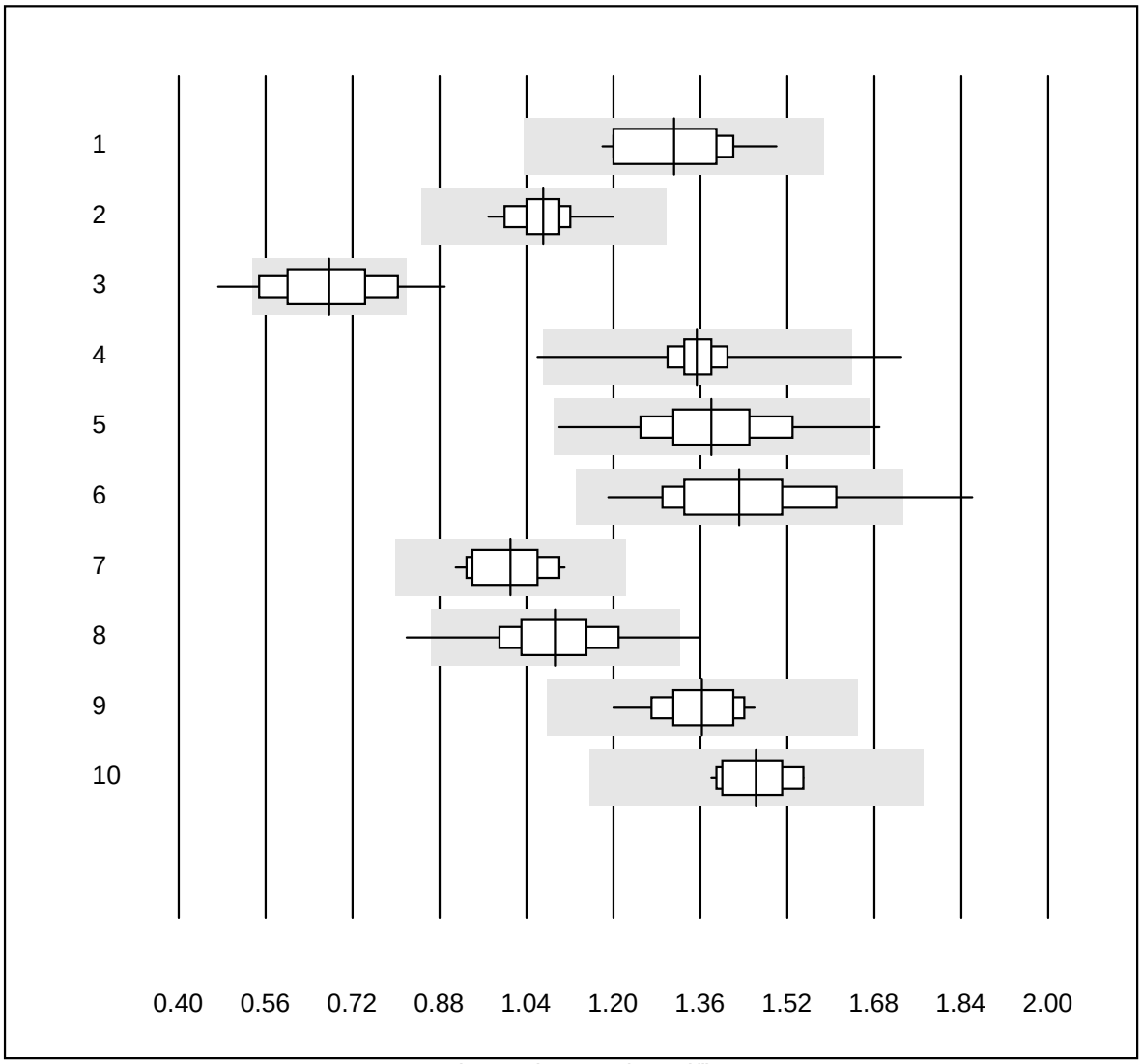
Colesterolo



Deviatione QUALAB : 10 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	22	95.5	4.5	0.0	5.1	6.3	e*
2	Cobas	15	100.0	0.0	0.0	5.0	2.0	e
3	Reflotron	769	95.4	3.8	0.8	5.3	4.8	e
4	Fuji Dri-Chem	683	97.8	1.5	0.7	5.4	3.5	e
5	Spotchem/Ready	142	85.2	12.0	2.8	5.1	6.1	e
6	Spotchem D-Concept	137	97.1	2.9	0.0	5.2	4.2	e
7	Piccolo	21	100.0	0.0	0.0	5.0	1.0	e
8	Cholestech LDX	187	94.1	4.8	1.1	5.0	4.8	e
9	Abx Mira	20	90.0	10.0	0.0	4.9	5.2	e
10	Hitachi S40/M40	13	100.0	0.0	0.0	5.0	3.0	e

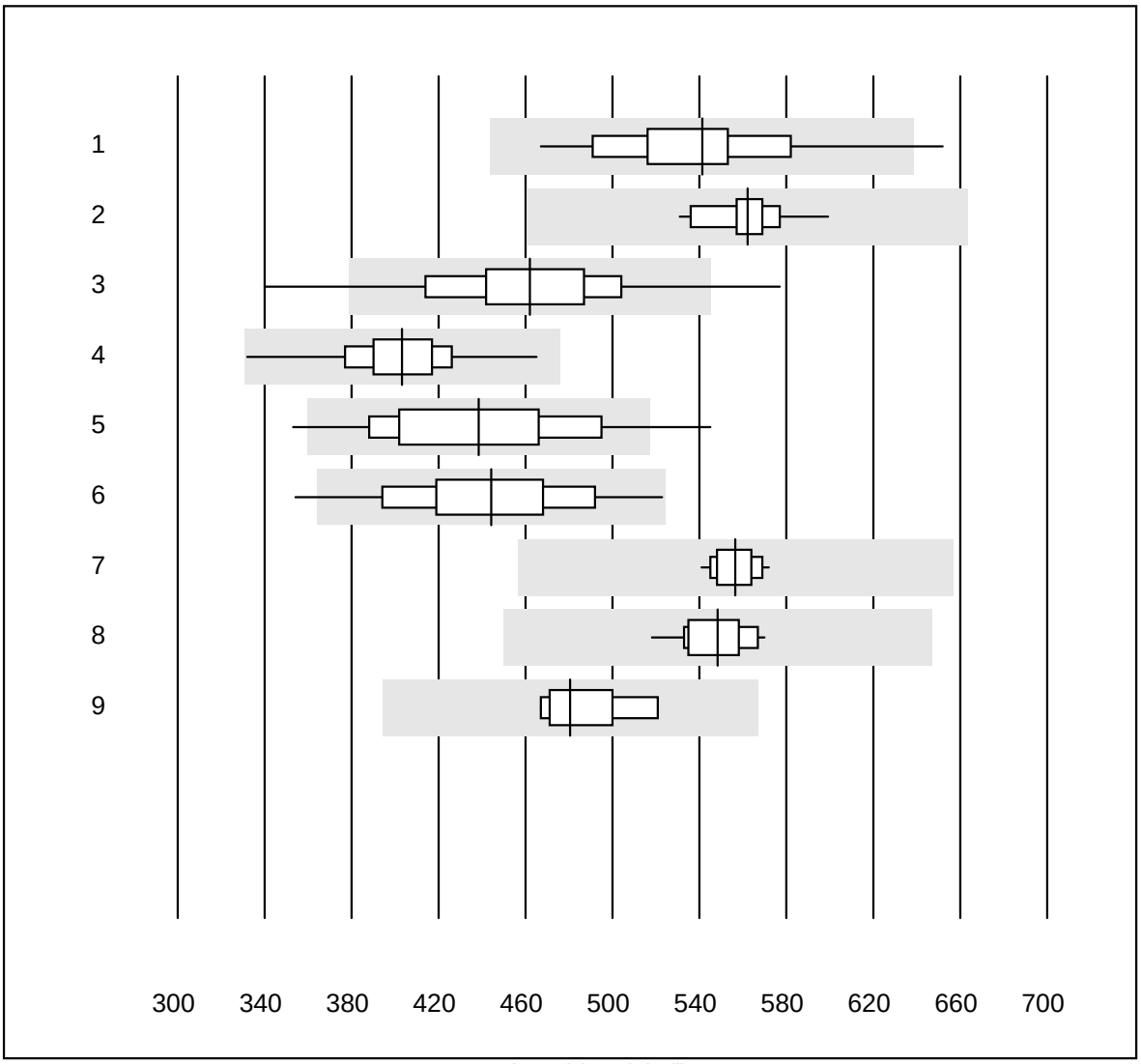
Colesterolo HDL



Deviatione QUALAB : 21 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	umida, diretto	18	100.0	0.0	0.0	1.31	7.1	e
2	Cobas	13	100.0	0.0	0.0	1.07	5.5	e
3	Reflotron	576	74.6	13.4	12.0	0.68	14.4	e
4	Fuji Dri-Chem	646	99.0	0.5	0.5	1.35	3.8	e
5	Spotchem/Ready	128	96.9	2.3	0.8	1.38	8.6	e
6	Spotchem D-Concept	133	96.2	3.0	0.8	1.43	9.4	e
7	Piccolo	22	100.0	0.0	0.0	1.01	6.4	e
8	Cholestech LDX	187	95.2	3.2	1.6	1.09	8.8	e
9	Abx Mira	19	100.0	0.0	0.0	1.36	4.9	e
10	Hitachi S40/M40	12	100.0	0.0	0.0	1.46	4.1	e

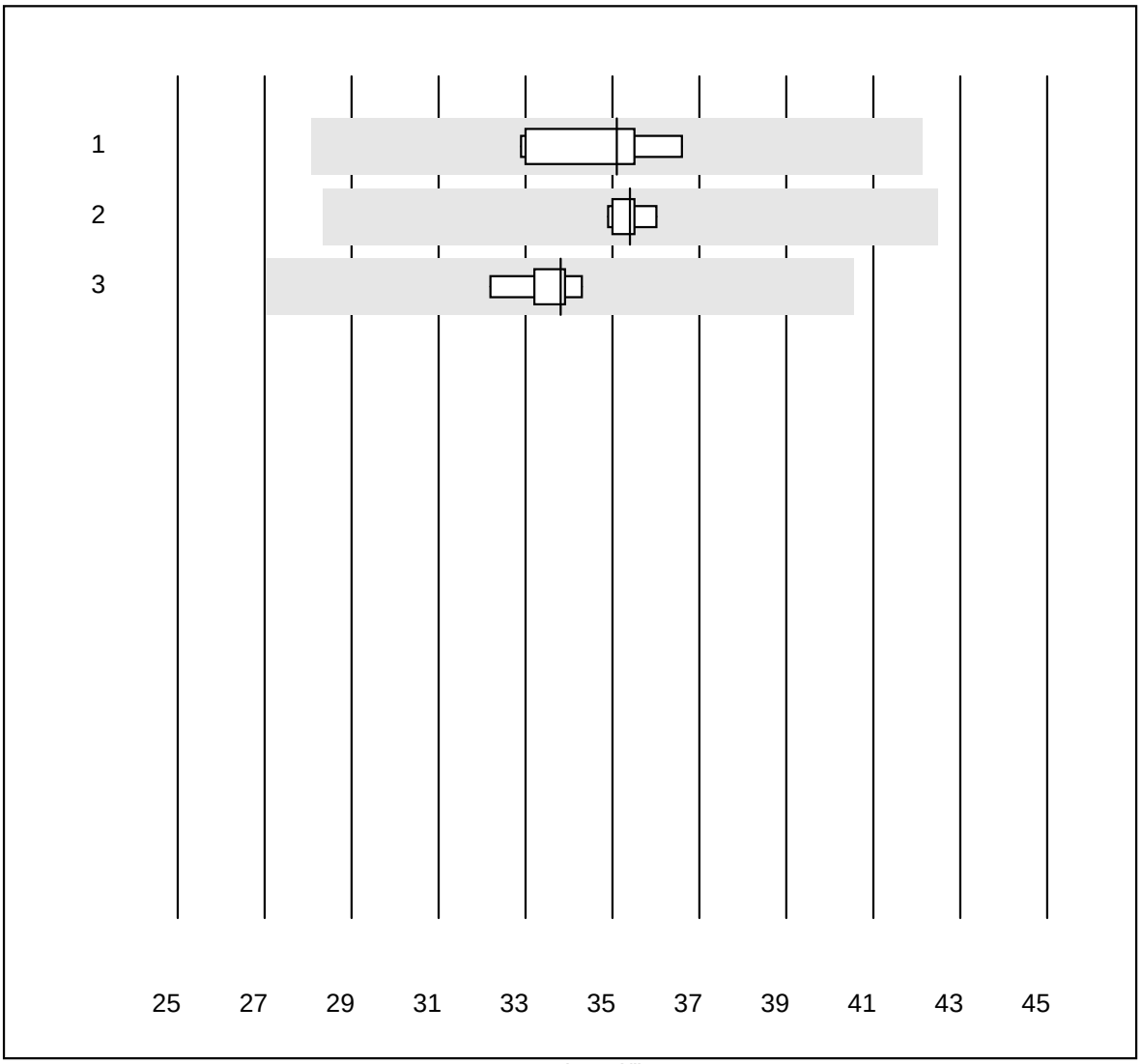
Creatina chinasi



Deviatione QUALAB : 18 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	IFCC/SGKC/SFBC 37'C	20	95.0	5.0	0.0	541	7.3	e
2	Cobas	15	100.0	0.0	0.0	562	2.9	e
3	Reflotron	412	92.0	4.4	3.6	462	8.2	e
4	Fuji Dri-Chem	426	98.6	0.0	1.4	403	5.0	e
5	Spotchem/Ready	54	90.7	5.6	3.7	438	9.7	e
6	Spotchem D-Concept	82	95.2	2.4	2.4	444	8.3	e
7	Piccolo	12	100.0	0.0	0.0	557	1.8	e
8	Abx Mira	16	100.0	0.0	0.0	549	2.7	e
9	Hitachi S40/M40	8	100.0	0.0	0.0	481	4.0	e

Ferro

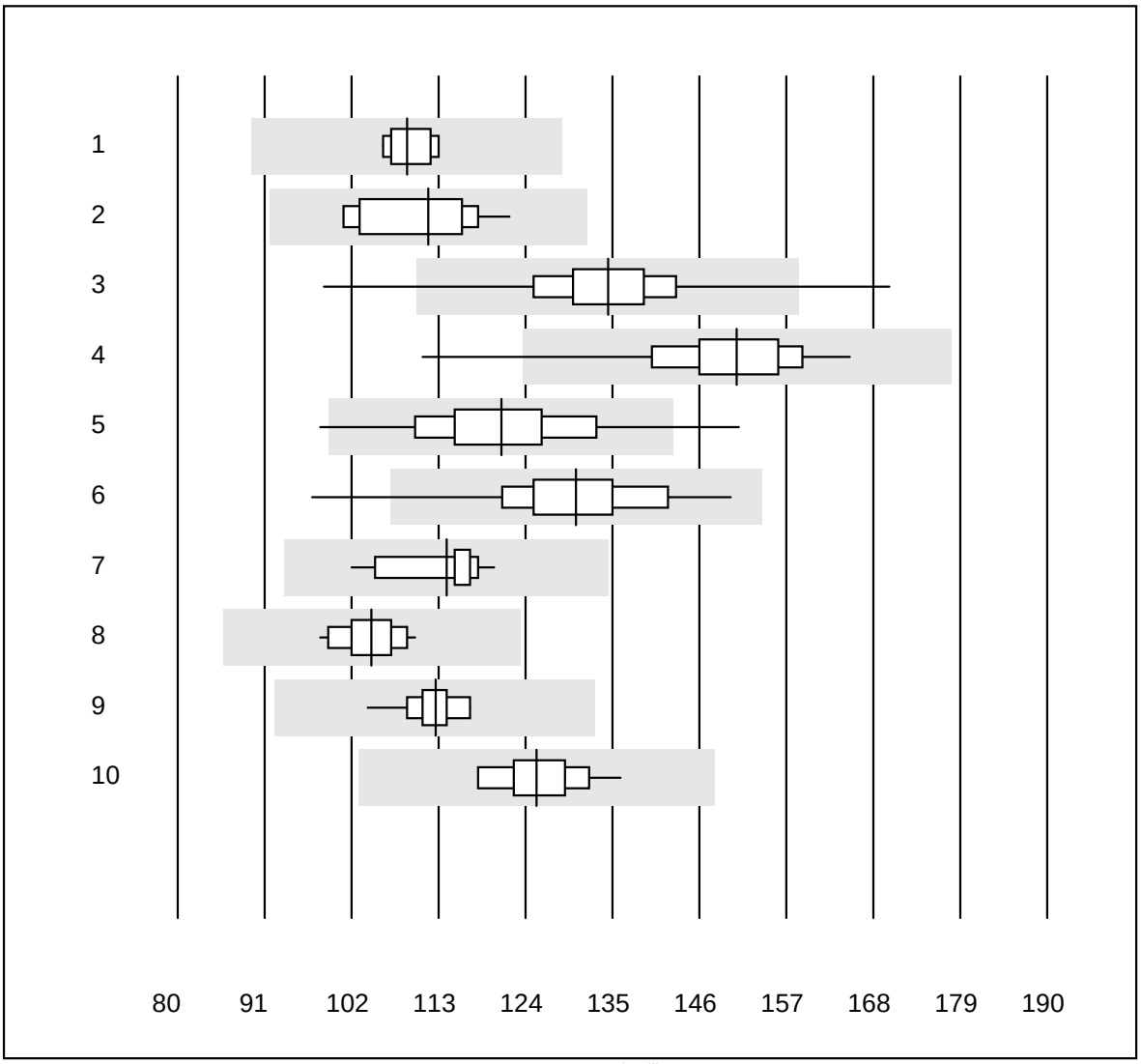


Deviatione QUALAB : 20 %

Ferro (µmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	9	88.9	0.0	11.1	35	4.0	e
2	Cobas	9	100.0	0.0	0.0	35	1.1	e
3	Abx Mira	5	100.0	0.0	0.0	34	2.4	e

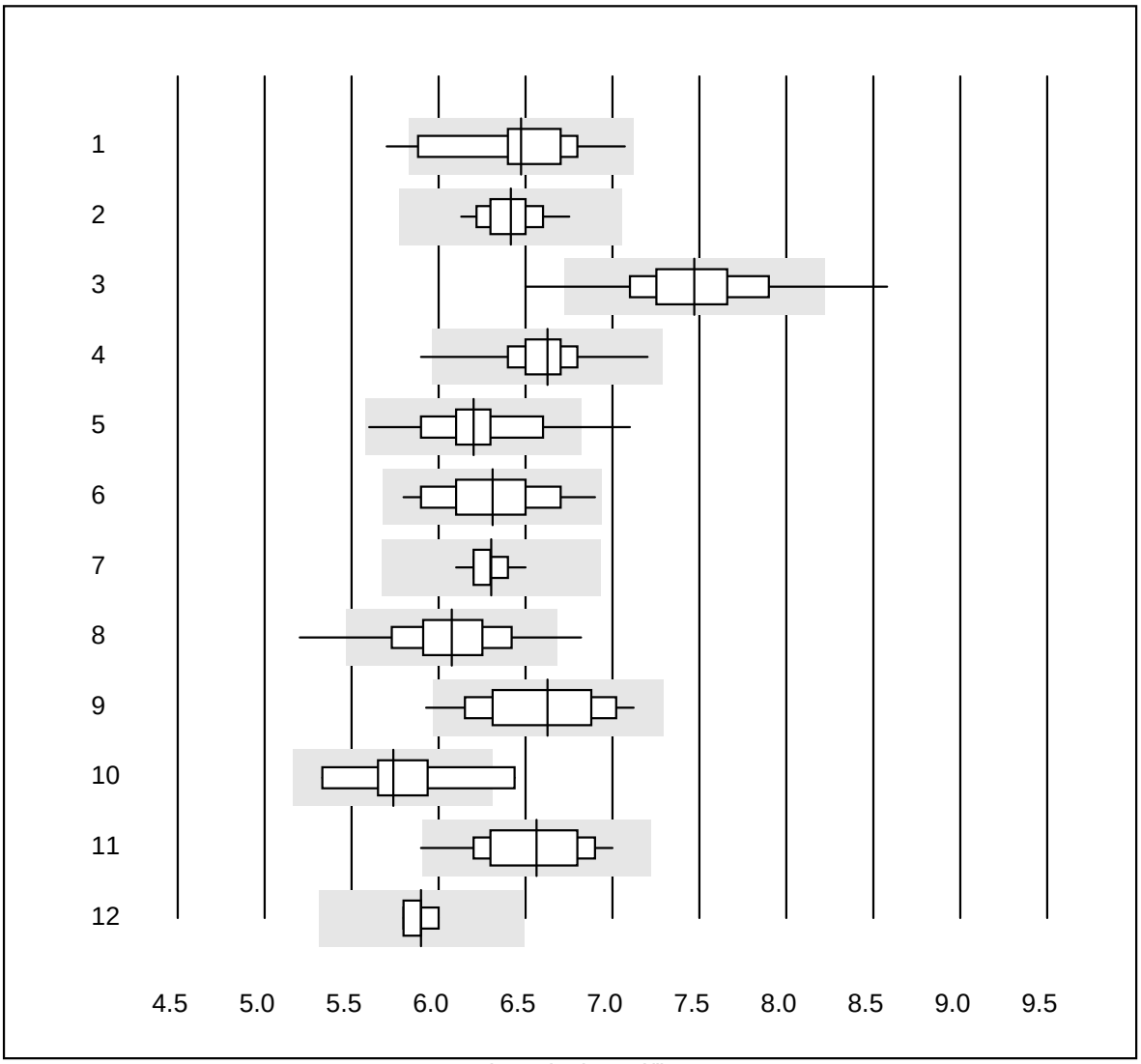
Gamma-GT



Deviatione QUALAB : 18 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	IFCC/SGKC/SFBC 37'C	7	100.0	0.0	0.0	109	2.4	e
2	Cobas	16	100.0	0.0	0.0	112	6.3	e
3	Reflotron	882	97.5	1.9	0.6	134	6.0	e
4	Fuji Dri-Chem	722	99.0	0.6	0.4	151	5.2	e
5	Spotchem/Ready	146	97.2	1.4	1.4	121	7.4	e
6	Spotchem D-Concept	149	98.6	0.7	0.7	130	6.3	e
7	Mètodo standard, 37'	13	100.0	0.0	0.0	114	4.7	e
8	Piccolo	28	100.0	0.0	0.0	105	3.0	e
9	Abx Mira	21	100.0	0.0	0.0	113	3.0	e
10	Hitachi S40/M40	15	100.0	0.0	0.0	125	4.1	e

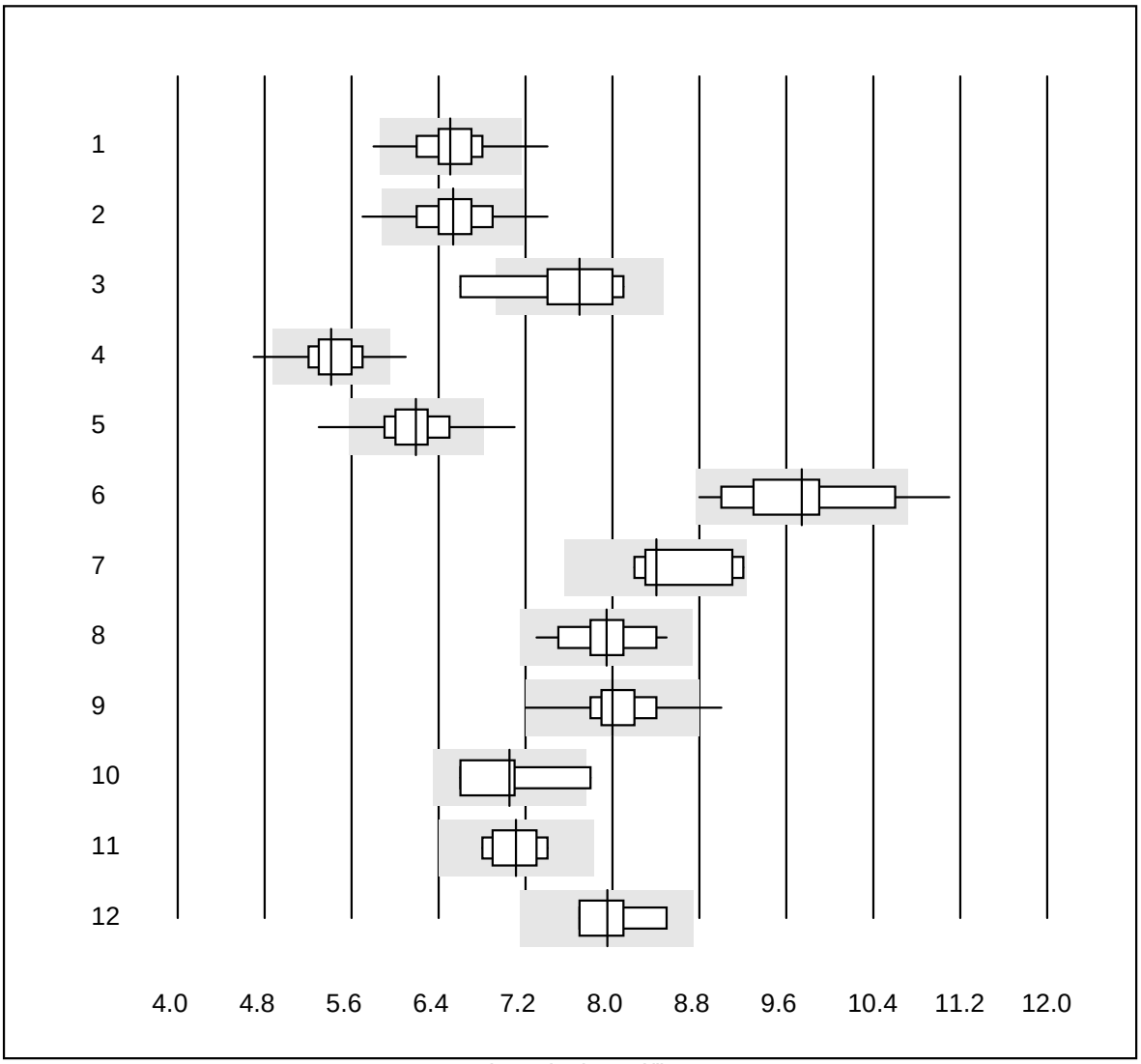
Glucosio



Deviatione QUALAB : 10 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	27	92.6	7.4	0.0	6.5	5.3	e
2	Cobas	16	100.0	0.0	0.0	6.4	2.5	e
3	Reflotron	899	95.3	2.7	2.0	7.5	4.3	e
4	Fuji Dri-Chem	685	99.5	0.1	0.4	6.6	2.4	e
5	Spotchem/Ready	132	97.7	1.5	0.8	6.2	4.3	e
6	Spotchem D-Concept	139	99.3	0.0	0.7	6.3	4.1	e
7	Piccolo	36	100.0	0.0	0.0	6.3	1.5	e
8	Cholestech LDX	150	95.4	3.3	1.3	6.1	4.5	e
9	Abx Mira	21	90.4	4.8	4.8	6.6	4.9	e
10	Lange	6	66.6	16.7	16.7	5.7	7.0	e*
11	Hitachi S40/M40	16	93.7	6.3	0.0	6.6	4.5	e
12	iStat Chem8	4	100.0	0.0	0.0	5.9	1.4	e

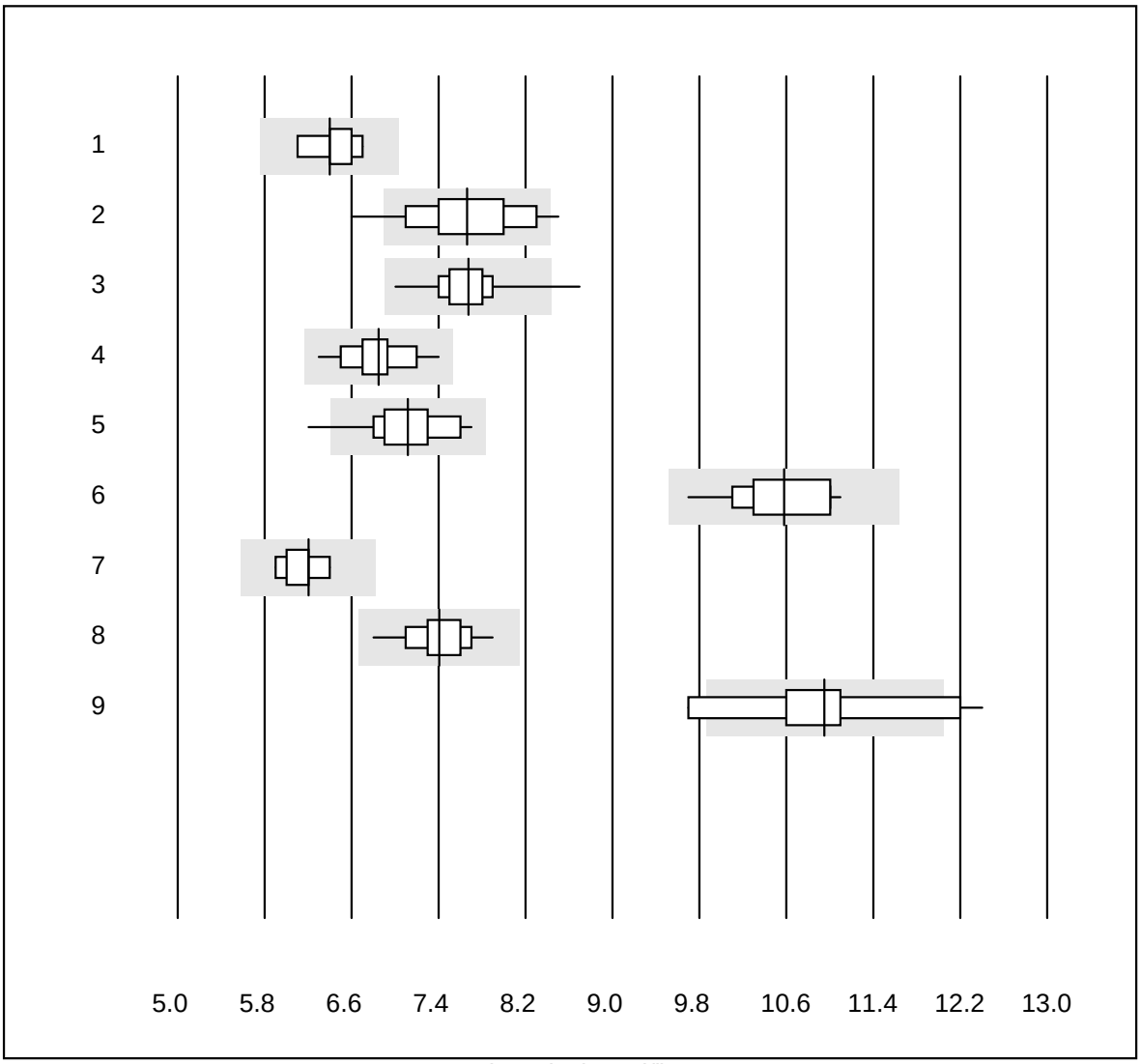
Glucosio



Deviatione QUALAB : 10 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Accu-Chek Aviva	390	96.9	1.8	1.3	6.5	3.6	e
2	Accu-Chek Inform 2	229	97.8	2.2	0.0	6.5	4.0	e
3	Accu-Chek Mobile	5	80.0	20.0	0.0	7.7	8.0	e*
4	Bayer Contour 2 (5s)	49	91.8	4.1	4.1	5.4	4.6	e
5	Bayer Contour XT/NEX	1091	97.3	2.0	0.7	6.2	4.1	e
6	Bayer Breeze 2	17	94.1	5.9	0.0	9.7	5.9	e*
7	Glucocard	9	77.8	0.0	22.2	8.4	4.6	e*
8	Hemocue 201+ P-equiv	83	94.0	0.0	6.0	7.9	3.7	e
9	Hemocue 201RT P-equi	36	88.8	5.6	5.6	8.0	3.7	e
10	FreeStyle Precision	6	50.0	16.7	33.3	7.1	7.0	e*
11	Freestyle Freedom li	10	90.0	0.0	10.0	7.1	3.2	e
12	Sanofi BG Star	6	83.3	0.0	16.7	8.0	4.3	e*

Glucosio

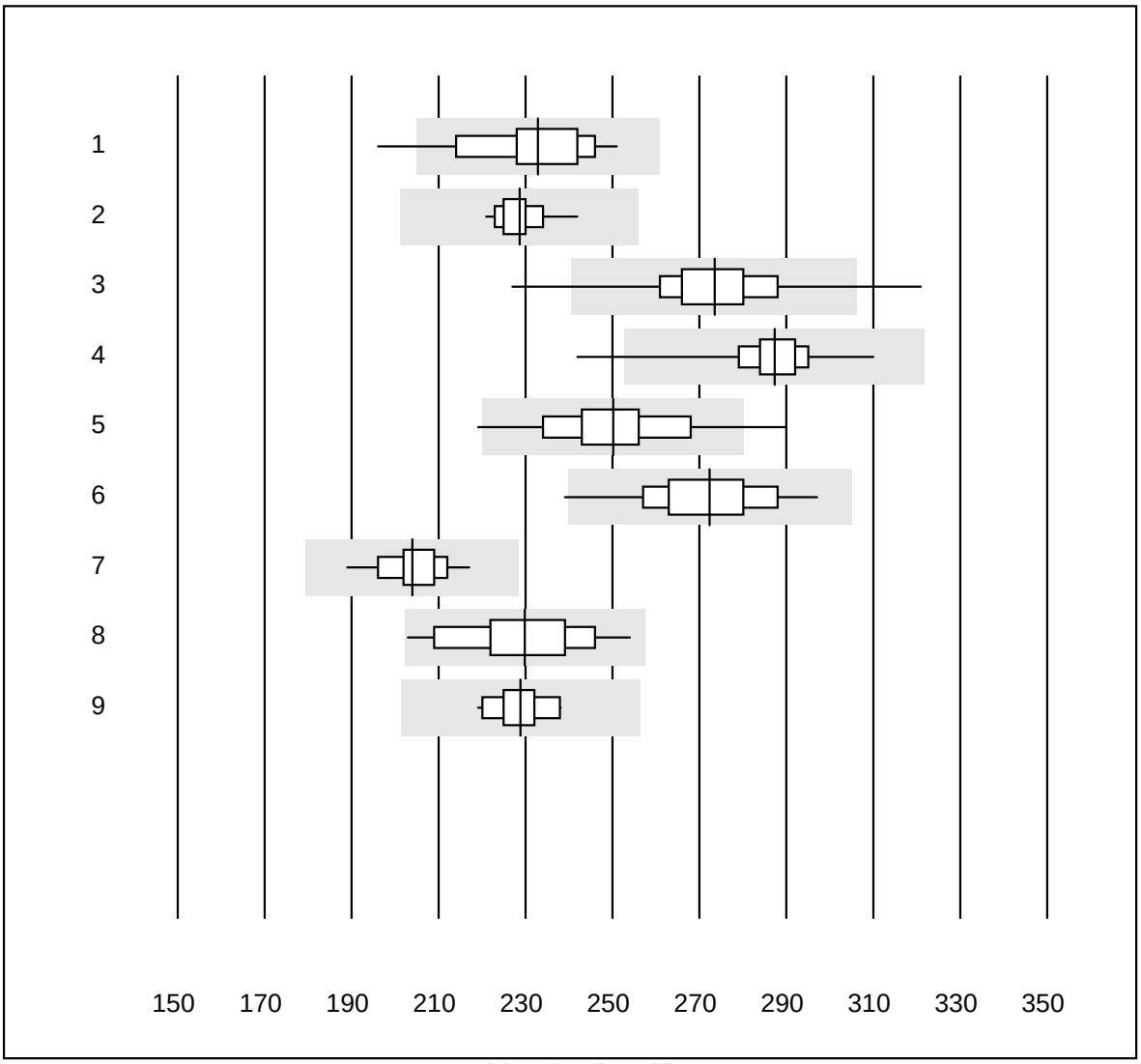


Deviatione QUALAB : 10 %

Glucosio (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Bayer Elite	9	100.0	0.0	0.0	6.4	2.8	e
2	Hemocue 201+ (alt)	50	92.0	6.0	2.0	7.7	5.6	e
3	OneTouch Ultra	19	94.7	5.3	0.0	7.7	4.3	e
4	OneTouch Verio	16	100.0	0.0	0.0	6.8	3.9	e
5	Bayer Contour (15s)	45	97.8	2.2	0.0	7.1	4.3	e
6	Healthpro	14	100.0	0.0	0.0	10.6	4.1	e
7	Mylife UNIO	5	100.0	0.0	0.0	6.2	3.2	e*
8	mylife Pura	65	100.0	0.0	0.0	7.4	3.0	e
9	Omnitest	17	70.6	29.4	0.0	11.0	7.6	e*

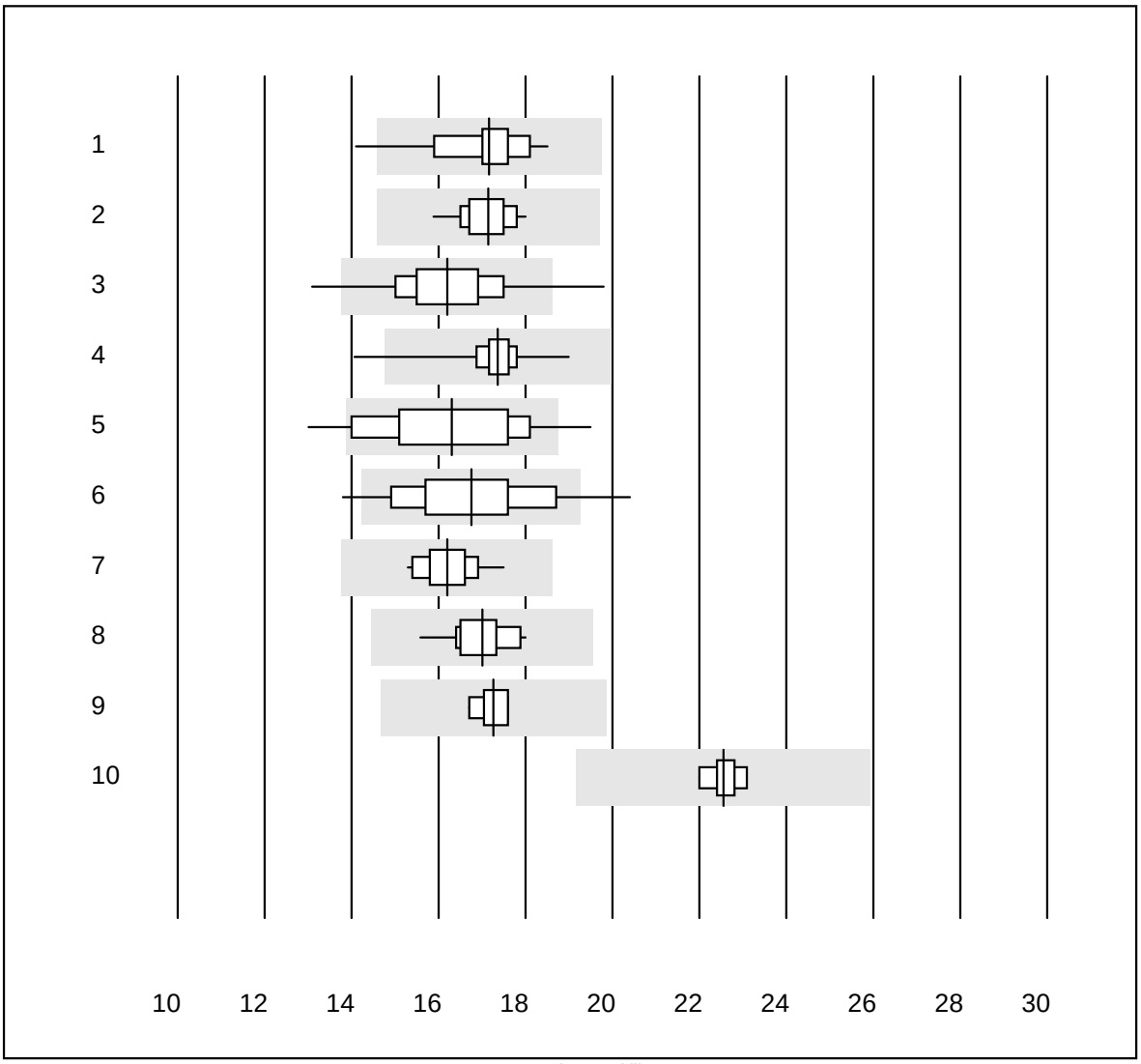
Acido urico



Deviatione QUALAB : 12 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	22	95.5	4.5	0.0	233	5.7	e
2	Cobas	12	100.0	0.0	0.0	229	2.4	e
3	Reflotron	781	96.7	1.0	2.3	273	4.1	e
4	Fuji Dri-Chem	681	99.0	0.4	0.6	287	2.5	e
5	Spotchem/Ready	123	98.4	1.6	0.0	250	5.0	e
6	Spotchem D-Concept	132	99.2	0.8	0.0	272	4.5	e
7	Piccolo	24	95.8	0.0	4.2	204	3.3	e
8	Abx Mira	19	100.0	0.0	0.0	230	5.4	e
9	Hitachi S40/M40	14	100.0	0.0	0.0	229	2.6	e

Urea

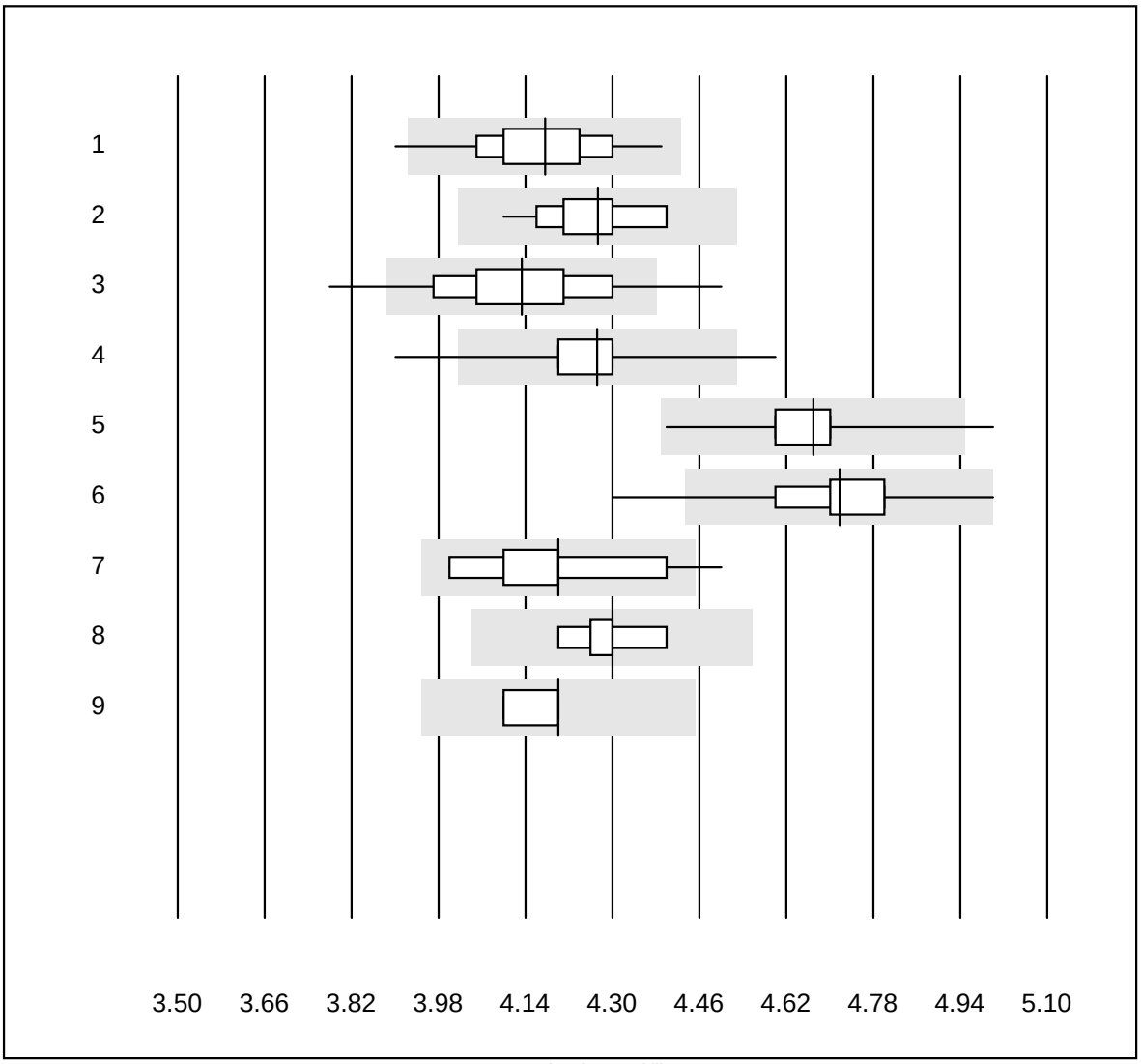


Deviatione QUALAB : 15 %

Urea (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	21	80.9	4.8	14.3	17.2	5.6	e
2	Cobas	15	100.0	0.0	0.0	17.1	3.1	e
3	Reflotron	341	94.7	3.2	2.1	16.2	6.4	e
4	Fuji Dri-Chem	423	99.6	0.2	0.2	17.4	2.4	e
5	Spotchem/Ready	81	84.0	12.3	3.7	16.3	9.7	e
6	Spotchem D-Concept	82	85.3	9.8	4.9	16.8	8.7	e
7	Piccolo	34	97.1	0.0	2.9	16.2	3.5	e
8	Abx Mira	12	100.0	0.0	0.0	17.0	4.0	e
9	Hitachi S40/M40	10	100.0	0.0	0.0	17.3	1.8	e
10	iStat Chem8	6	100.0	0.0	0.0	22.6	1.7	e

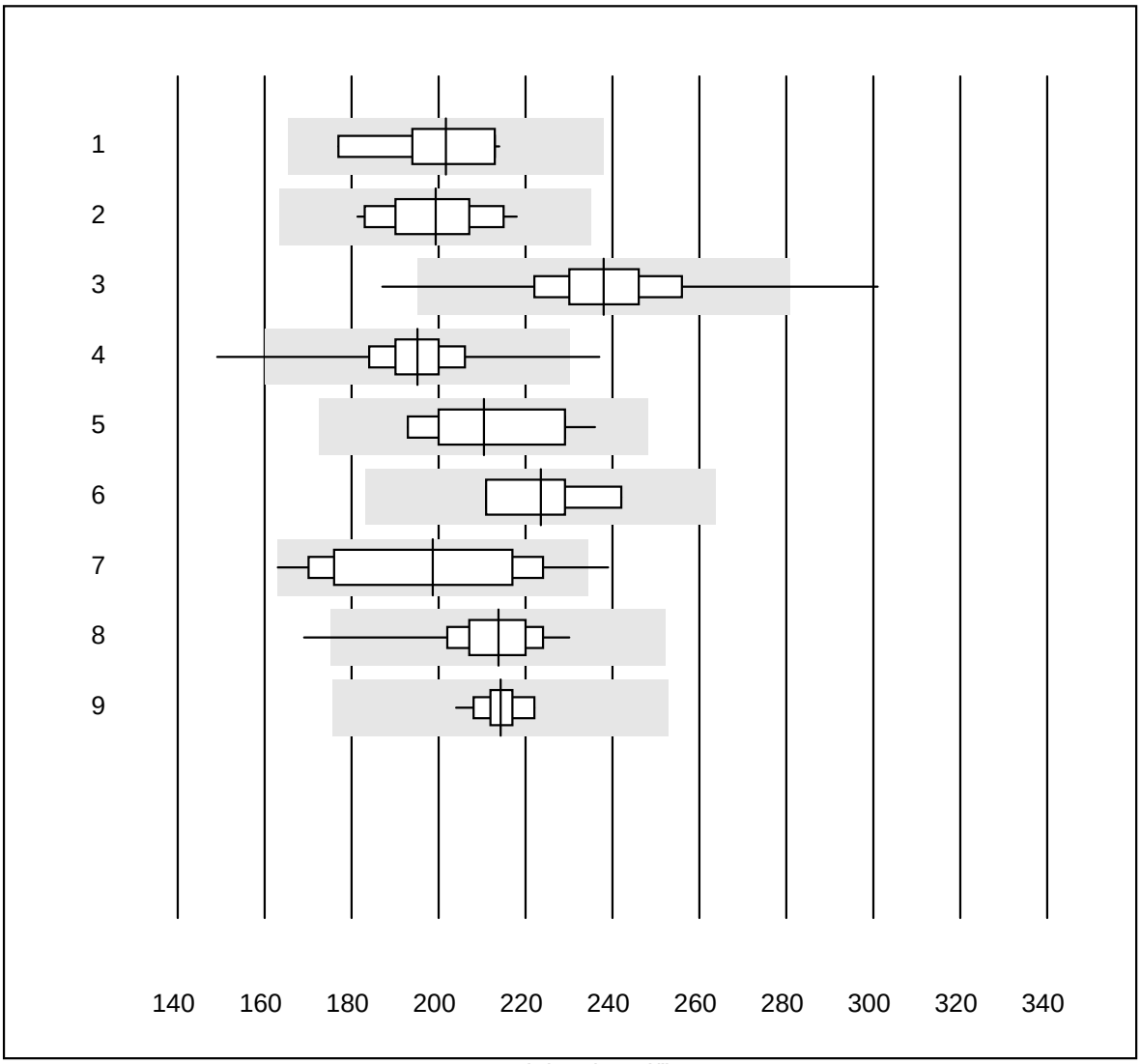
Potassio



Deviatione QUALAB : 6 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ISE	28	92.8	3.6	3.6	4.18	2.5	e
2	Cobas	16	100.0	0.0	0.0	4.27	1.9	e
3	Reflotron	808	89.9	6.3	3.8	4.13	3.2	e
4	Fuji Dri-Chem	715	97.7	2.0	0.3	4.27	2.0	e
5	Spotchem D-Concept	136	97.8	0.7	1.5	4.67	1.8	e
6	Spotchem EL-SE 1520	115	95.7	1.7	2.6	4.72	2.4	e
7	Piccolo	21	71.4	4.8	23.8	4.20	3.3	e*
8	Abx Mira	5	100.0	0.0	0.0	4.30	1.7	e*
9	iStat Chem8	7	100.0	0.0	0.0	4.20	1.3	e

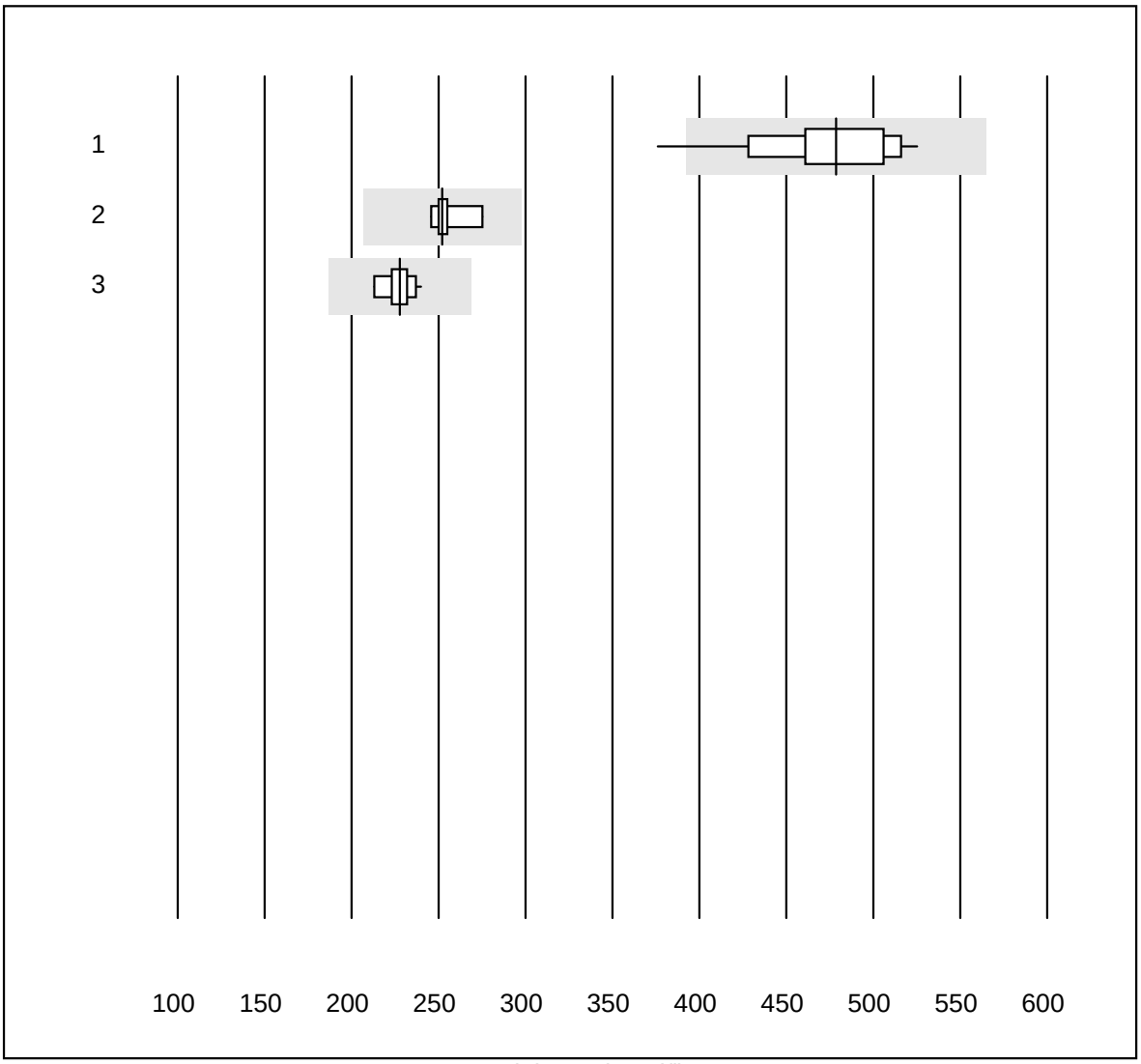
Creatinina



Deviatione QUALAB : 18 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	11	90.9	0.0	9.1	202	6.4	e
2	Cobas	17	100.0	0.0	0.0	199	6.1	e
3	Reflotron	990	98.0	0.9	1.1	238	5.8	e
4	Fuji Dri-Chem	748	97.7	0.8	1.5	195	4.8	e
5	Jaffé	10	100.0	0.0	0.0	210	7.2	e*
6	Enzymatisch	4	100.0	0.0	0.0	224	6.0	e*
7	Piccolo	34	97.1	2.9	0.0	199	11.0	e
8	Abx Mira	21	95.2	4.8	0.0	214	6.4	e
9	Hitachi S40/M40	15	100.0	0.0	0.0	214	2.4	e

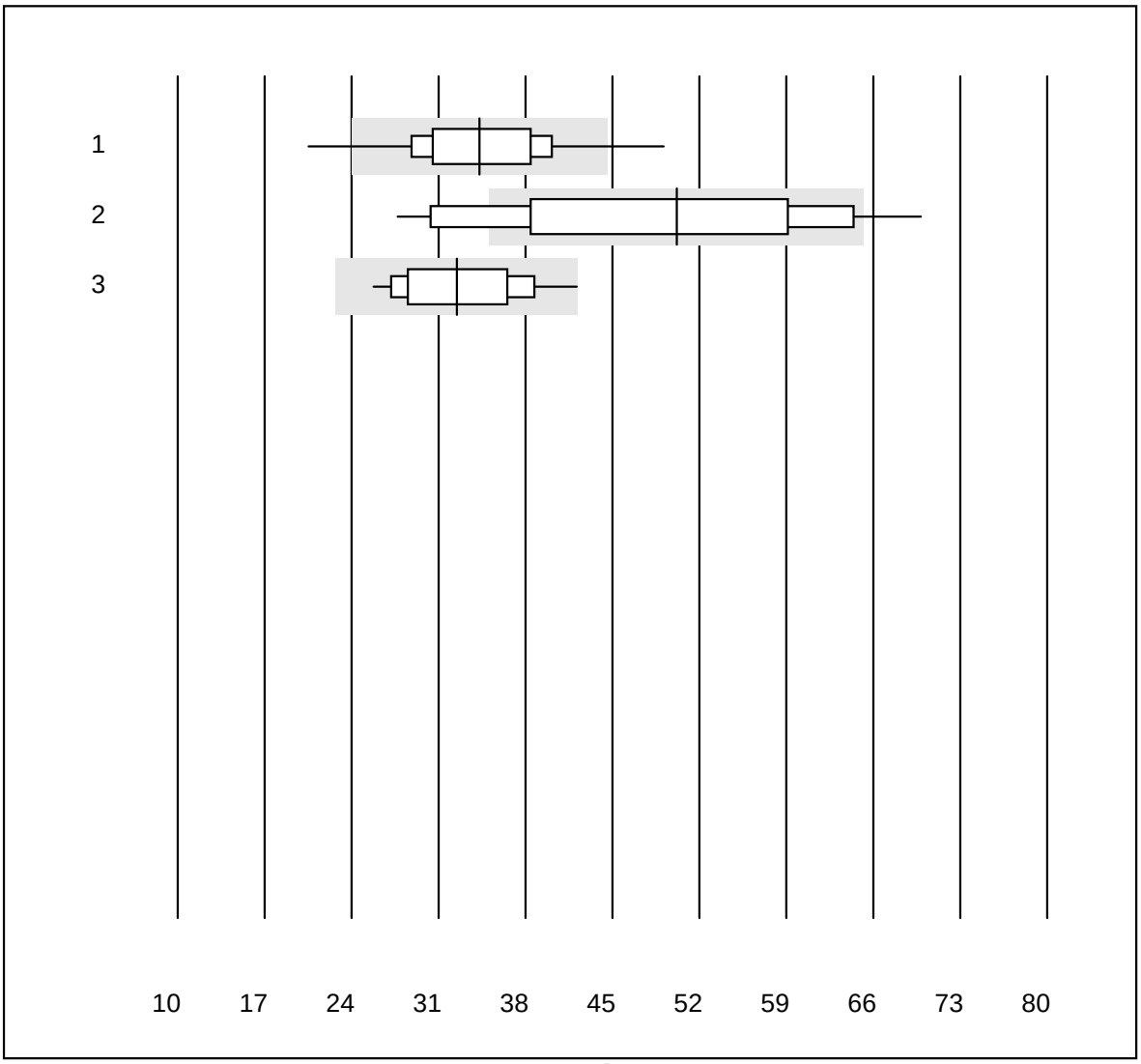
Creatinina E



Deviatione QUALAB : 18 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Statsensor i / Nova	28	96.4	3.6	0.0	479	7.5	e
2	iStat Chem8	9	100.0	0.0	0.0	252	3.4	e
3	ABL700/800 Radiomete	10	100.0	0.0	0.0	228	3.5	e

eGFR

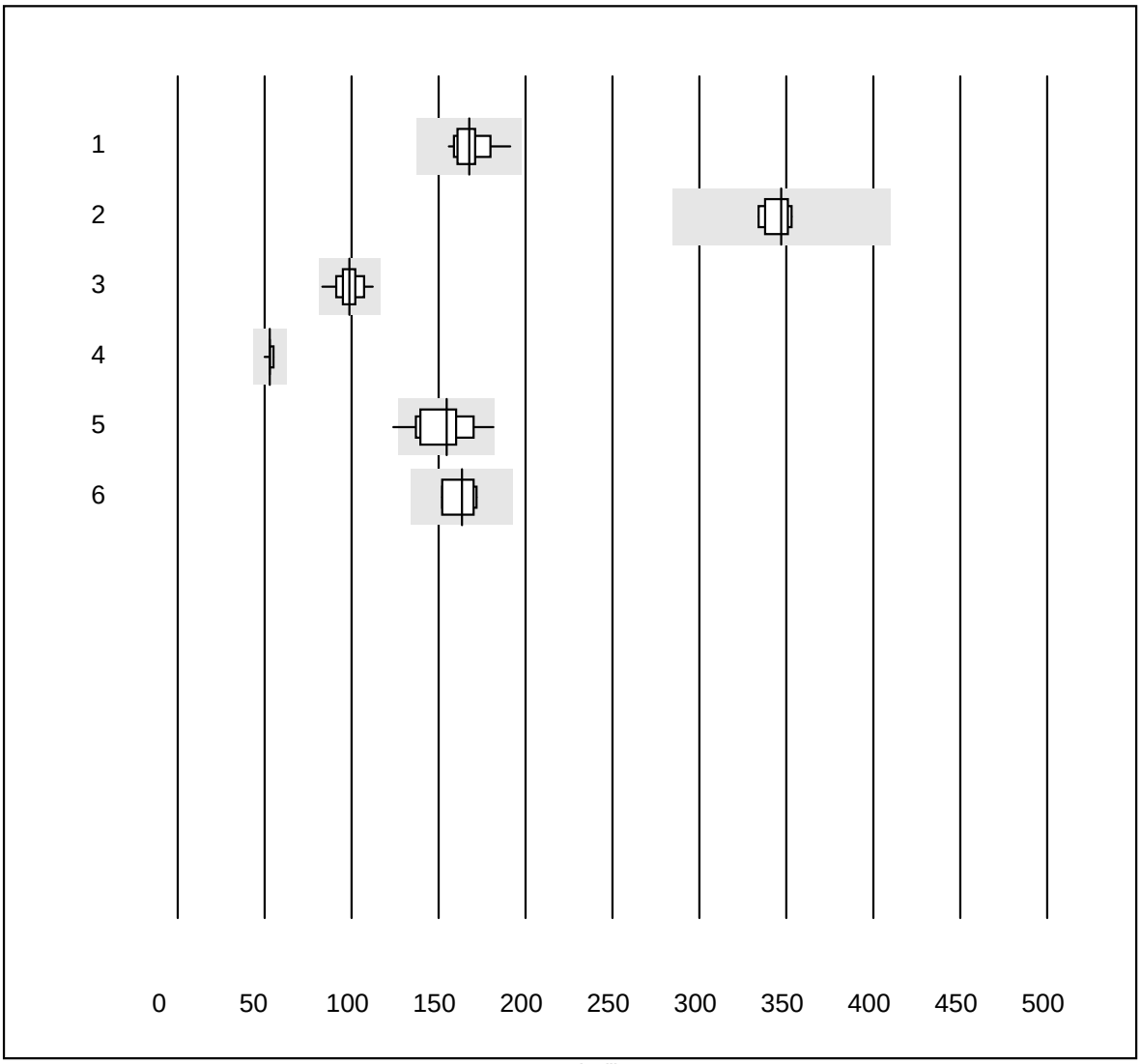


Deviatione QUALAB : 30 %

eGFR ()

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	CKD-EPI	720	94.2	0.8	5.0	34	13.5	e
2	Cockcroft-Gault	57	73.7	22.8	3.5	50	25.0	e
3	MDRD	30	80.0	0.0	20.0	32	14.2	e

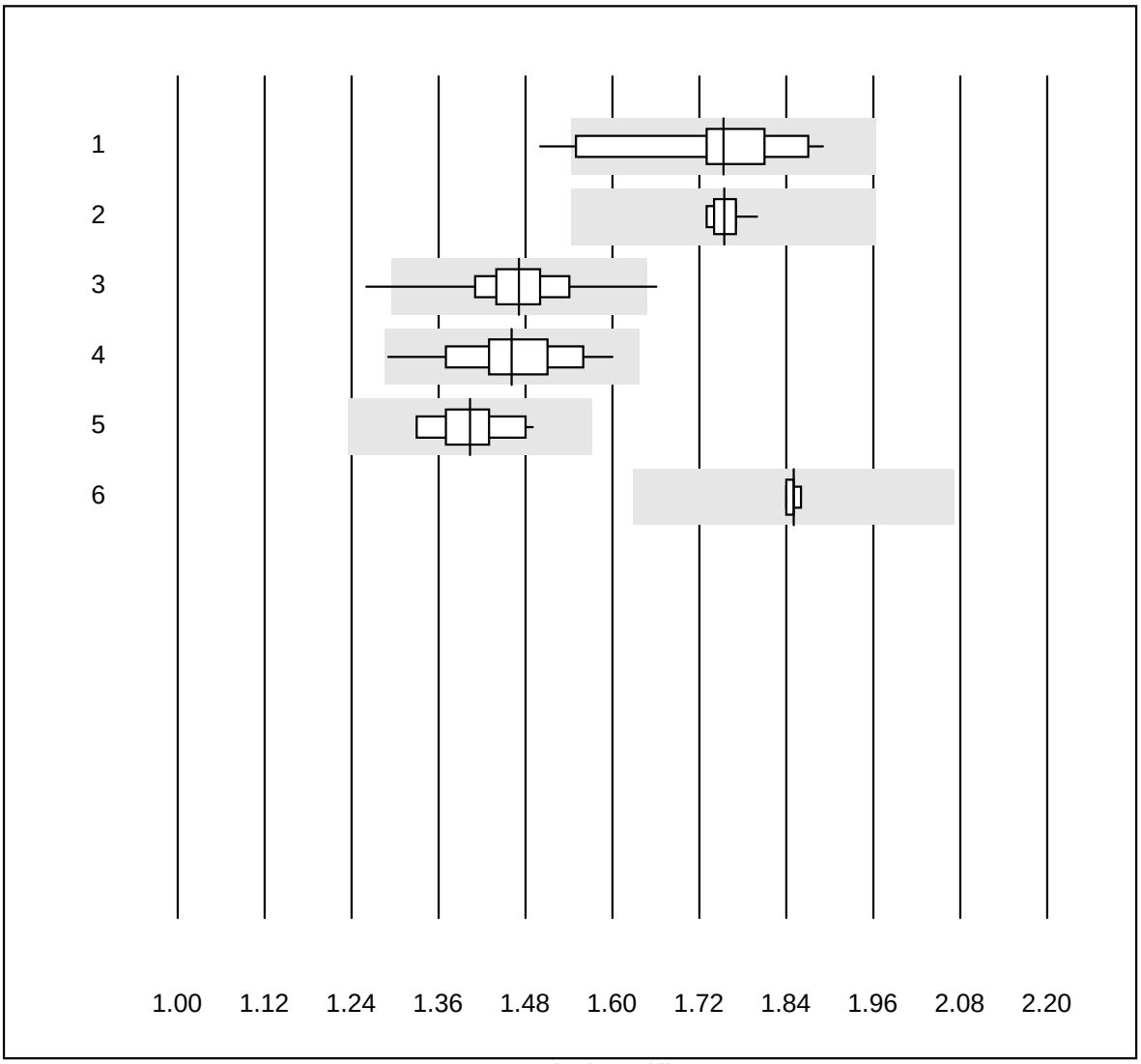
LDH



Deviatione QUALAB : 18 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	IFCC	18	100.0	0.0	0.0	168	5.0	e
2	Cobas	9	100.0	0.0	0.0	347	2.1	e
3	Fuji Dri-Chem	137	99.3	0.0	0.7	99	6.1	e
4	Spotchem/Ready	35	74.3	0.0	25.7	53	1.7	e
5	Abx Mira	12	91.7	8.3	0.0	155	10.0	e*
6	Hitachi S40/M40	5	100.0	0.0	0.0	164	5.9	e*

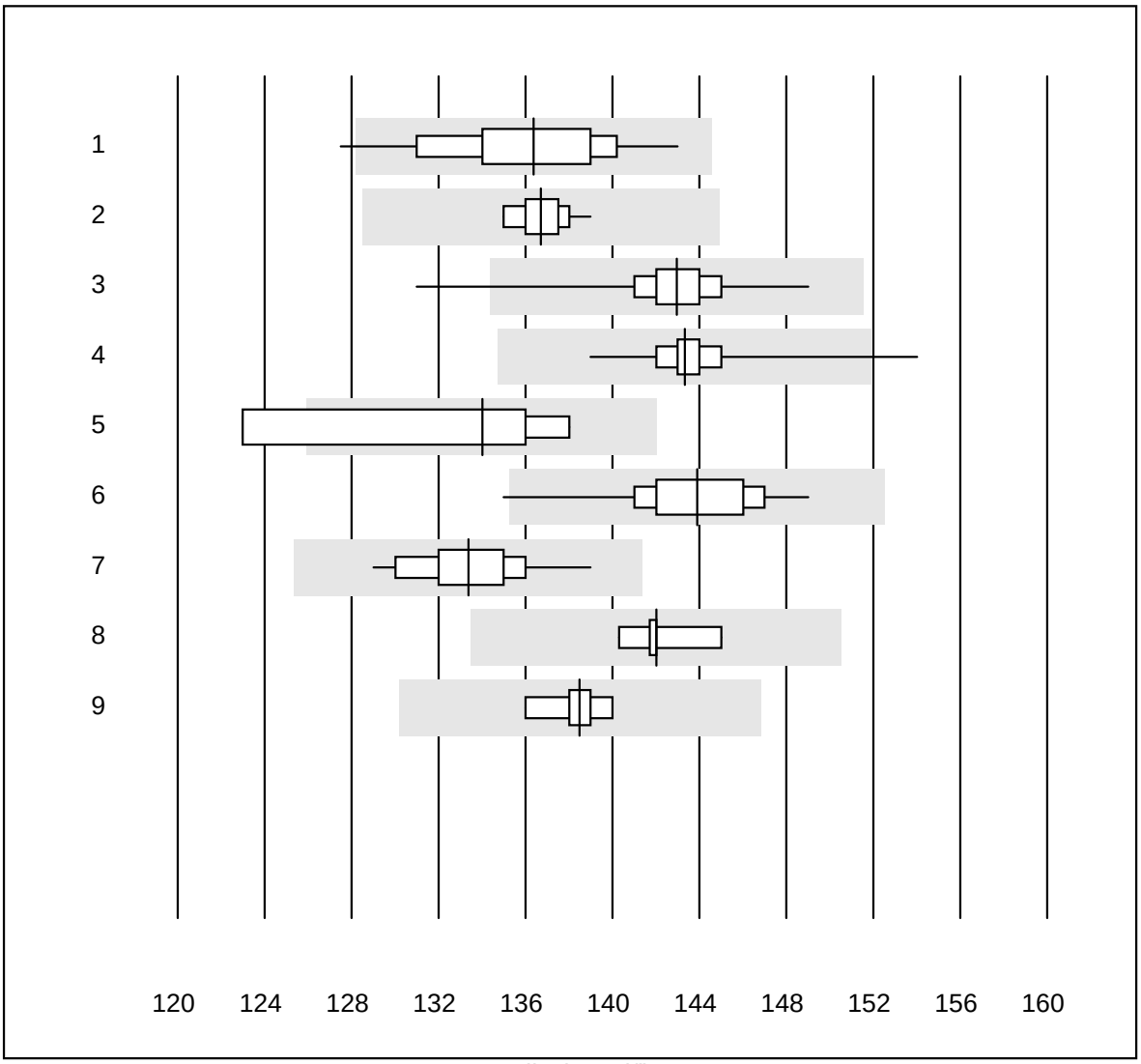
Magnesio



Deviatione QUALAB : 12 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	16	93.7	6.3	0.0	1.75	6.2	e*
2	Cobas	10	100.0	0.0	0.0	1.75	1.2	e
3	Fuji Dri-Chem	116	96.5	2.6	0.9	1.47	4.2	e
4	Spotchem D-Concept	25	100.0	0.0	0.0	1.46	5.3	e
5	Spotchem/Ready	18	100.0	0.0	0.0	1.40	3.4	e
6	Piccolo	4	100.0	0.0	0.0	1.85	0.4	e

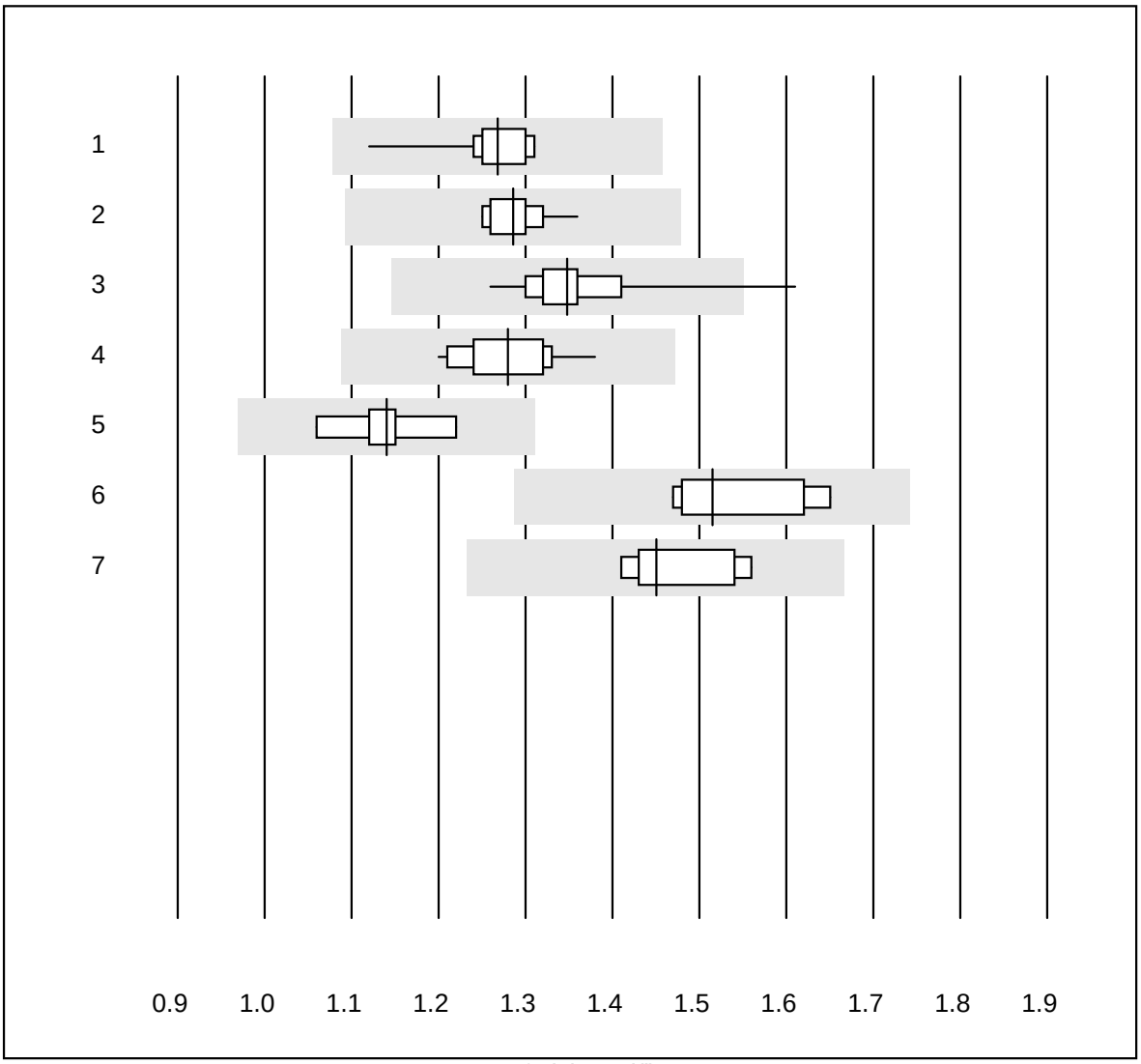
Sodio



Deviatione QUALAB : 6 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ISE	26	92.4	3.8	3.8	136	2.8	e
2	Cobas	15	100.0	0.0	0.0	137	0.8	e
3	Fuji Dri-Chem	660	98.6	1.1	0.3	143	1.5	e
4	Spotchem D-Concept	130	98.4	0.8	0.8	143	1.2	e
5	Chimica umida	4	75.0	25.0	0.0	134	5.0	e*
6	Spotchem EL-SE 1520	115	98.3	1.7	0.0	144	1.8	e
7	Piccolo	23	100.0	0.0	0.0	133	2.0	e
8	Abx Mira	6	100.0	0.0	0.0	142	1.1	e
9	iStat Chem8	6	100.0	0.0	0.0	139	1.0	e

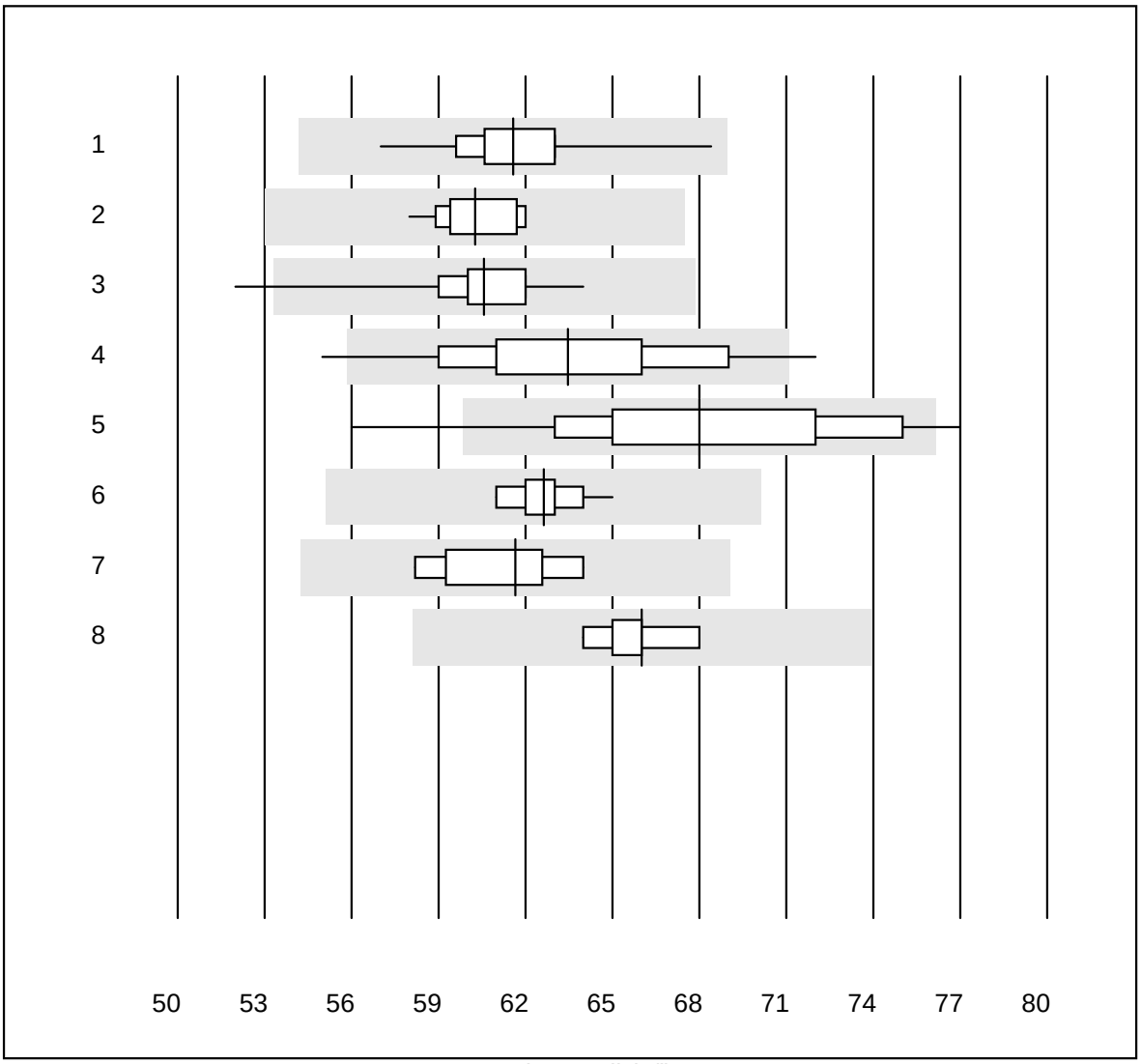
Fosfati



Deviatione QUALAB : 15 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	15	100.0	0.0	0.0	1.3	3.7	e
2	Cobas	10	100.0	0.0	0.0	1.3	2.7	e
3	Fuji Dri-Chem	75	97.3	2.7	0.0	1.3	4.3	e
4	Spotchem D-Concept	15	100.0	0.0	0.0	1.3	4.1	e
5	Spotchem/Ready	8	100.0	0.0	0.0	1.1	4.0	e
6	Piccolo	6	100.0	0.0	0.0	1.5	4.9	e*
7	Abx Mira	5	100.0	0.0	0.0	1.5	4.6	e*

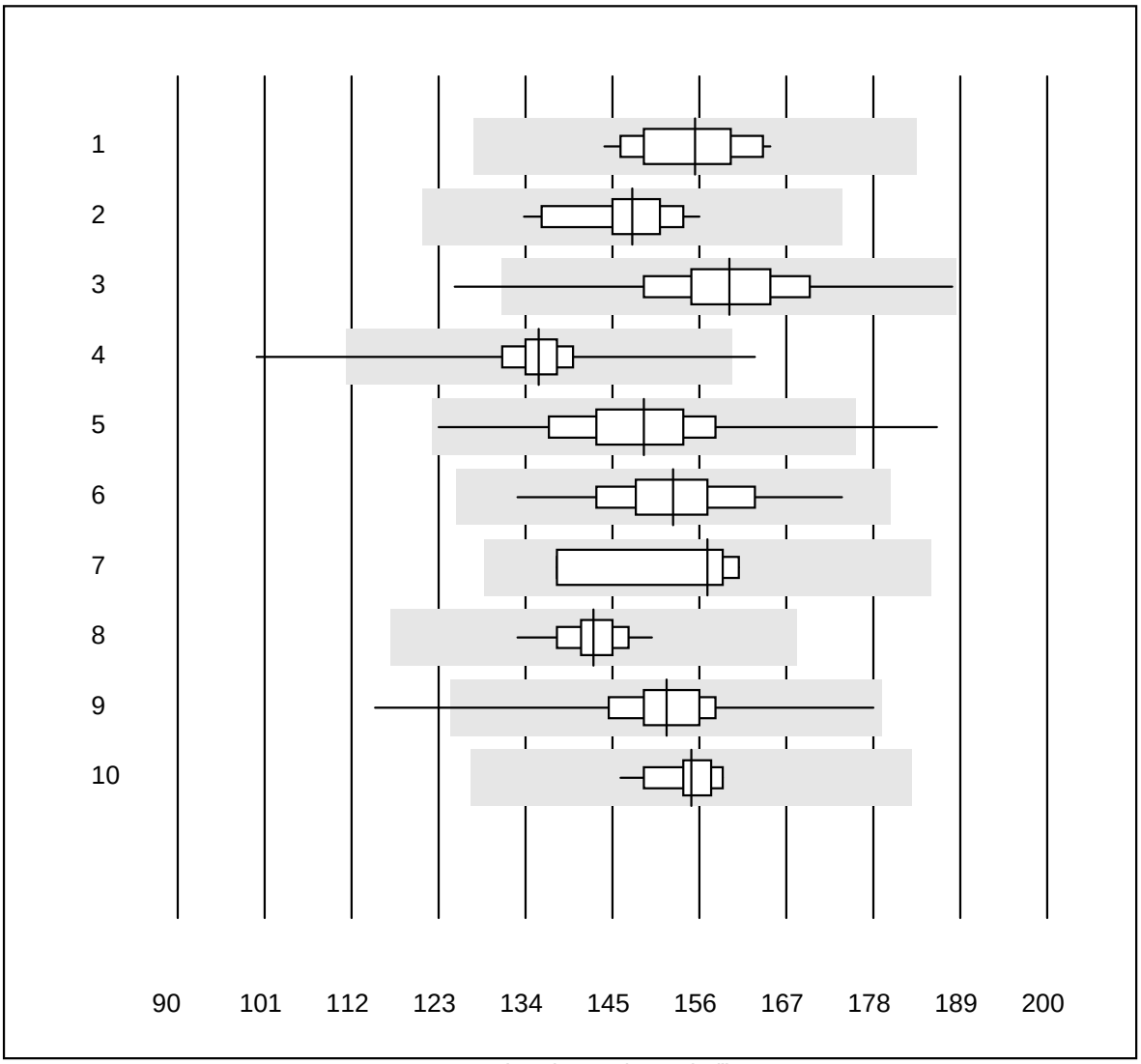
Proteine totali



Deviatione QUALAB : 12 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	21	100.0	0.0	0.0	61.6	3.6	e
2	Cobas	11	100.0	0.0	0.0	60.3	2.1	e
3	Fuji Dri-Chem	188	98.4	0.5	1.1	60.6	2.6	e
4	Spotchem/Ready	40	92.5	7.5	0.0	63.5	6.4	e
5	Spotchem D-Concept	64	92.2	4.7	3.1	68.0	6.8	e
6	Piccolo	24	100.0	0.0	0.0	62.6	1.5	e
7	Abx Mira	8	100.0	0.0	0.0	61.7	3.2	e
8	Hitachi S40/M40	5	100.0	0.0	0.0	66.0	2.3	e

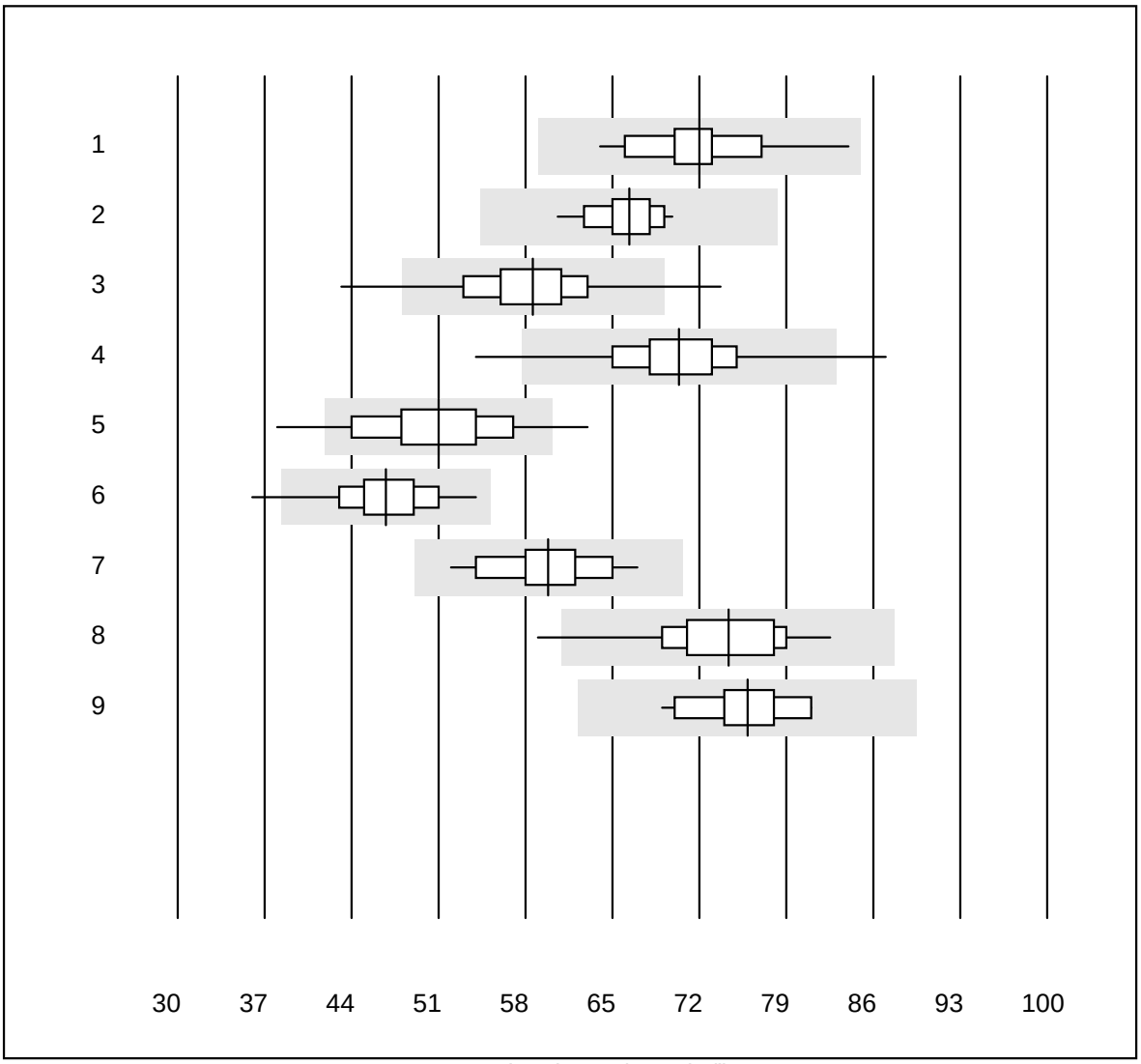
Transaminasi GOT/AST



Deviatione QUALAB : 18 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	IFCC con Pyridox 37'	18	100.0	0.0	0.0	155	4.3	e
2	Cobas	16	100.0	0.0	0.0	147	4.1	e
3	Reflotron	890	98.0	0.4	1.6	160	5.6	e
4	Fuji Dri-Chem	721	99.3	0.6	0.1	136	3.4	e
5	Spotchem/Ready	154	98.7	1.3	0.0	149	6.3	e
6	Spotchem D-Concept	143	100.0	0.0	0.0	153	4.9	e
7	IFCC senza Pyridox 3	4	100.0	0.0	0.0	157	6.8	e*
8	Piccolo	33	100.0	0.0	0.0	143	2.6	e
9	Abx Mira	21	95.2	4.8	0.0	152	7.8	e
10	Hitachi S40/M40	17	100.0	0.0	0.0	155	2.2	e

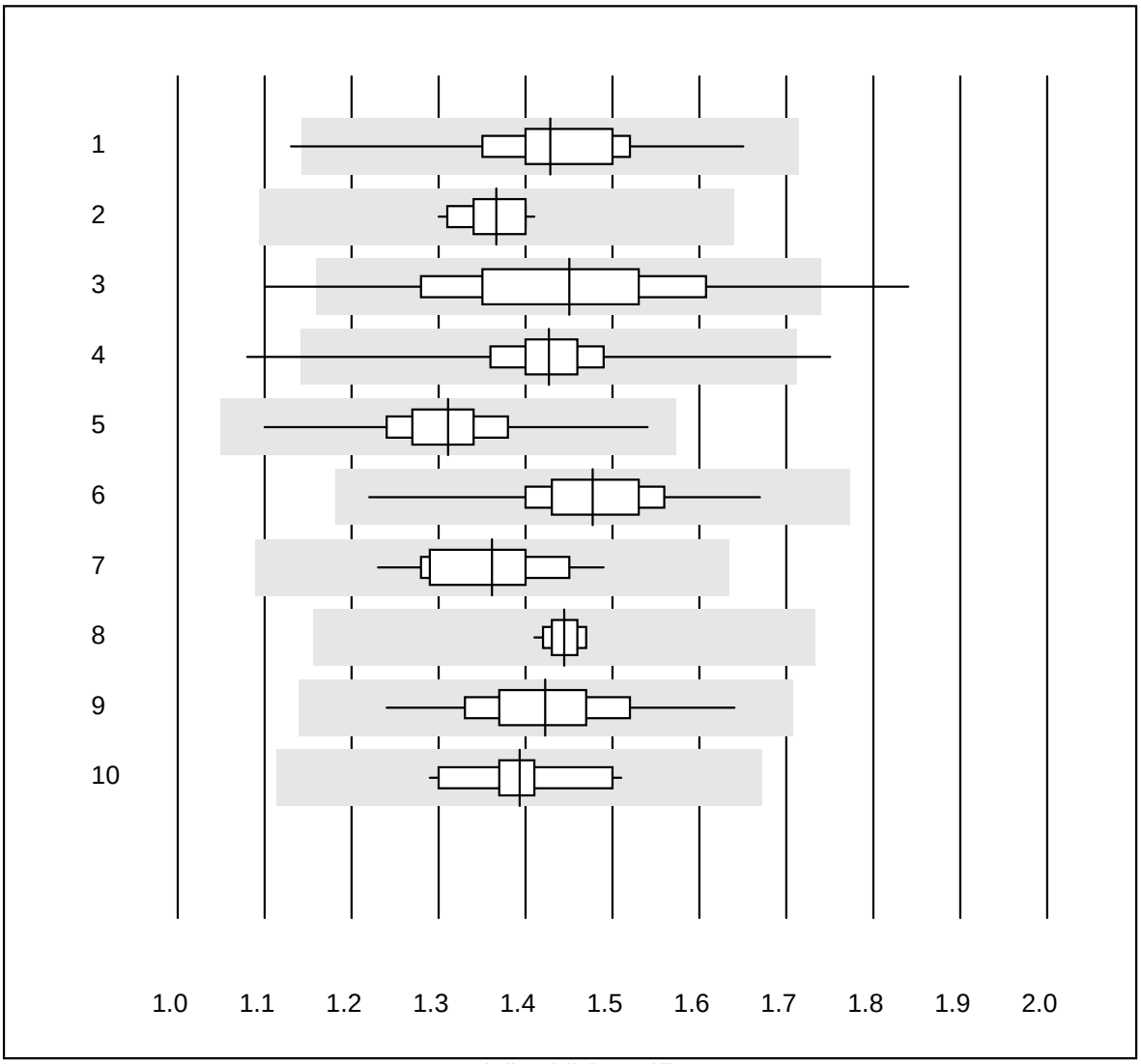
Transaminasi GPT/ALT



Deviatione QUALAB : 18 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	IFCC con Pyridox 37'	19	94.7	0.0	5.3	72	6.0	e
2	Cobas	17	100.0	0.0	0.0	66	3.5	e
3	Reflotron	927	97.2	1.3	1.5	59	6.9	e
4	Fuji Dri-Chem	737	98.5	0.8	0.7	70	5.8	e
5	Spotchem/Ready	157	96.8	1.9	1.3	51	9.1	e
6	Spotchem D-Concept	148	96.6	2.7	0.7	47	7.0	e
7	Piccolo	34	100.0	0.0	0.0	60	6.2	e
8	Abx Mira	21	95.2	4.8	0.0	74	7.0	e
9	Hitachi S40/M40	16	100.0	0.0	0.0	76	4.9	e

Trigliceridi

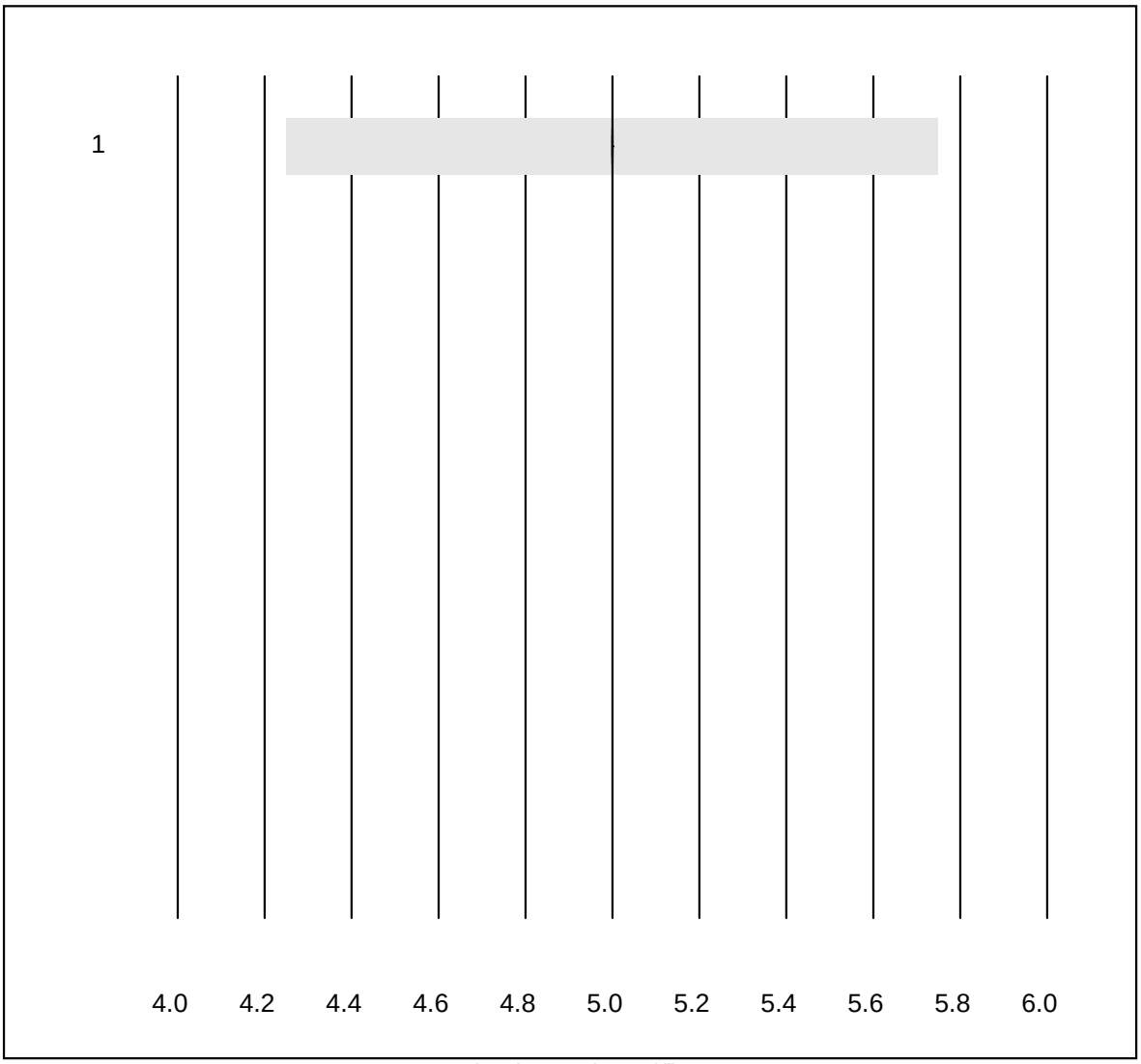


Deviatione QUALAB : 20 %

Trigliceridi (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	22	95.5	4.5	0.0	1.43	7.3	e
2	Cobas	16	100.0	0.0	0.0	1.37	2.7	e
3	Reflotron	670	95.1	2.5	2.4	1.45	9.1	e
4	Fuji Dri-Chem	661	98.6	0.6	0.8	1.43	4.2	e
5	Spotchem/Ready	136	100.0	0.0	0.0	1.31	5.0	e
6	Spotchem D-Concept	133	99.2	0.0	0.8	1.48	4.7	e
7	Hitachi S40/M40	12	100.0	0.0	0.0	1.36	5.7	e
8	Piccolo	19	100.0	0.0	0.0	1.44	1.2	e
9	Cholestech LDX	187	99.5	0.0	0.5	1.42	5.1	e
10	Abx Mira	19	100.0	0.0	0.0	1.39	4.1	e

Bicarbonat

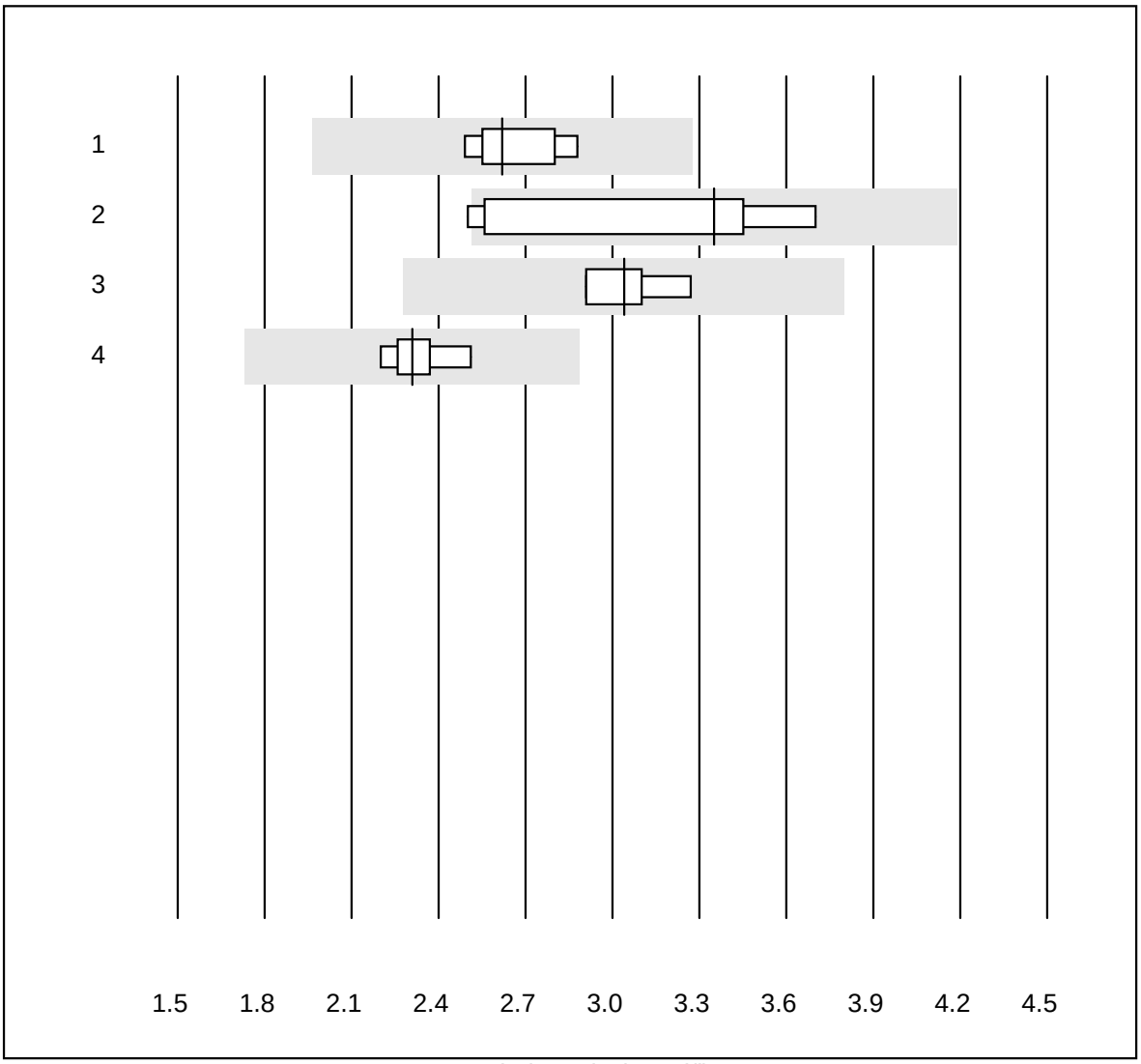


Deviatione QUALAB : 15 %

Bicarbonat (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Piccolo	5	100.0	0.0	0.0	5	0.0	e

LDL Cholesterin

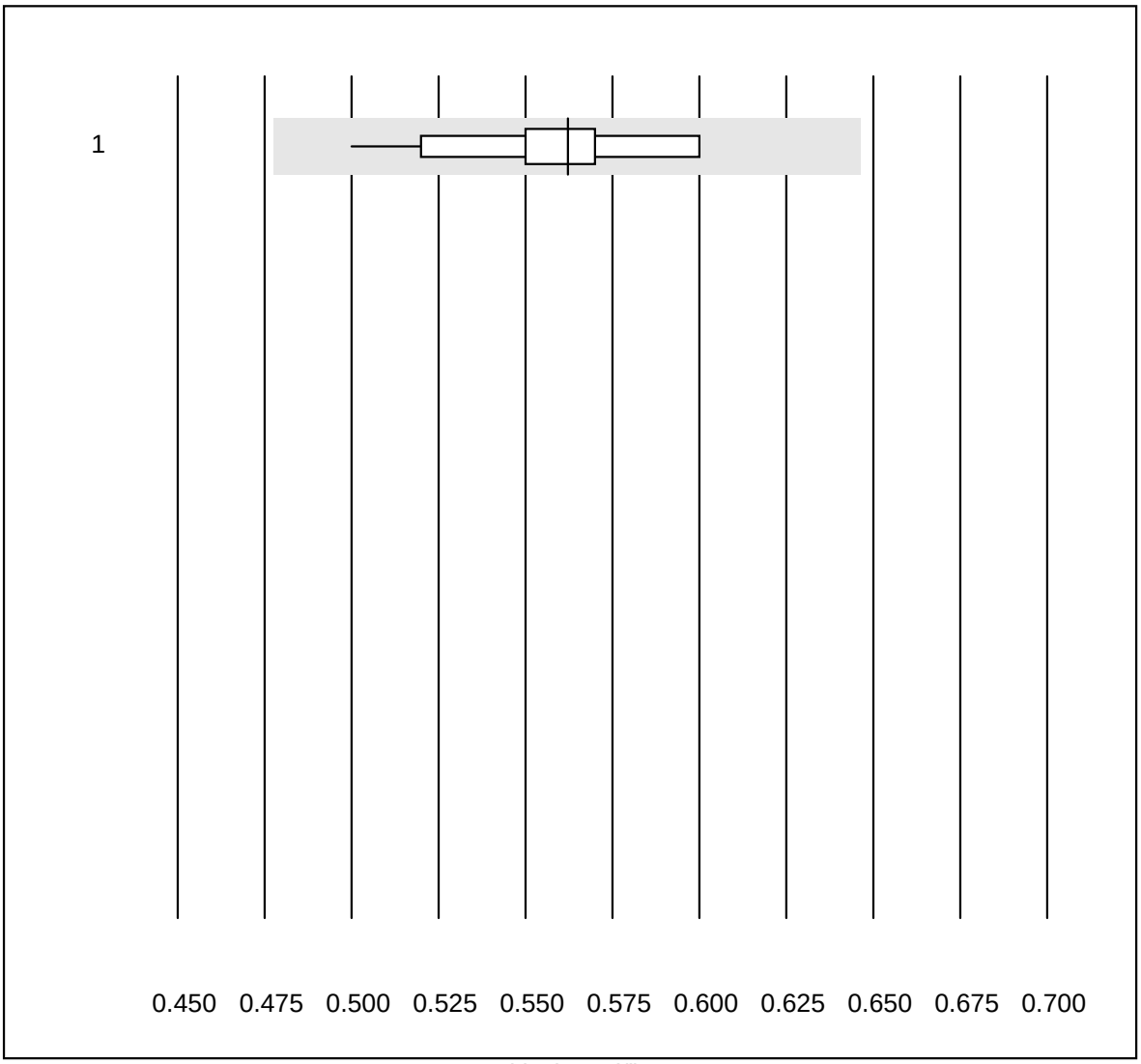


Deviatione QUALAB : 25 %

LDL Cholesterin (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Abx Mira	9	100.0	0.0	0.0	2.6	5.8	e
2	Chimica umida	8	87.5	12.5	0.0	3.4	14.2	e*
3	Roche, Cobas	4	100.0	0.0	0.0	3.0	5.1	e
4	Hitachi S40/M40	5	100.0	0.0	0.0	2.3	5.1	e

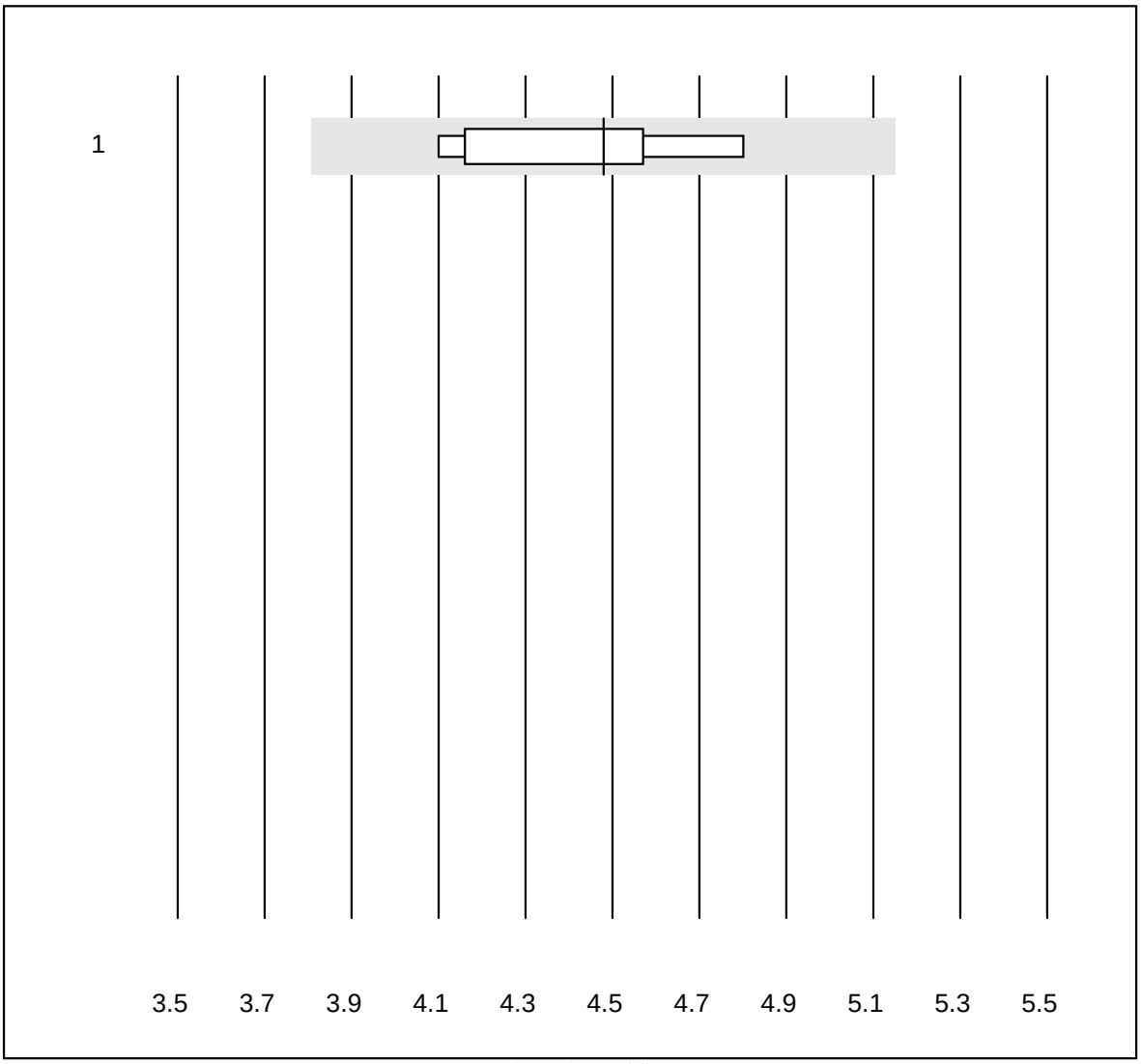
Litio



Deviatione QUALAB : 15 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	tutti	14	100.0	0.0	0.0	0.56	4.9	e

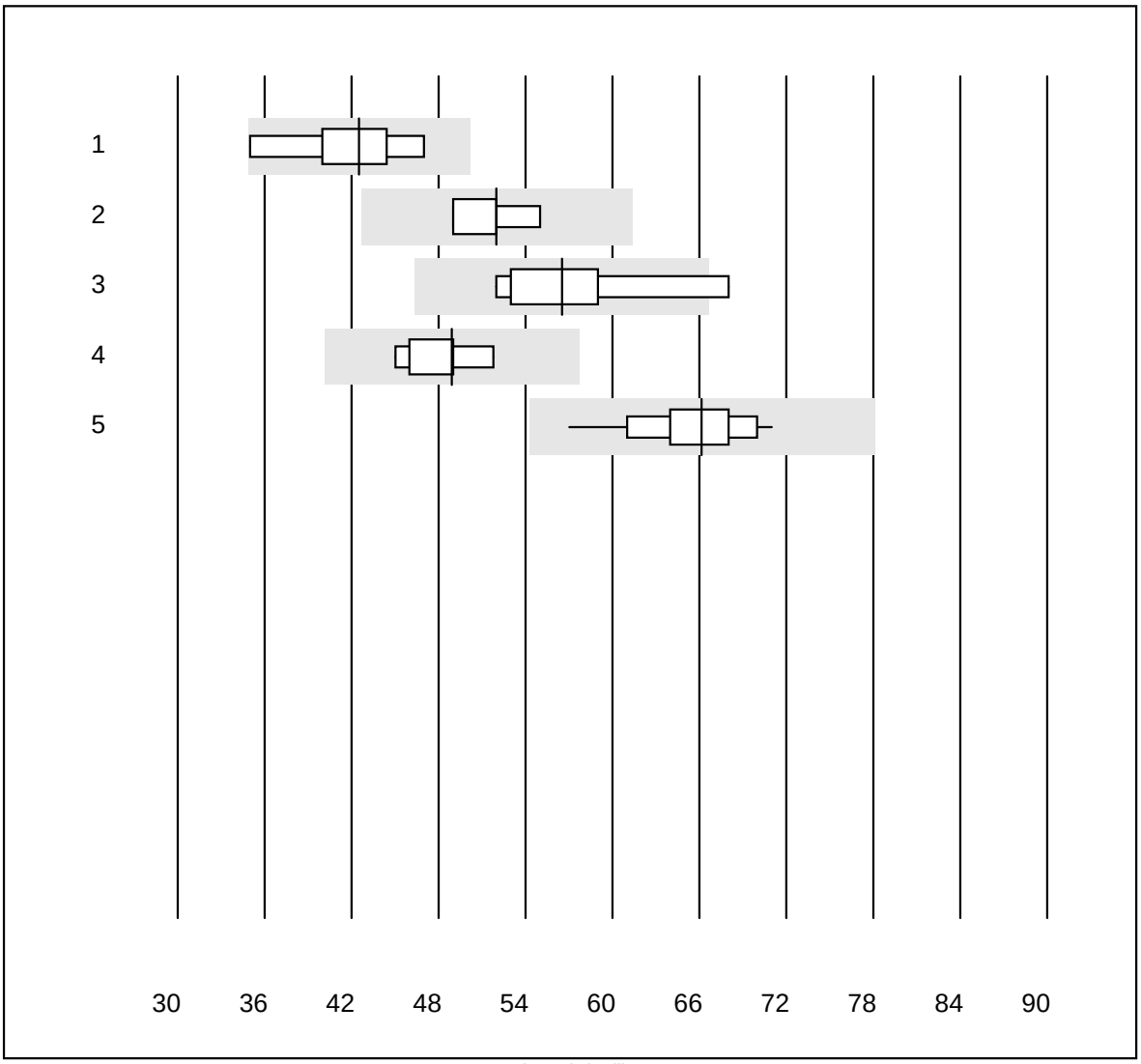
Laktat



Deviatione QUALAB : 15 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	tutti	7	100.0	0.0	0.0	4.48	5.7	e*

Lipasi

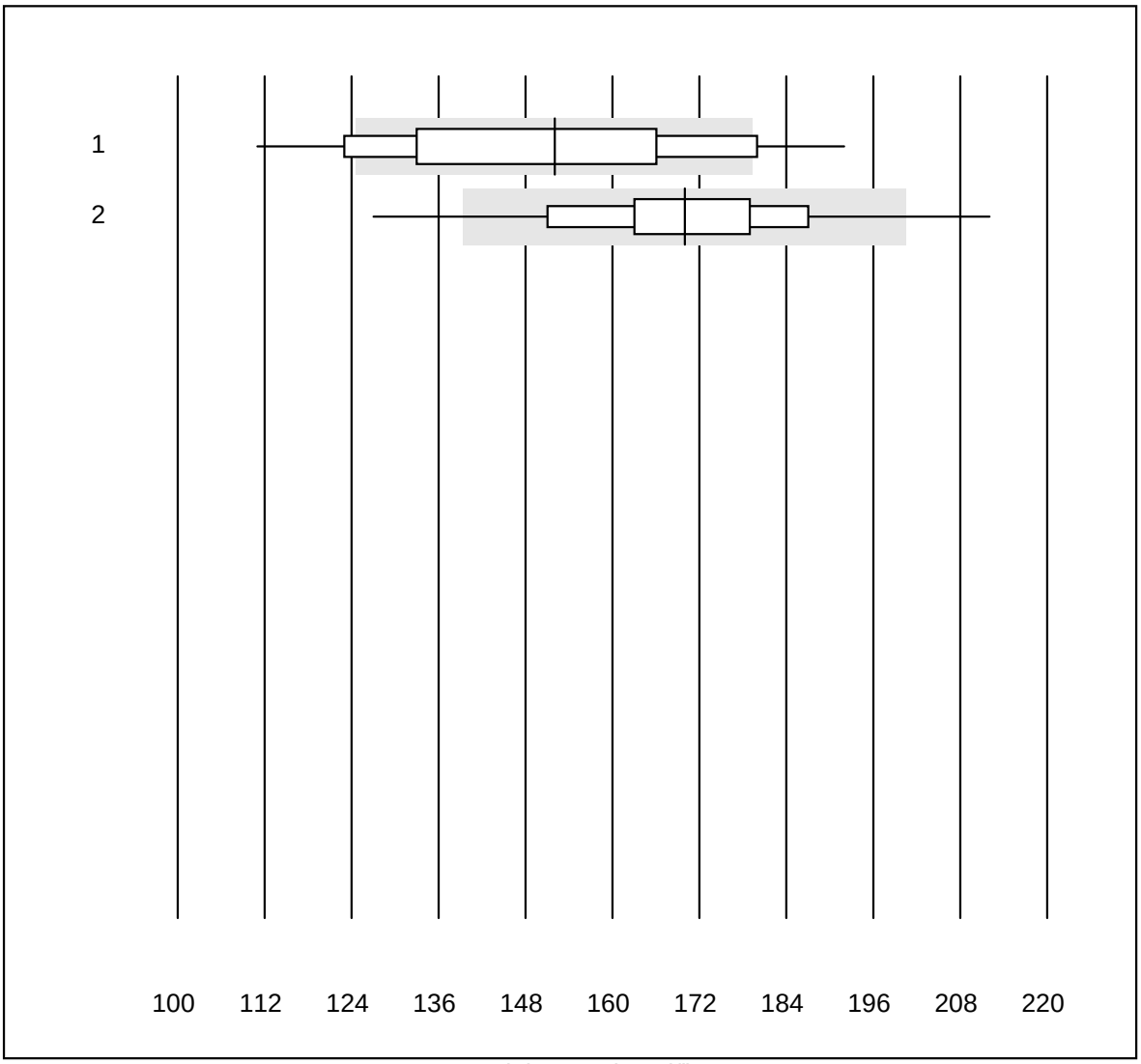


Deviatione QUALAB : 18 %

Lipasi (U/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Abx Mira	6	100.0	0.0	0.0	42.5	9.8	e*
2	Architect	4	100.0	0.0	0.0	52.0	4.7	e*
3	Beckman/Olympus	6	83.3	16.7	0.0	56.5	10.1	e*
4	Cobas	8	100.0	0.0	0.0	48.9	4.4	e
5	Fuji Dri-Chem	63	96.8	0.0	3.2	66.1	4.9	e

Creatinina SP

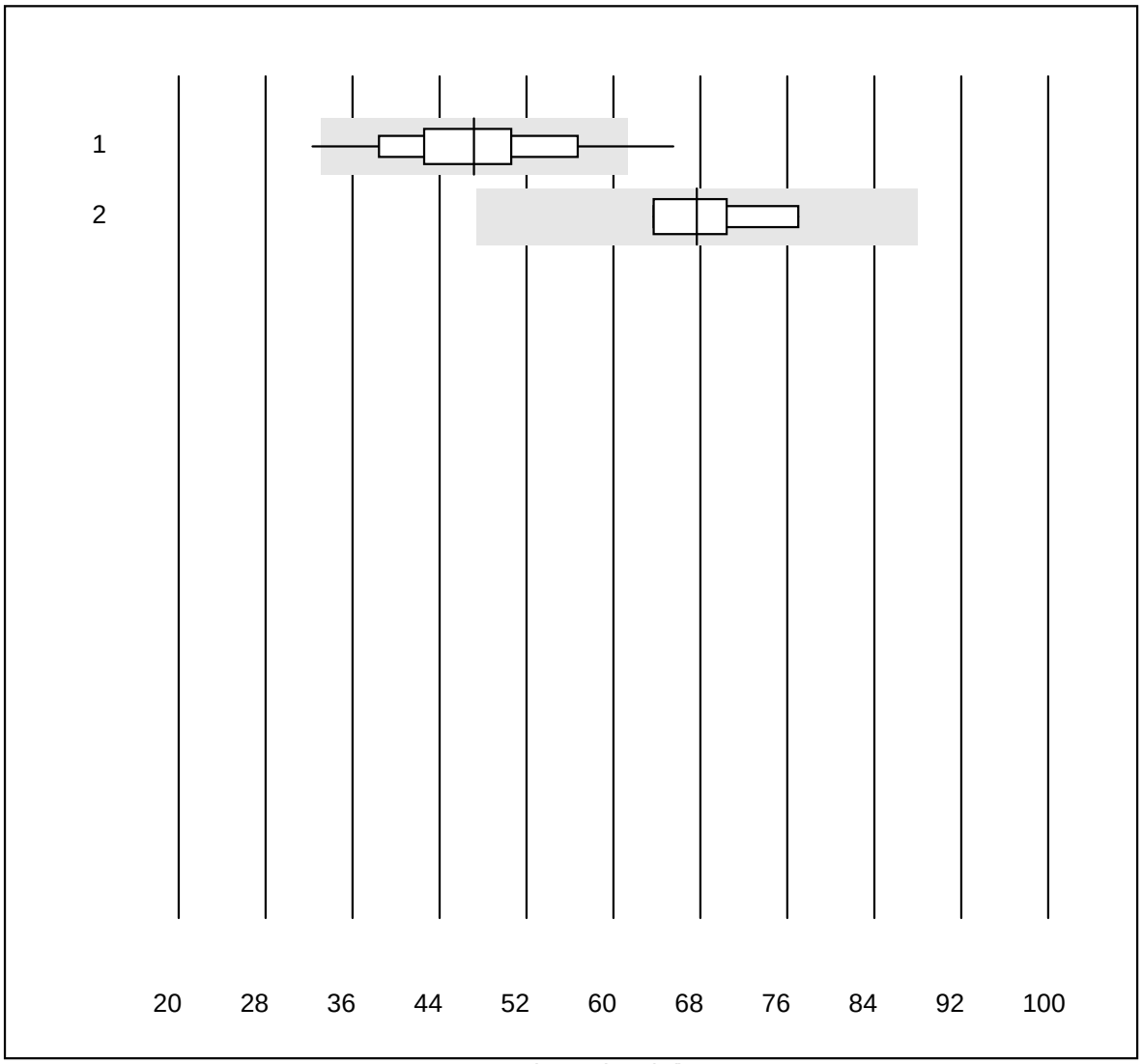


Deviatione QUALAB : 18 %

Creatinina SP (µmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Spotchem/Ready	156	65.4	19.2	15.4	152	13.7	e
2	Spotchem D-Concept	142	93.0	7.0	0.0	170	8.7	e

eGFR (Spotchem)

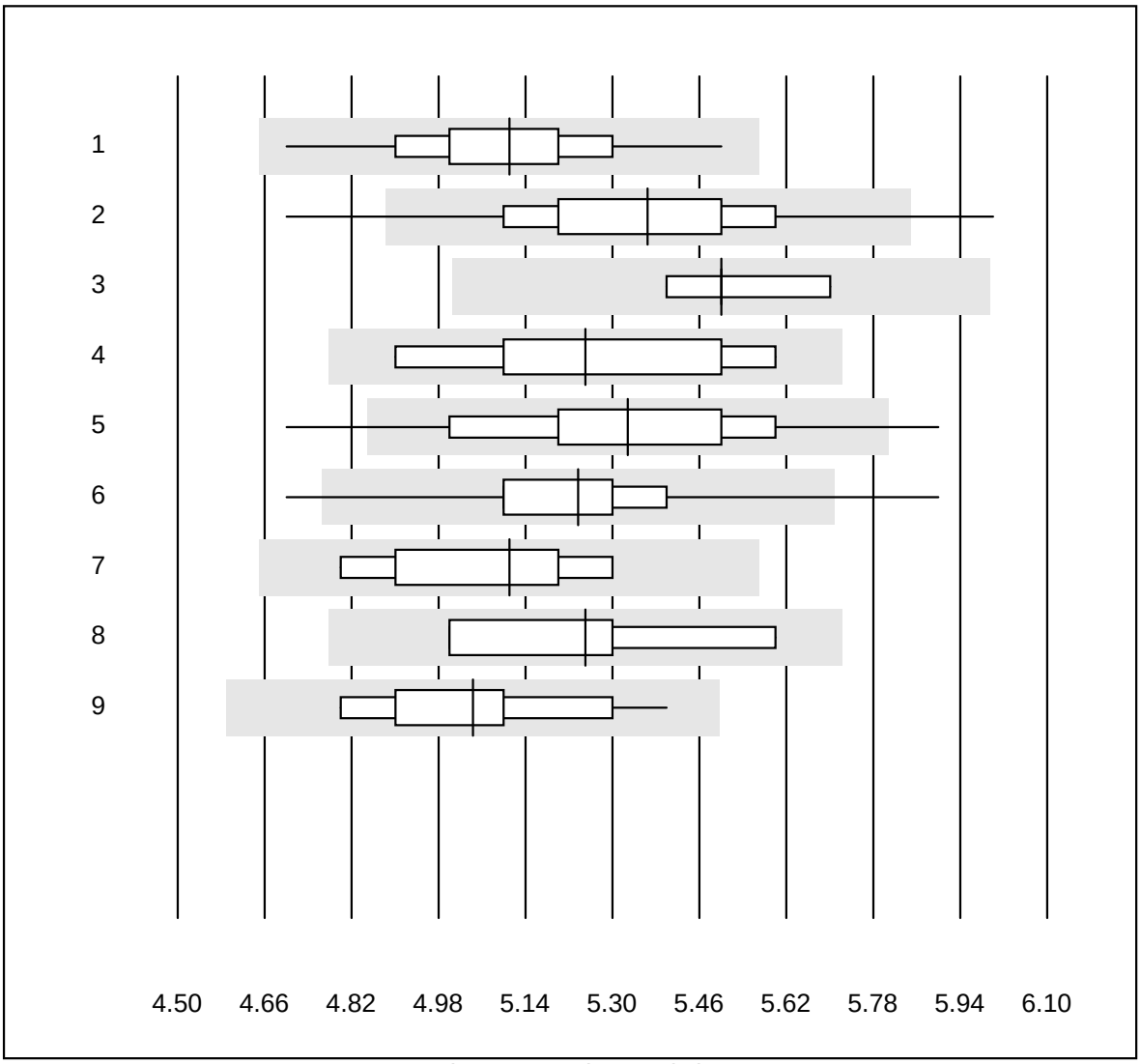


Deviatione QUALAB : 30 %

eGFR (Spotchem) ()

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	CKD-EPI	95	81.1	6.3	12.6	47	15.3	e
2	Cockcroft-Gault	5	80.0	0.0	20.0	68	8.0	e

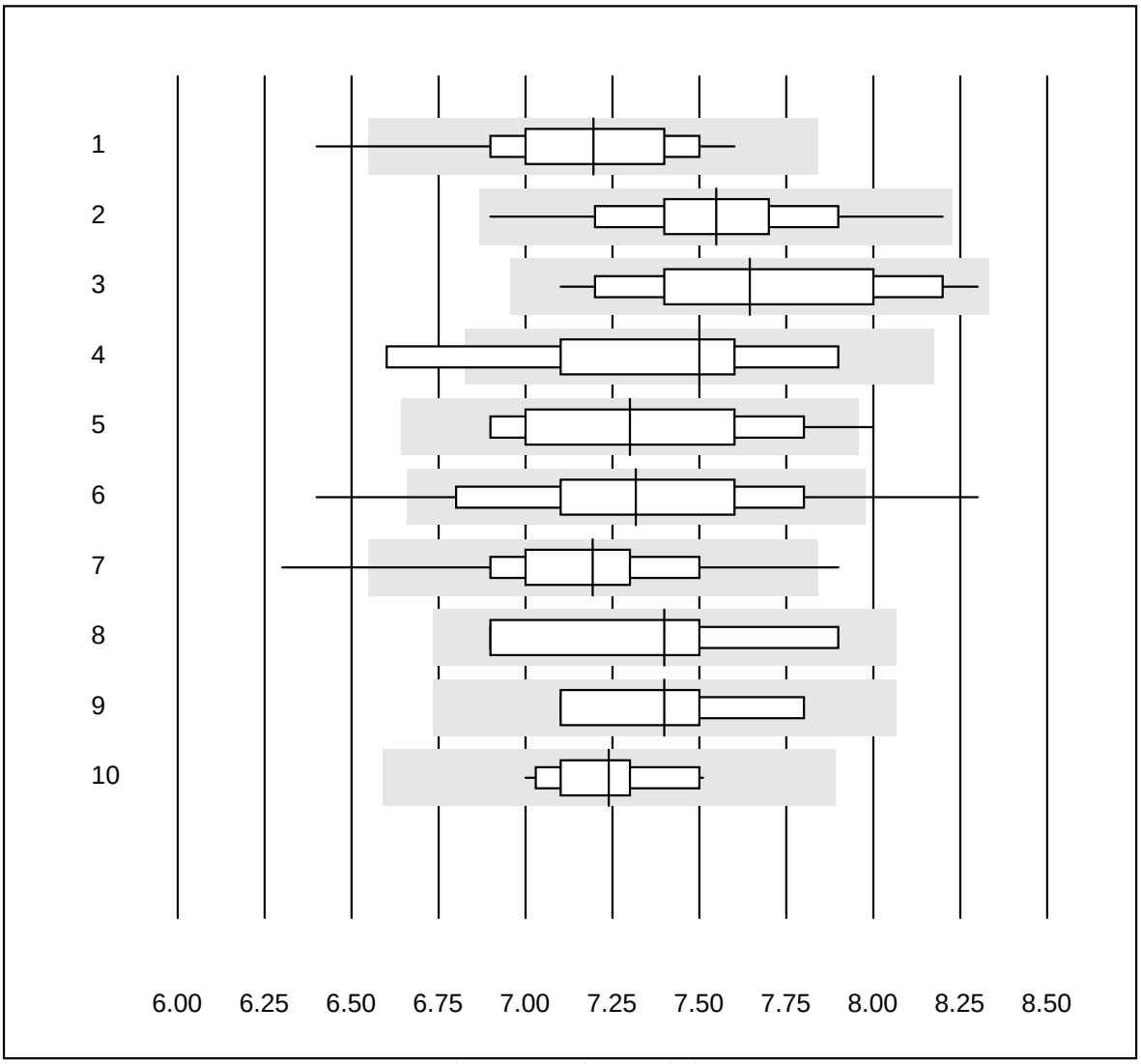
HbA1c campione A



Deviatione QUALAB : 9 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas b101	31	100.0	0.0	0.0	5.1	3.3	e
2	Afinion	596	98.0	1.5	0.5	5.4	3.7	e
3	Eurolyser	9	100.0	0.0	0.0	5.5	1.7	e
4	Hemocue HbA1c 501	8	87.5	0.0	12.5	5.3	4.5	e*
5	NycoCard	95	84.2	5.3	10.5	5.3	4.9	e
6	DCA2000/Vantage	226	97.8	1.8	0.4	5.2	3.4	e
7	Andere	8	100.0	0.0	0.0	5.1	3.4	e*
8	HPLC	6	100.0	0.0	0.0	5.3	4.3	e*
9	Roche, Cobas	16	93.7	0.0	6.3	5.0	3.6	e

HbA1c campione B

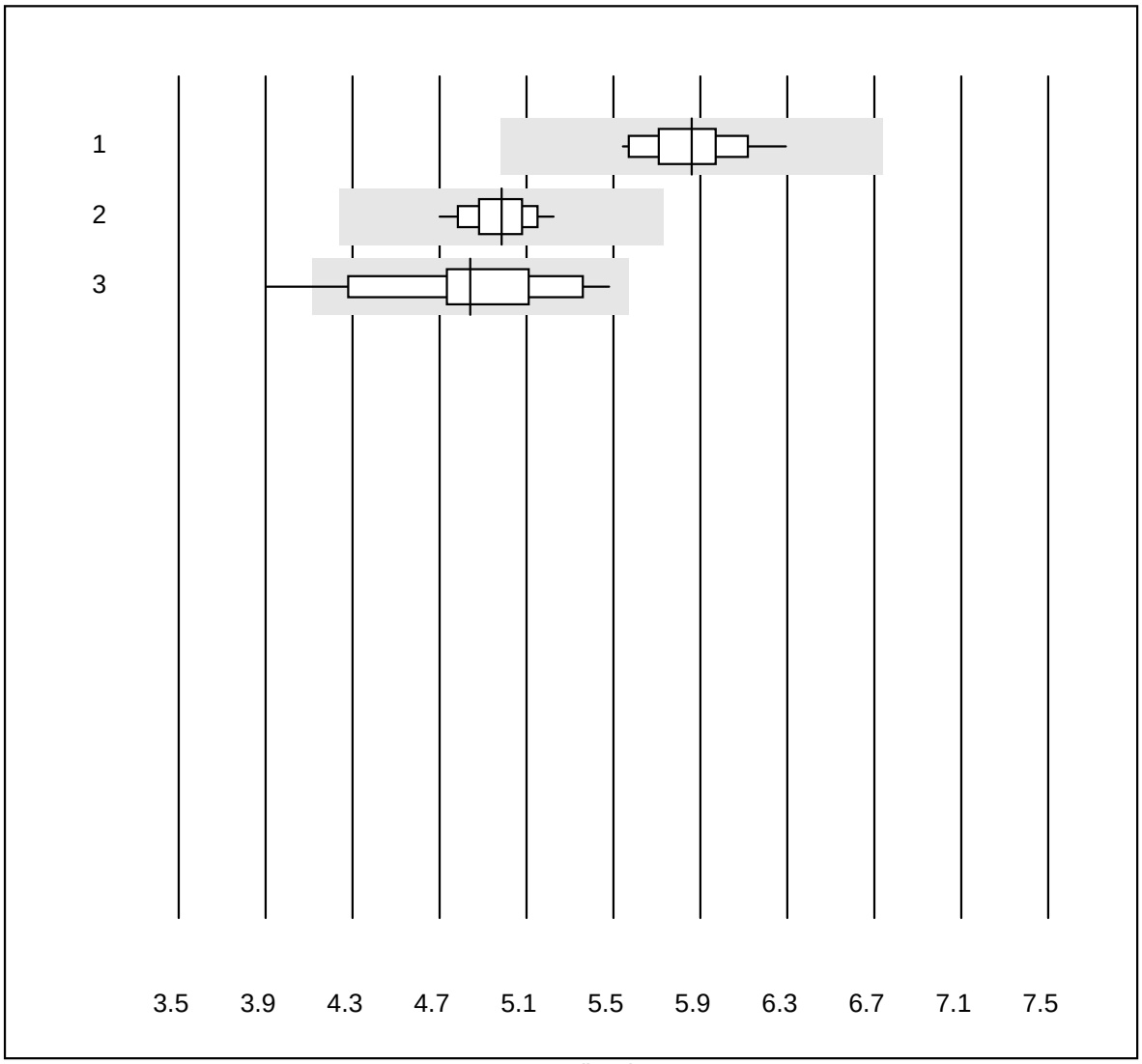


Deviatione QUALAB : 9 %

HbA1c campione B (%)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas b101	23	95.7	4.3	0.0	7.2	3.9	e
2	Afinion	599	99.8	0.0	0.2	7.5	3.3	e
3	Eurolyser	18	100.0	0.0	0.0	7.6	4.7	e*
4	A1c Now	5	80.0	20.0	0.0	7.5	6.9	e*
5	Hemocue HbA1c 501	11	90.9	9.1	0.0	7.3	5.3	e*
6	NycoCard	93	81.7	8.6	9.7	7.3	5.3	e
7	DCA2000/Vantage	194	98.0	1.0	1.0	7.2	3.2	e
8	Andere	7	100.0	0.0	0.0	7.4	4.8	e*
9	HPLC	4	100.0	0.0	0.0	7.4	4.0	e*
10	Roche, Cobas	15	100.0	0.0	0.0	7.2	2.1	e

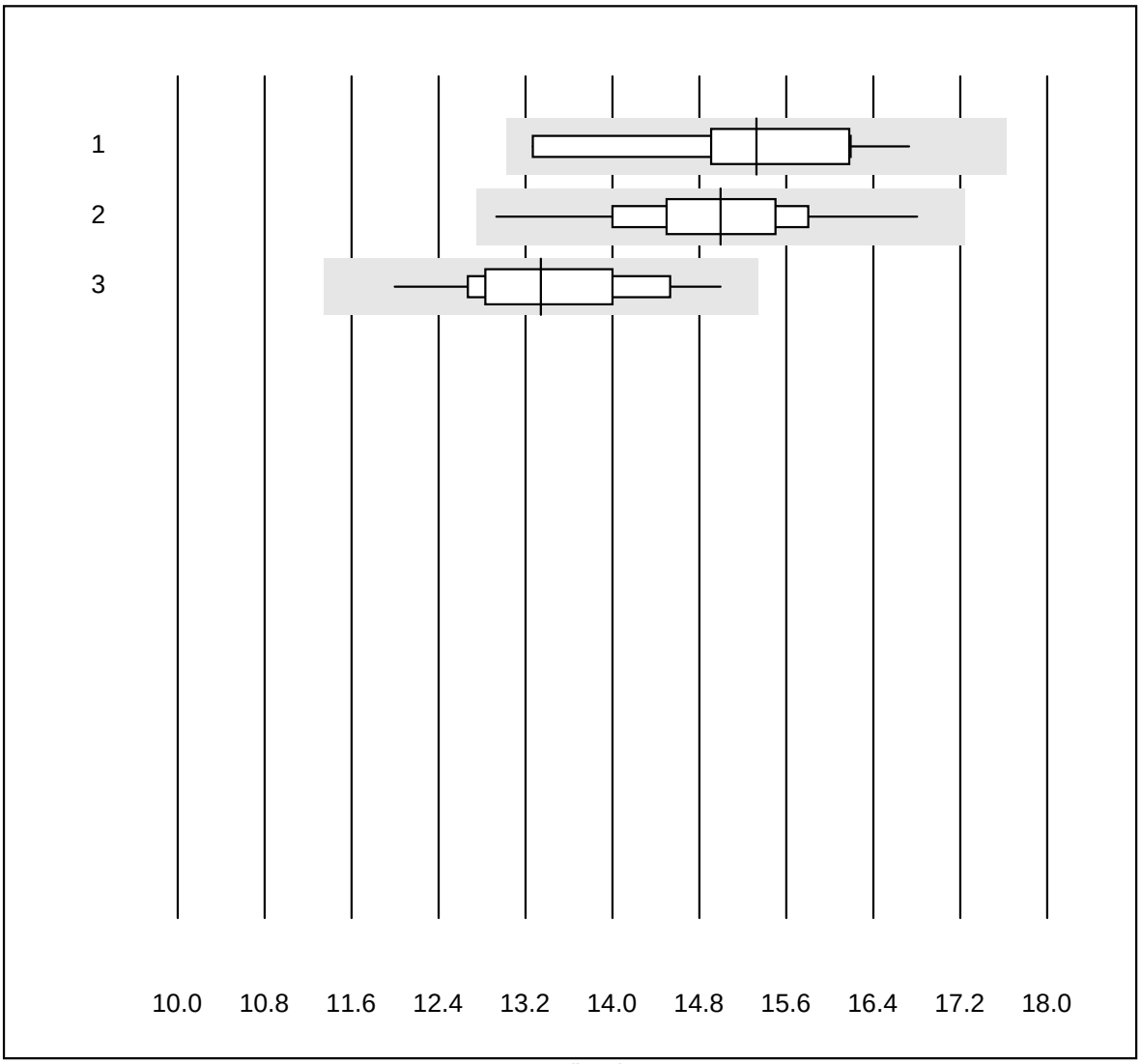
pCO2



Deviatione QUALAB : 15 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas b121/123/221	14	100.0	0.0	0.0	5.86	3.6	e
2	iStat	34	100.0	0.0	0.0	4.99	2.8	e
3	EPOC	26	92.3	7.7	0.0	4.84	8.2	e

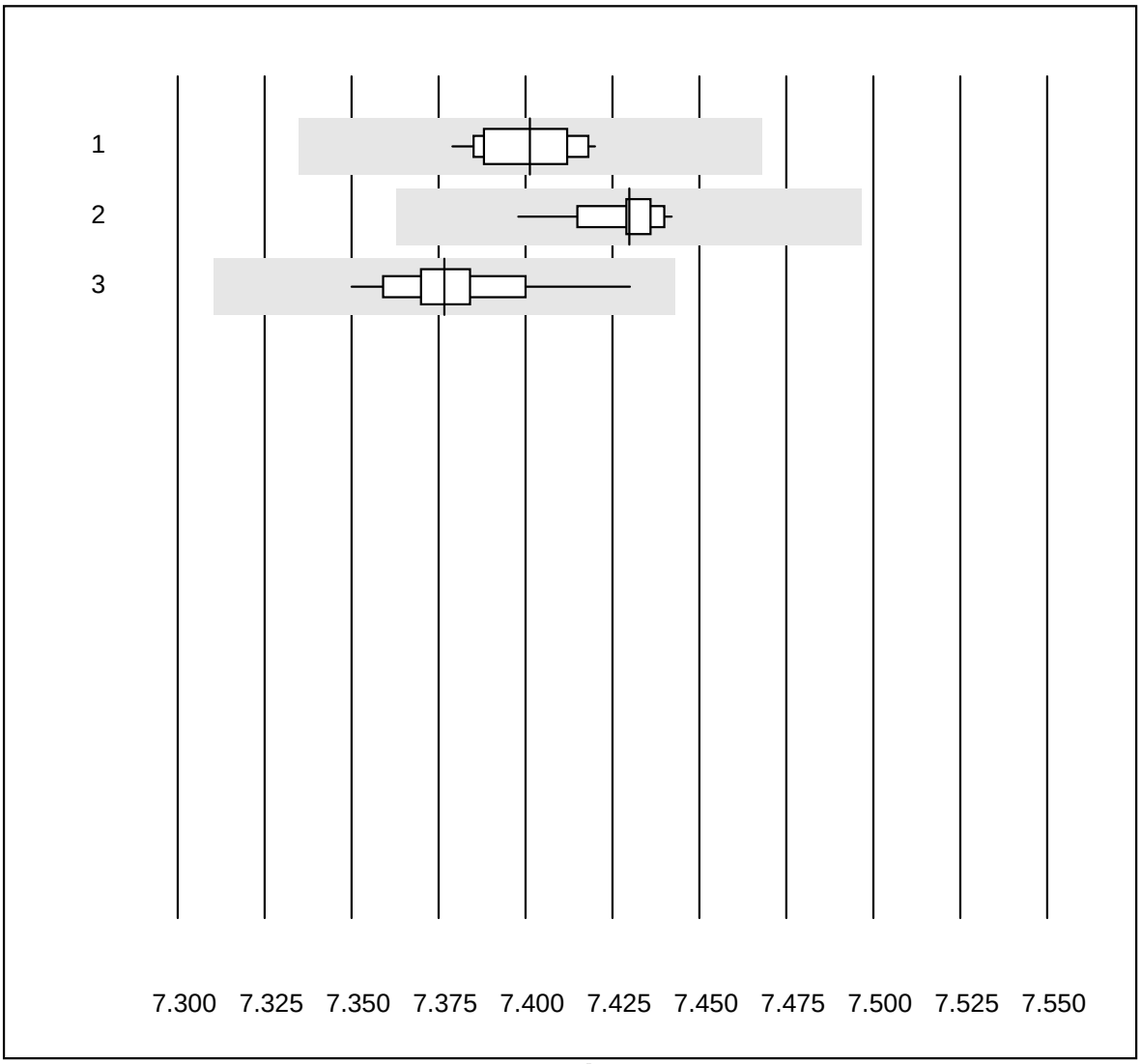
pO2



Deviatione QUALAB : 15 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas b121/123/221	11	90.9	0.0	9.1	15.32	7.2	e*
2	iStat	33	100.0	0.0	0.0	15.00	5.3	e
3	EPOC	26	84.6	0.0	15.4	13.34	5.6	e

pH

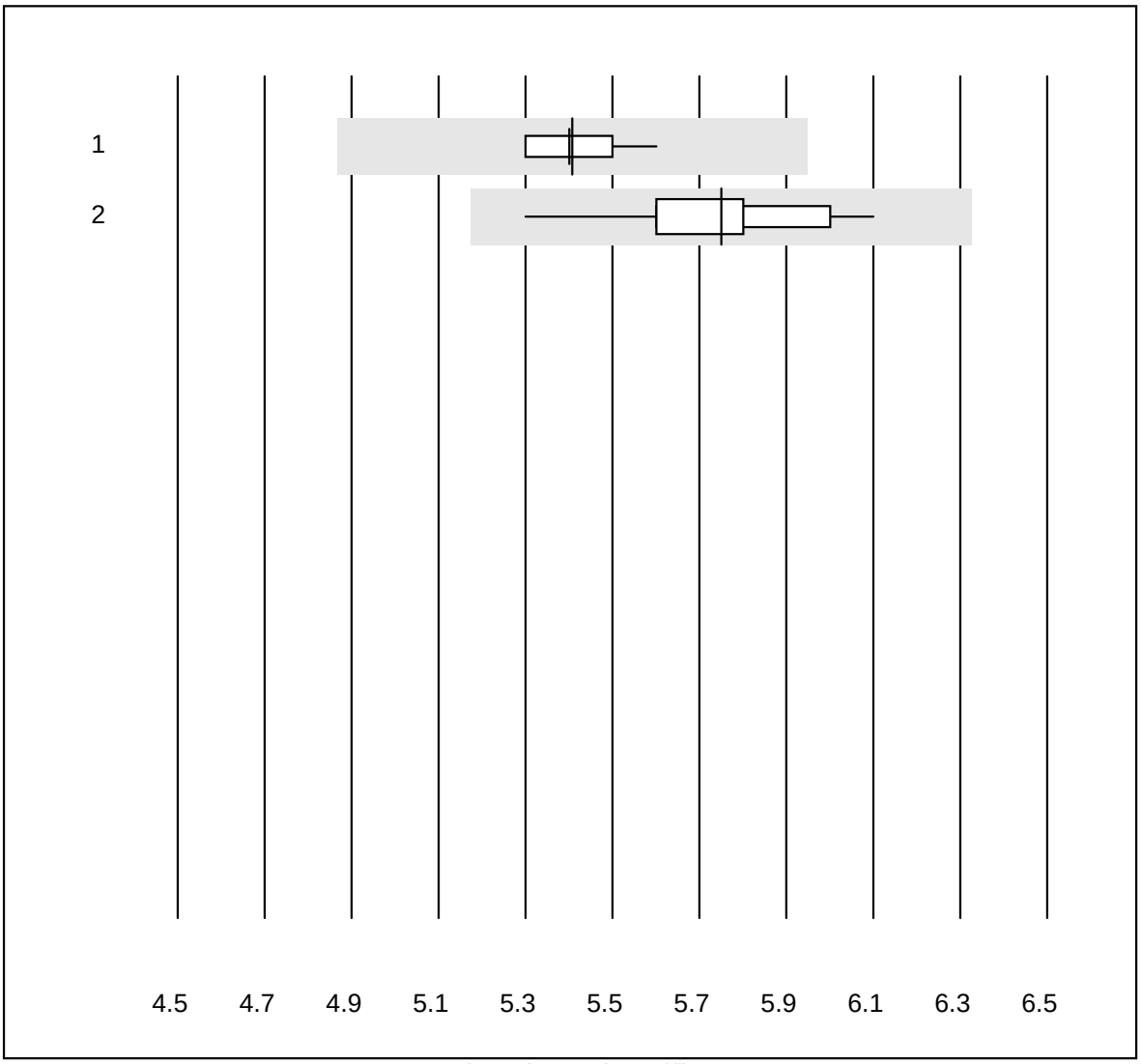


Deviatione QUALAB : 1 %

pH ()

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas b121/123/221	13	100.0	0.0	0.0	7.40	0.2	e
2	iStat	34	100.0	0.0	0.0	7.43	0.1	e
3	EPOC	26	100.0	0.0	0.0	7.38	0.2	e

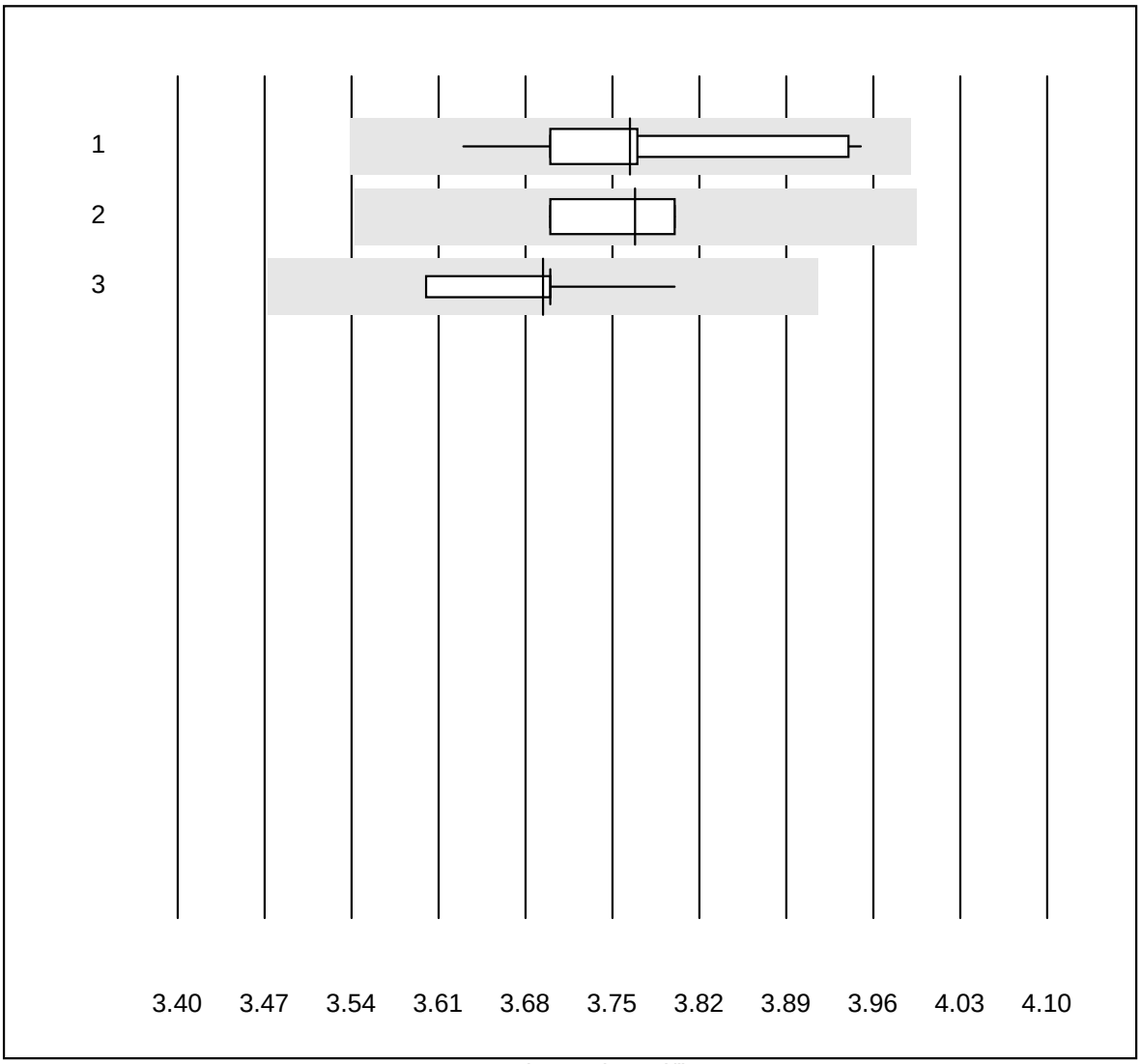
Glucosio GS



Deviatione QUALAB : 10 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	iStat	12	100.0	0.0	0.0	5.4	1.5	e
2	EPOC	18	100.0	0.0	0.0	5.7	3.1	e

Potassio BG

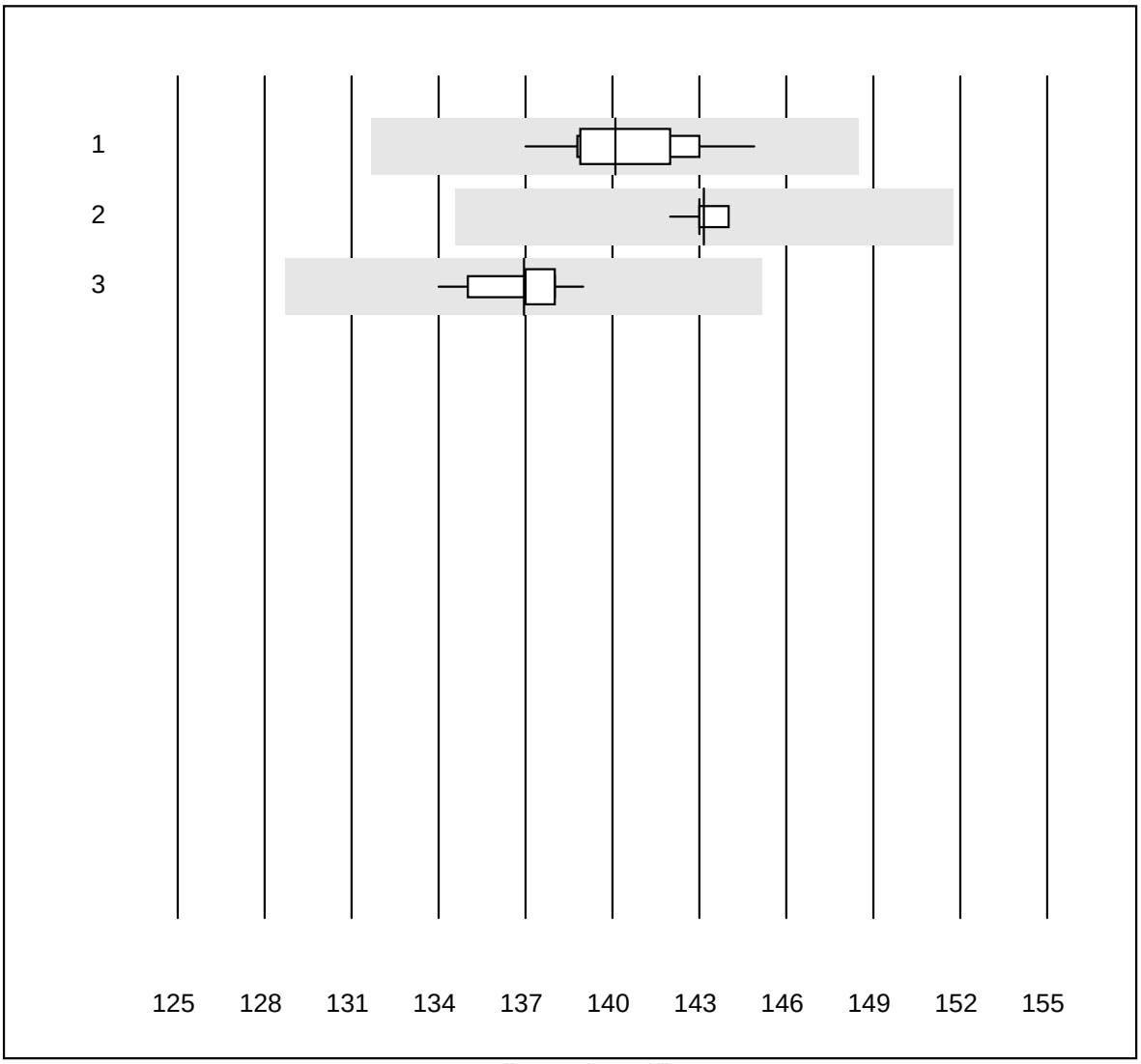


Deviatione QUALAB : 6 %

Potassio BG (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas b121/123/221	12	100.0	0.0	0.0	3.8	2.5	e
2	iStat	19	100.0	0.0	0.0	3.8	1.3	e
3	EPOC	22	100.0	0.0	0.0	3.7	1.3	e

Sodio BG

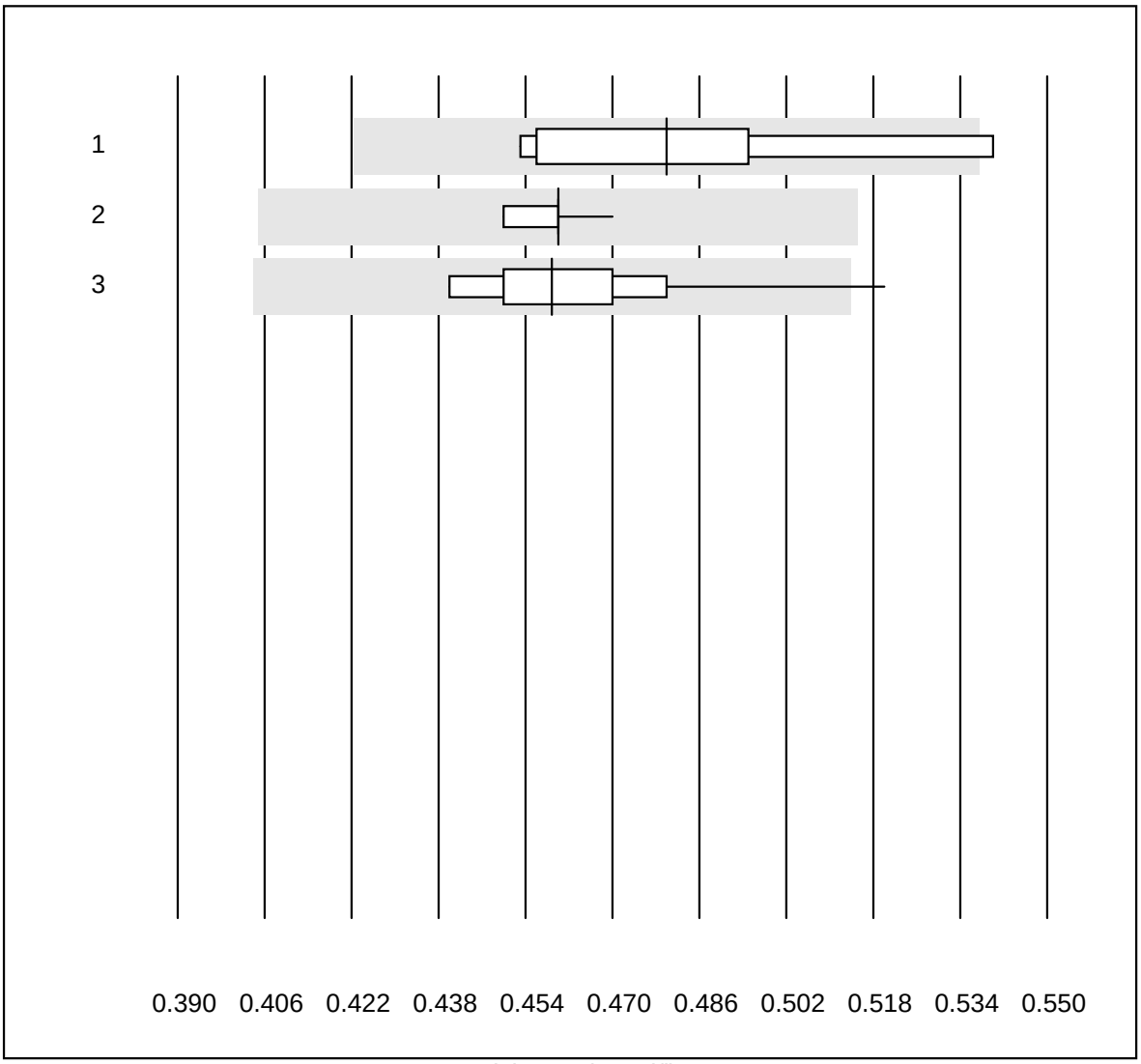


Deviatione QUALAB : 6 %

Sodio BG (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas b121/123/221	13	100.0	0.0	0.0	140.1	1.6	e
2	iStat	19	100.0	0.0	0.0	143.2	0.4	e
3	EPOC	21	100.0	0.0	0.0	136.9	1.0	e

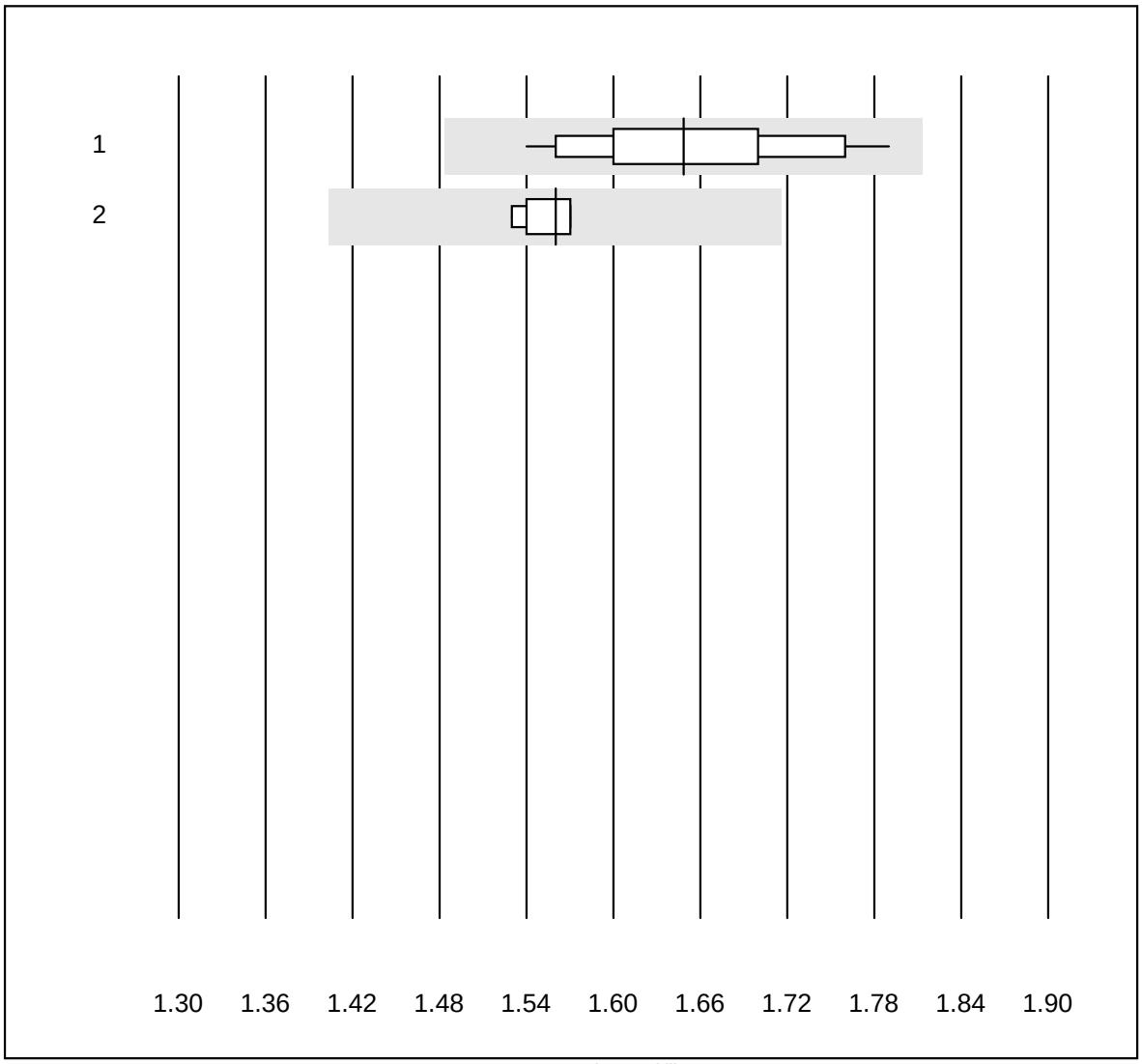
Calcio-BG



Deviatione QUALAB : 12 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas b121/123/221	8	87.5	12.5	0.0	0.48	6.1	e*
2	iStat	10	100.0	0.0	0.0	0.46	1.0	e
3	EPOC	21	95.2	4.8	0.0	0.46	4.6	e

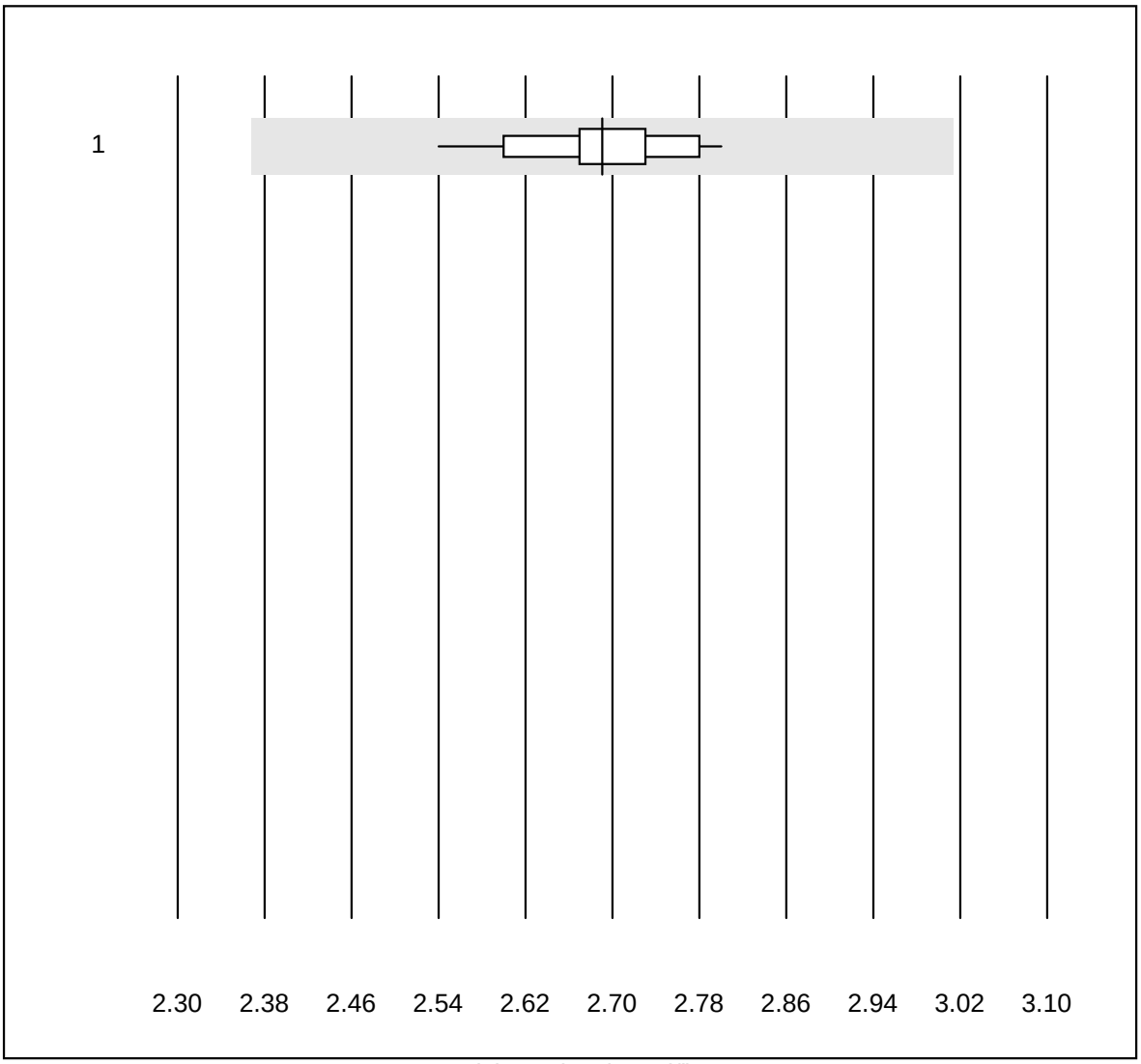
Lattato-BG



Deviatione QUALAB : 10 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	EPOC	22	100.0	0.0	0.0	1.65	4.4	e
2	iStat	7	85.7	0.0	14.3	1.56	1.1	e

Calcio - urine

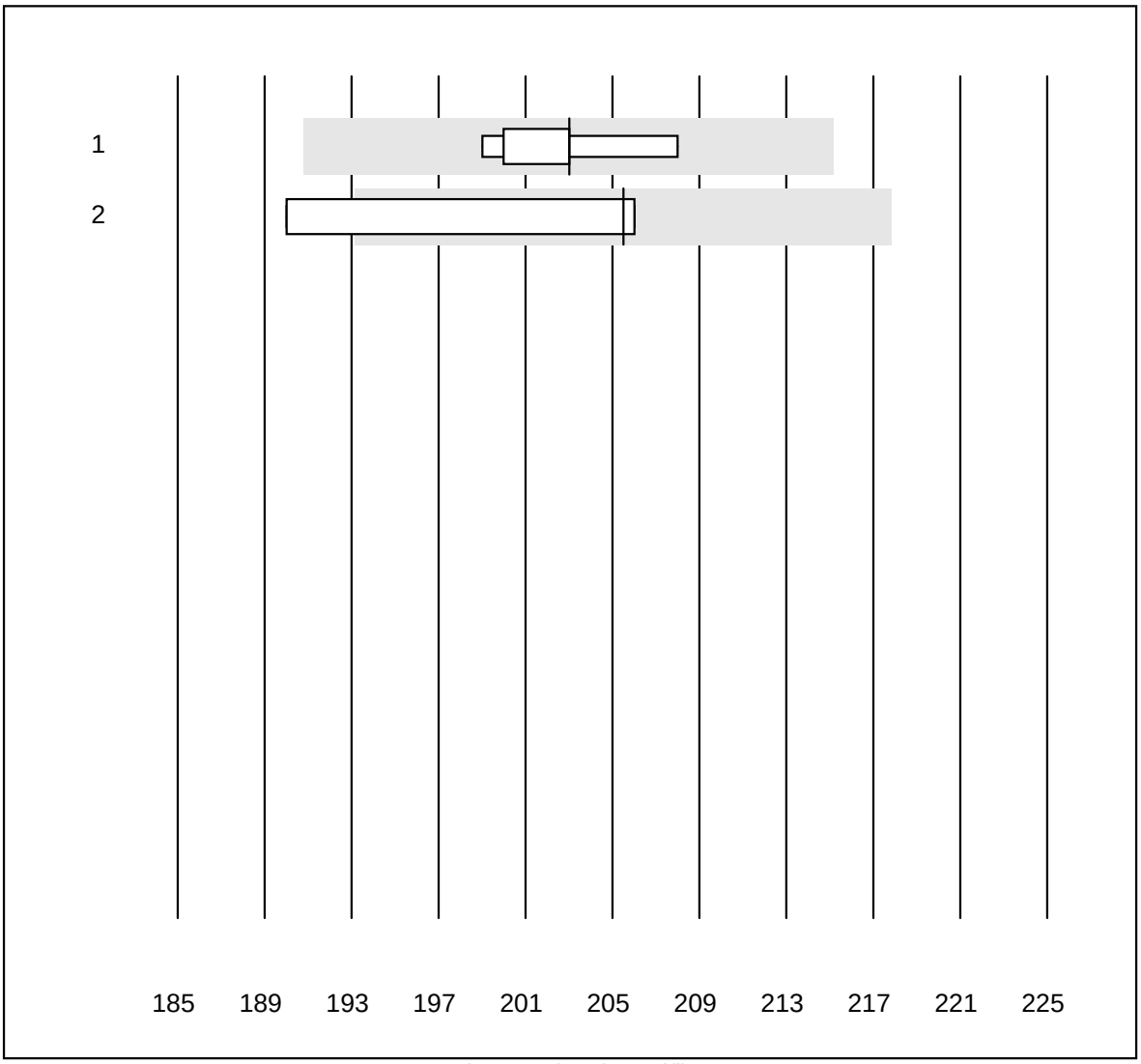


Deviatione QUALAB : 12 %

Calcio - urine (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	14	100.0	0.0	0.0	2.69	2.5	e

Cloro - urine

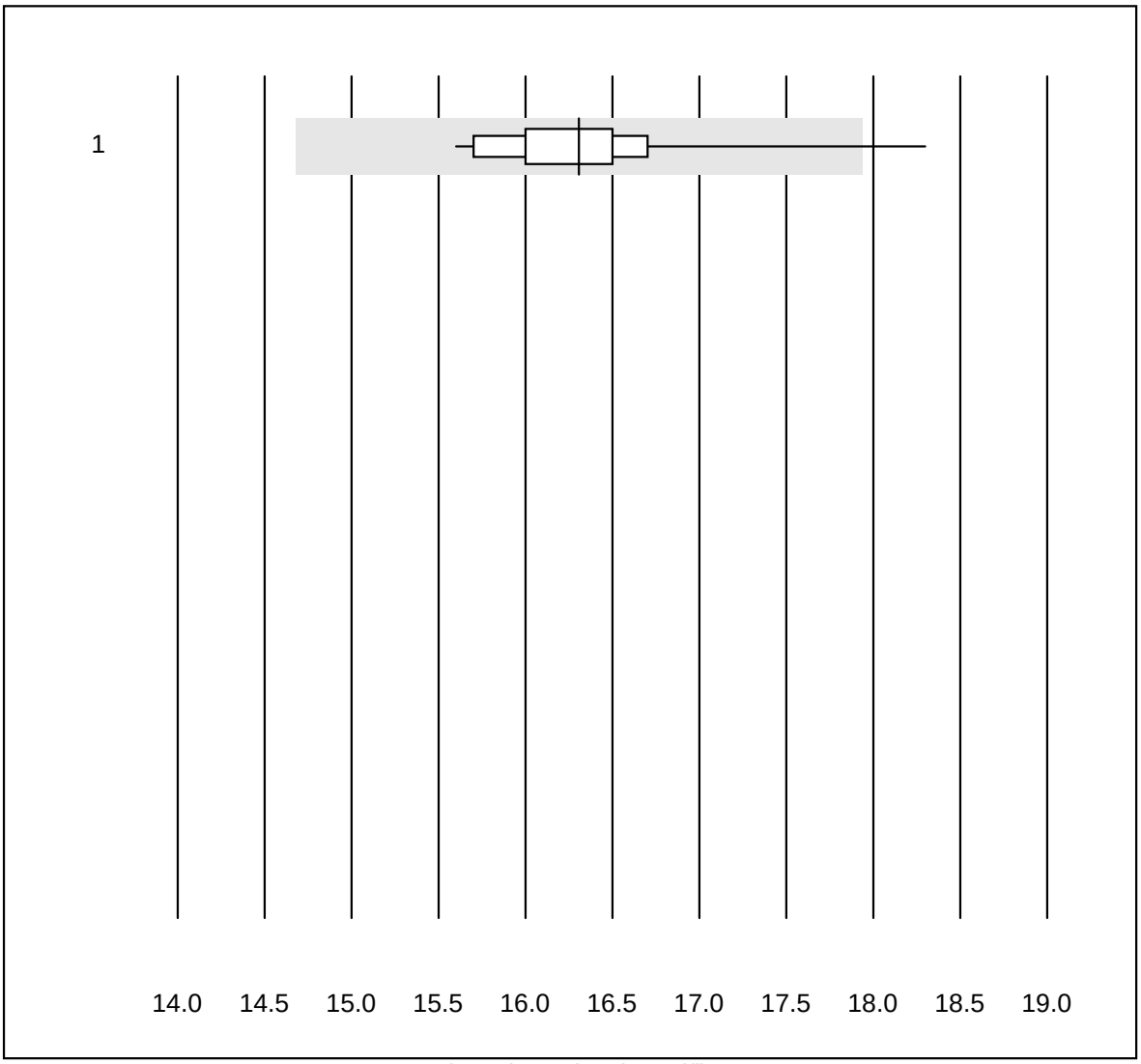


Deviatione QUALAB : 6 %

Cloro - urine (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	6	100.0	0.0	0.0	203	1.5	e
2	ISE diretto	4	75.0	25.0	0.0	206	3.9	e*

Glucosio - urine

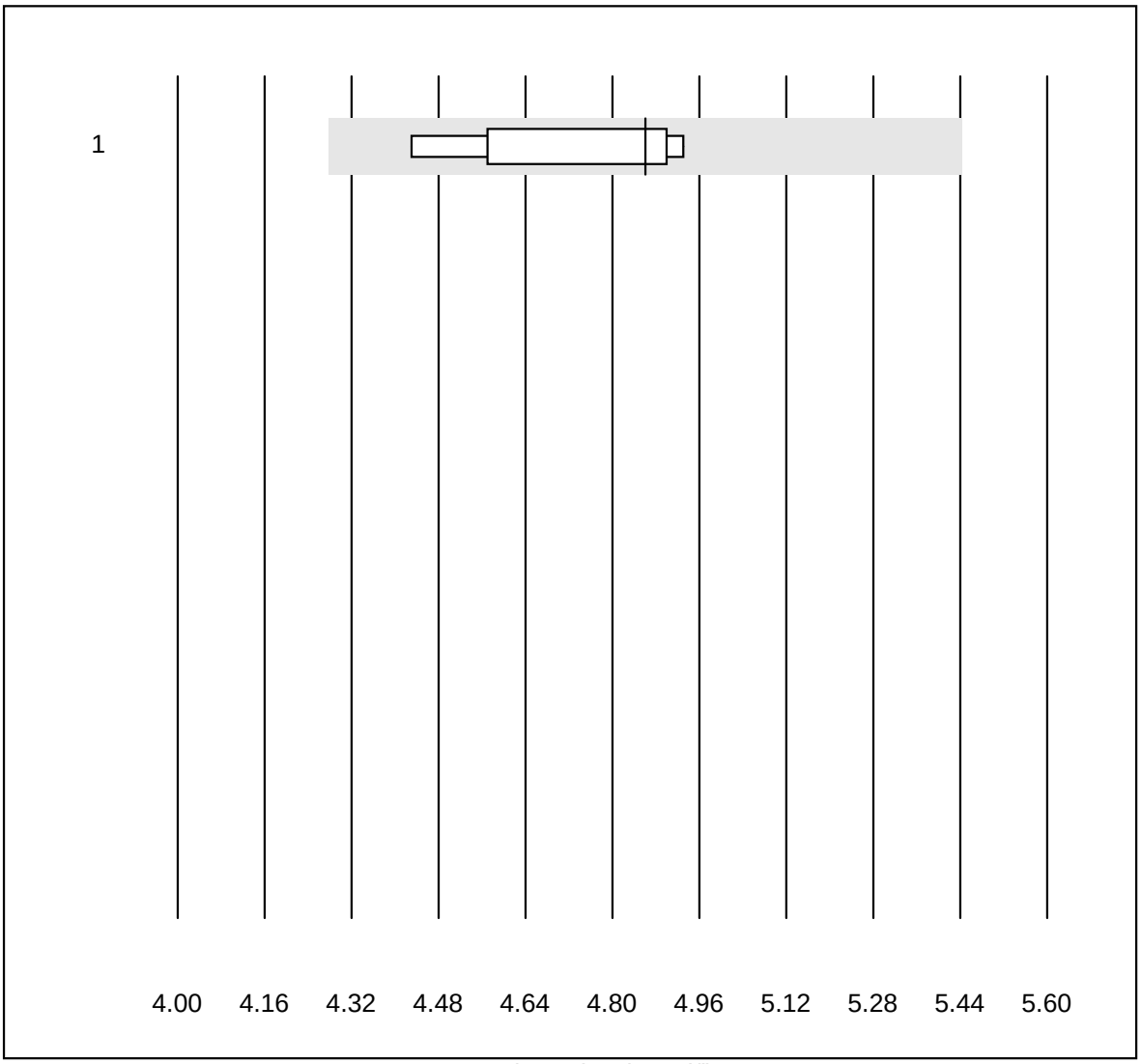


Deviatione QUALAB : 10 %

Glucosio - urine (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	13	92.3	7.7	0.0	16.3	4.2	e

Magnesio - urine

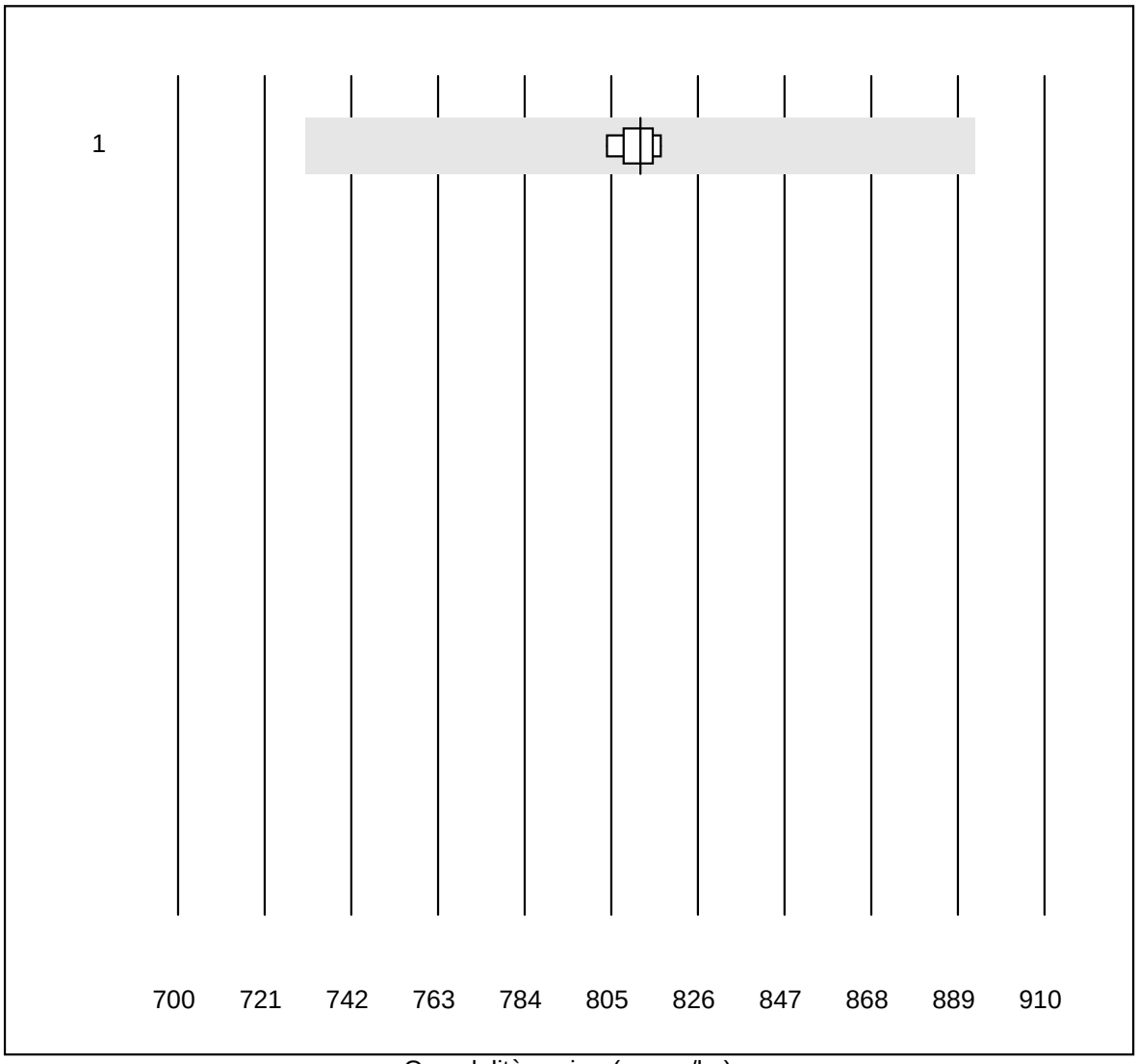


Deviatione QUALAB : 12 %

Magnesio - urine (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	9	100.0	0.0	0.0	4.9	4.1	e

Osmolalità - urine

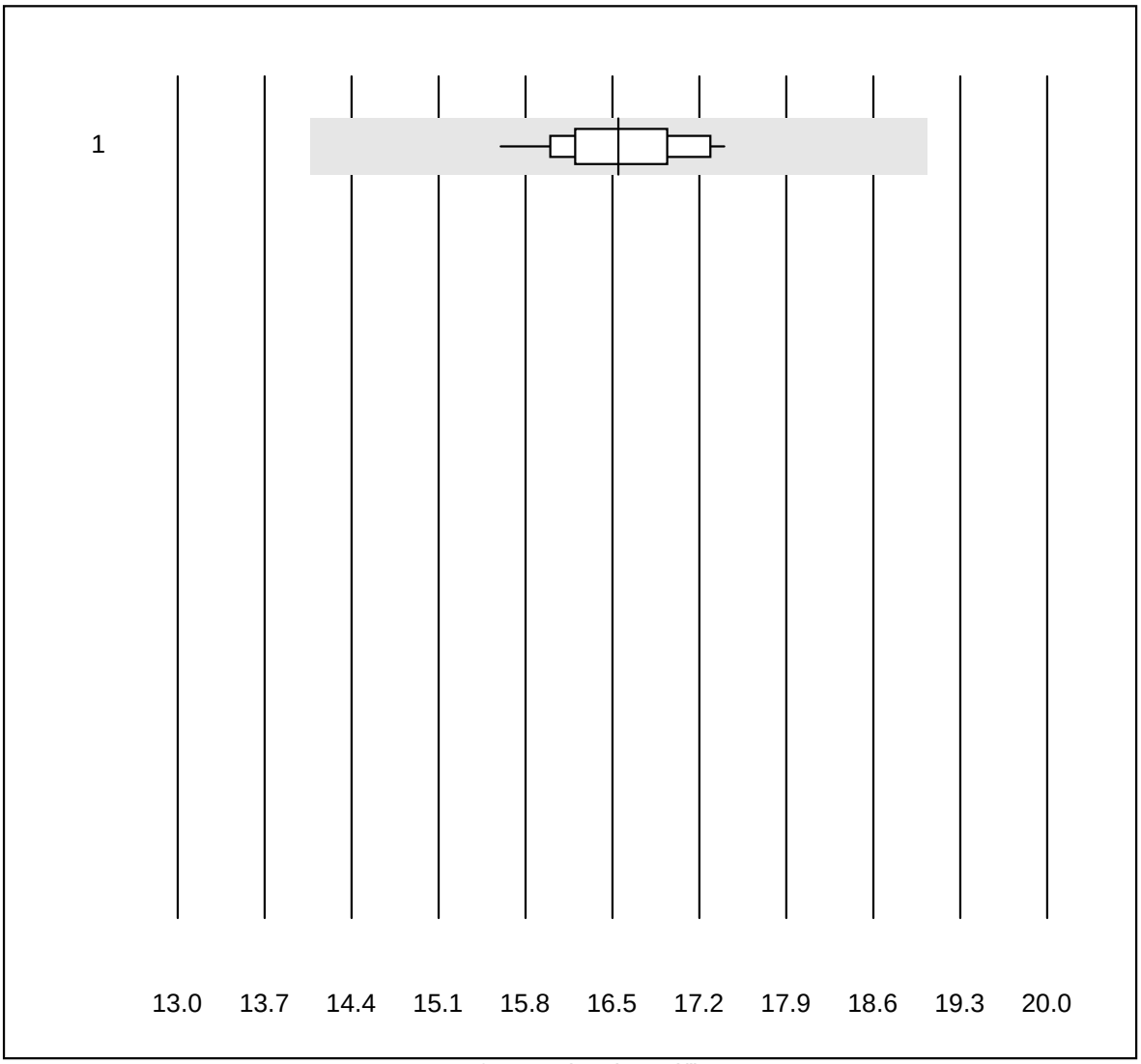


Deviatione QUALAB : 10 %

Osmolalità - urine (mosm/kg)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cryoscopie	6	100.0	0.0	0.0	812	0.6	e

Fosforo - urine

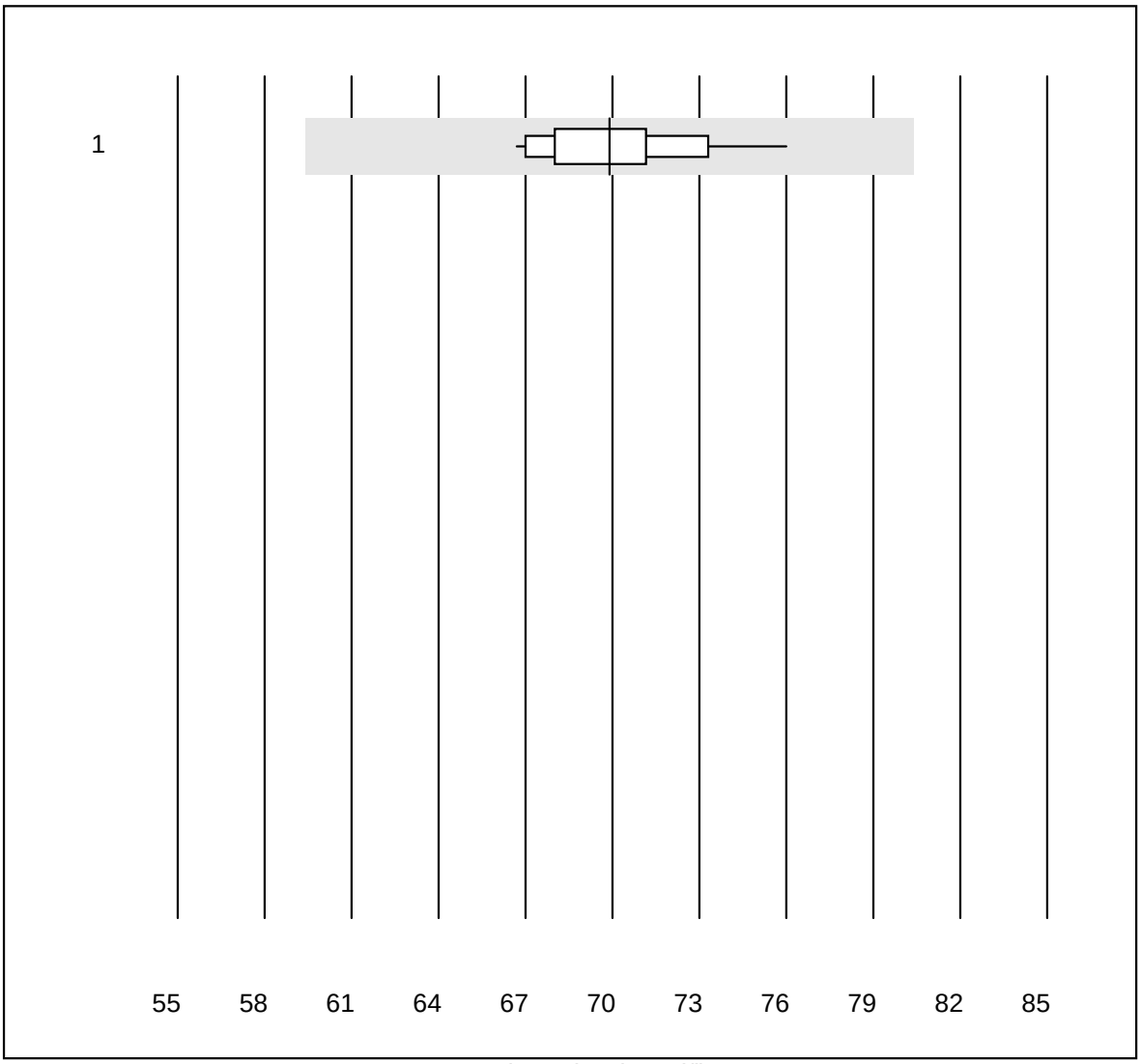


Deviatione QUALAB : 15 %

Fosforo - urine (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	15	100.0	0.0	0.0	16.5	3.1	e

Potassio - urine

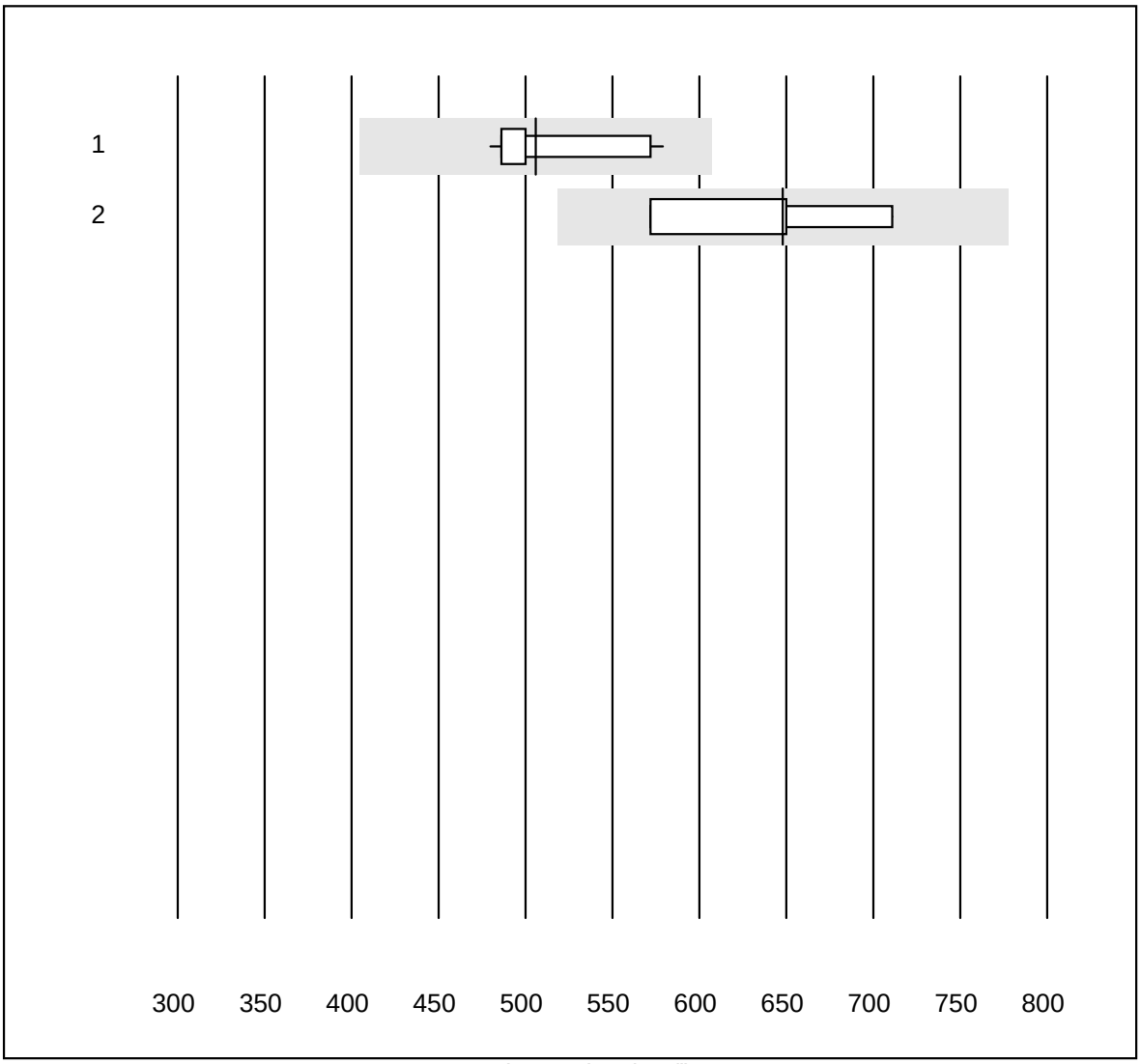


Deviatione QUALAB : 15 %

Potassio - urine (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	tutti	19	100.0	0.0	0.0	70	3.3	e

Proteina - urina

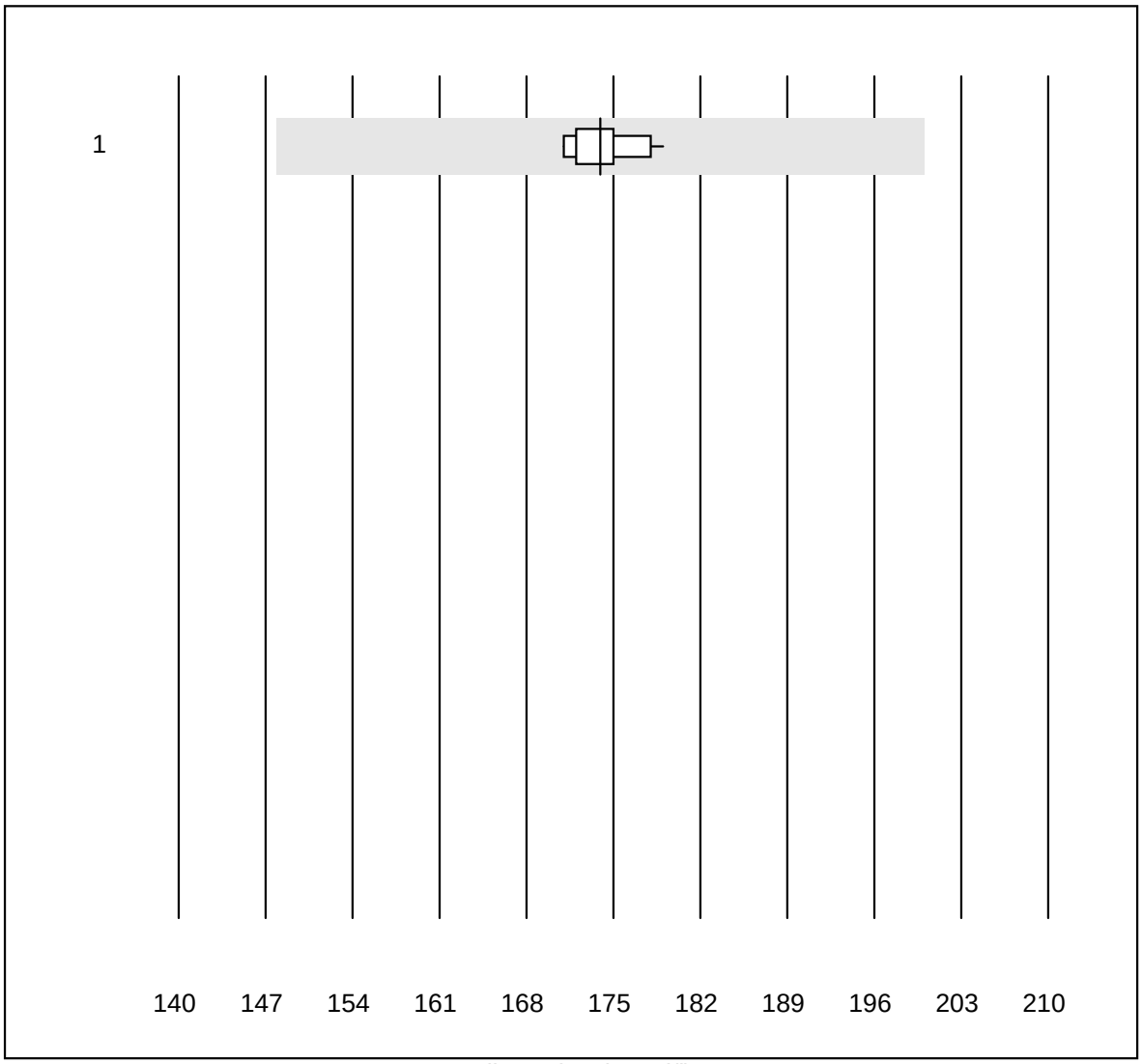


Deviatione QUALAB : 20 %

Proteina - urina (mg/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas/Roche	12	100.0	0.0	0.0	506.0	6.5	e
2	altro	4	100.0	0.0	0.0	648.0	8.8	e*

Sodio - urine



Deviatione QUALAB : 15 %

Sodio - urine (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	tutti	19	100.0	0.0	0.0	174	1.4	e

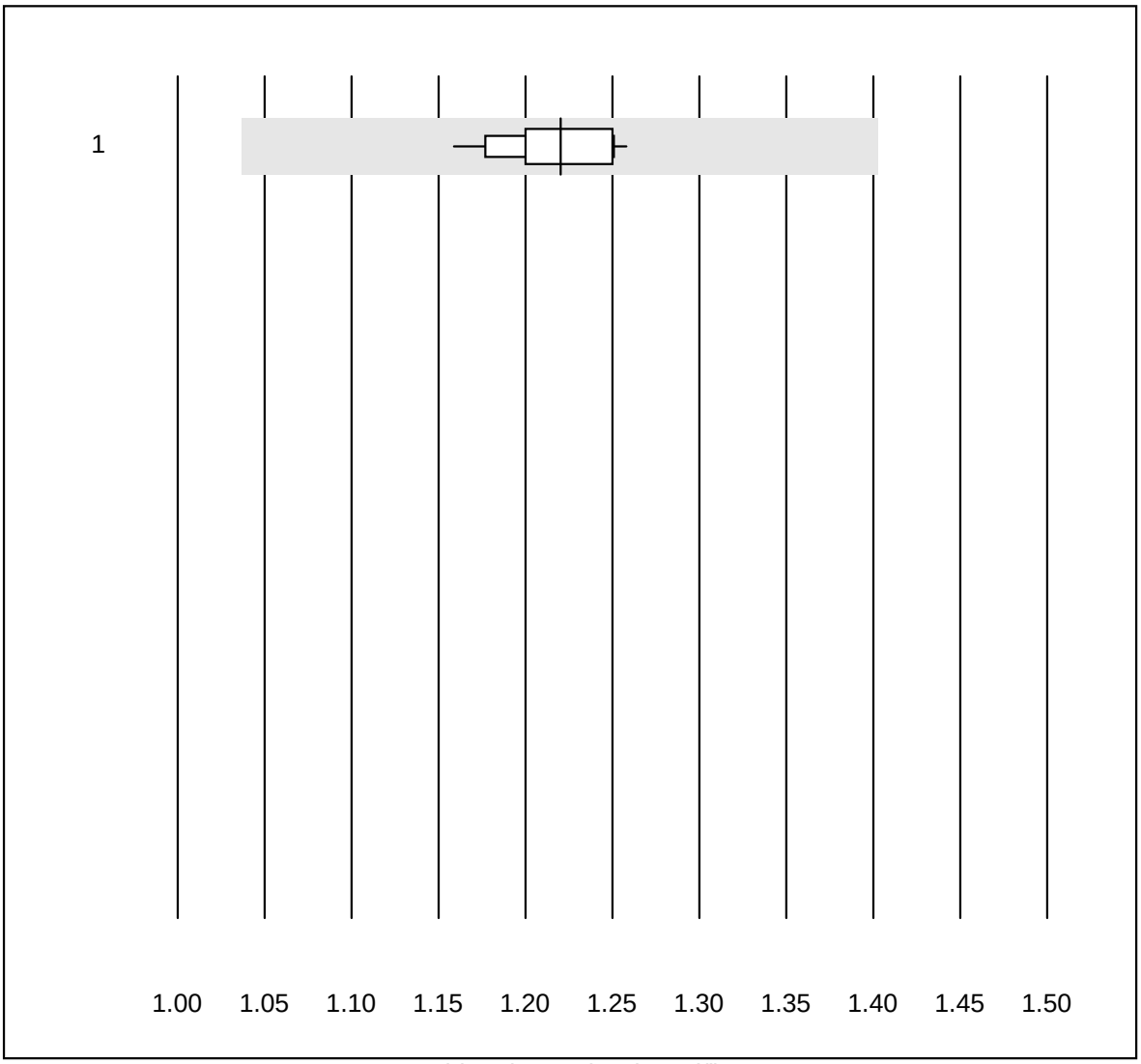
Urea - urine

Deviatione QUALAB : 15 %

Urea - urine (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	15	100.0	0.0	0.0	267	4.0	e

Acido urico - urine

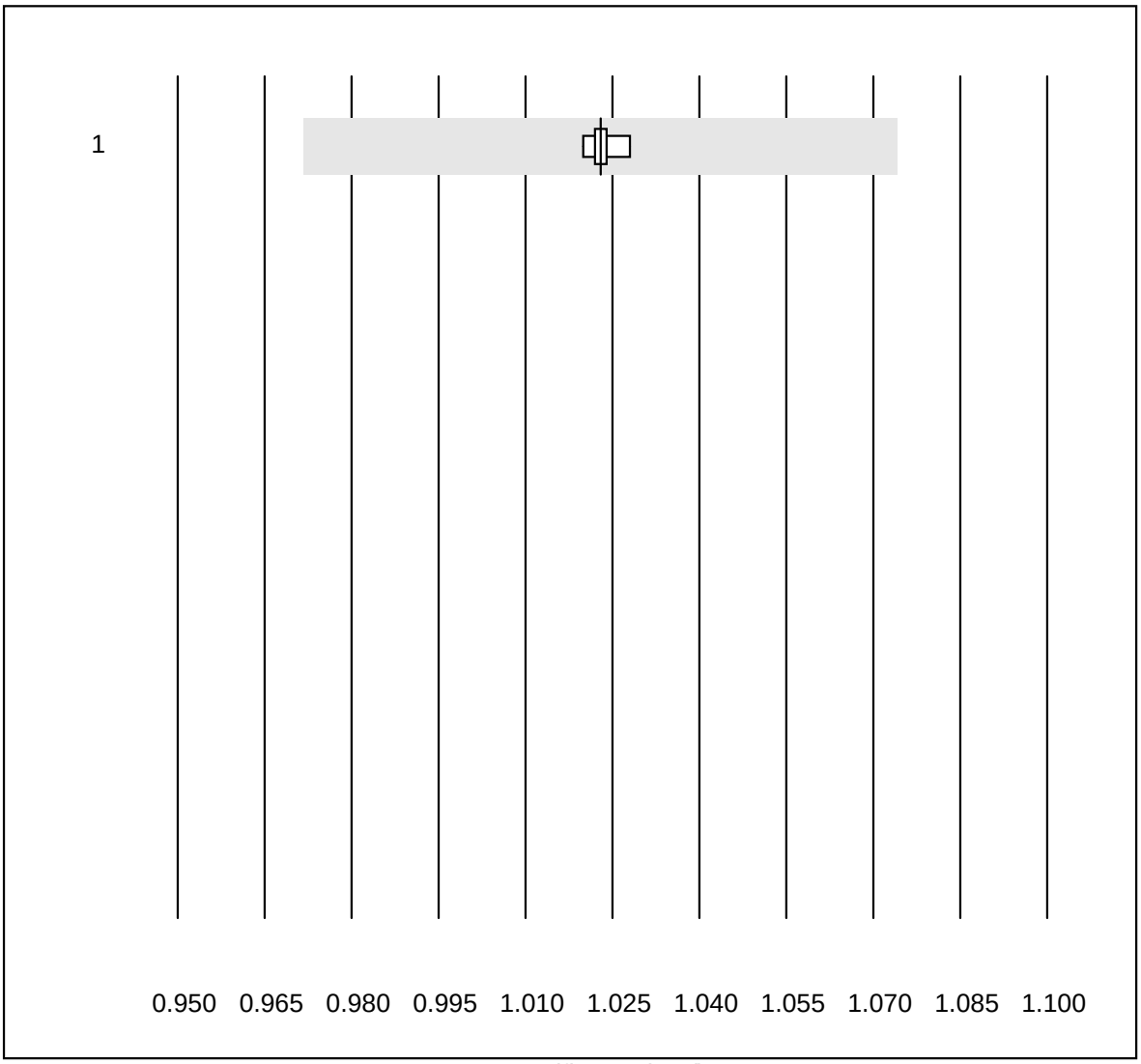


Deviatione QUALAB : 15 %

Acido urico - urine (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	14	100.0	0.0	0.0	1.22	2.5	e

Peso Specifico - urine

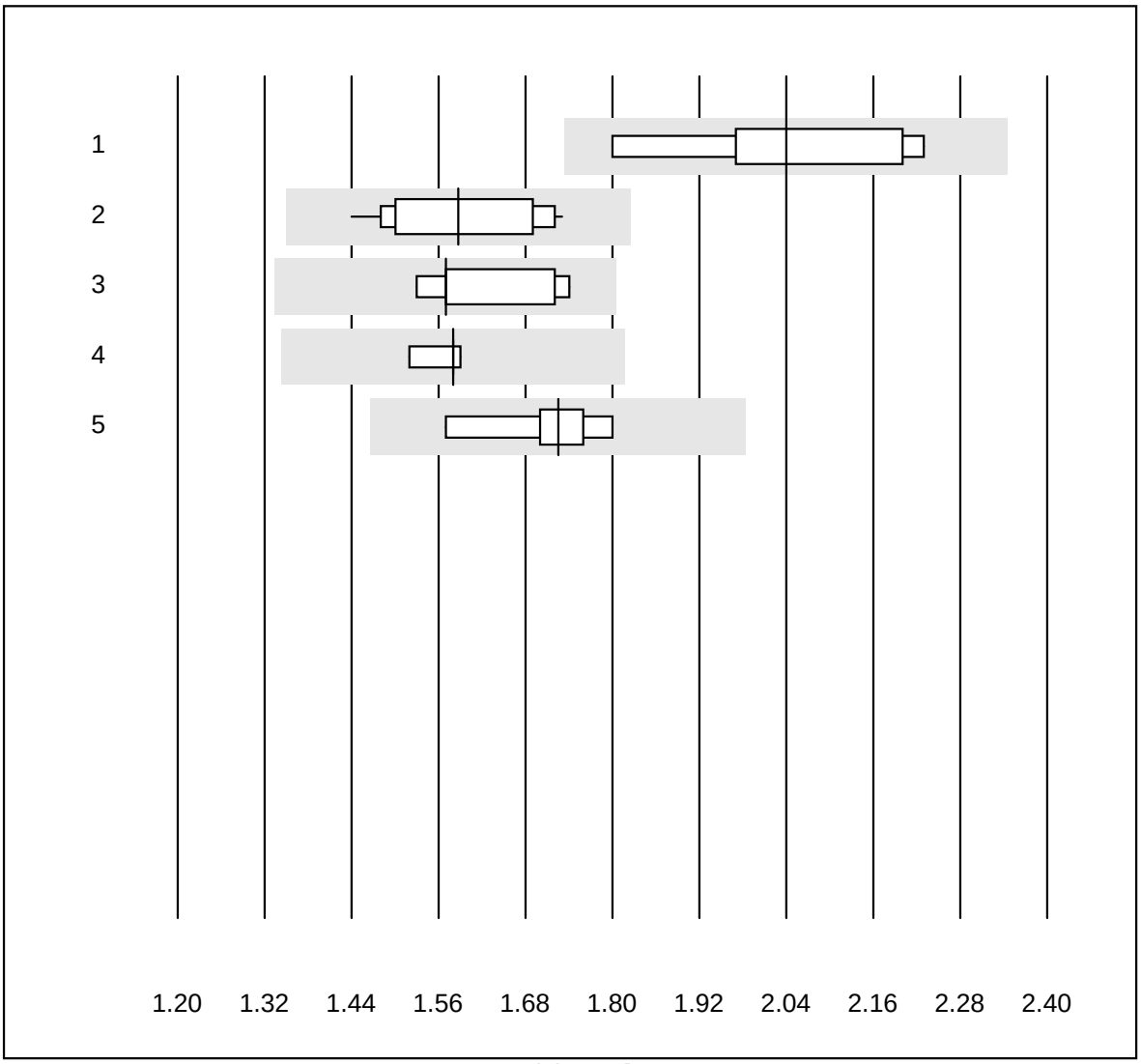


Deviatione QUALAB : 5 %

Peso Specifico - urine ()

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Refraktometer	6	100.0	0.0	0.0	1.023	0.3	e

Quick OA

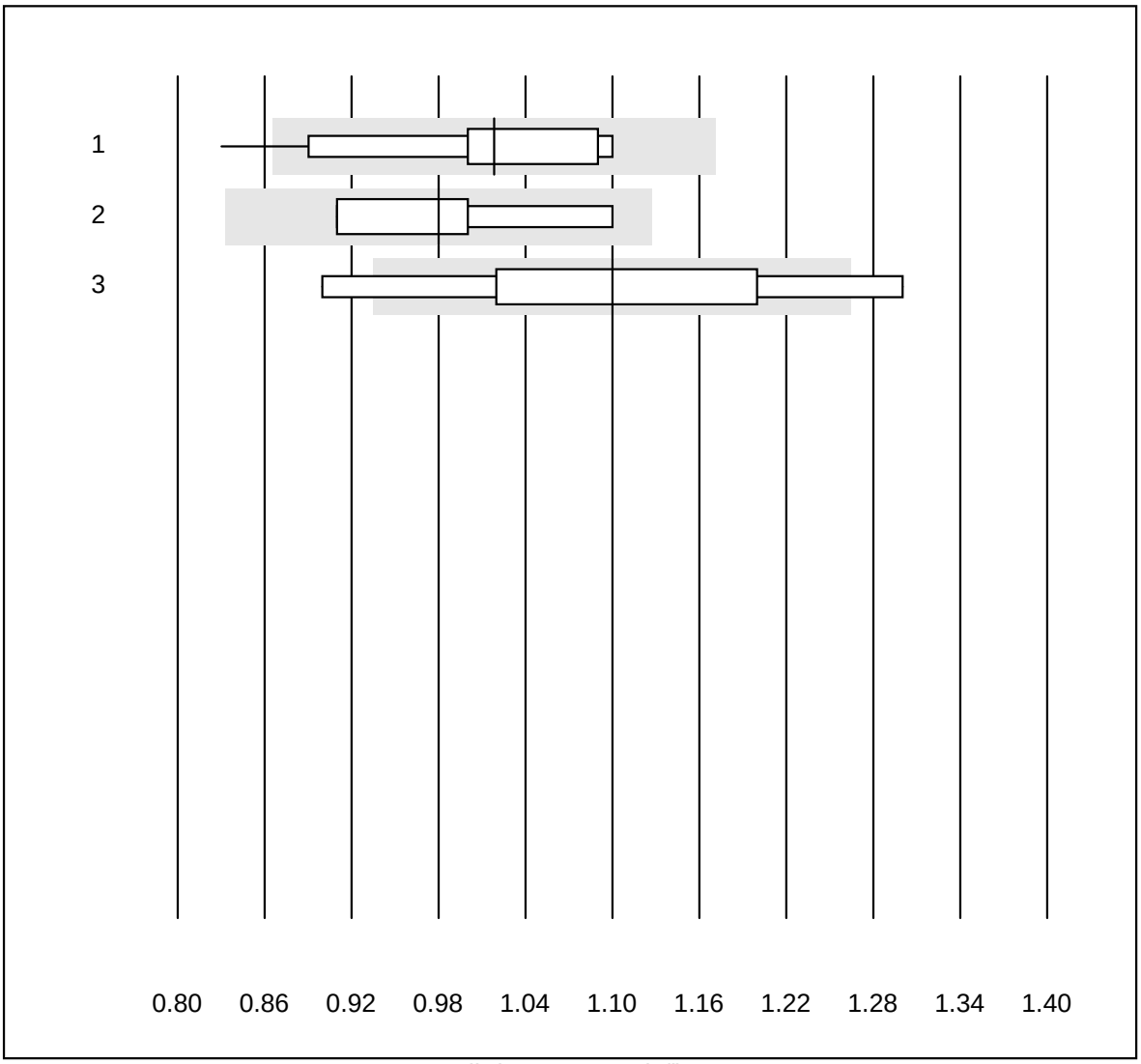


Deviatione QUALAB : 15 %

Quick OA ()

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Neoplastin Plus	5	100.0	0.0	0.0	2.04	8.6	e*
2	Innovin	18	100.0	0.0	0.0	1.59	5.8	e
3	Recombiplastin 2G	5	100.0	0.0	0.0	1.57	5.9	e*
4	altro	5	100.0	0.0	0.0	1.58	1.8	e
5	Neoplastin R	8	87.5	0.0	12.5	1.73	4.2	e

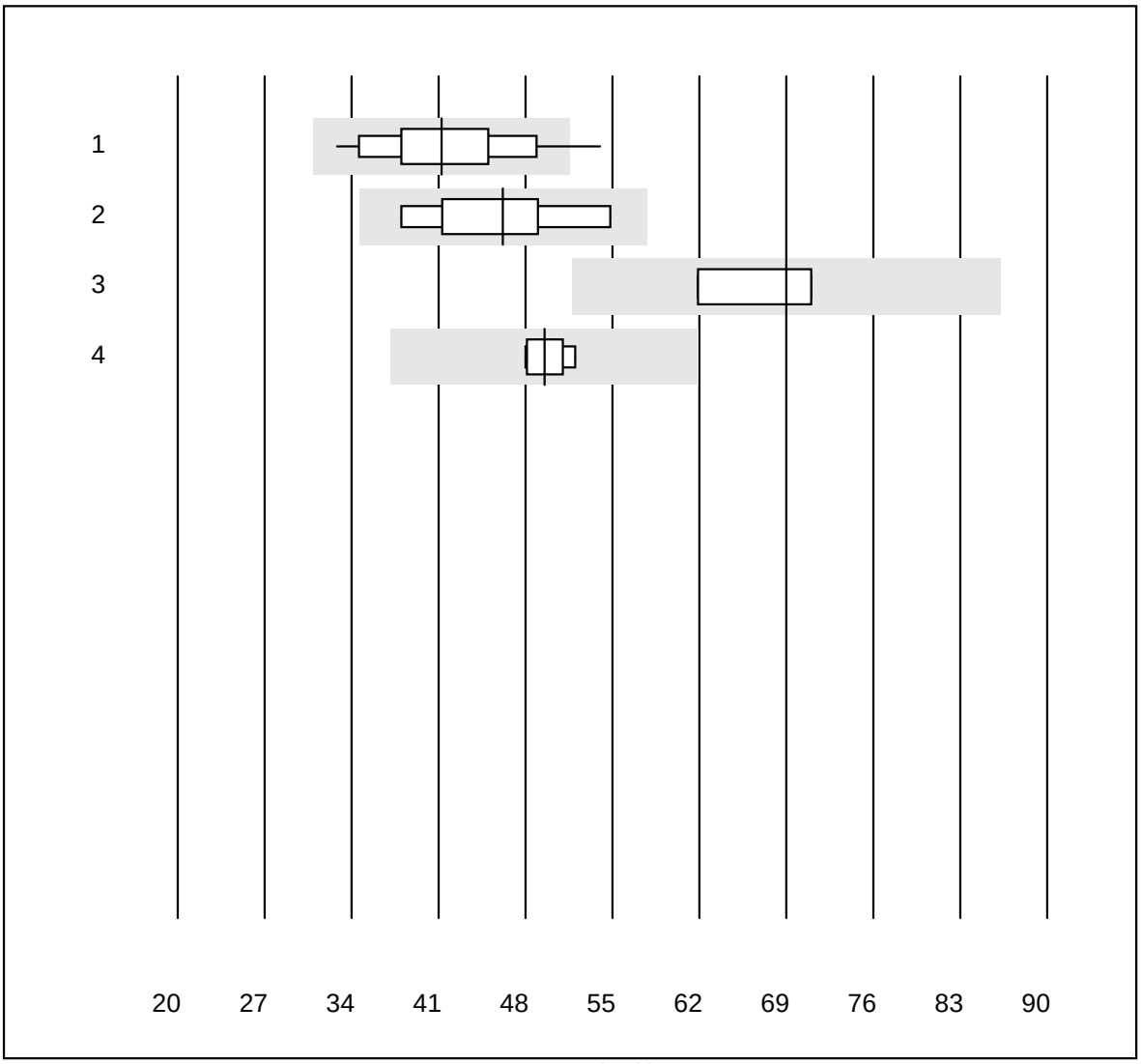
Fibrinogeno OA



Deviatione QUALAB : 15 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	altro	11	90.9	9.1	0.0	1.02	8.5	e*
2	Siemens Thrombin	4	100.0	0.0	0.0	0.98	8.1	e*
3	Stago/STA	7	71.4	28.6	0.0	1.10	11.6	e*

aPTT OA

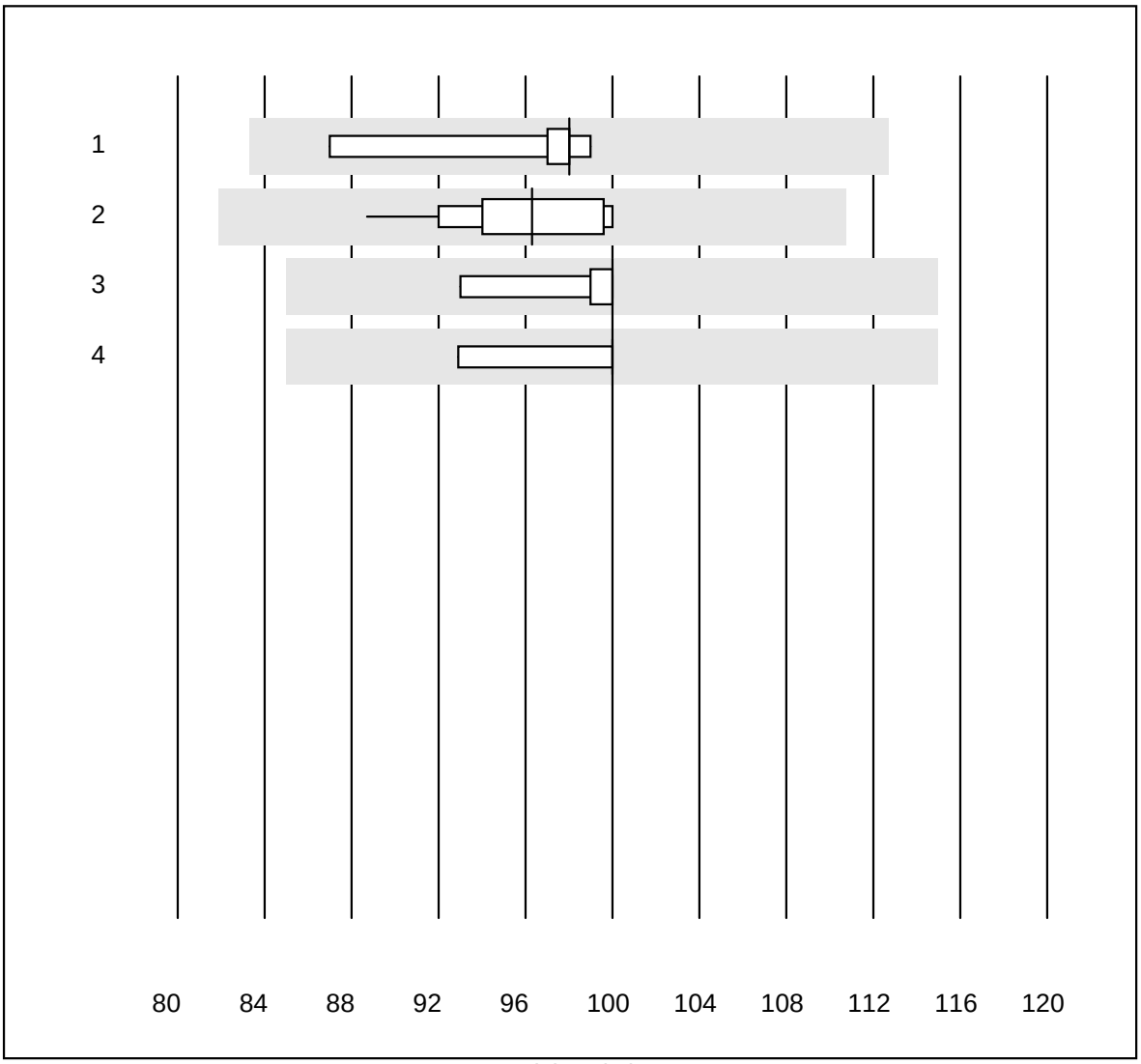


Deviatione QUALAB : 25 %

aPTT OA (Sek)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	altro	13	92.3	7.7	0.0	41.3	14.8	e*
2	Actin FS	8	100.0	0.0	0.0	46.2	12.4	e*
3	Pathromtin SL	4	100.0	0.0	0.0	69.0	6.4	e*
4	Stago/STA	6	100.0	0.0	0.0	49.5	3.2	e

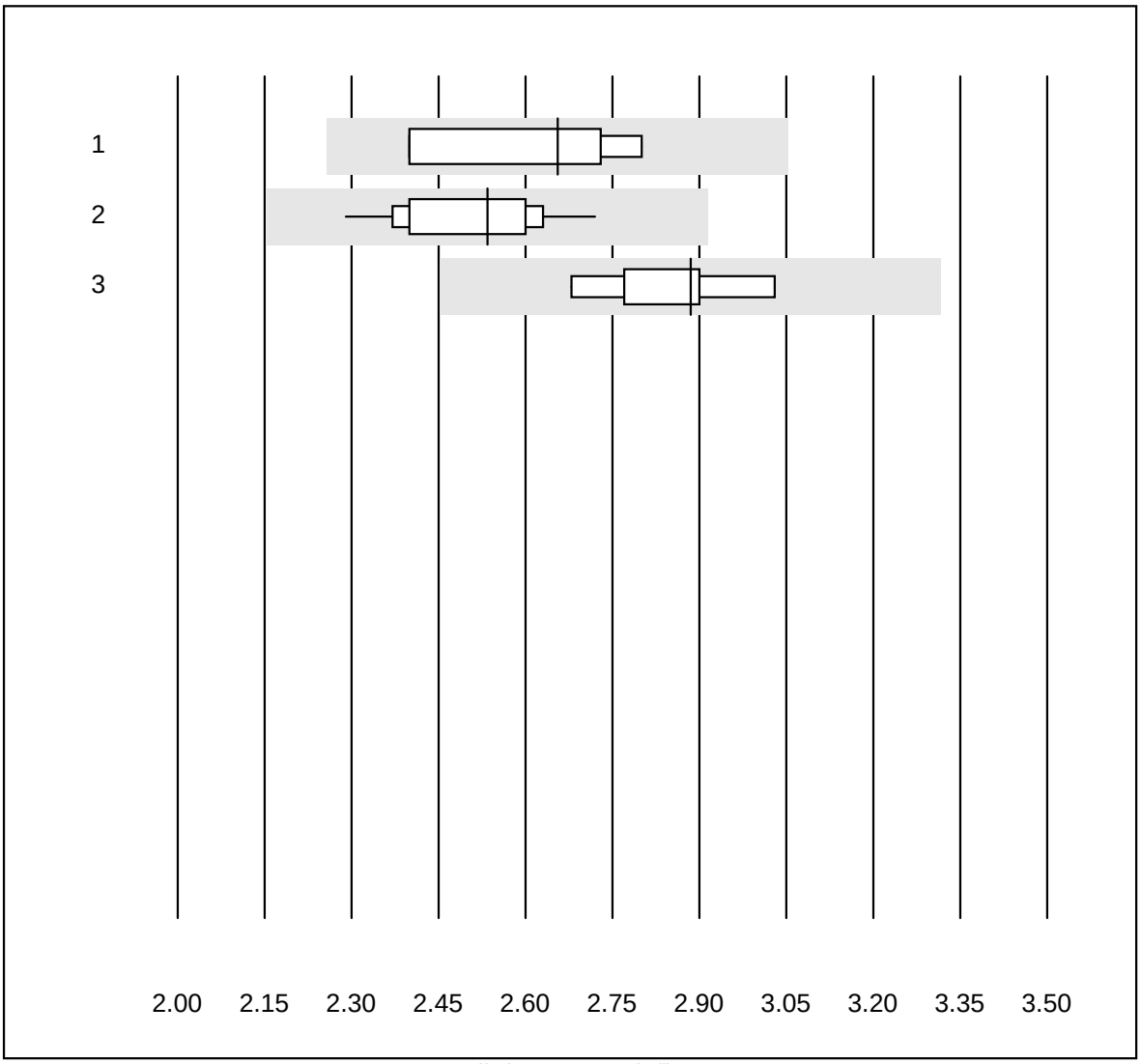
Quick N



Deviatione QUALAB : 15 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Neoplastin R	5	100.0	0.0	0.0	98	5.2	e*
2	Innovin	11	100.0	0.0	0.0	96	3.8	e
3	tutti	6	100.0	0.0	0.0	100	2.8	e
4	Recombiplastin 2G	6	100.0	0.0	0.0	100	2.9	e

Fibrinogeno N

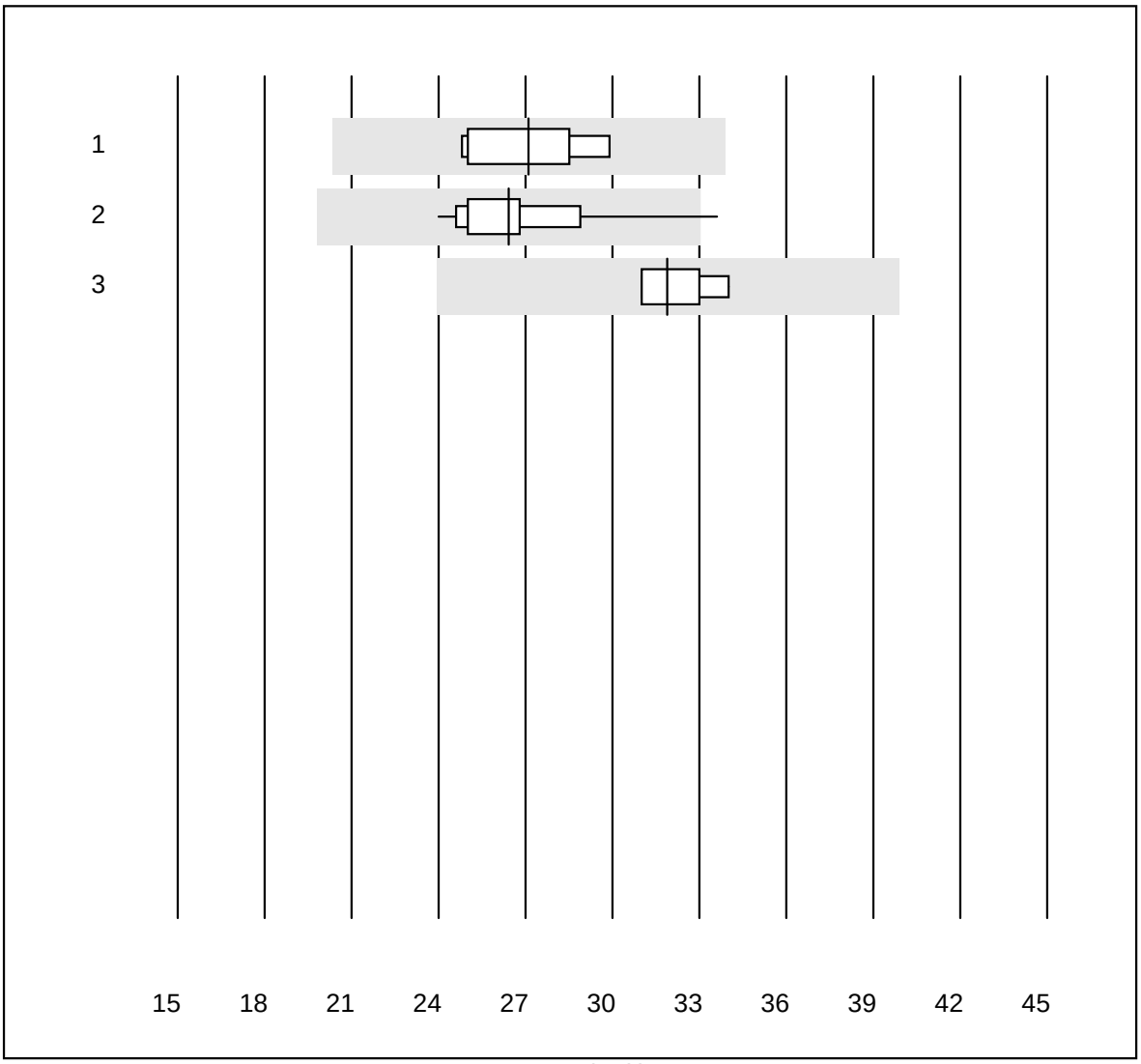


Deviatione QUALAB : 15 %

Fibrinogeno N (g/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Siemens Thrombin	4	100.0	0.0	0.0	2.66	6.7	e*
2	altro	12	91.7	0.0	8.3	2.53	5.1	e
3	Stago/STA	8	100.0	0.0	0.0	2.89	3.7	e

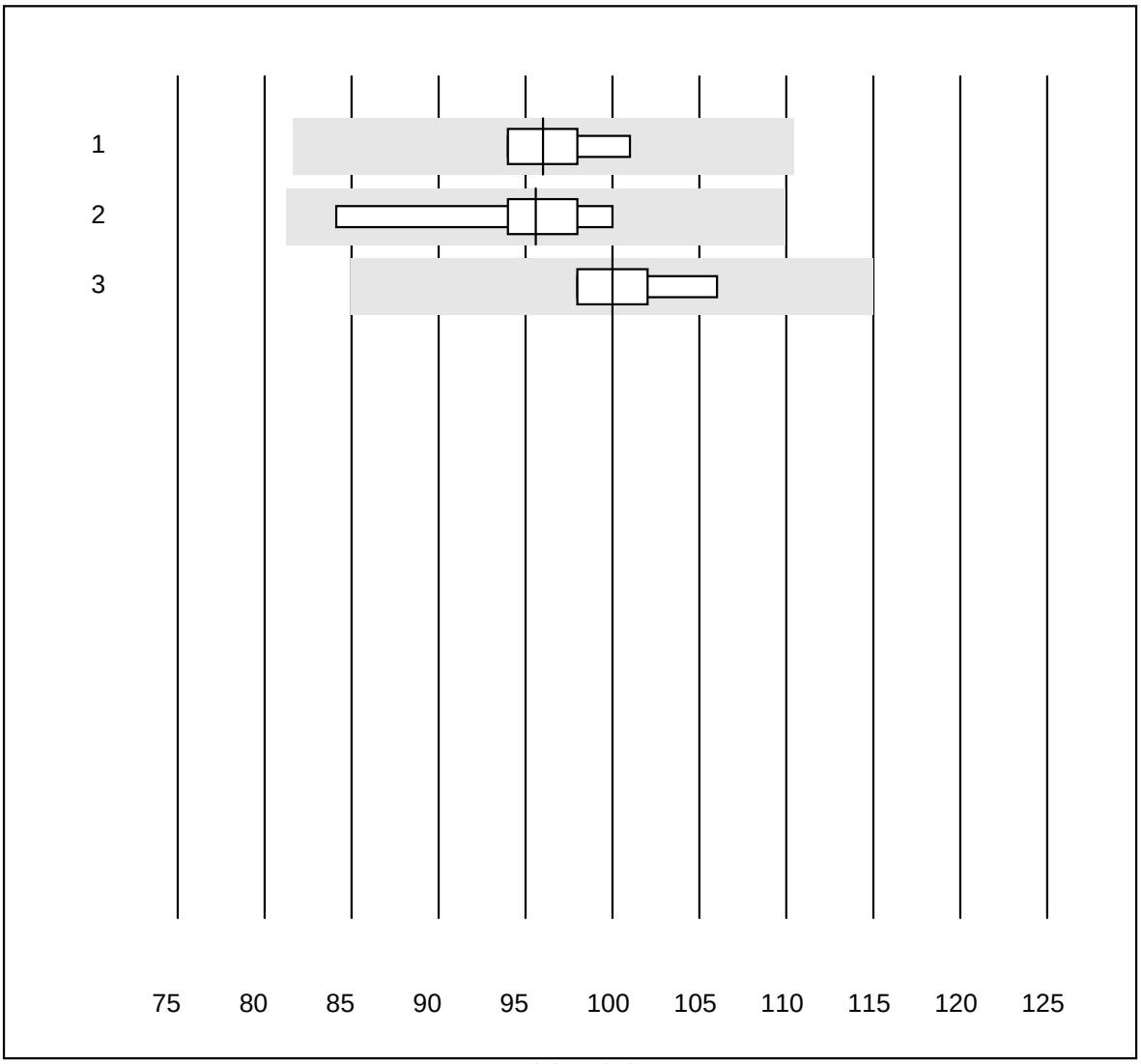
aPTT N



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Actin FS	8	100.0	0.0	0.0	27.1	7.0	e
2	altro	13	92.3	7.7	0.0	26.4	9.4	e
3	Stago/STA	7	100.0	0.0	0.0	31.9	3.6	e

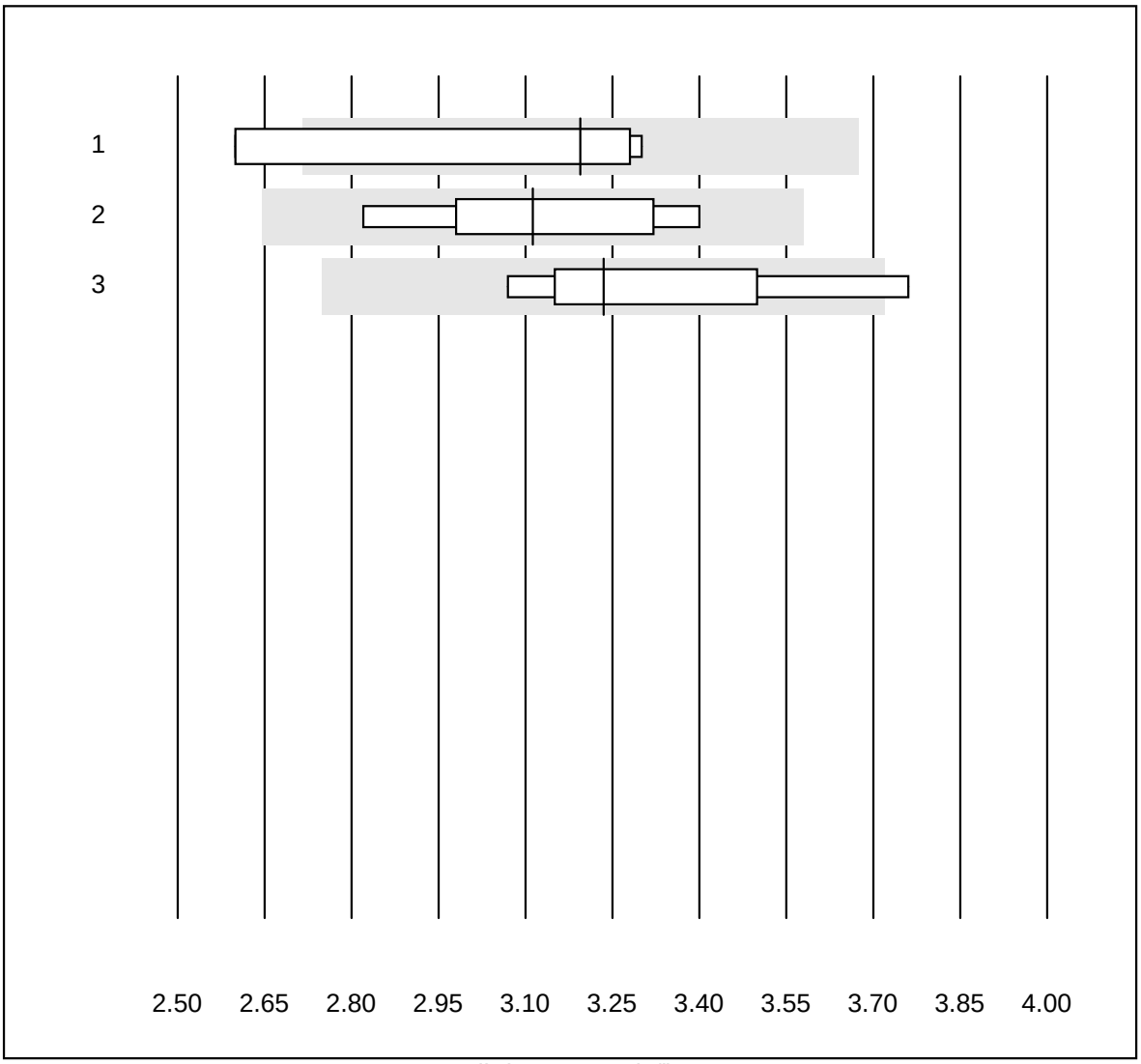
Quick H



Deviatione QUALAB : 15 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Neoplastin R	5	100.0	0.0	0.0	96	3.1	e
2	Innovin	9	100.0	0.0	0.0	96	5.0	e
3	Recombiplastin 2G	5	100.0	0.0	0.0	100	3.3	e

Fibrinogeno H

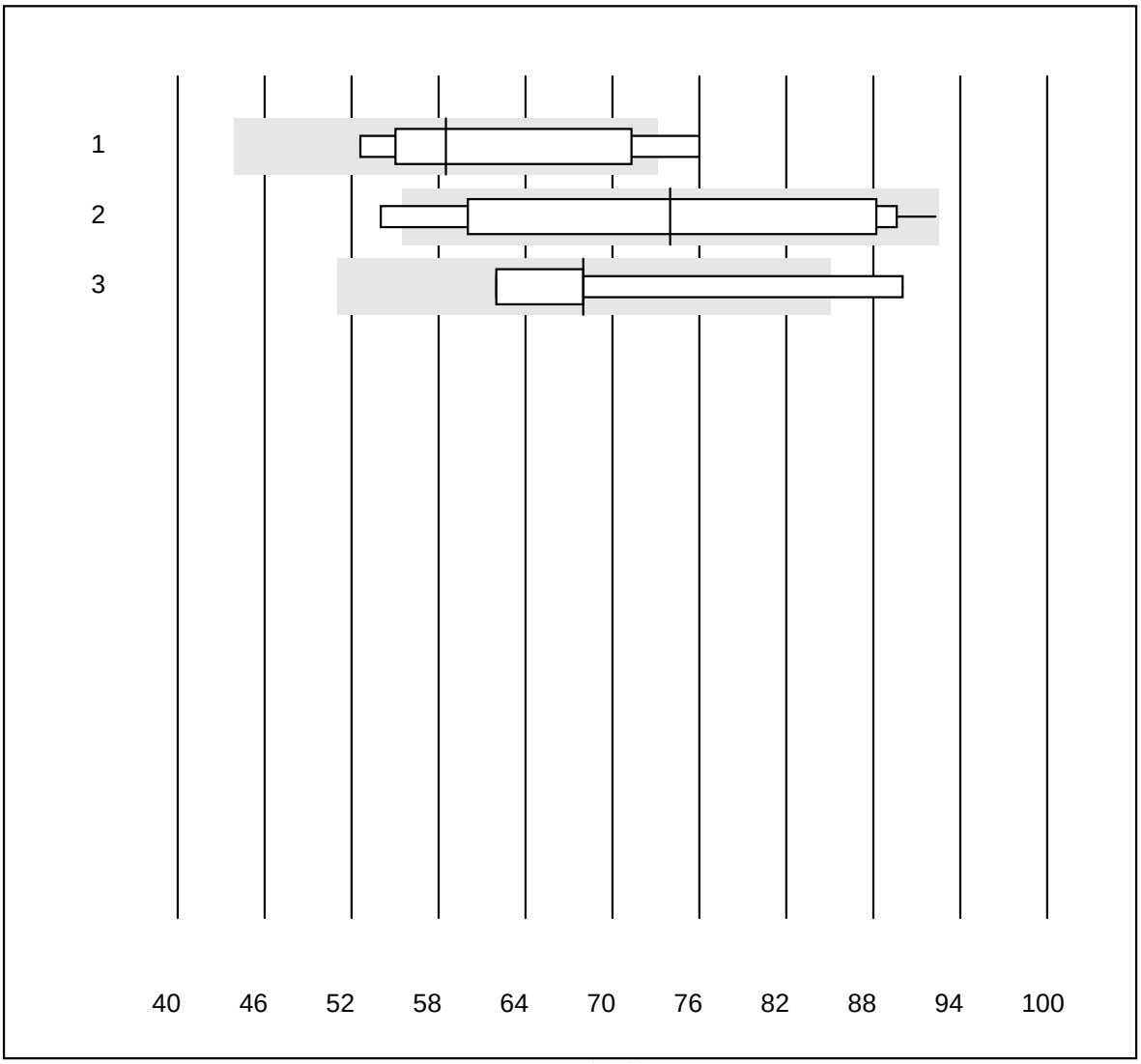


Deviatione QUALAB : 15 %

Fibrinogeno H (g/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Siemens Thrombin	4	75.0	25.0	0.0	3.20	10.6	e*
2	altro	10	100.0	0.0	0.0	3.11	6.4	e*
3	Stago/STA	6	83.3	16.7	0.0	3.24	7.8	e*

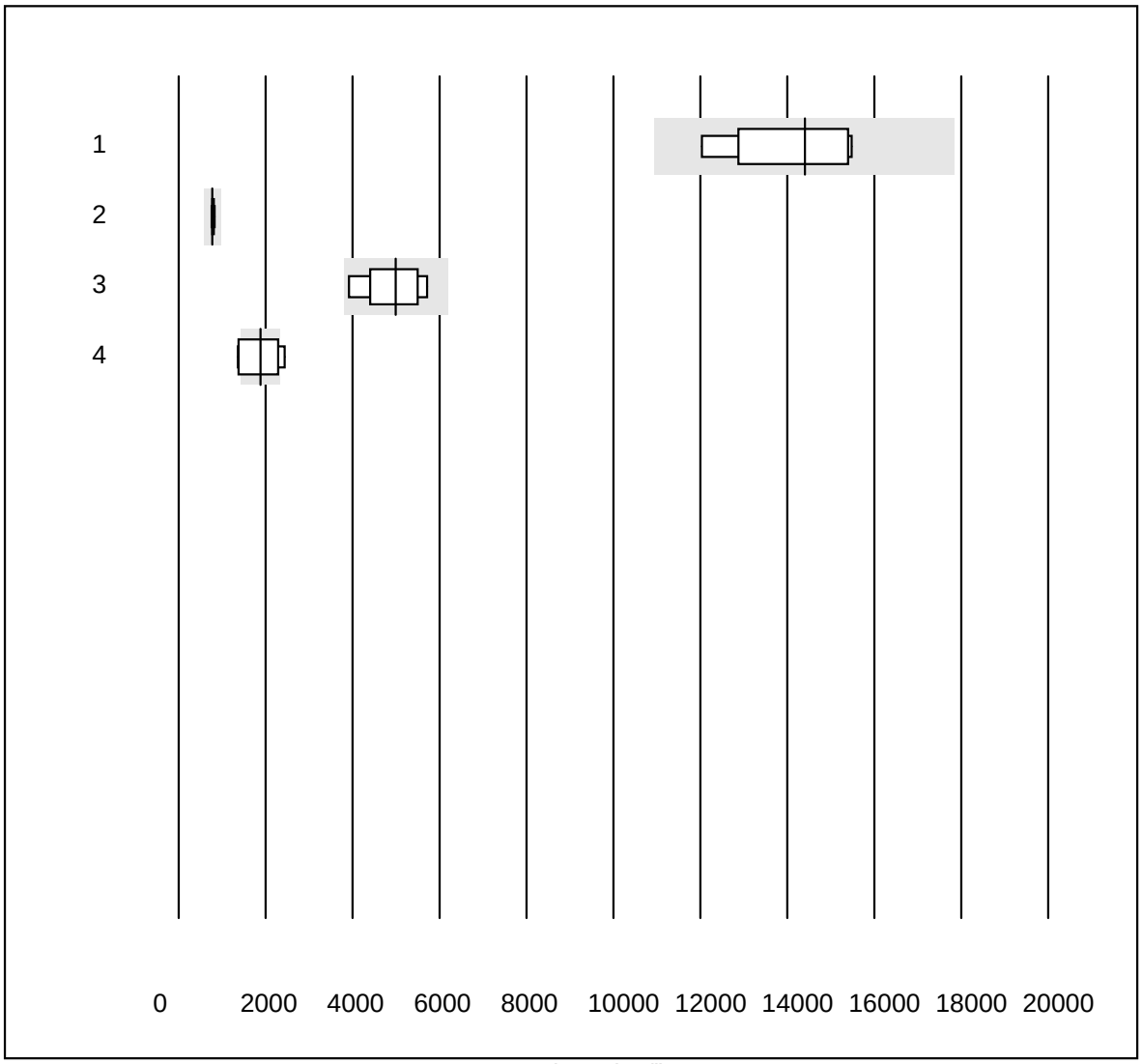
aPTT H



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Actin FS	7	85.7	14.3	0.0	58.5	13.9	e*
2	altro	10	90.0	10.0	0.0	74.0	18.8	e*
3	Stago/STA	5	60.0	20.0	20.0	68.0	18.2	e*

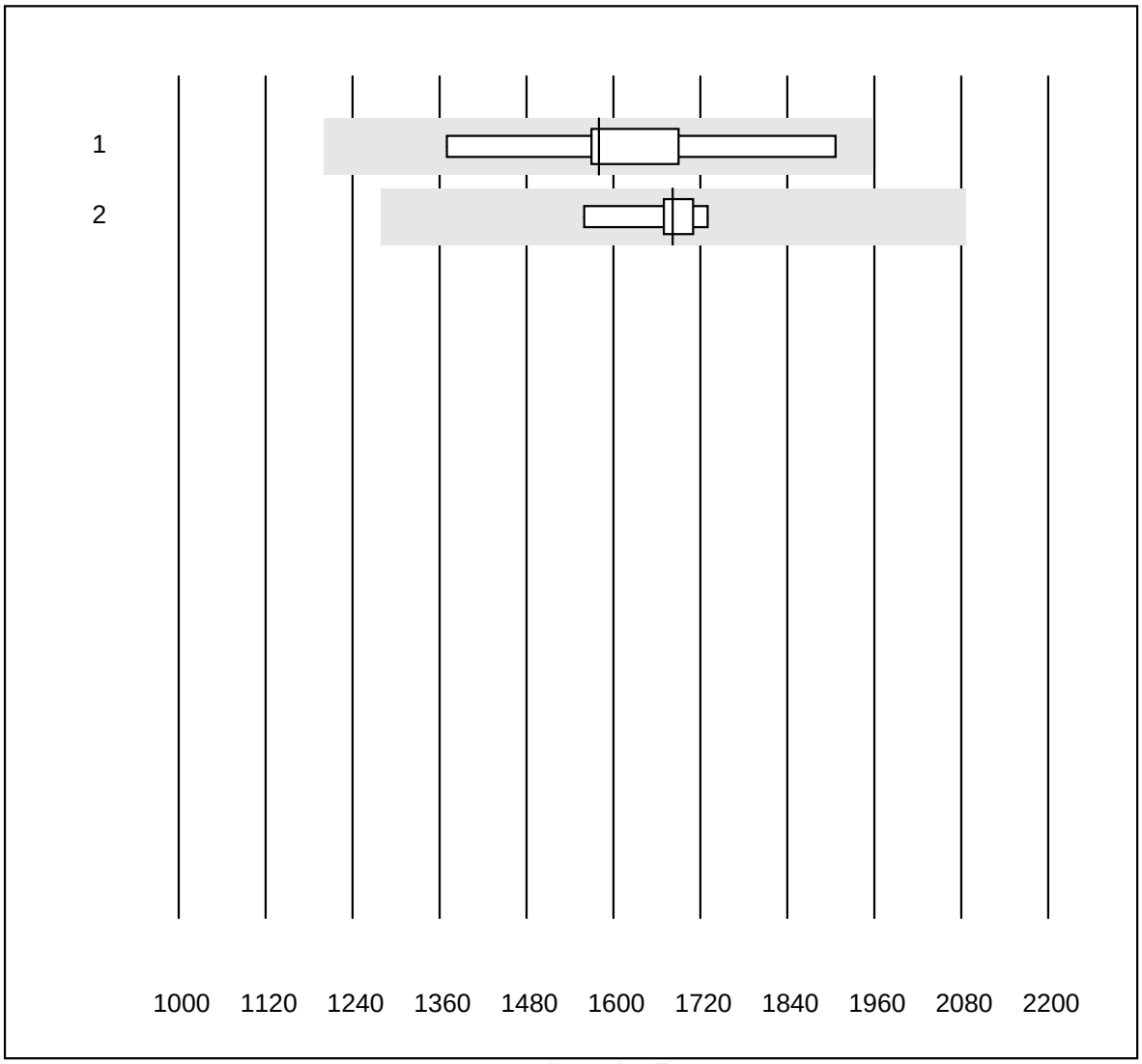
Troponina I



Deviatione QUALAB : 24 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Vidas	6	100.0	0.0	0.0	14395.0	9.9	e*
2	AQT 90 FLEX	5	100.0	0.0	0.0	780.0	3.8	e
3	ADVIA Centaur XP/CP	6	100.0	0.0	0.0	4992.5	14.2	e*
4	Eurolyser	13	38.4	30.8	30.8	1877.0	24.0	e*

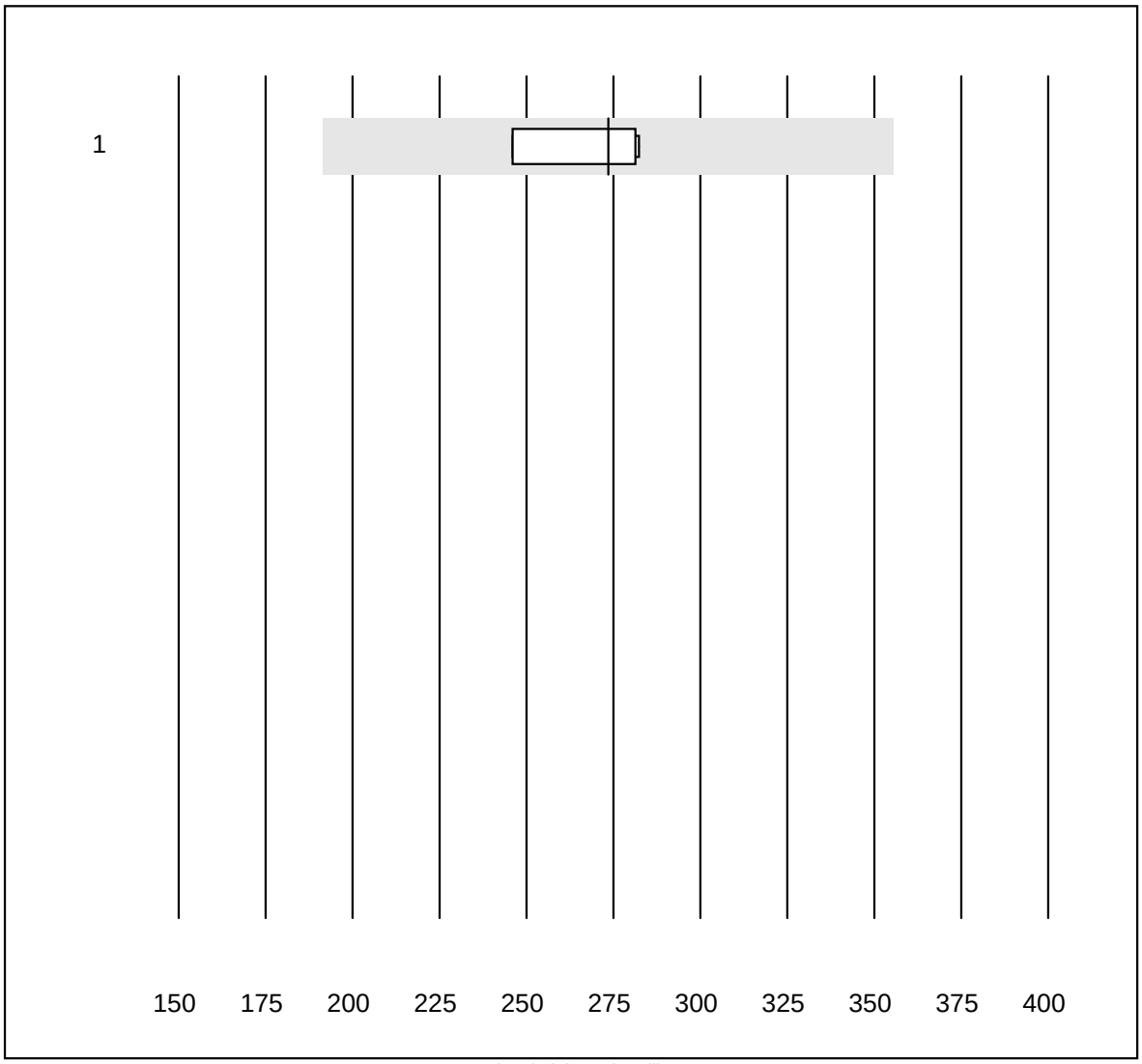
Troponina T



Deviatione QUALAB : 24 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas hs	5	100.0	0.0	0.0	1580.00	12.1	e*
2	Cobas hs STAT	6	100.0	0.0	0.0	1682.00	3.6	e

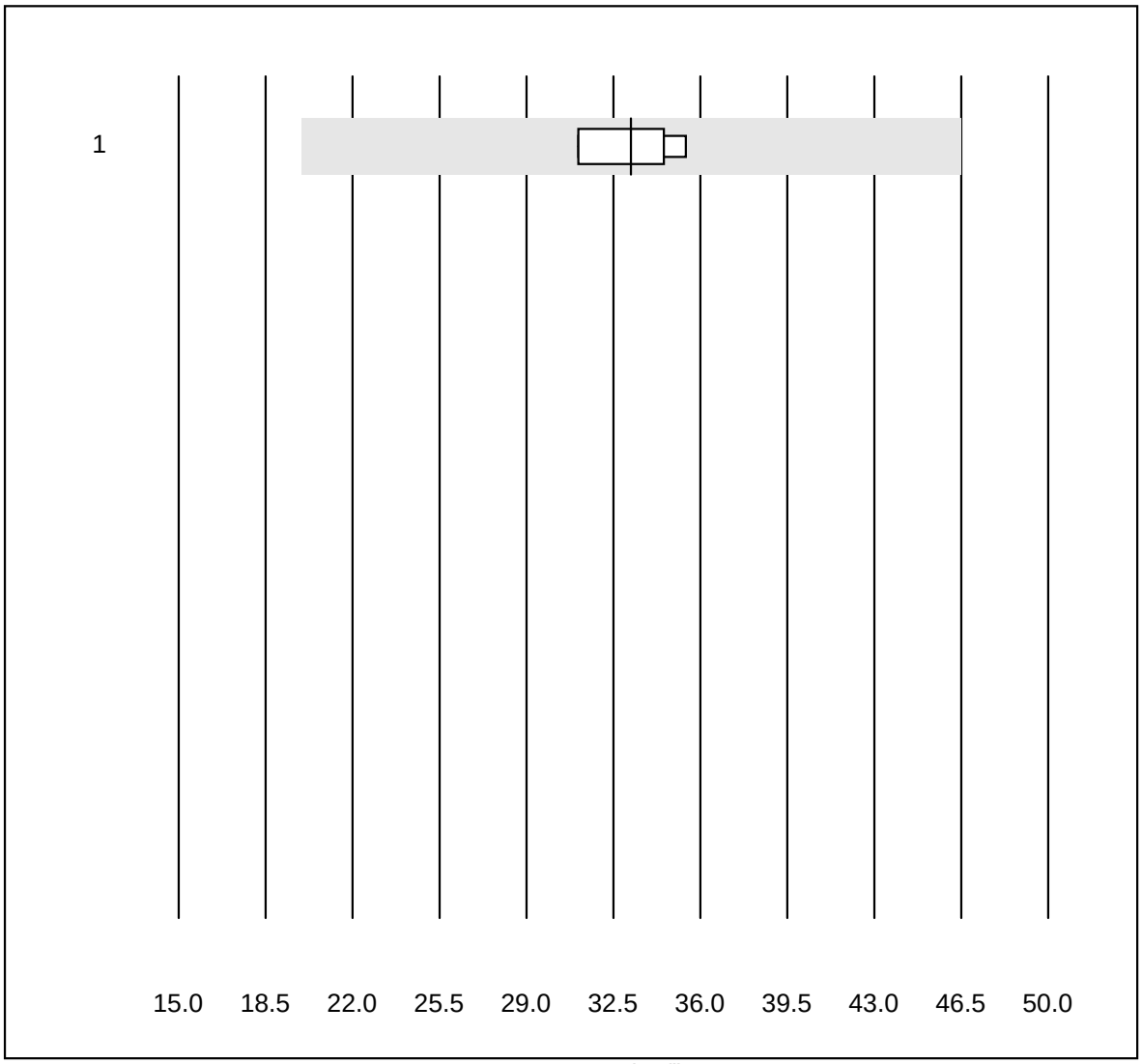
Mioglobina



Deviatione QUALAB : 30 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas E / Elecsys	4	100.0	0.0	0.0	273.6	6.3	e

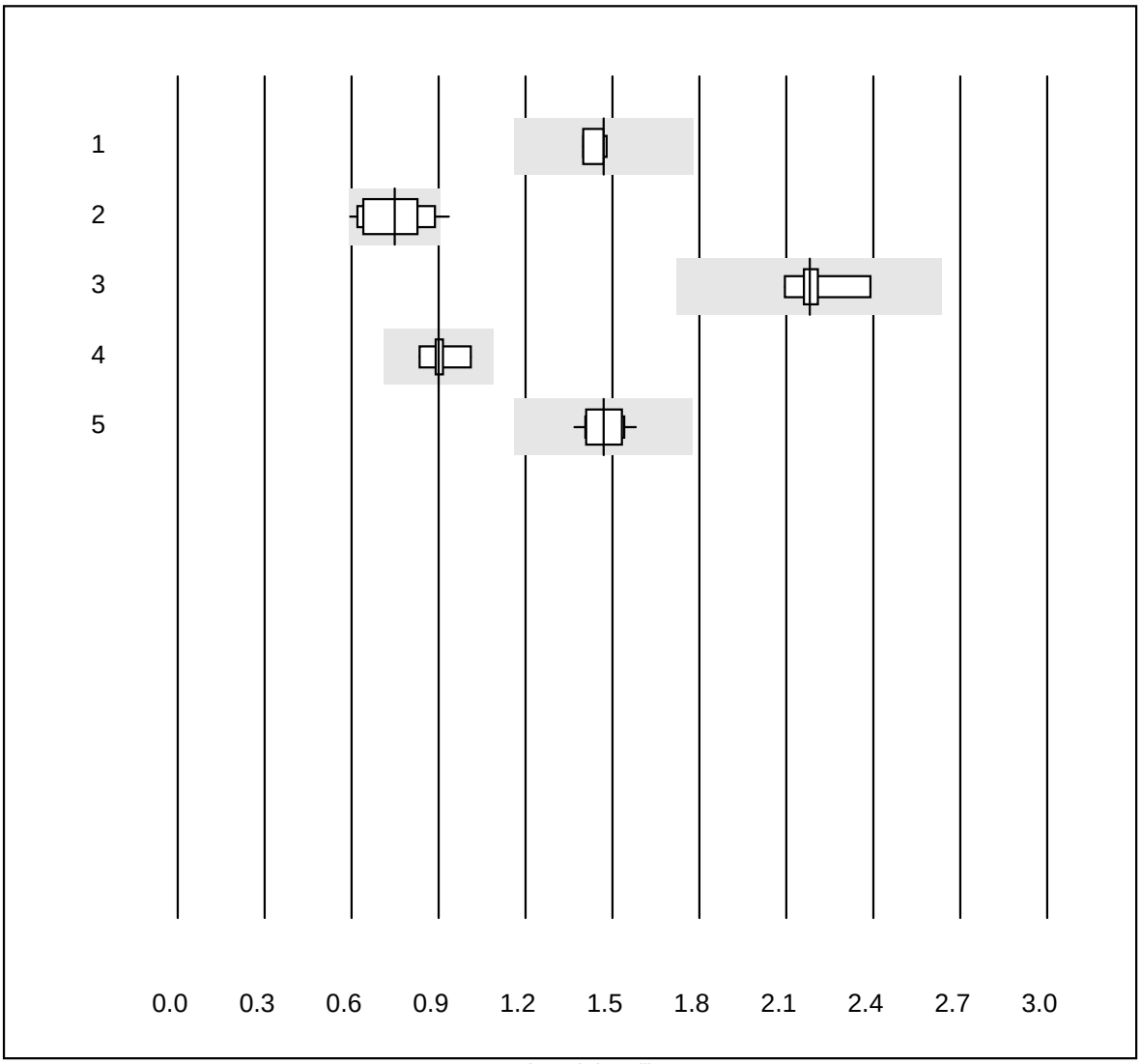
CK-MB massa



Deviatione QUALAB : 40 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	altro	4	100.0	0.0	0.0	33.2	6.2	e

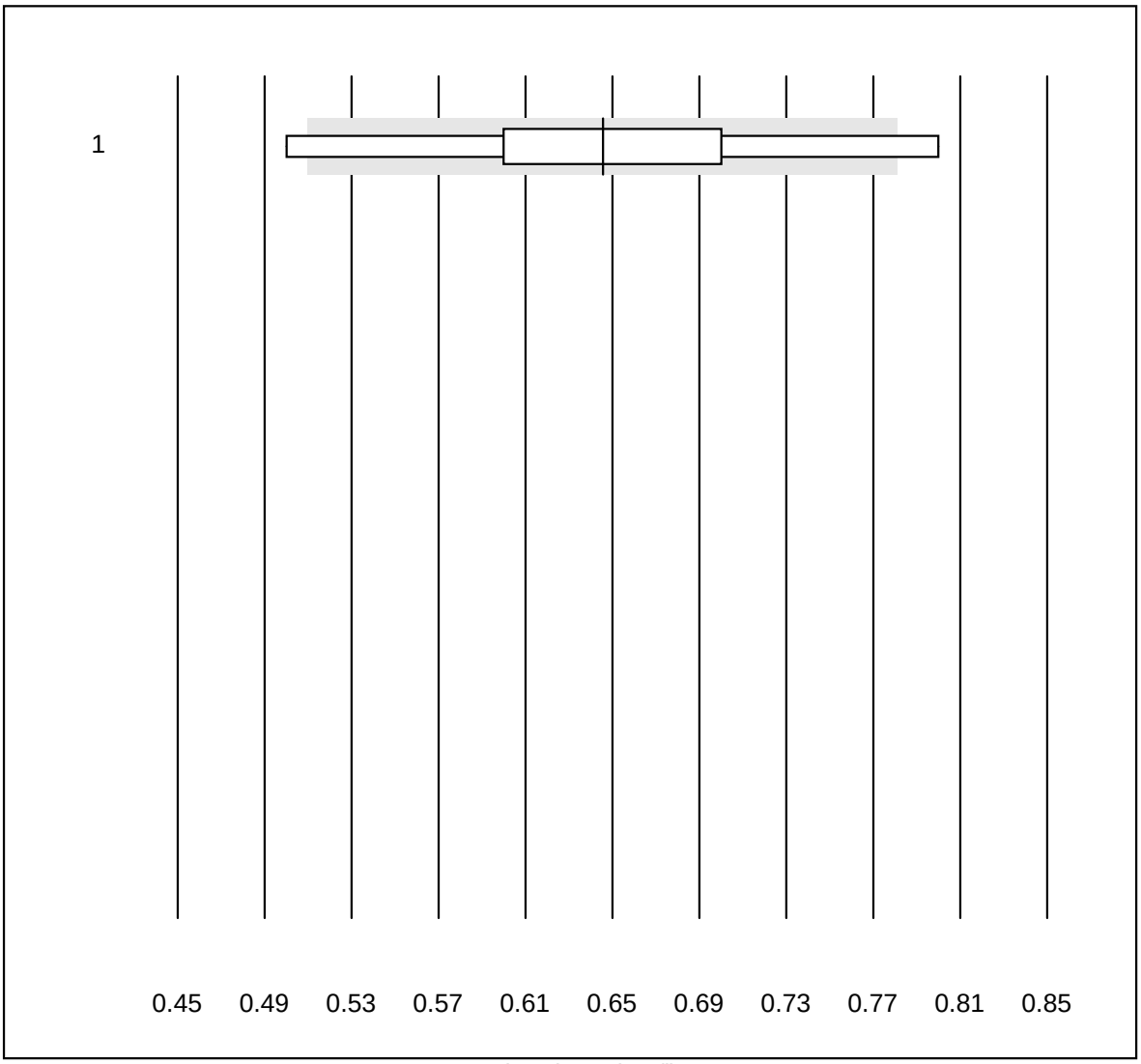
D-Dimeri



Deviatione QUALAB : 21 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	STA Liatest	8	87.5	0.0	12.5	1.47	2.3	e
2	Eurolyser	27	81.5	3.7	14.8	0.75	13.0	e
3	ACL	5	100.0	0.0	0.0	2.18	5.0	e
4	AQT 90 FLEX	6	100.0	0.0	0.0	0.90	6.3	e*
5	Vidas	11	100.0	0.0	0.0	1.47	4.5	e

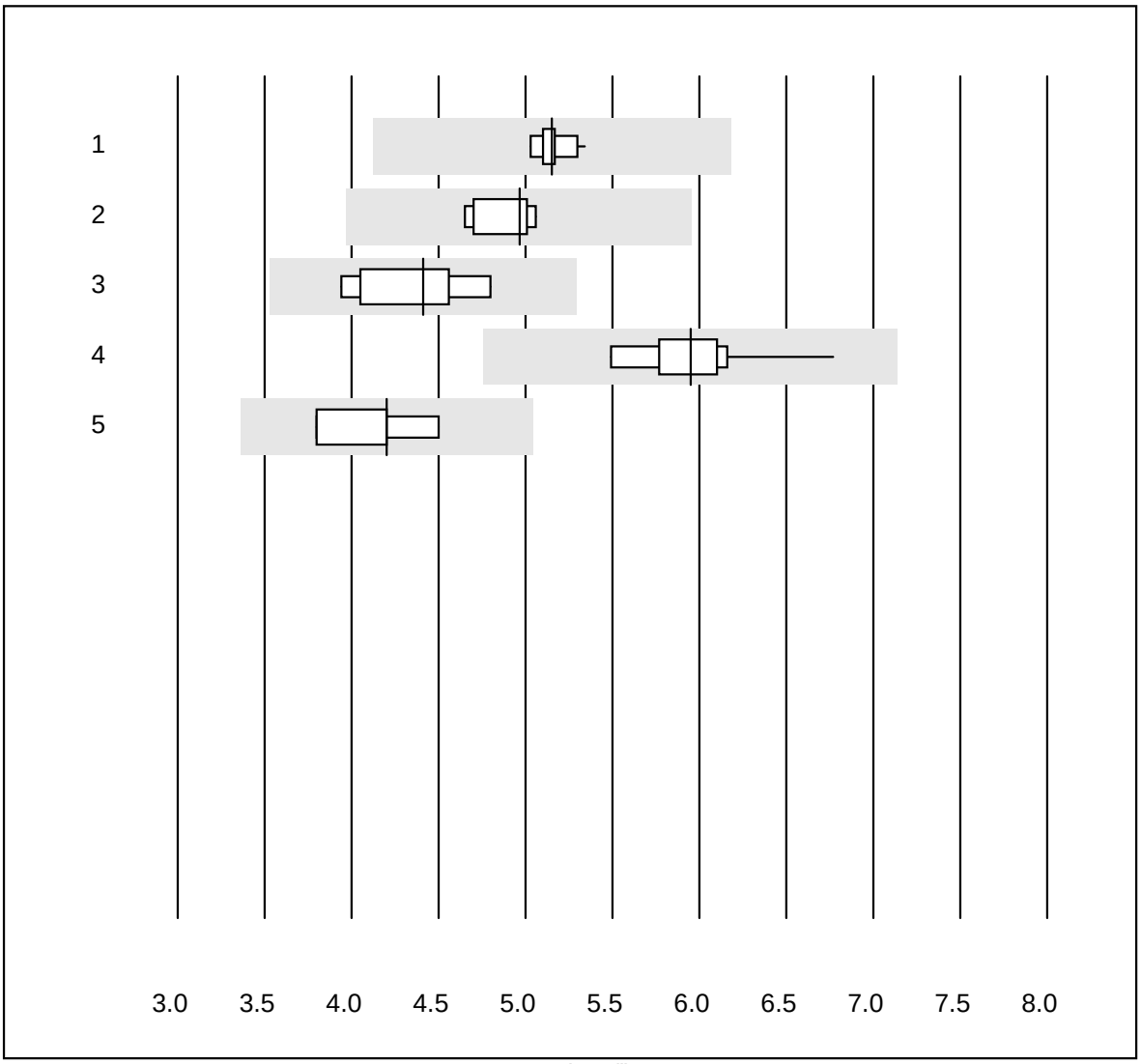
D-Dimeri NC



Deviatione QUALAB : 21 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	NycoCard	35	40.0	22.9	37.1	0.65	15.7	e*

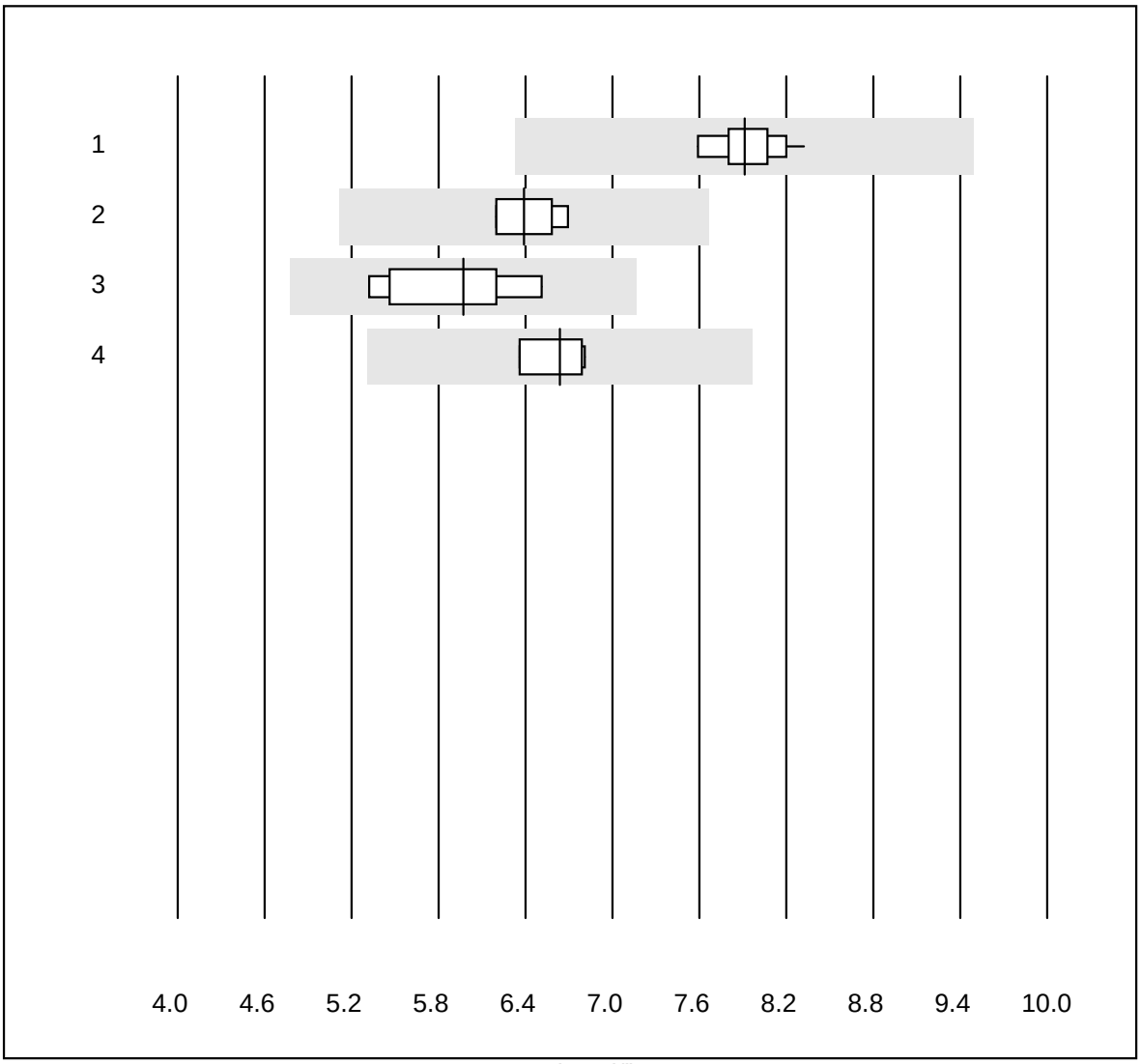
TSH



Deviatione QUALAB : 20 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas E / Elecsys	10	100.0	0.0	0.0	5.2	1.9	e
2	ADVIA Centaur XP/CP	5	100.0	0.0	0.0	5.0	3.9	e
3	Architect	8	100.0	0.0	0.0	4.4	6.7	e
4	Vidas	10	100.0	0.0	0.0	5.9	6.2	e
5	Qualigen	5	80.0	0.0	20.0	4.2	7.0	e*

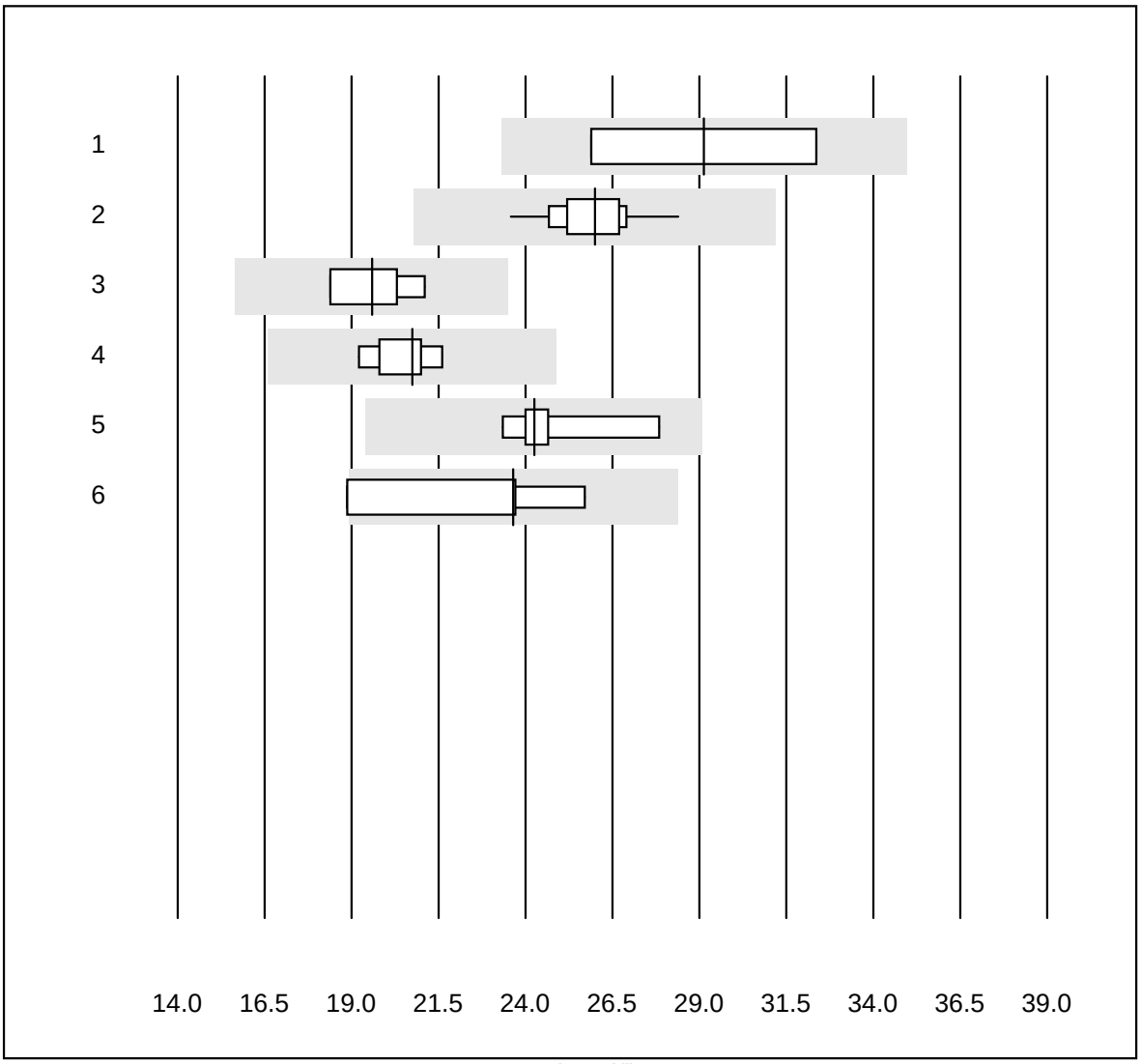
FT3



Deviatione QUALAB : 20 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas E / Elecsys	10	100.0	0.0	0.0	7.9	3.0	e
2	ADVIA Centaur XP/CP	4	100.0	0.0	0.0	6.4	4.0	e
3	Architect	7	100.0	0.0	0.0	6.0	7.1	e*
4	Vidas	4	100.0	0.0	0.0	6.6	3.4	e

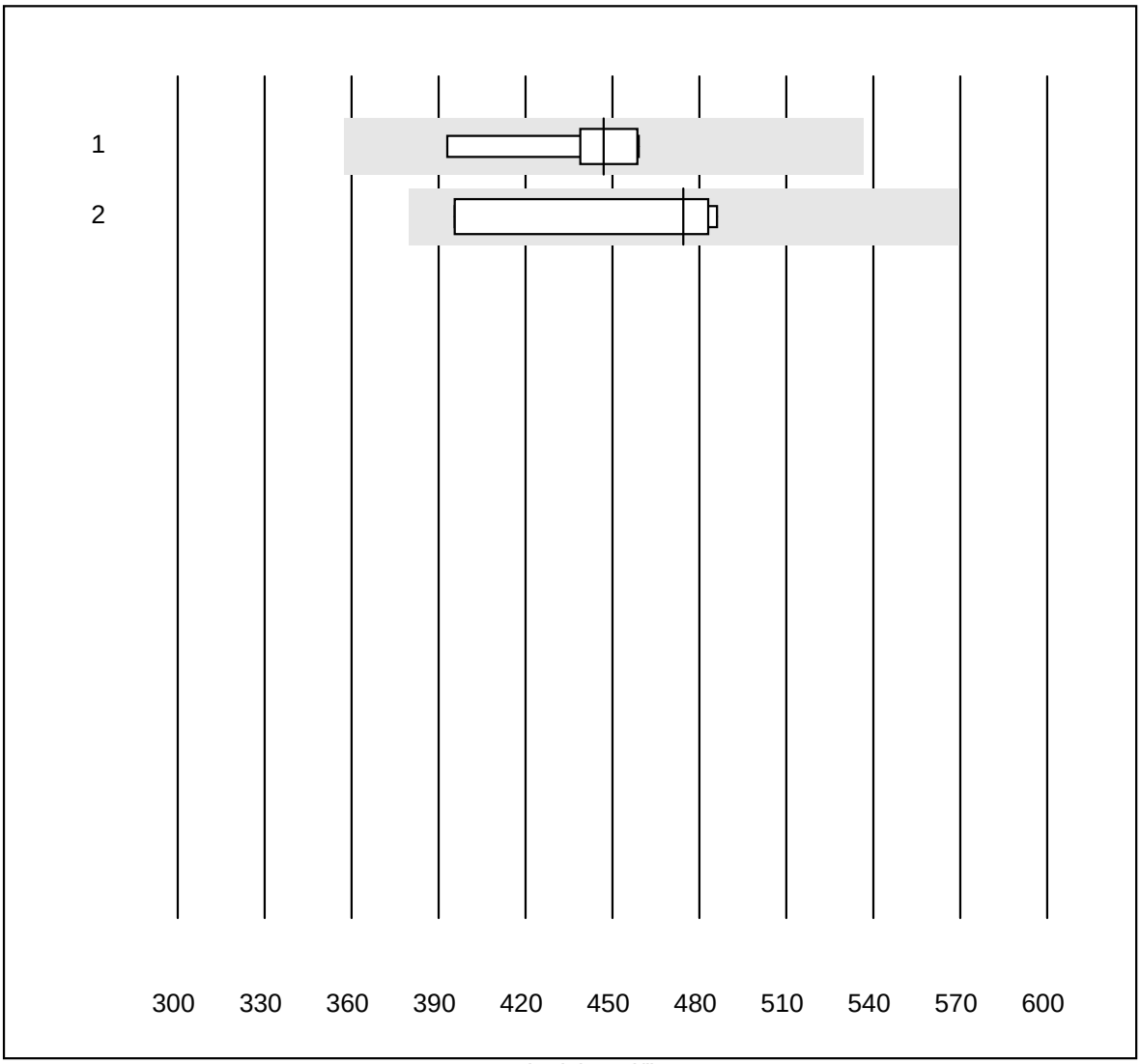
FT4



Deviatione QUALAB : 20 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Qualigen	4	75.0	0.0	25.0	29.1	13.3	e*
2	Cobas E / Elecsys	11	100.0	0.0	0.0	26.0	4.8	e
3	ADVIA Centaur XP	4	100.0	0.0	0.0	19.6	6.4	e*
4	Architect	8	100.0	0.0	0.0	20.7	3.8	e
5	Vidas	6	100.0	0.0	0.0	24.2	6.5	e*
6	altro	4	75.0	25.0	0.0	23.7	12.6	e*

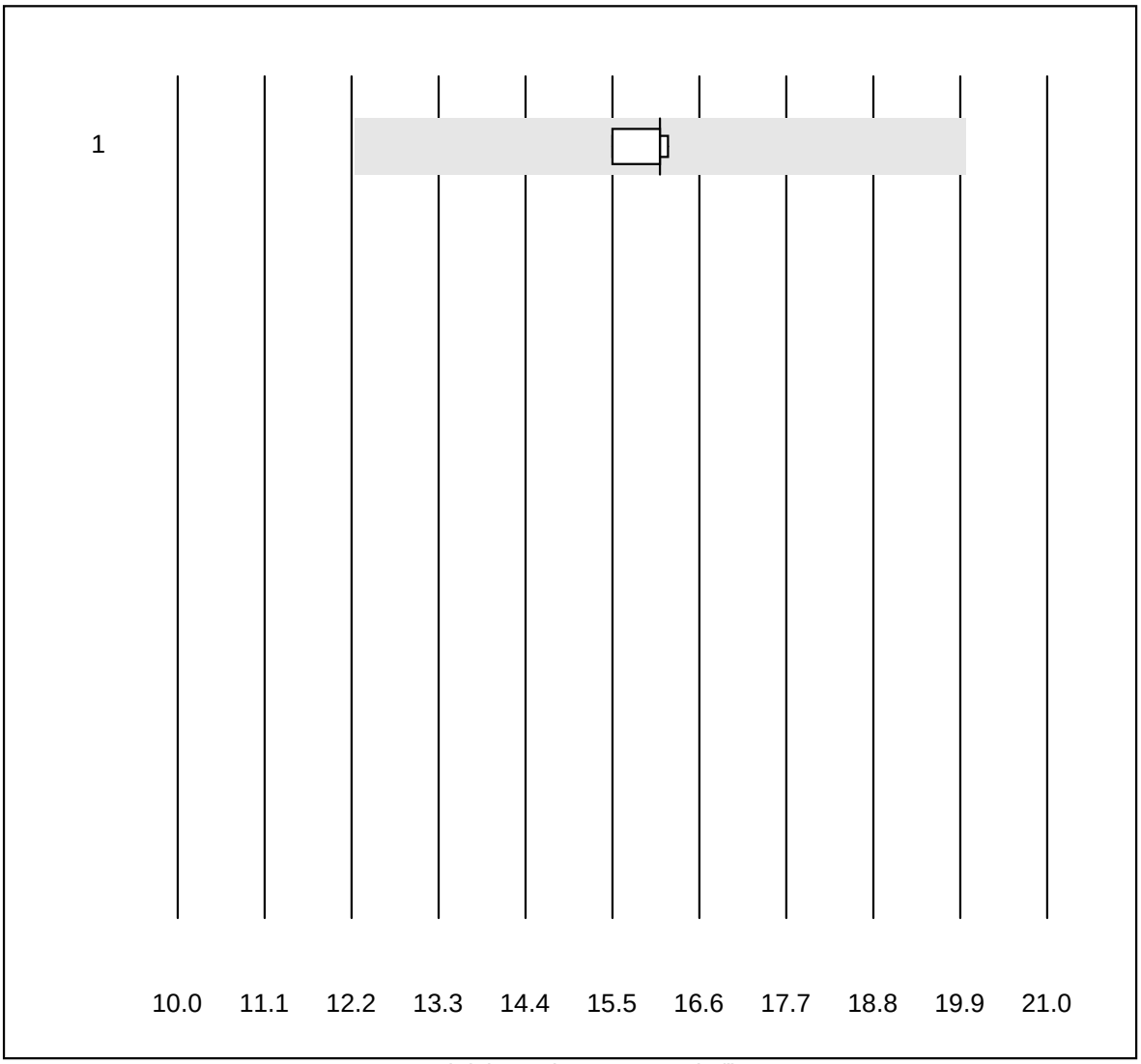
Cortisol



Deviatione QUALAB : 20 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas E / Elecsys	7	100.0	0.0	0.0	447	5.2	e
2	ADVIA Centaur XP/CP	4	100.0	0.0	0.0	475	9.2	e*

Luteinisierendes Hormon

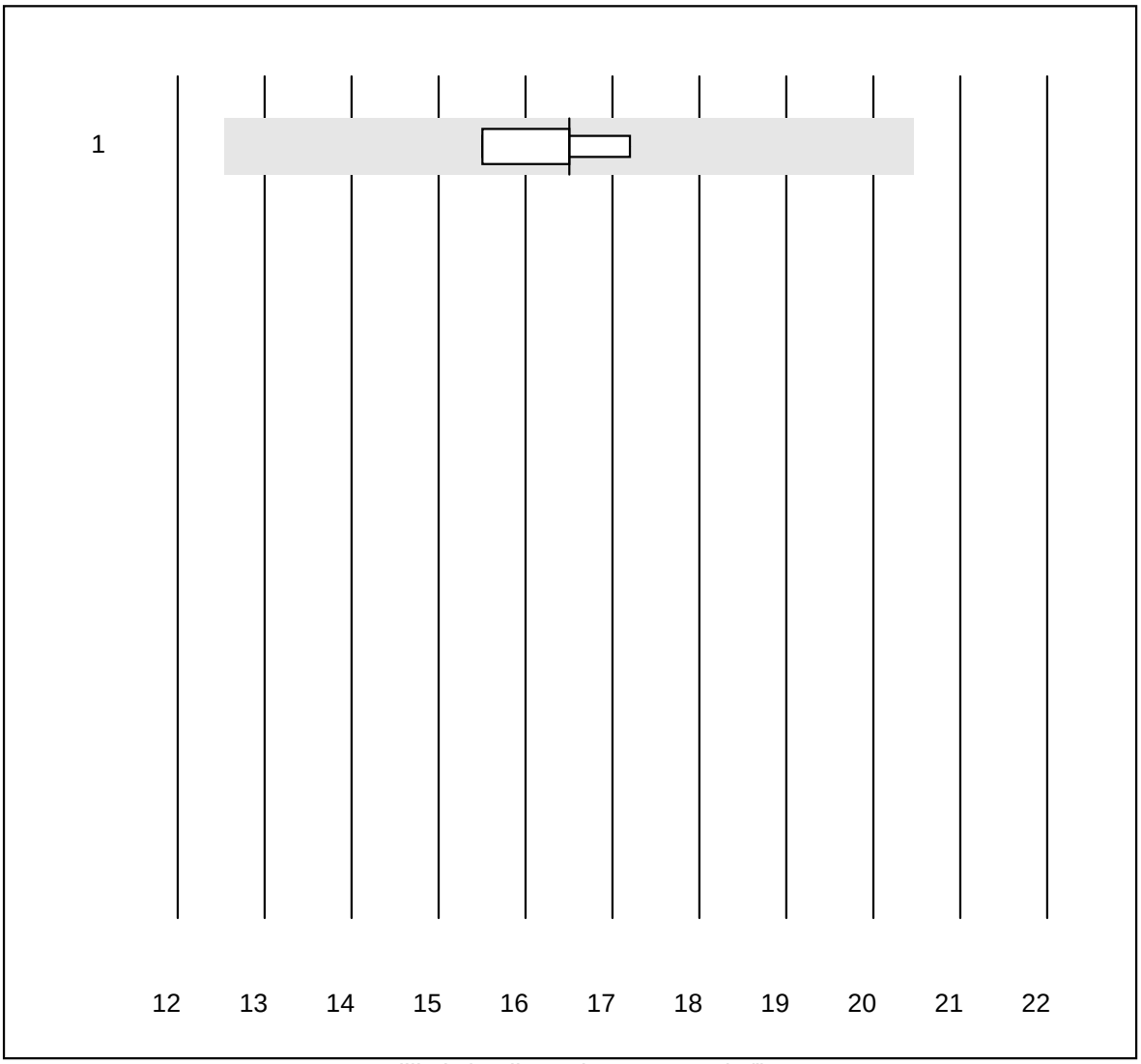


Deviatione QUALAB : 24 %

Luteinisierendes Hormon (U/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ADVIA Centaur XP/CP	4	100.0	0.0	0.0	16.1	2.0	e

Follikelstimulierendes Hormon

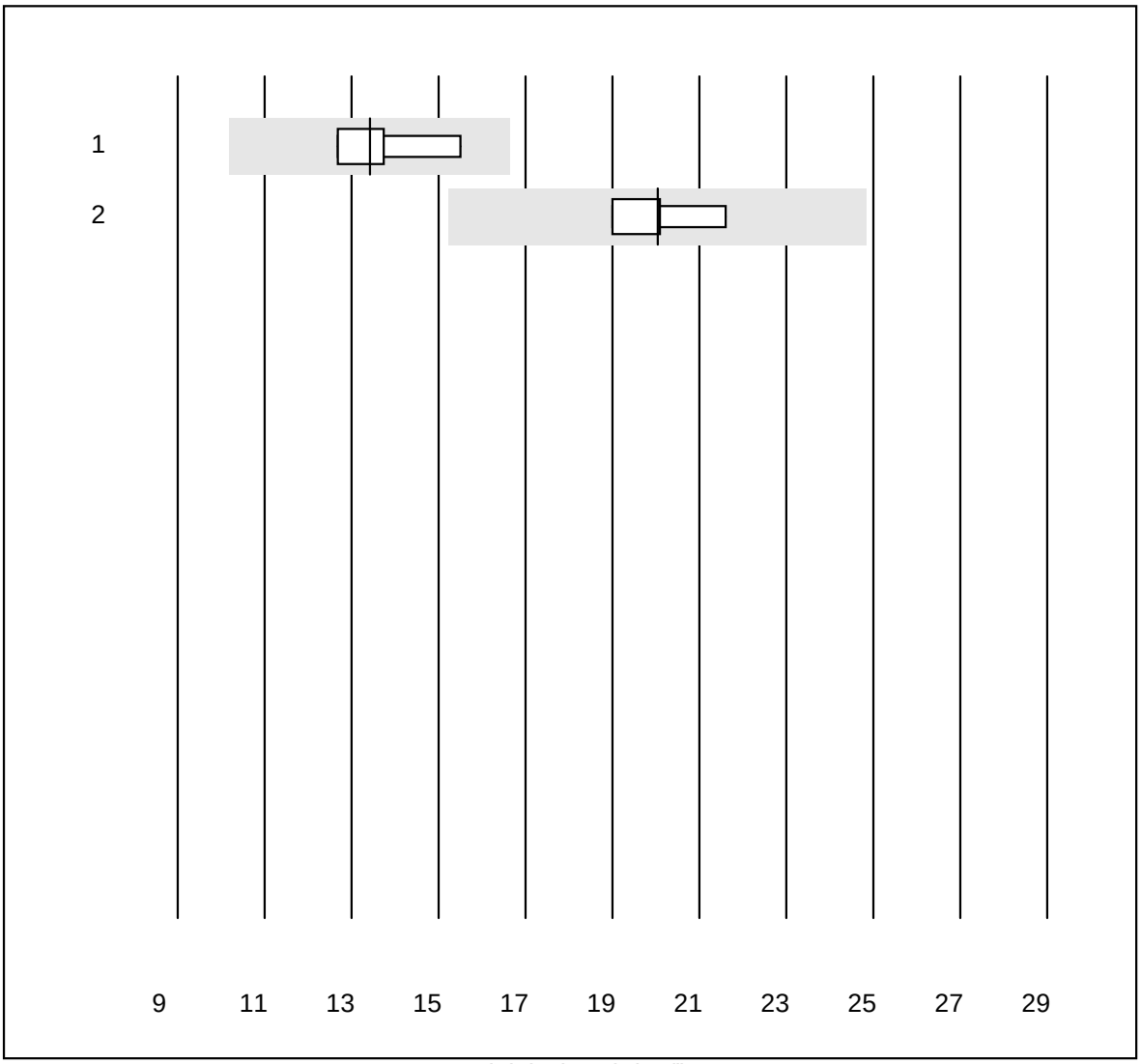


Deviatione QUALAB : 24 %

Follikelstimulierendes Hormon (U/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ADVIA Centaur XP/CP	4	100.0	0.0	0.0	16.5	4.3	e

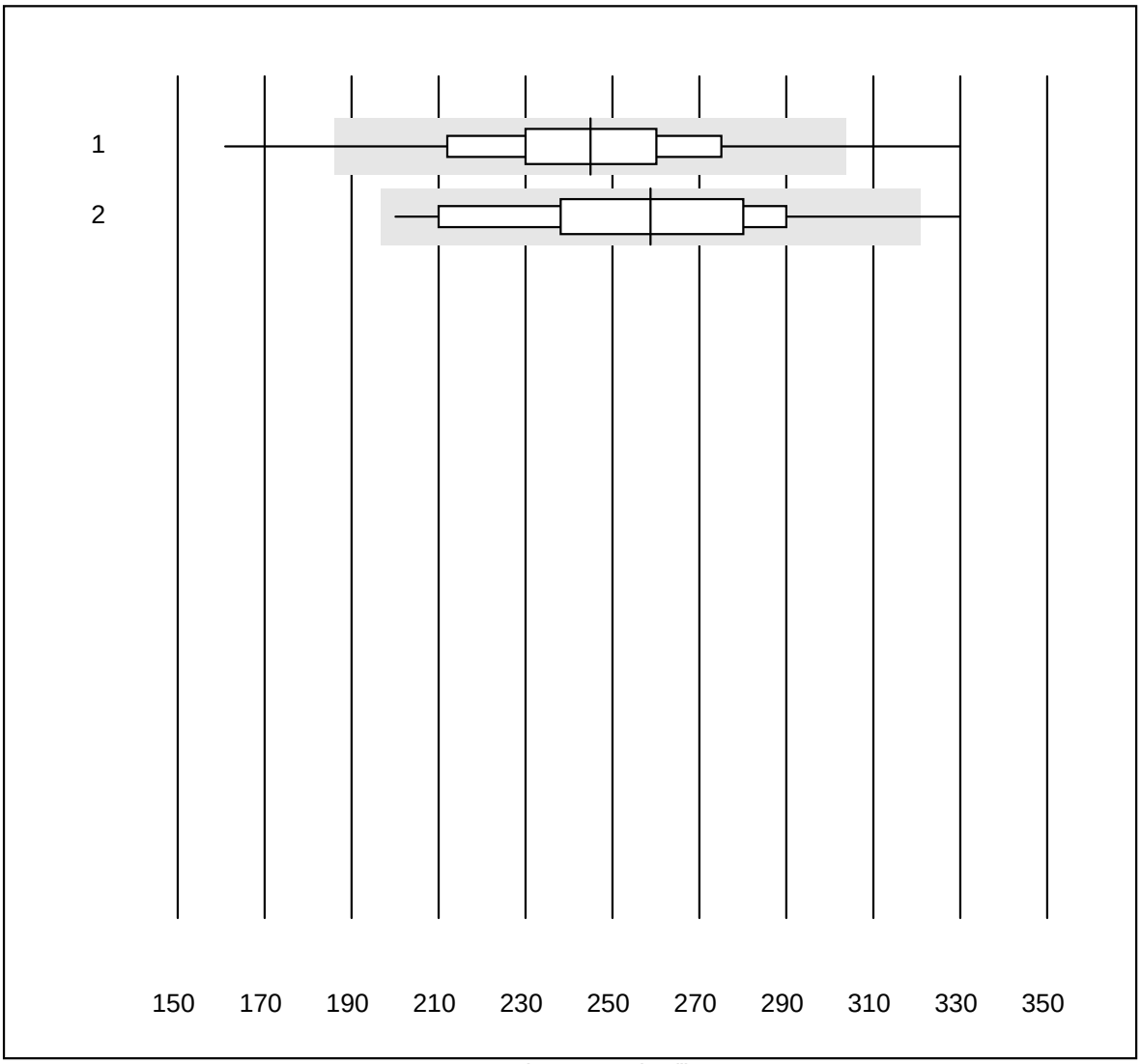
Prolaktin (PRL)



Deviatione QUALAB : 24 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ADVIA Centaur XP/CP	4	100.0	0.0	0.0	13.4	9.0	e*
2	Cobas/Roche	4	100.0	0.0	0.0	20.0	5.3	e

Troponina T CR

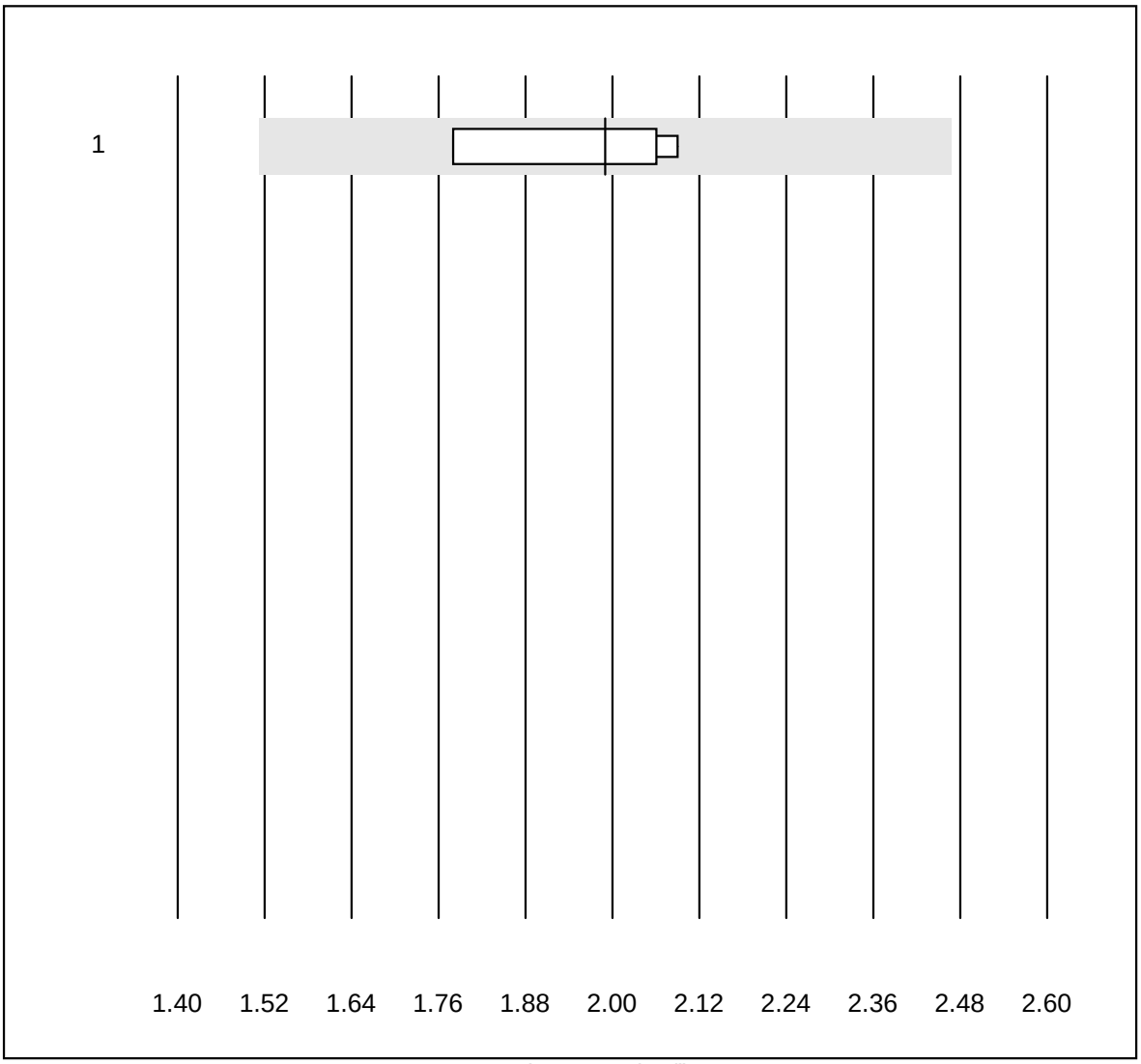


Deviatione QUALAB : 24 %

Troponina T CR (ng/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas h 232	782	97.1	2.3	0.6	244.93	10.0	e
2	Cardiac Reader	58	96.6	1.7	1.7	258.75	12.4	e

Troponina I WB

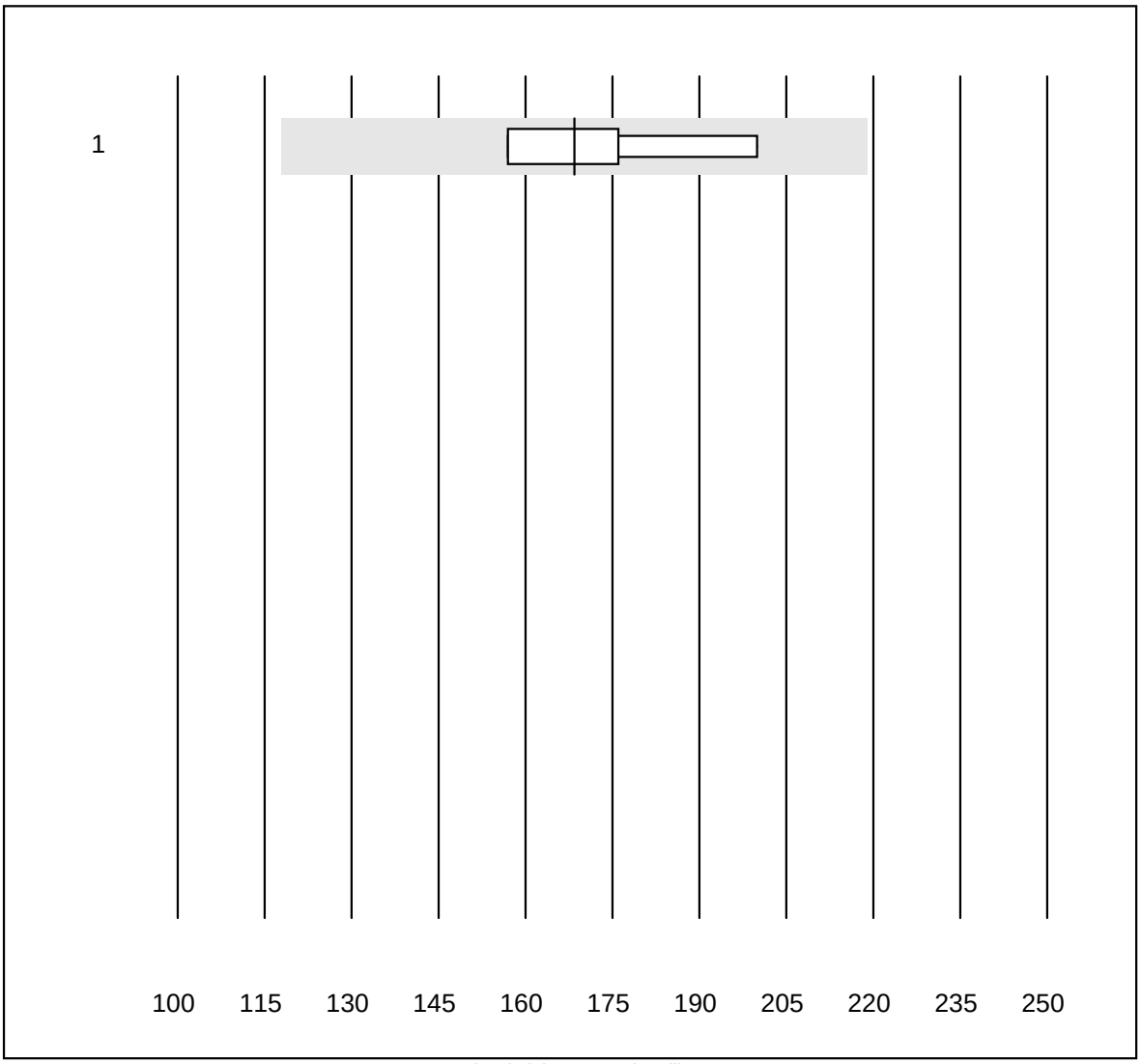


Deviatione QUALAB : 24 %

Troponina I WB (µg/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	iStat	4	100.0	0.0	0.0	1.99	7.3	e*

Mioglobina CR

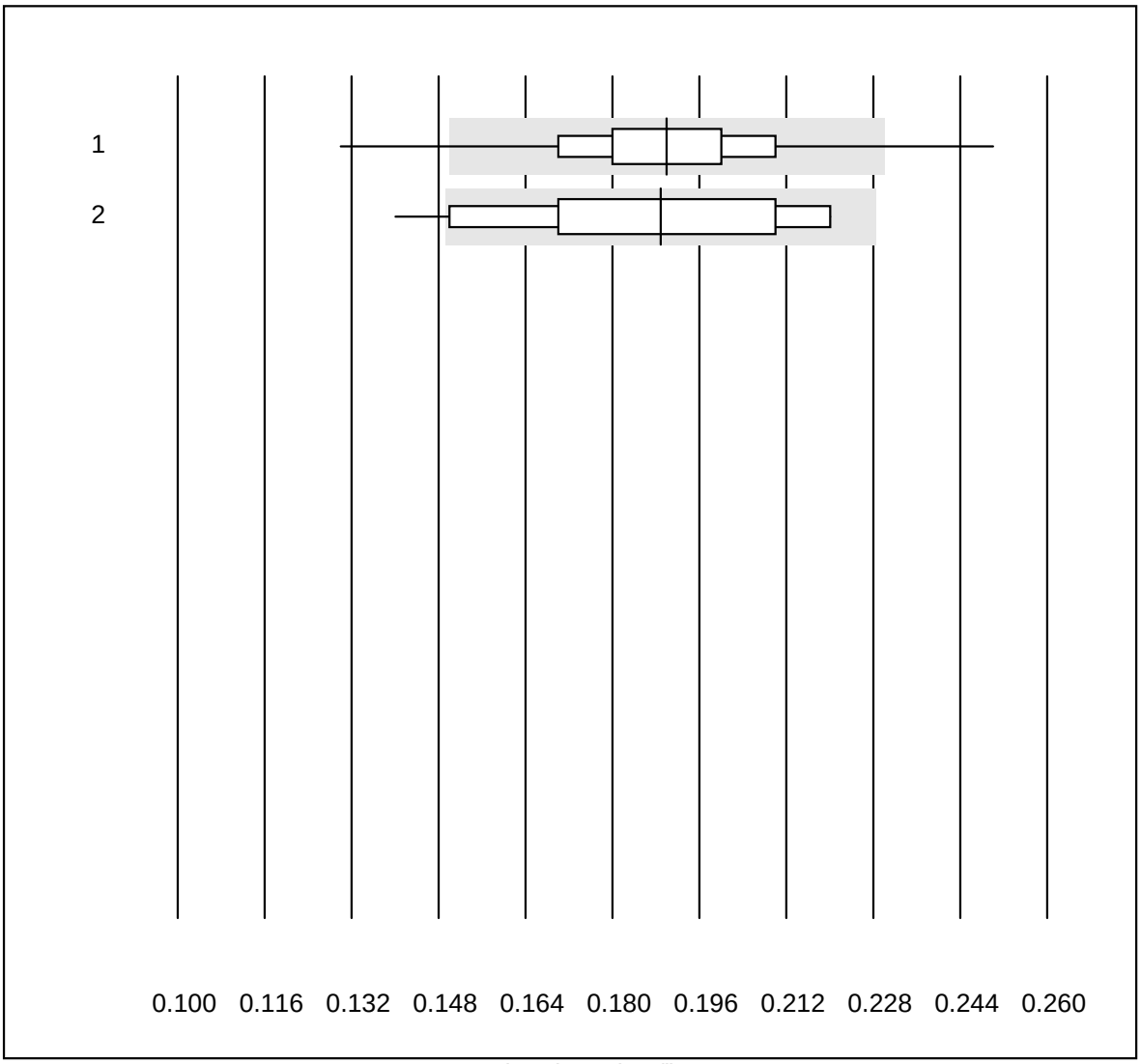


Deviatione QUALAB : 30 %

Mioglobina CR (µg/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas h 232	4	100.0	0.0	0.0	168.5	11.2	e*

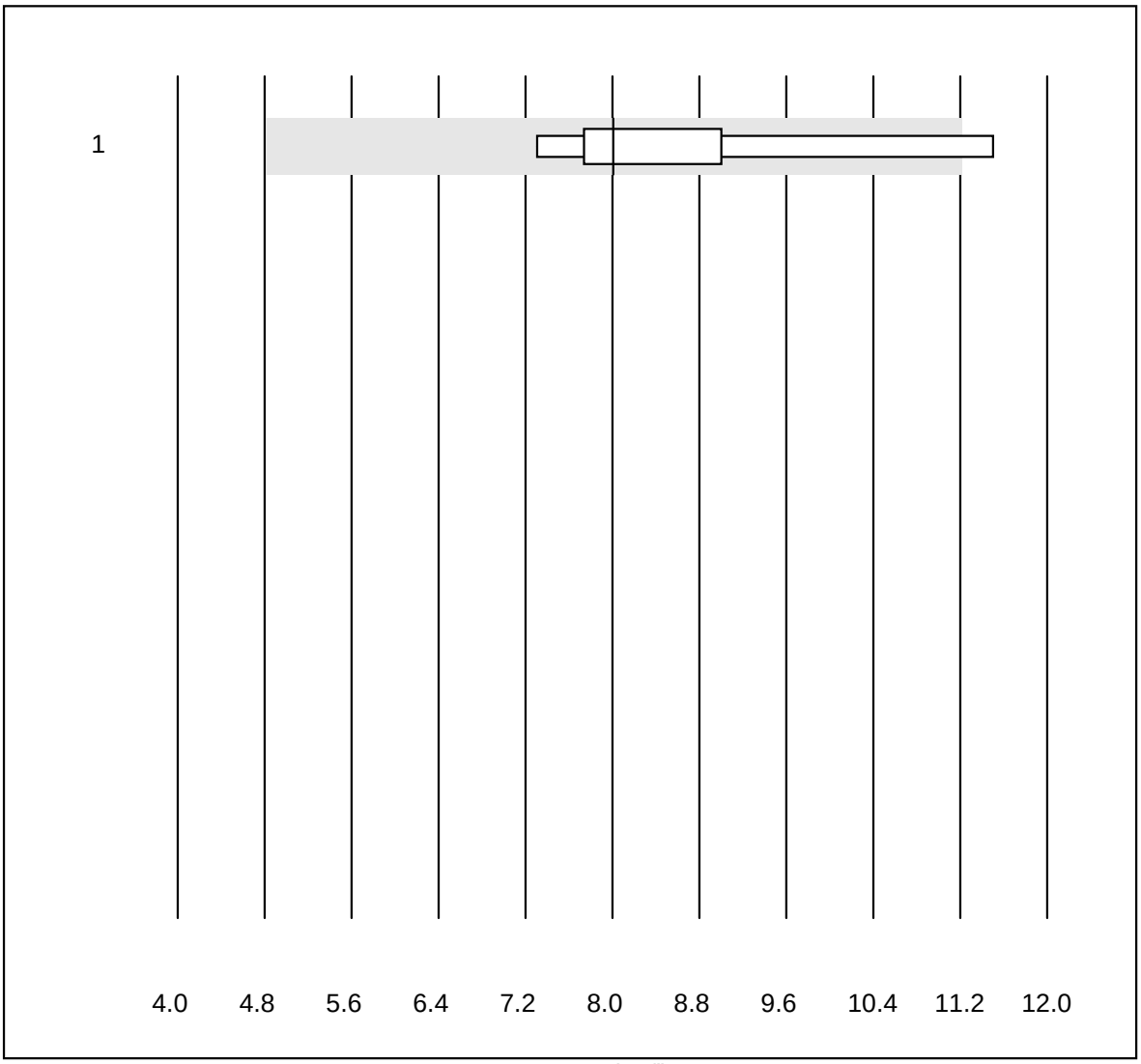
D-Dimeri CR



Deviatione QUALAB : 21 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas h 232	810	93.2	5.1	1.7	0.19	9.5	e
2	Cardiac Reader	51	90.2	3.9	5.9	0.19	12.5	e

CKMB - K8

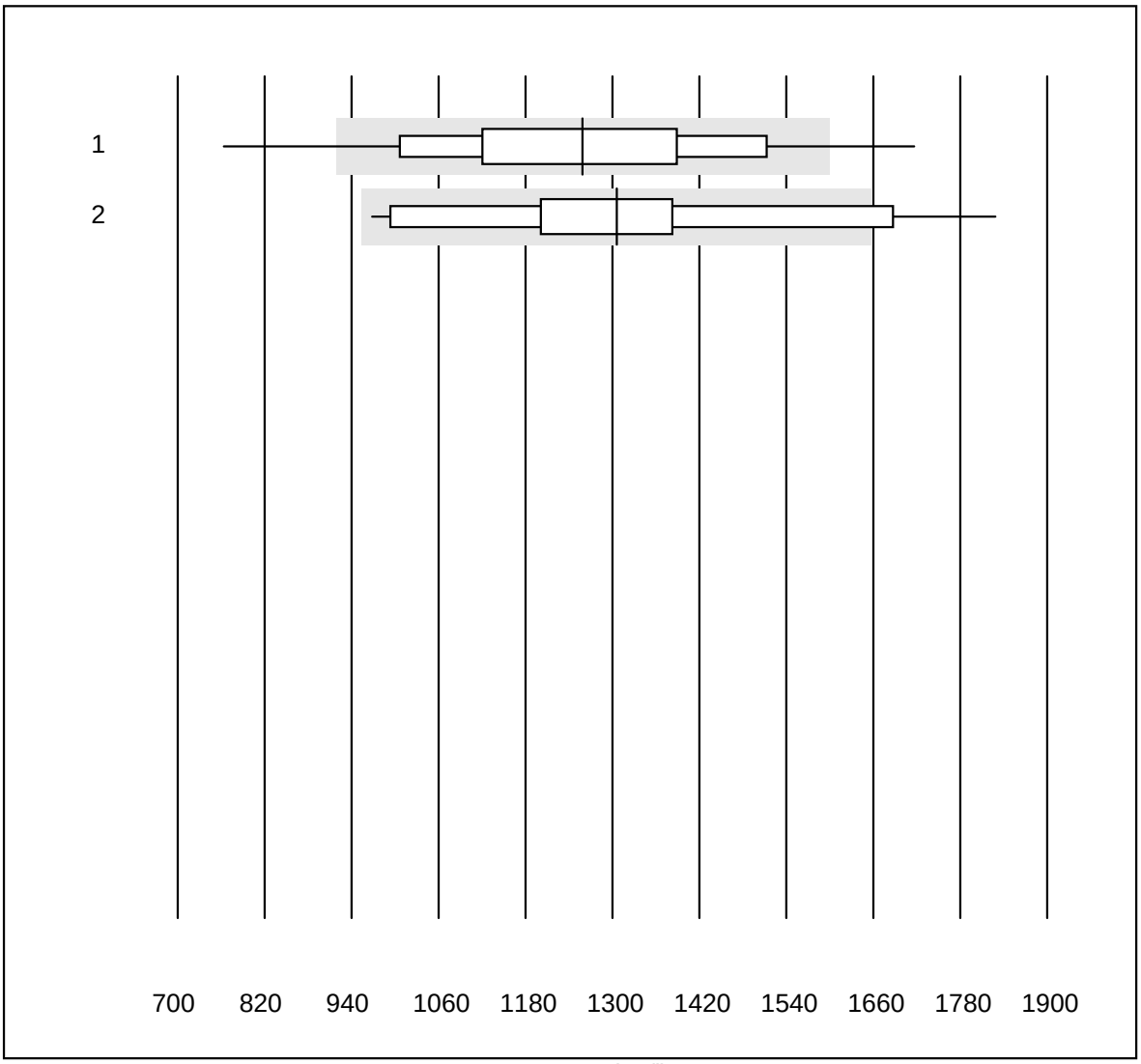


Deviatione QUALAB : 40 %

CKMB - K8 (µg/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas h 232	9	88.9	11.1	0.0	8.0	16.6	e*

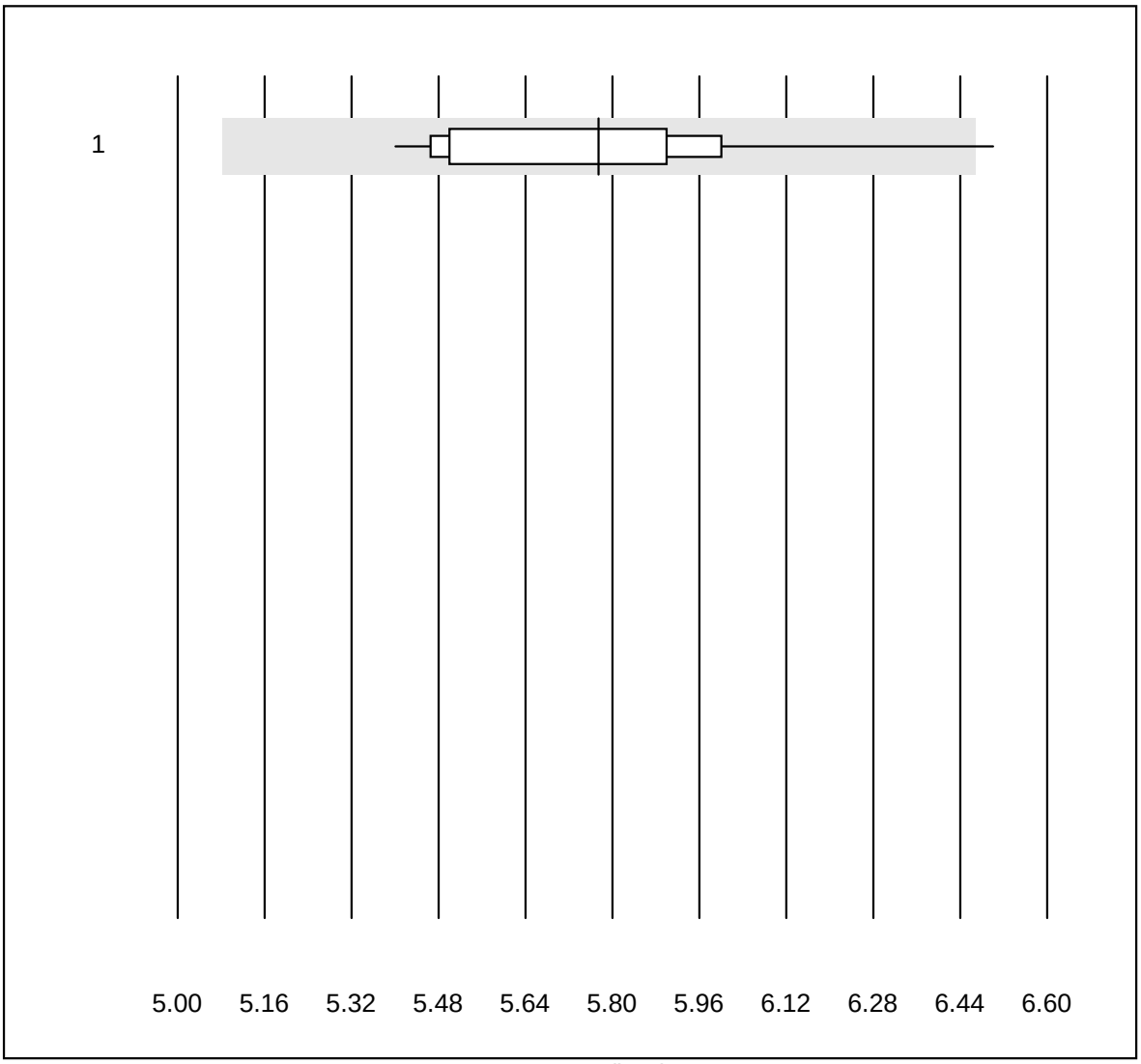
proBNP CR



Deviatione QUALAB : 27 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas h 232	506	91.3	5.5	3.2	1259	15.0	e
2	Cardiac Reader	17	76.4	11.8	11.8	1307	18.1	e*

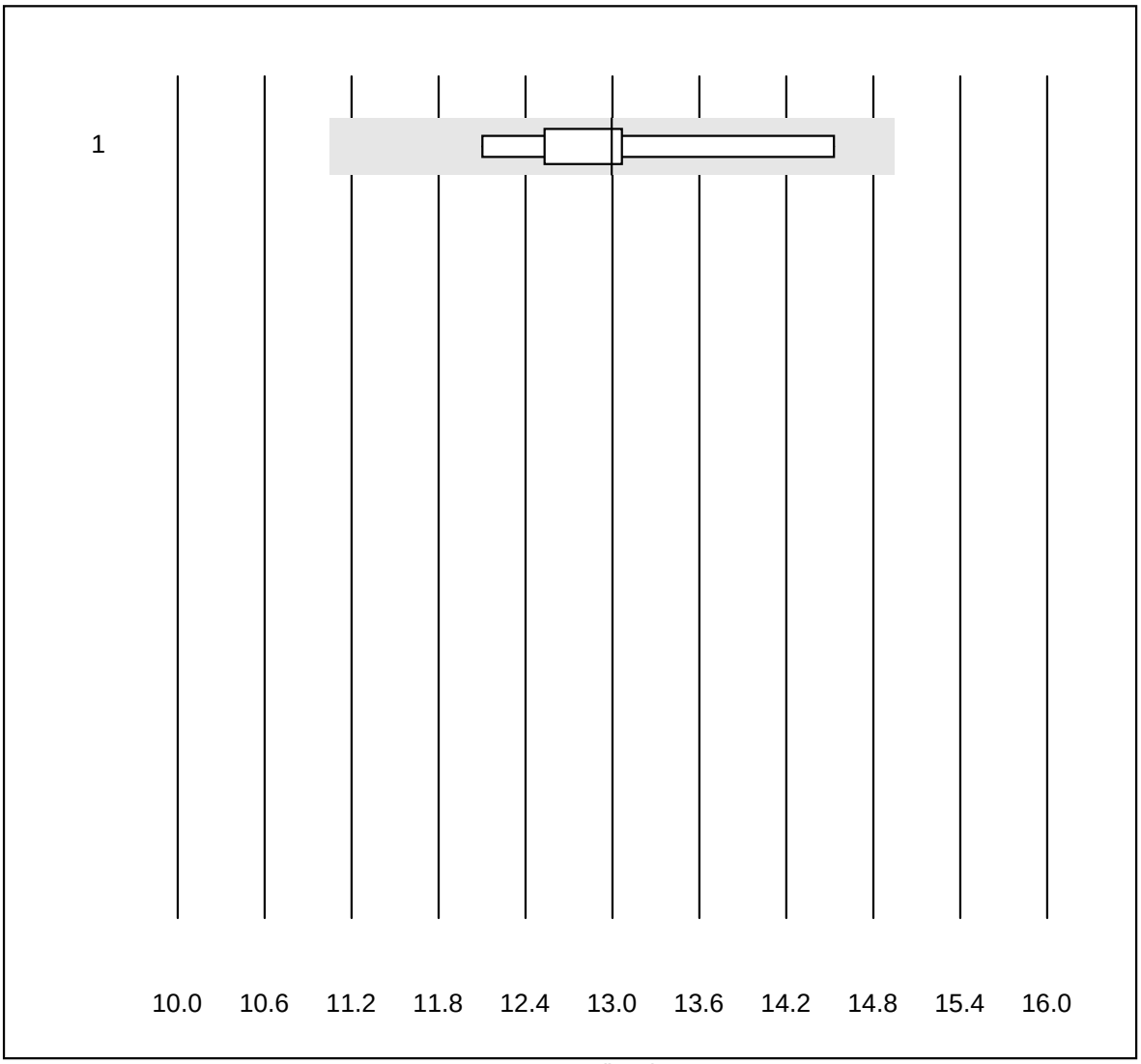
PCO2 CCA



Deviatione QUALAB : 12 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	OPTI CCA	11	90.9	9.1	0.0	5.77	5.4	e*

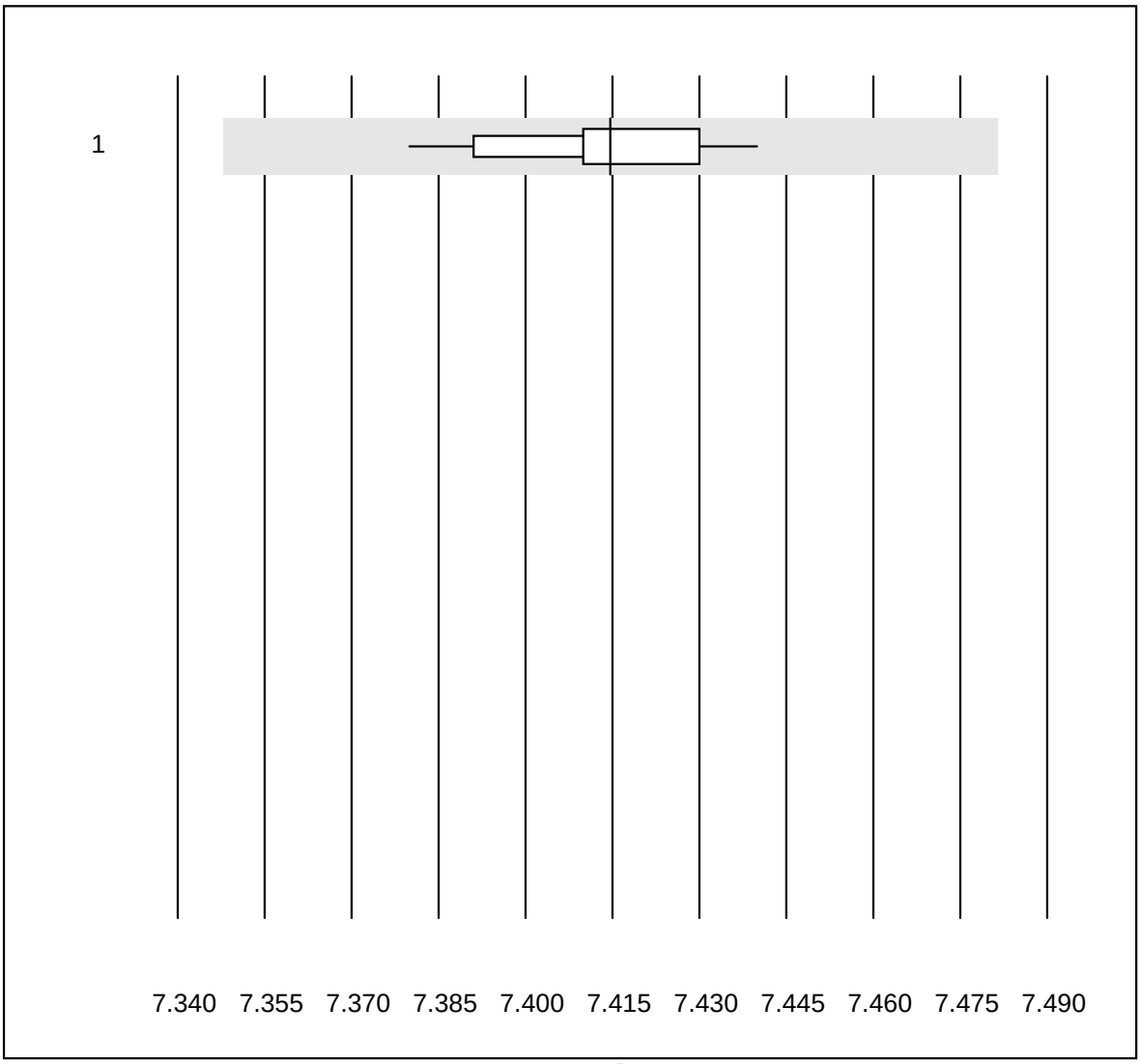
PO2 CCA



Deviatione QUALAB : 15 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	OPTI CCA	11	81.8	0.0	18.2	13.00	6.5	e*

pH CCA

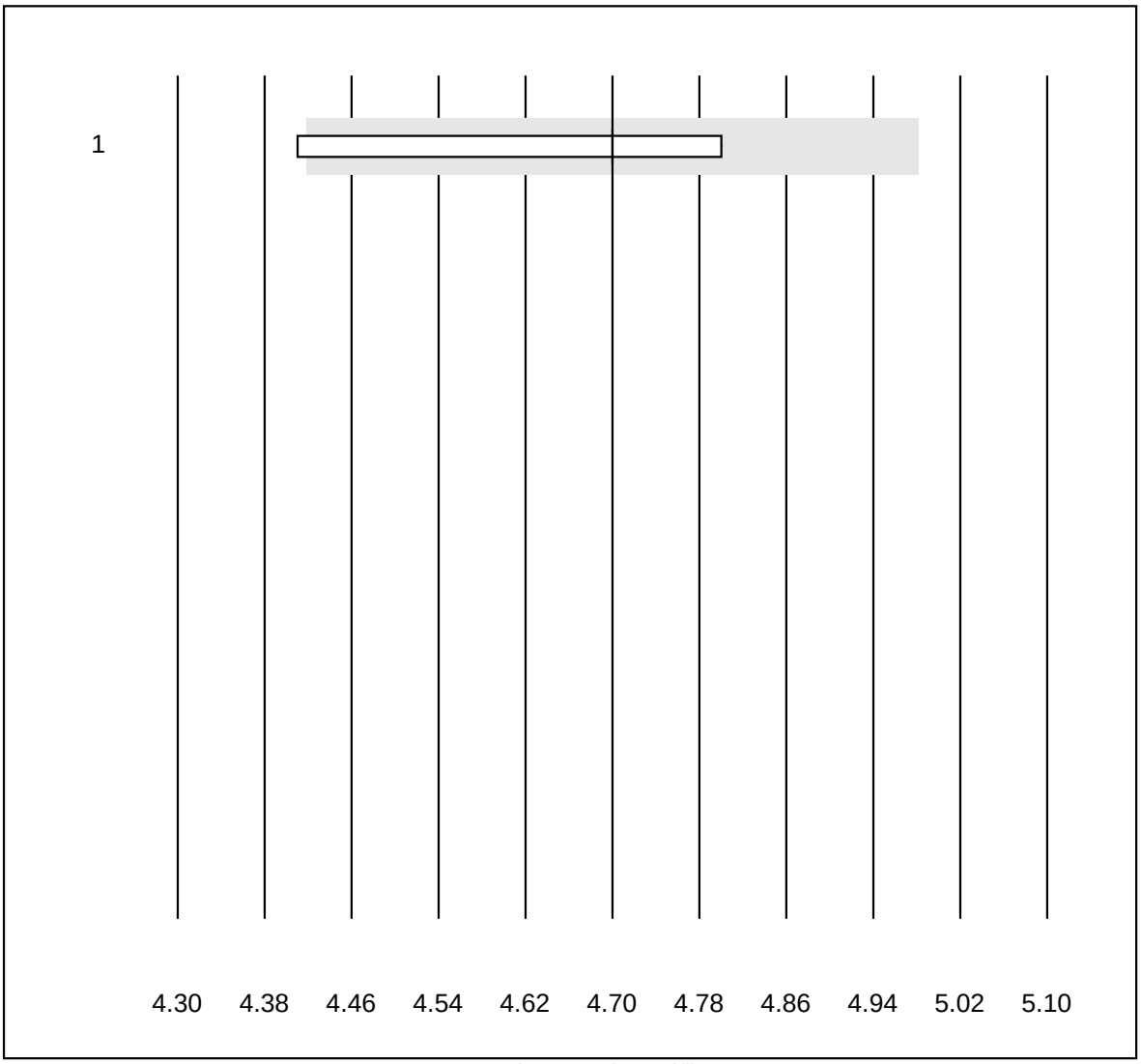


Deviatione QUALAB : 1 %

pH CCA ()

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	OPTI CCA	11	100.0	0.0	0.0	7.41	0.2	e

Potassio CCA

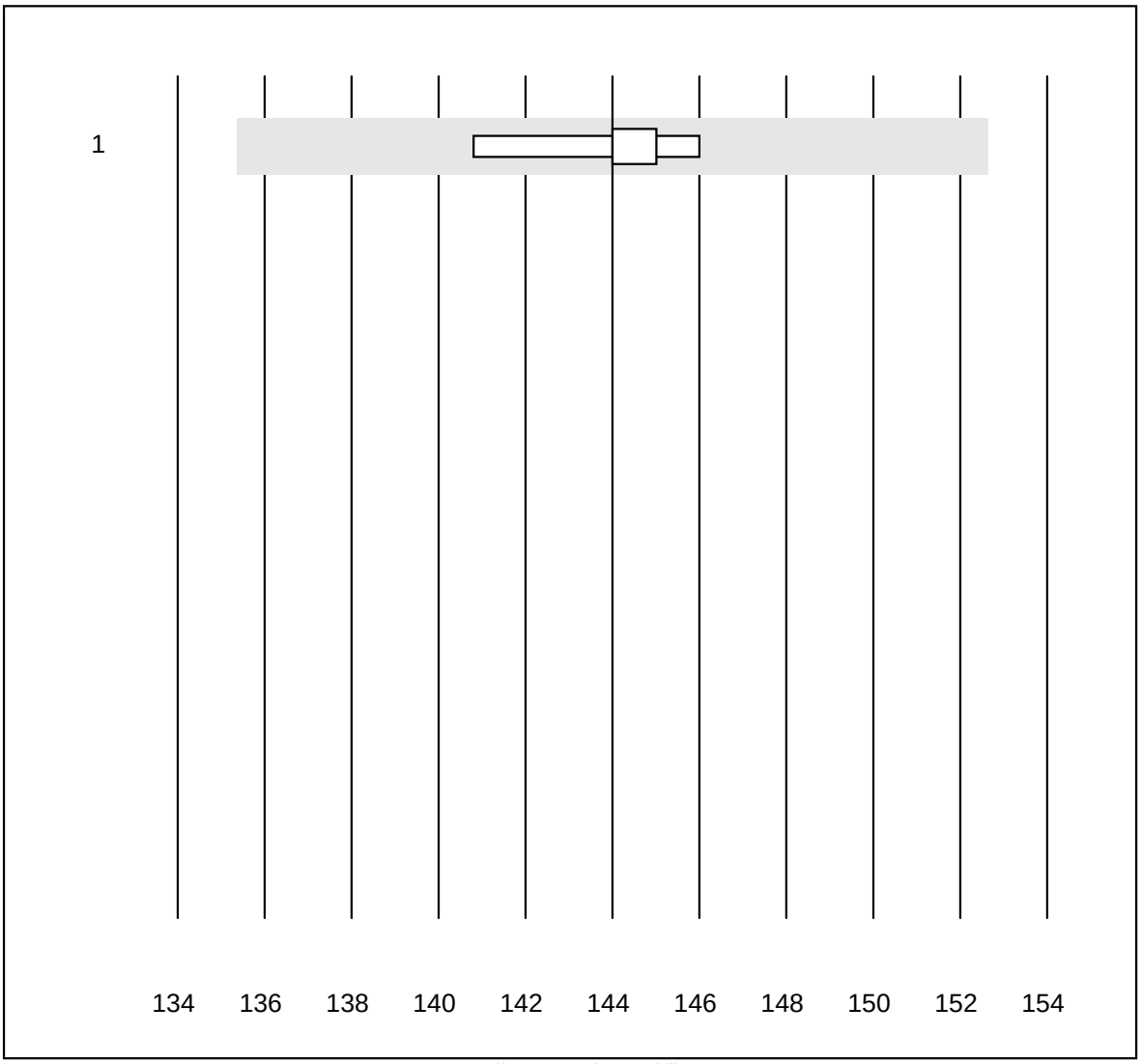


Deviatione QUALAB : 6 %

Potassio CCA (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	OPTI CCA	6	83.3	16.7	0.0	4.7	2.8	e*

Sodio CCA

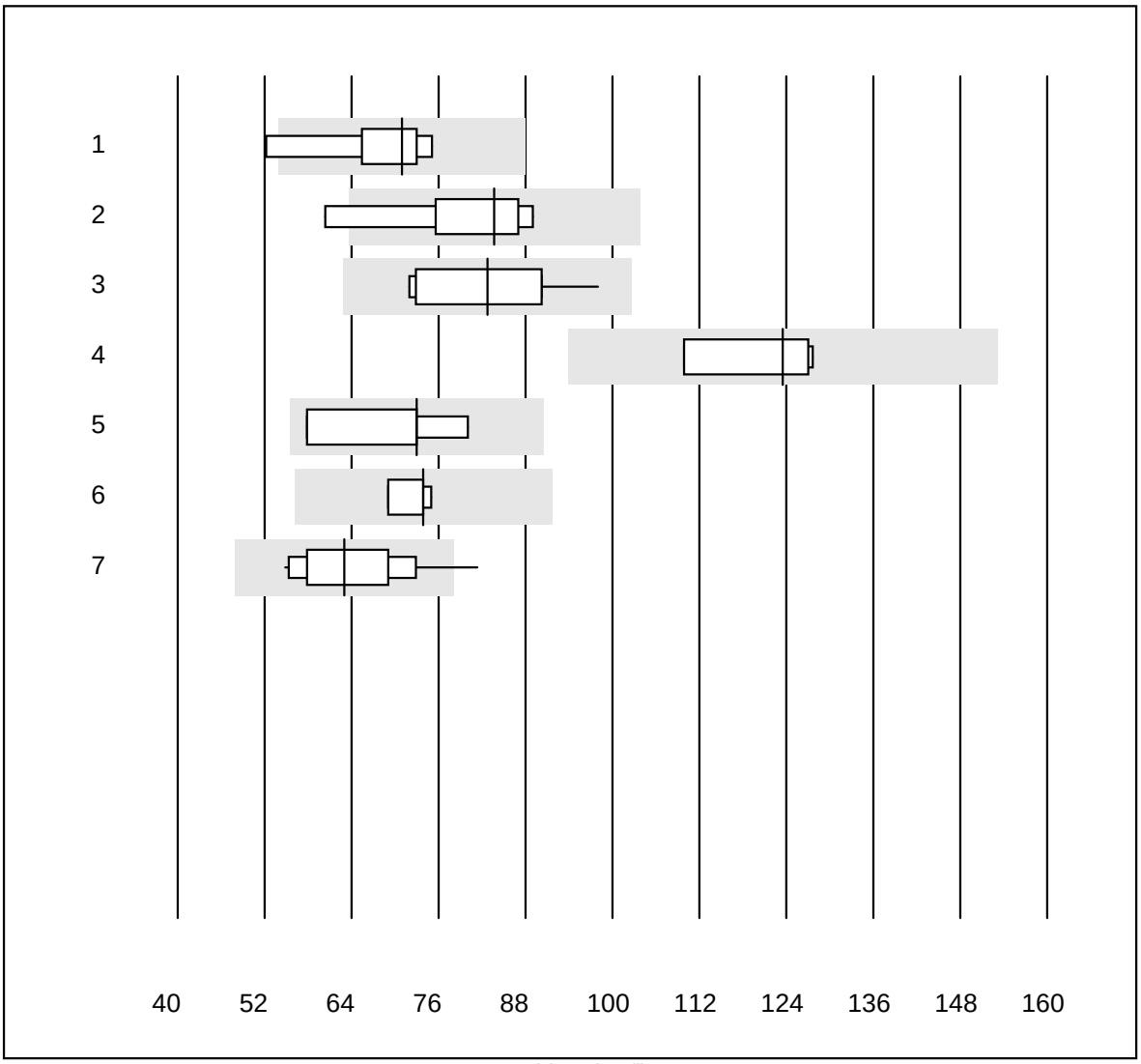


Deviatione QUALAB : 6 %

Sodio CCA (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	OPTI CCA	5	100.0	0.0	0.0	144.0	1.4	e

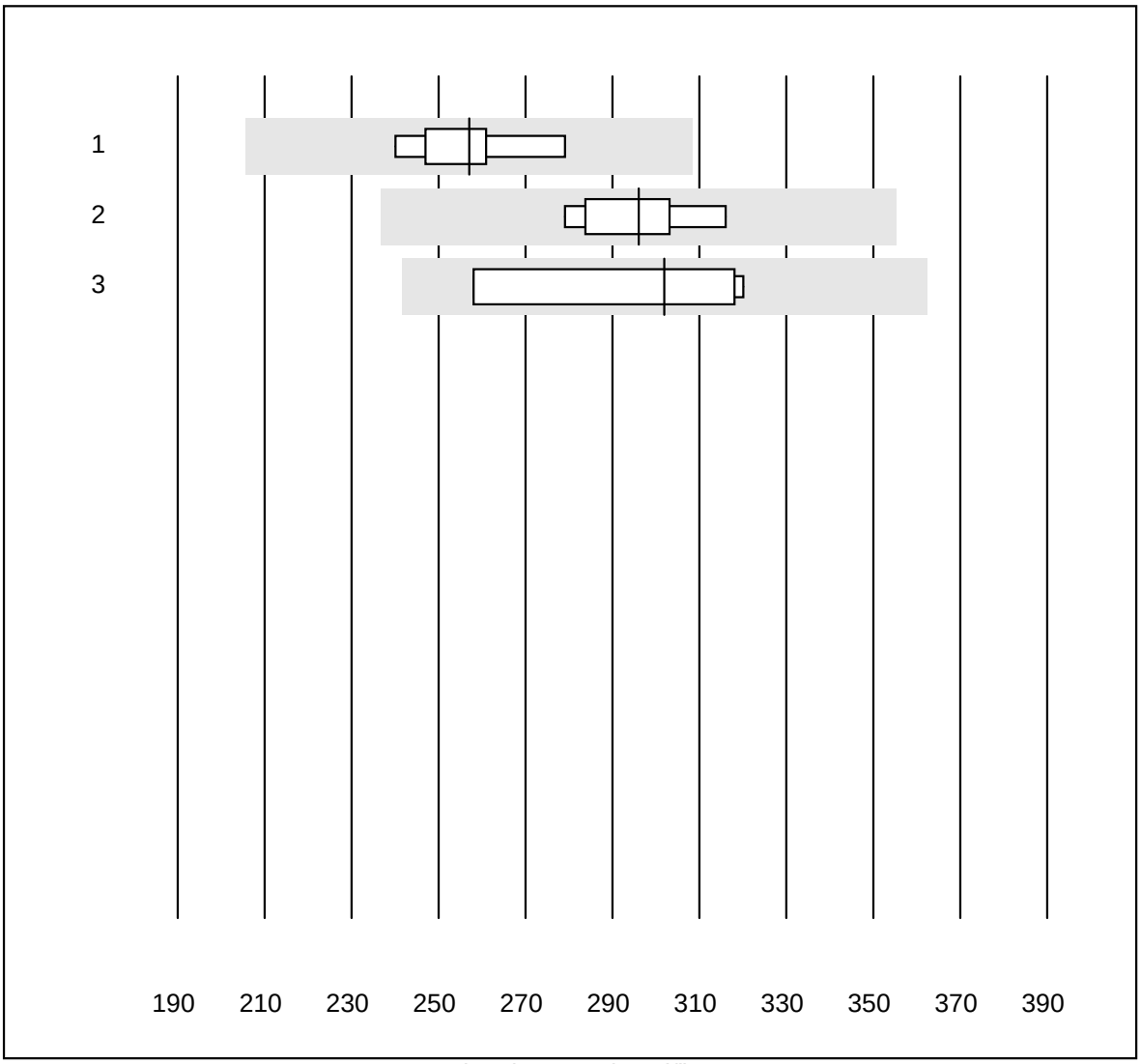
Ferritina



Deviatione QUALAB : 24 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Beckman	5	80.0	20.0	0.0	71.00	13.7	e*
2	tutti	5	80.0	20.0	0.0	83.70	14.7	e*
3	Cobas E / Elecsys	10	100.0	0.0	0.0	82.80	11.5	e*
4	Architect	4	100.0	0.0	0.0	123.54	6.8	e*
5	Mira/DiaSys	4	100.0	0.0	0.0	73.00	13.1	e*
6	Mini Vidas	4	100.0	0.0	0.0	73.87	3.7	e
7	Eurolyser	18	94.4	5.6	0.0	63.02	11.5	e

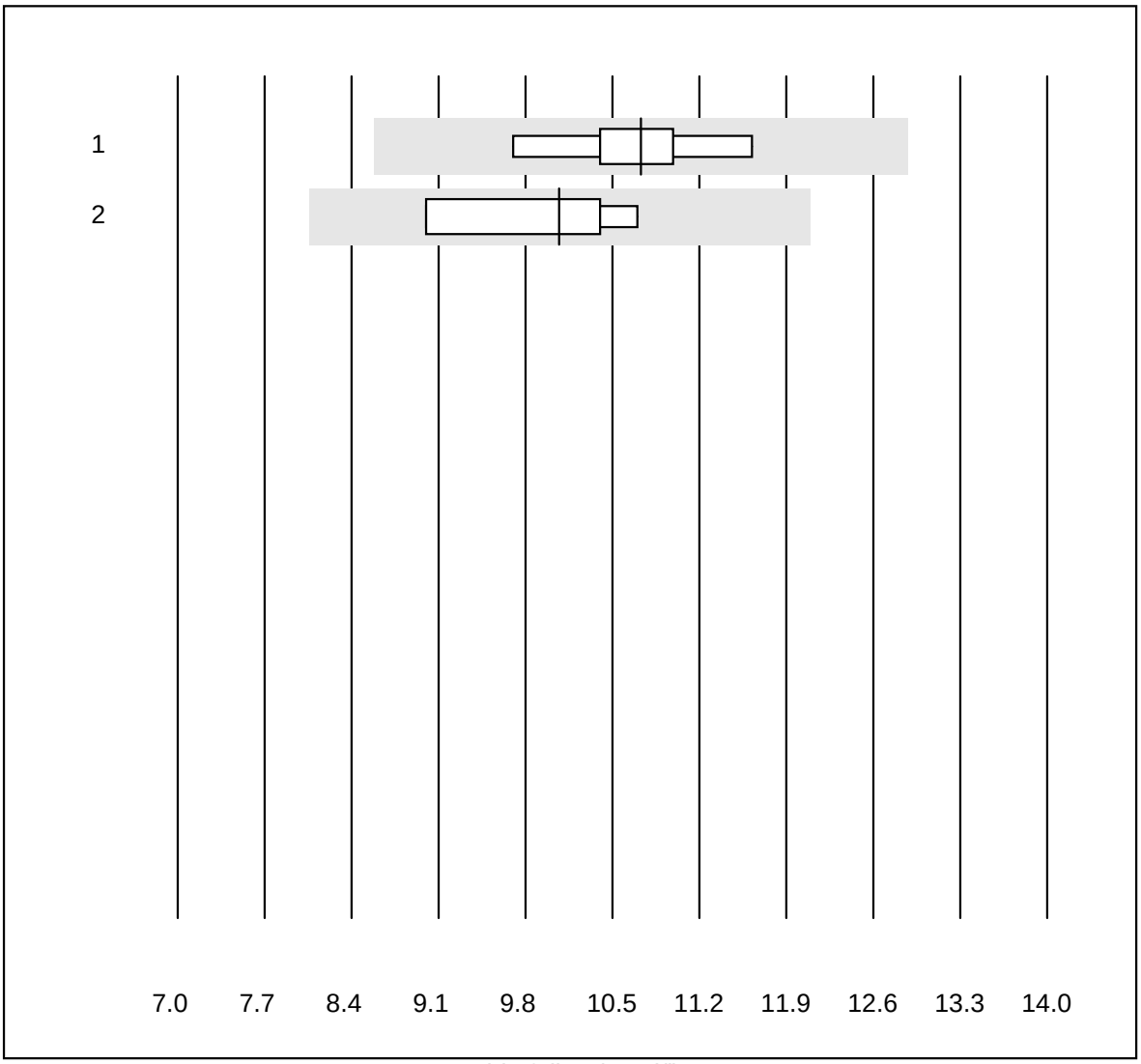
Vitamina B12



Deviatione QUALAB : 20 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ADVIA Centaur XP/CP	5	100.0	0.0	0.0	257.00	5.8	e*
2	Cobas E / Elecsys	8	100.0	0.0	0.0	296.00	4.6	e
3	Architect	4	100.0	0.0	0.0	301.96	10.0	e*

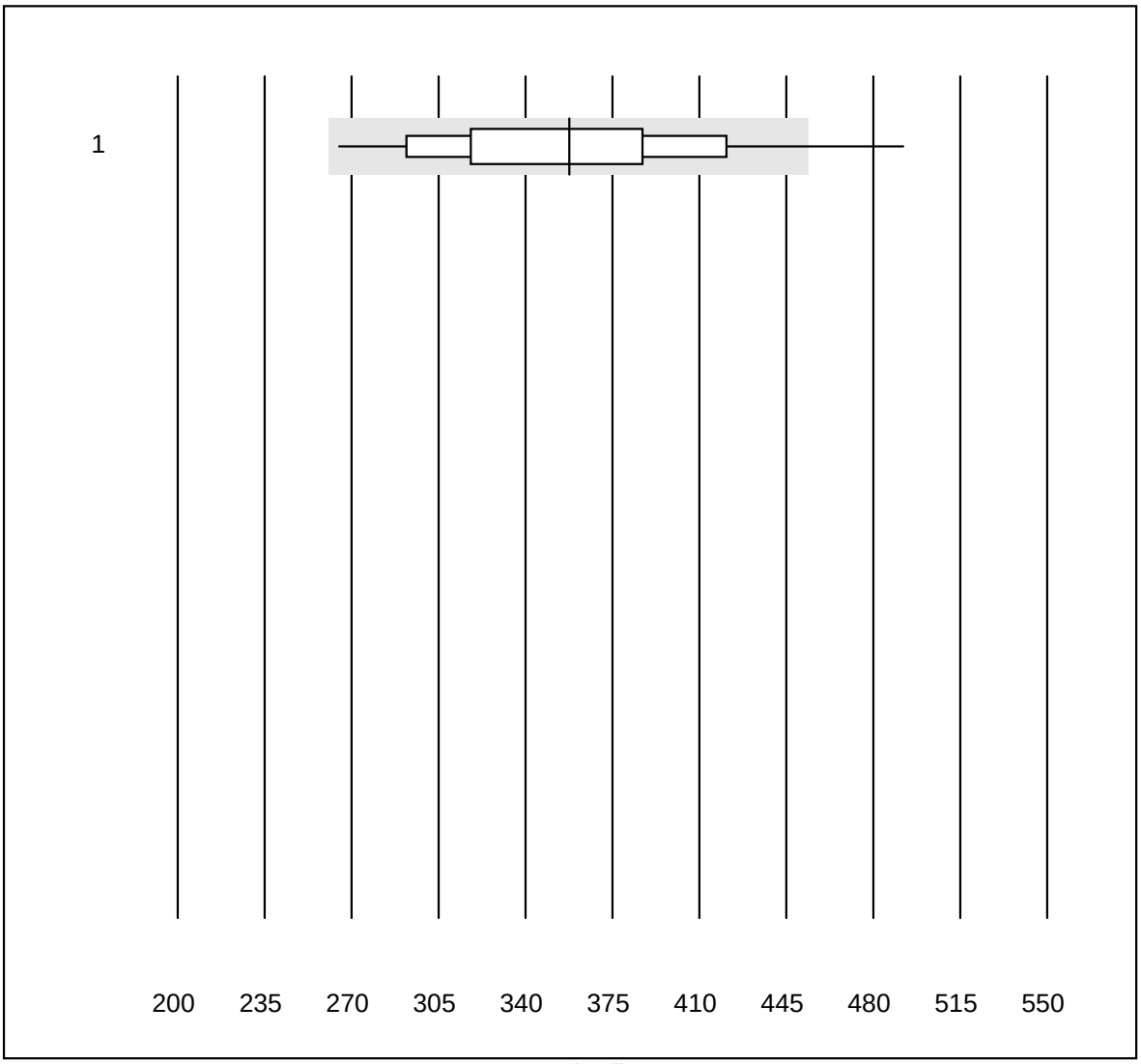
Acido folico



Deviatione QUALAB : 20 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas E / Elecsys	8	100.0	0.0	0.0	10.73	5.5	e
2	Architect	4	100.0	0.0	0.0	10.07	7.6	e*

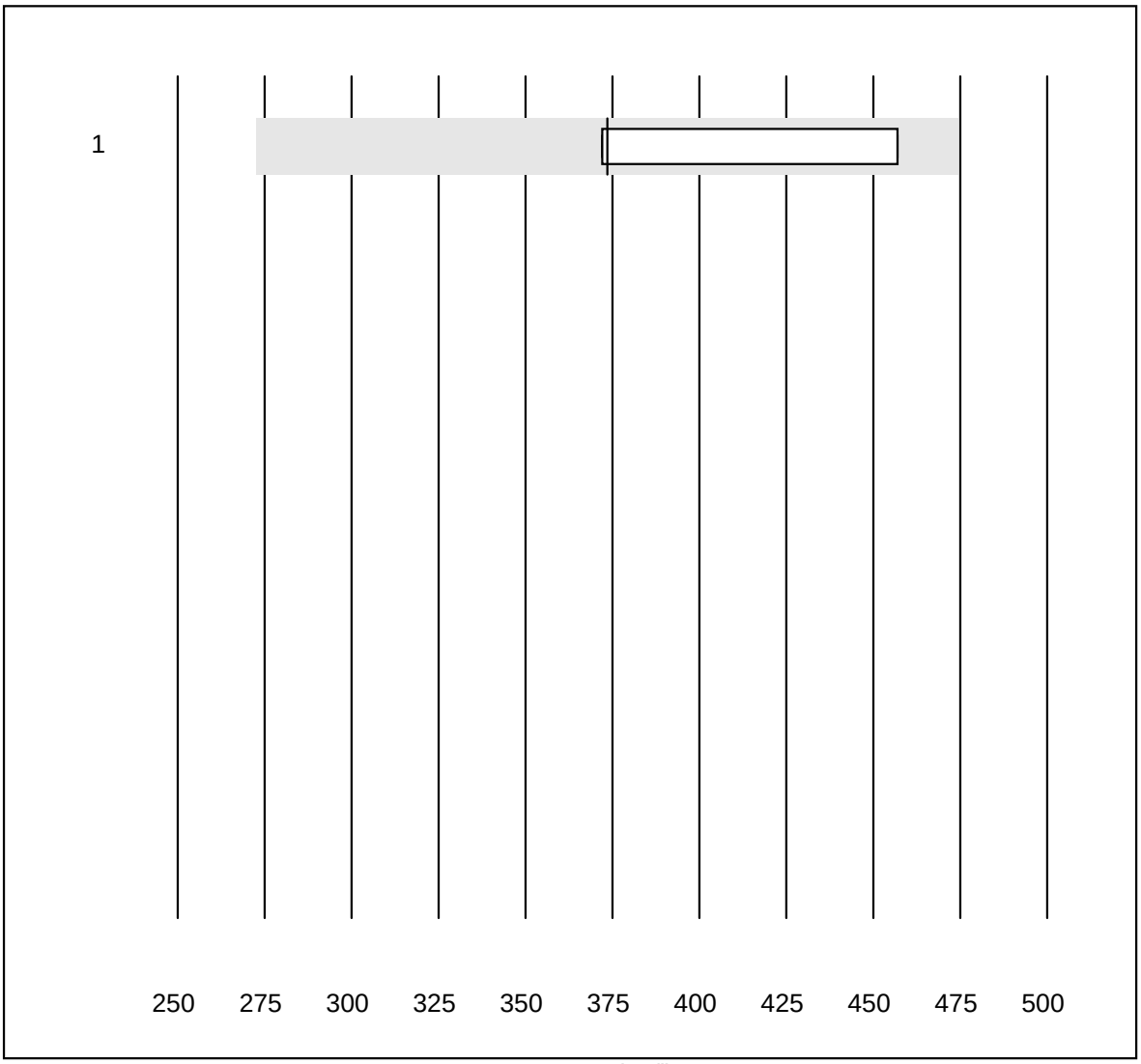
BNP



Deviatione QUALAB : 27 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Triage Meter	45	93.4	4.4	2.2	357.5	14.3	e

NT-Pro-BNP

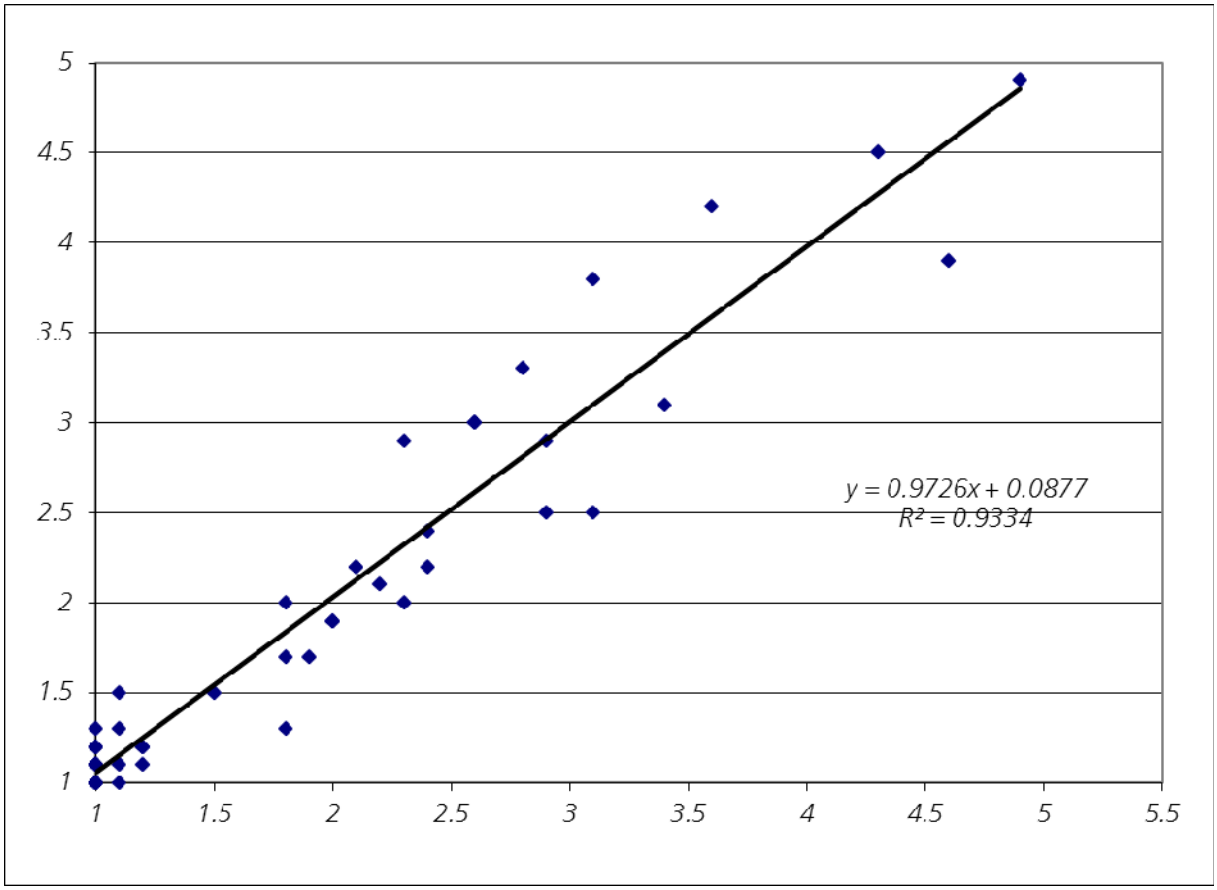


Deviatione QUALAB : 27 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Triage Meter	4	75.0	0.0	25.0	374	12.0	e*

Quick / INR WB

Ospedale universitario Zurigo

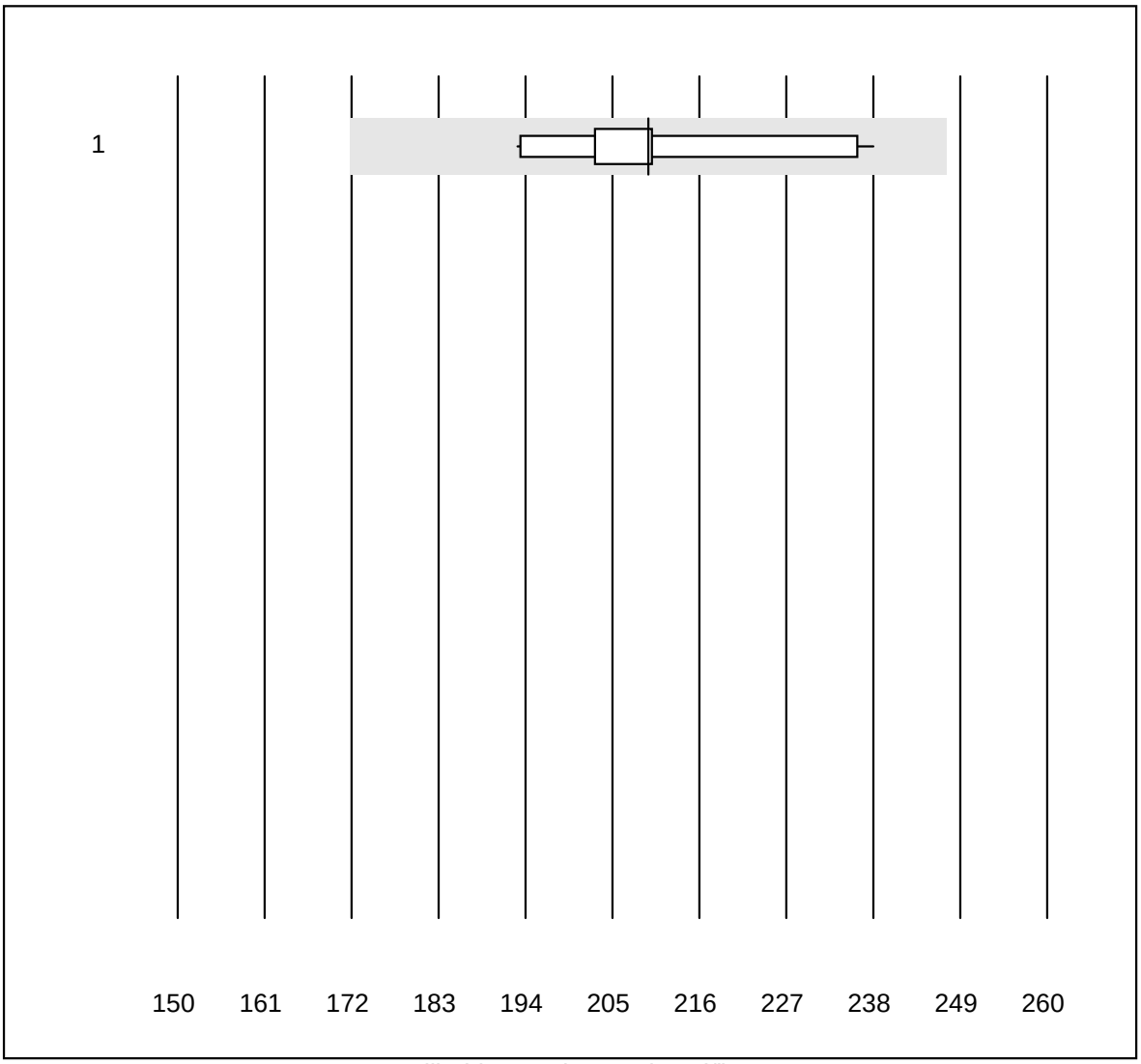


INRatio partecipanti

Nell'ambito del controllo circolare G10 vengono confrontati gli INR dei partecipanti con quelli dell'ospedale universitario di Zurigo.

Nr.	metodo	totale	% conforme	% insuff.	% outlier
1	INRatio	55	83.64	9.09	7.27

Bilirubina totale Neo

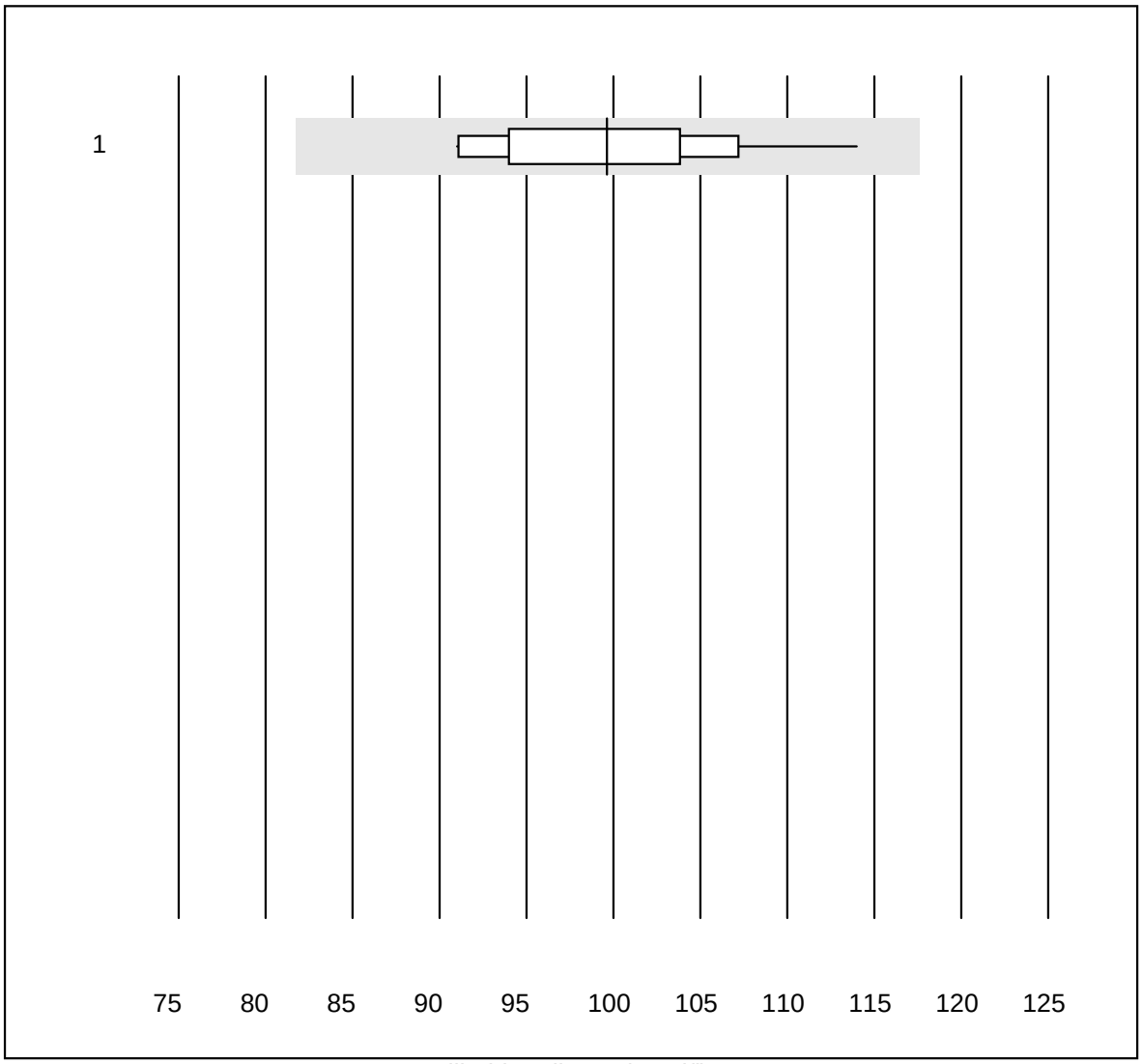


Deviatione QUALAB : 18 %

Bilirubina totale Neo (µmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	tutti	13	100.0	0.0	0.0	210	6.8	e

Bilirubina diretta

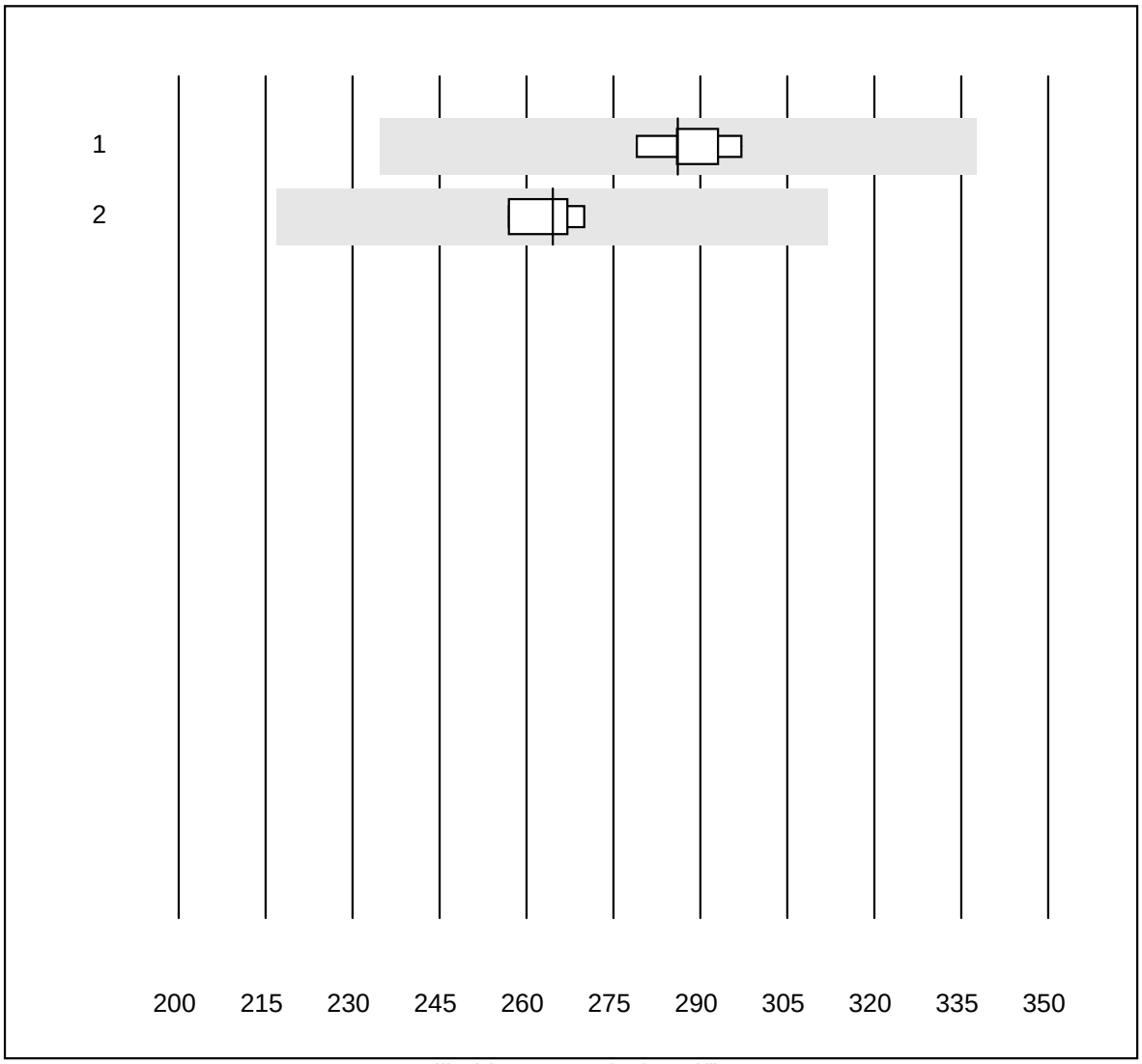


Deviatione QUALAB : 18 %

Bilirubina diretta (µmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	tutti	13	92.3	0.0	7.7	100	7.2	e

Bilirubin neonatale

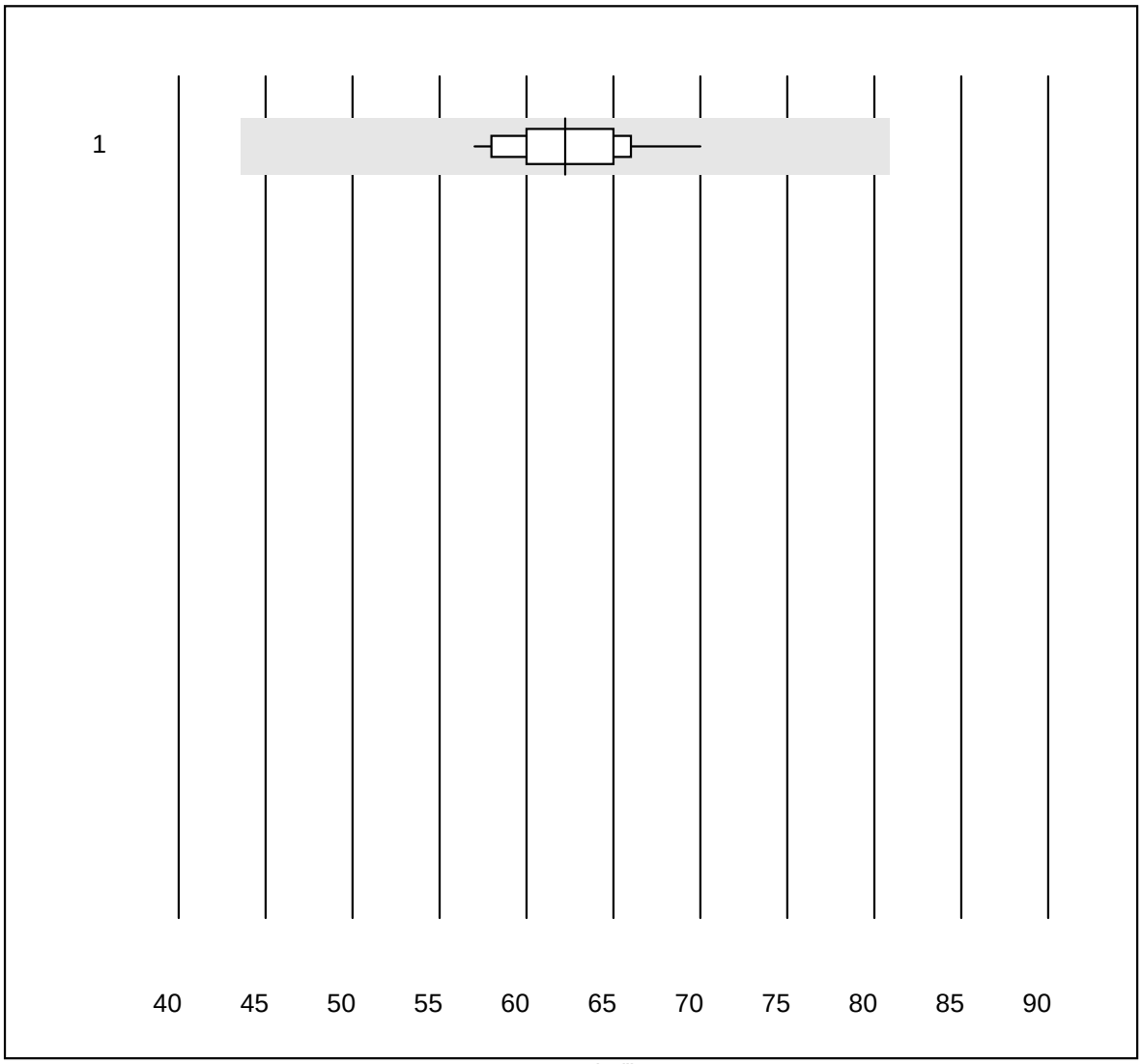


Deviatione QUALAB : 18 %

Bilirubin neonatale (µmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	tutti	5	100.0	0.0	0.0	286	2.4	e
2	ABL700/800 Radiomete	4	100.0	0.0	0.0	265	2.2	e

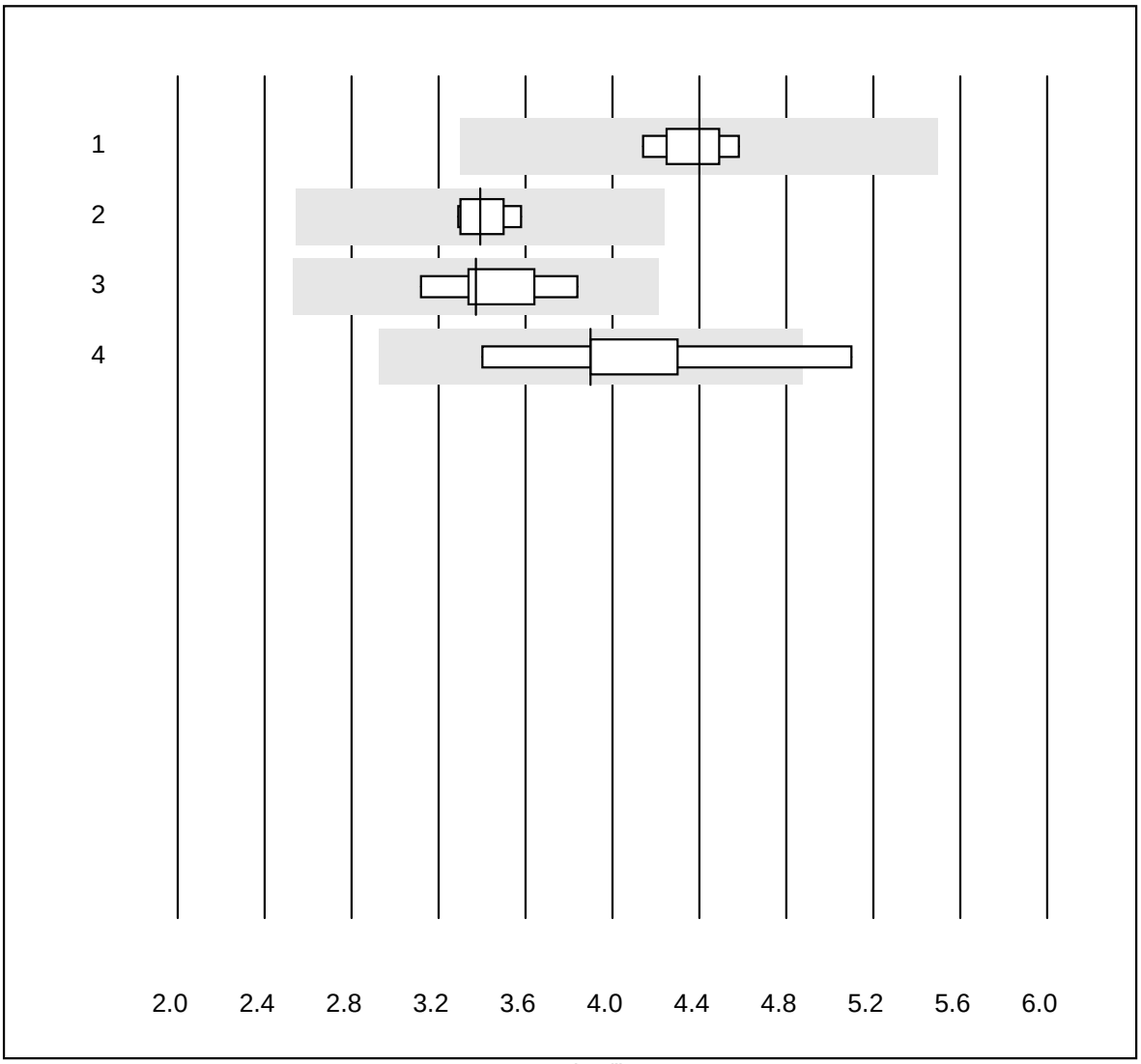
CK-MB



Deviatione QUALAB : 30 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Fuji Dri-Chem	38	97.4	0.0	2.6	62.2	5.2	e

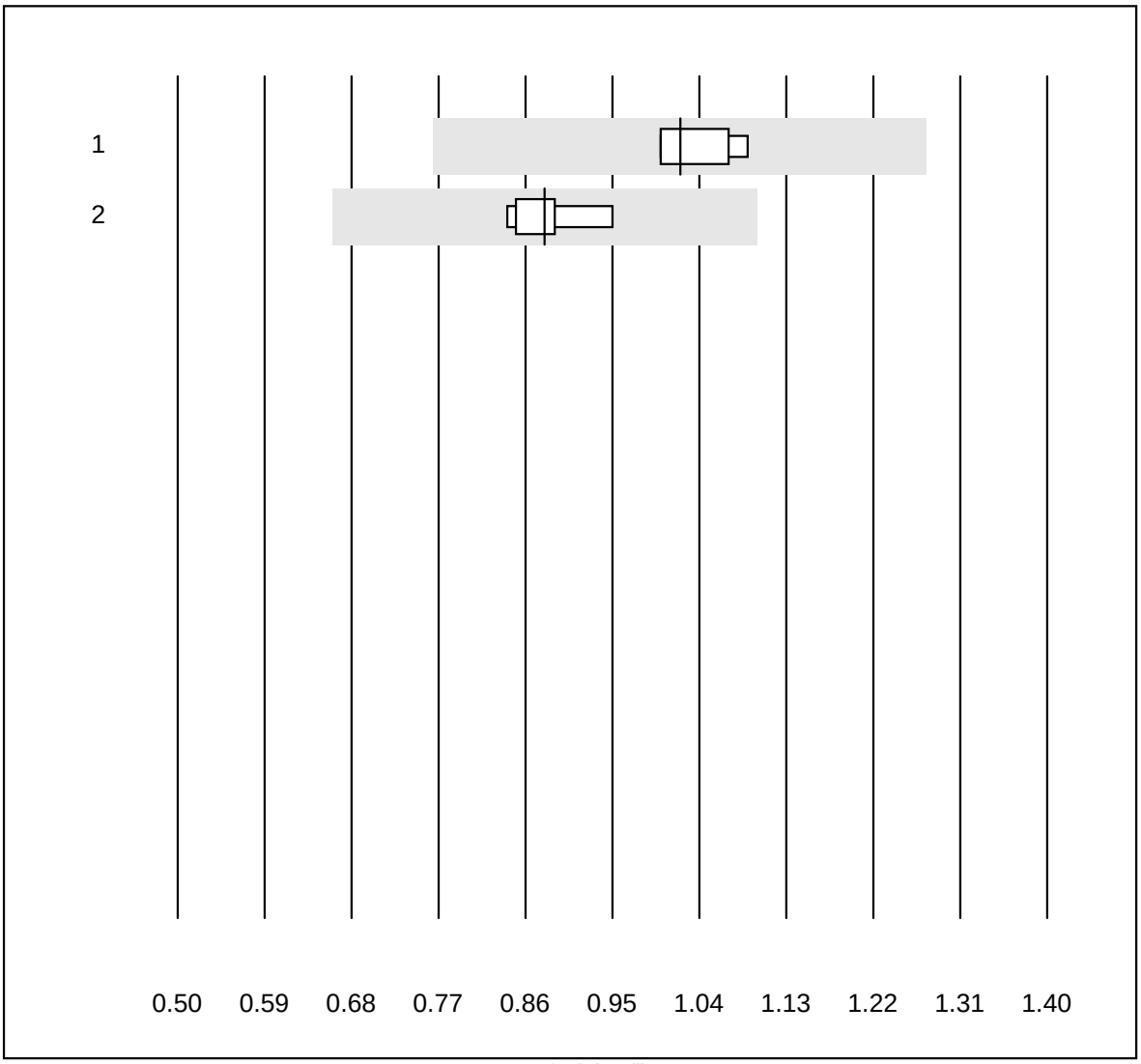
PSA



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas E / Elecsys	9	100.0	0.0	0.0	4.40	3.4	e
2	ADVIA Centaur XP/CP	5	100.0	0.0	0.0	3.39	3.7	e
3	Architect	6	100.0	0.0	0.0	3.37	7.4	e*
4	Qualigen	5	80.0	20.0	0.0	3.90	15.4	a

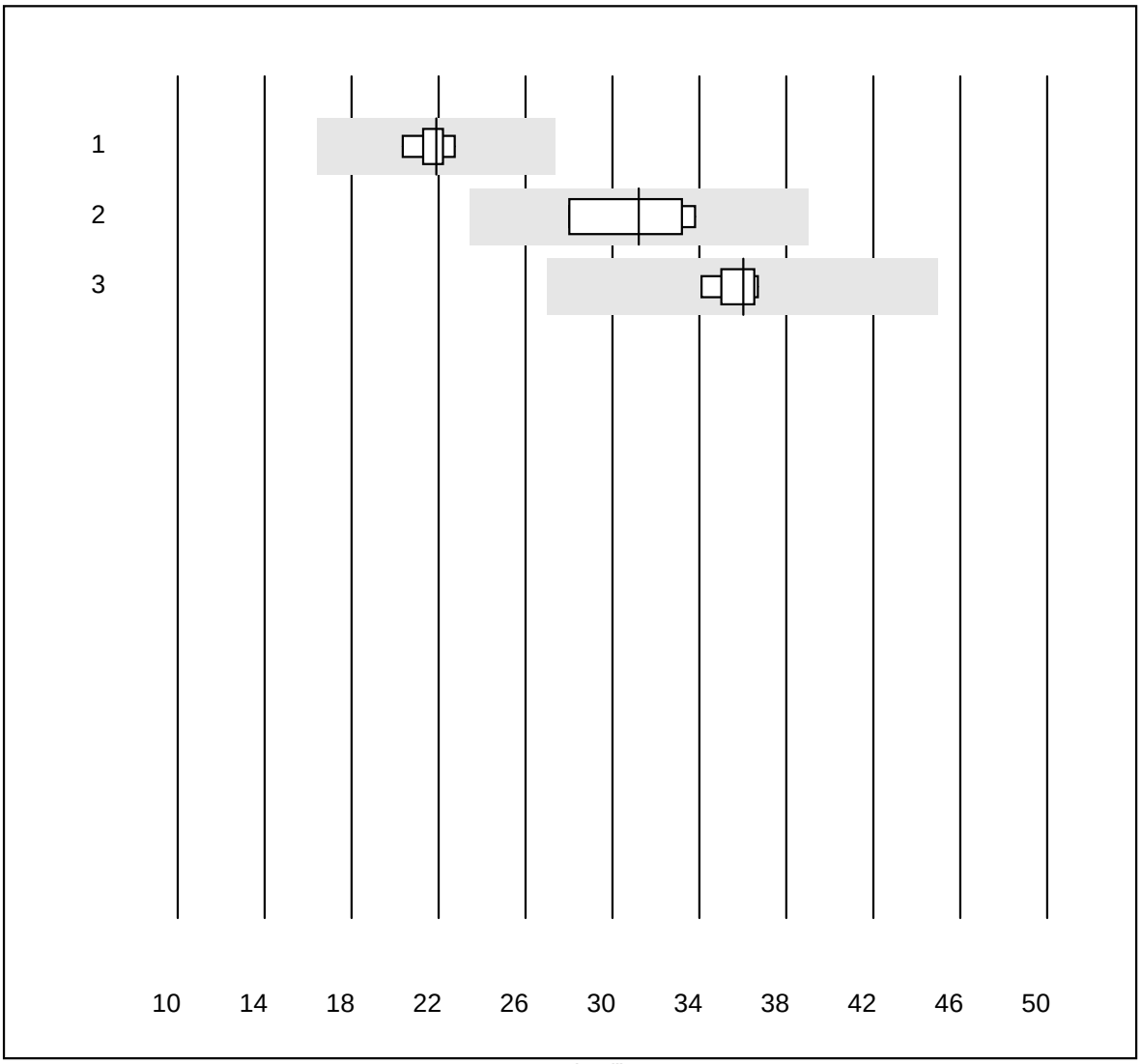
PSA frei



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas E / Elecsys	5	100.0	0.0	0.0	1.02	4.0	e
2	Architect	5	100.0	0.0	0.0	0.88	4.9	e

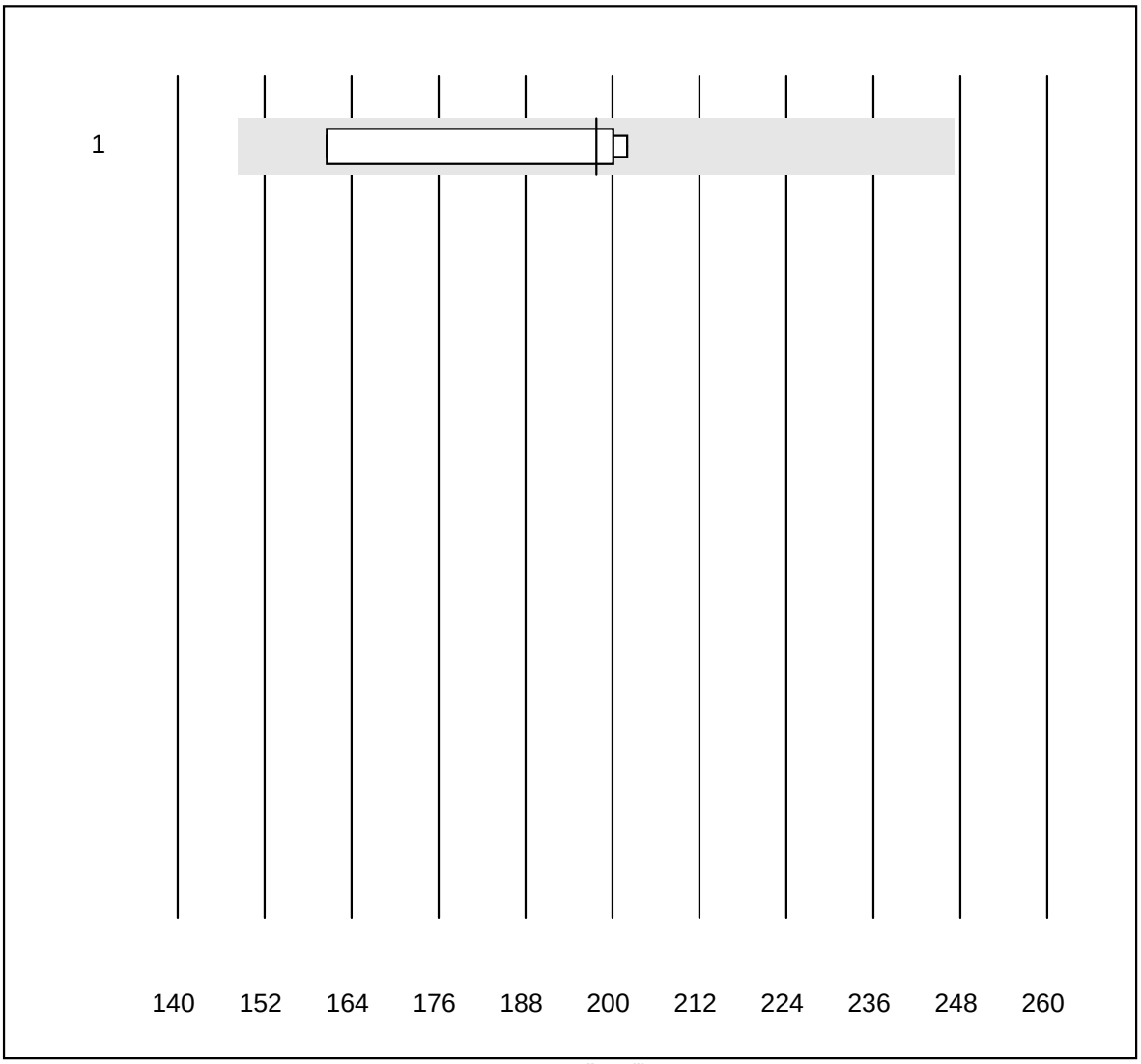
CEA



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas E / Elecsys	5	100.0	0.0	0.0	21.9	4.2	e
2	ADVIA Centaur XP/CP	4	100.0	0.0	0.0	31.2	9.3	e*
3	Architect	5	100.0	0.0	0.0	36.0	3.1	e

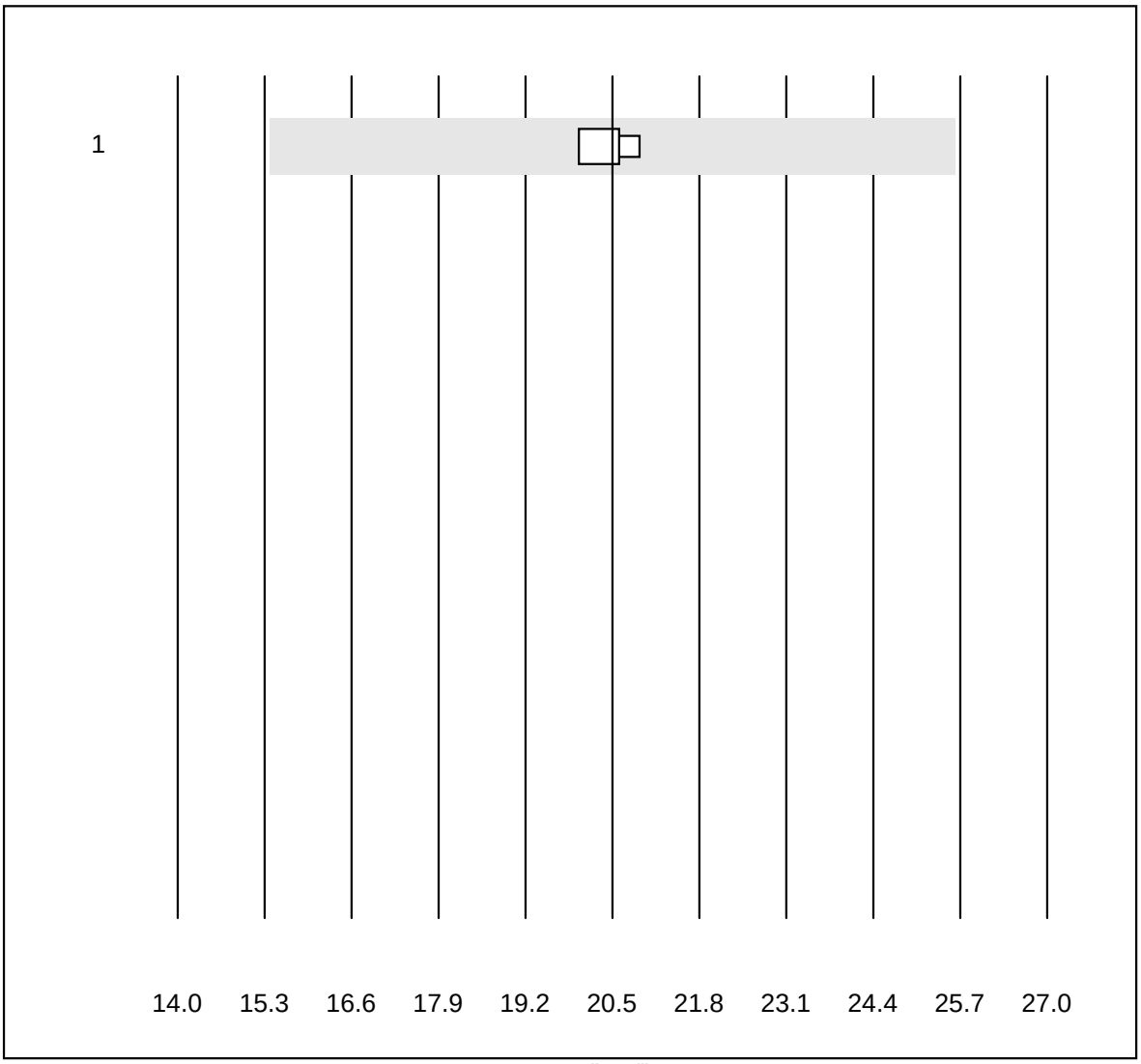
CA 125



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Architect	4	100.0	0.0	0.0	197.8	10.3	e*

CA 15-3

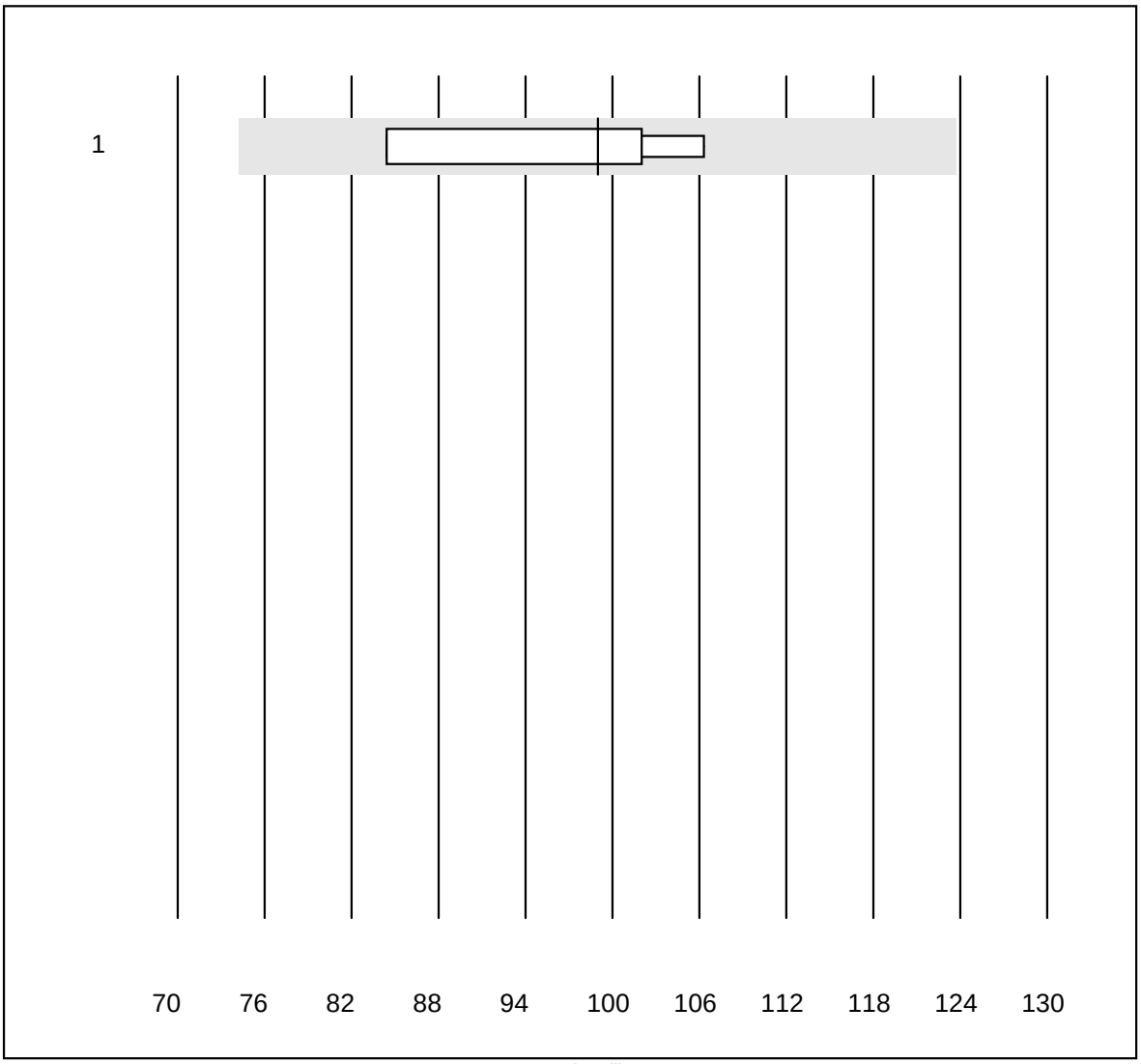


Deviatione QUALAB : 25 %

CA 15-3 (kIU/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Architect	4	100.0	0.0	0.0	20.5	1.8	e

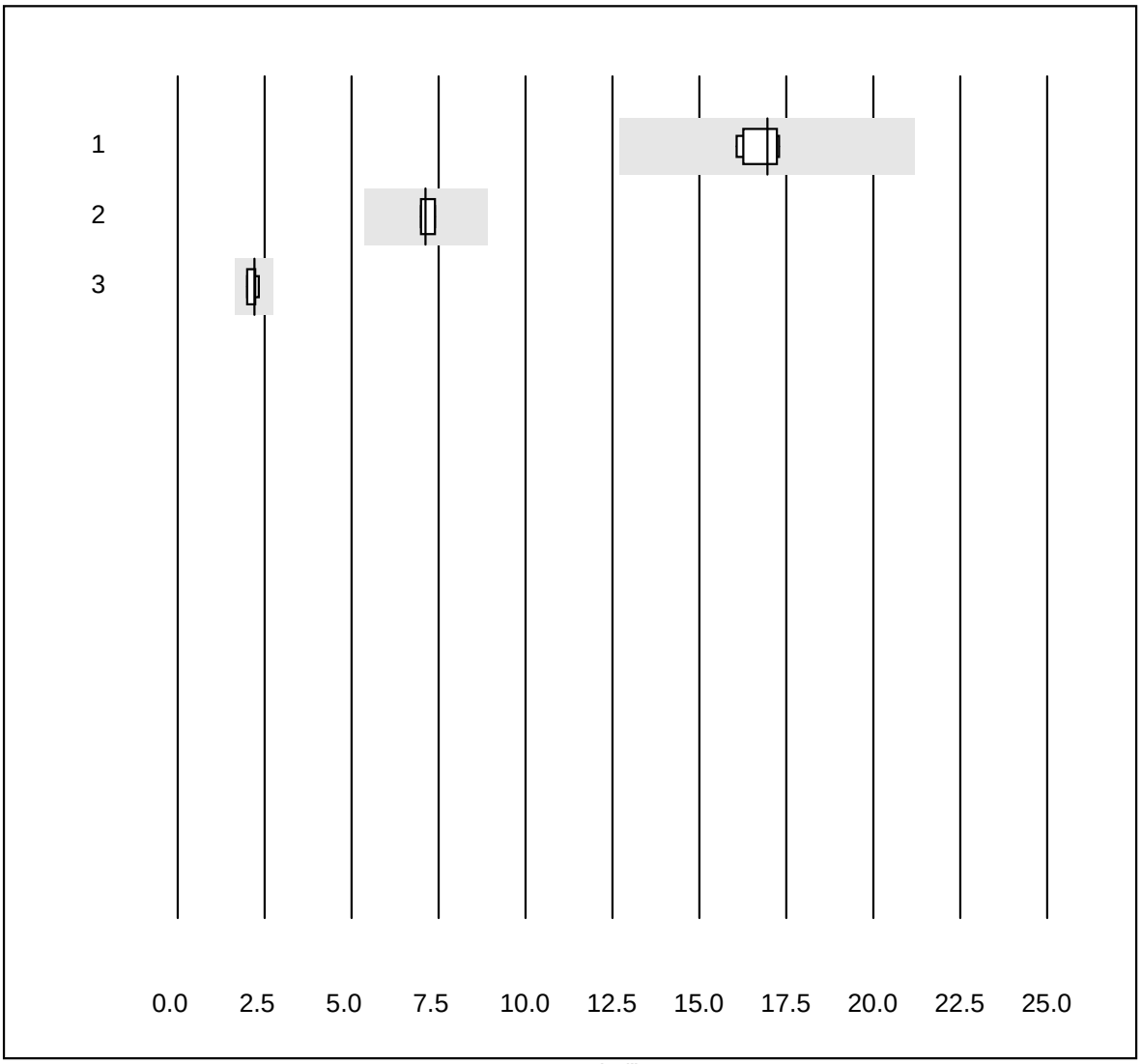
AFP



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas E / Elecsys	4	100.0	0.0	0.0	99	9.7	a

HCG qn

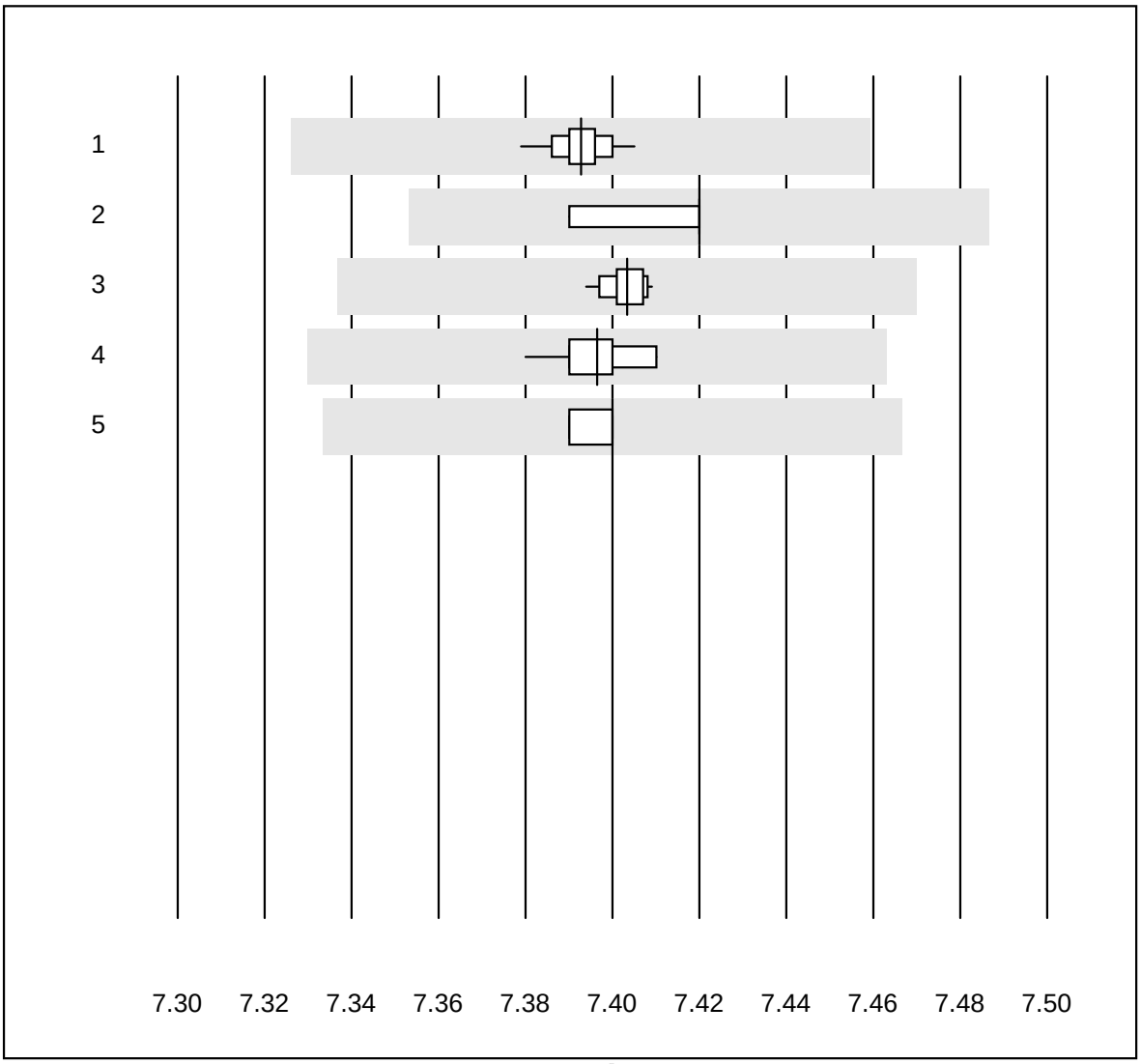


Deviatione QUALAB : 25 %

HCG qn (U/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas E / Elecsys	5	100.0	0.0	0.0	17	3.4	e
2	ADVIA Centaur XP/CP	4	75.0	0.0	25.0	7	3.2	a
3	Vidas	4	100.0	0.0	0.0	2	8.0	a

pH OR

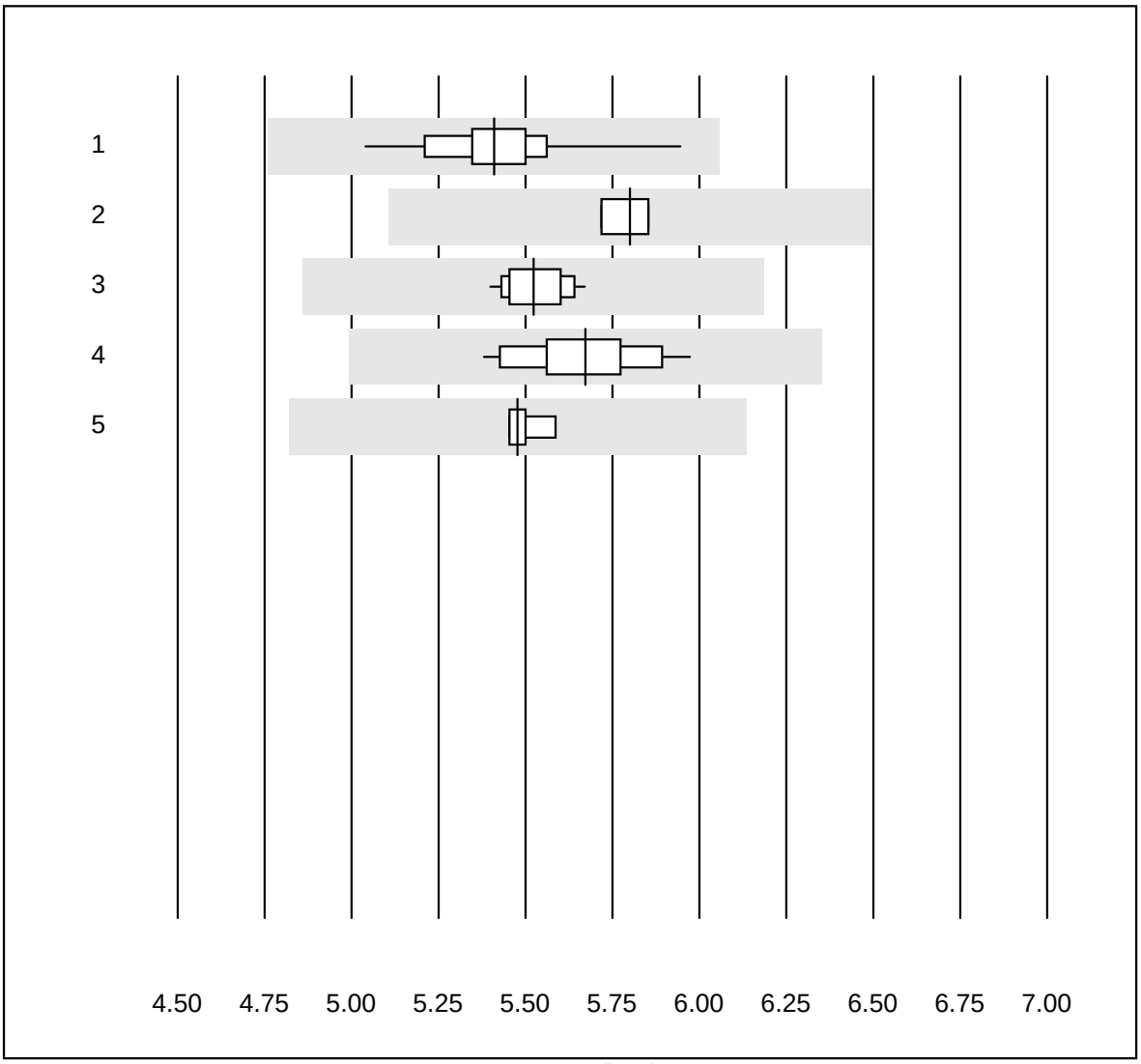


Deviatione QUALAB : 1 %

pH OR ()

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ABL700/800 Radiomete	81	100.0	0.0	0.0	7.39	0.1	e
2	Radiometer NPT-7	5	100.0	0.0	0.0	7.42	0.2	e
3	ABL 90	27	100.0	0.0	0.0	7.40	0.1	e
4	ABL 80 / Coox	14	100.0	0.0	0.0	7.40	0.1	e
5	ABL 5	6	100.0	0.0	0.0	7.40	0.1	e

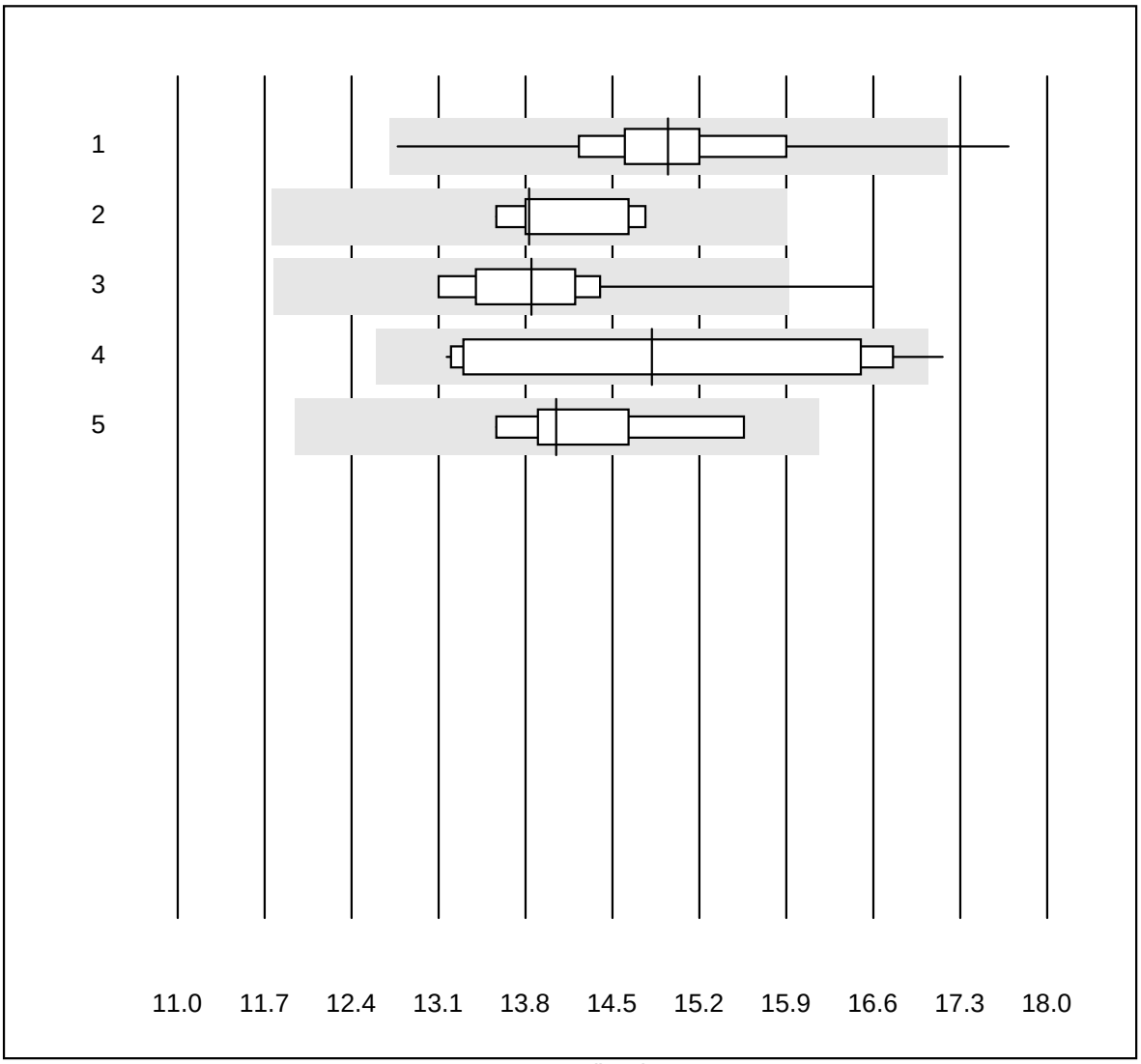
pCO2 OR



Deviatione QUALAB : 12 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ABL700/800 Radiomete	79	98.7	0.0	1.3	5.41	2.7	e
2	Radiometer NPT-7	5	100.0	0.0	0.0	5.80	1.2	e
3	ABL 90	27	100.0	0.0	0.0	5.52	1.4	e
4	ABL 80 / Coox	14	100.0	0.0	0.0	5.67	3.0	e
5	ABL 5	6	100.0	0.0	0.0	5.48	0.9	e

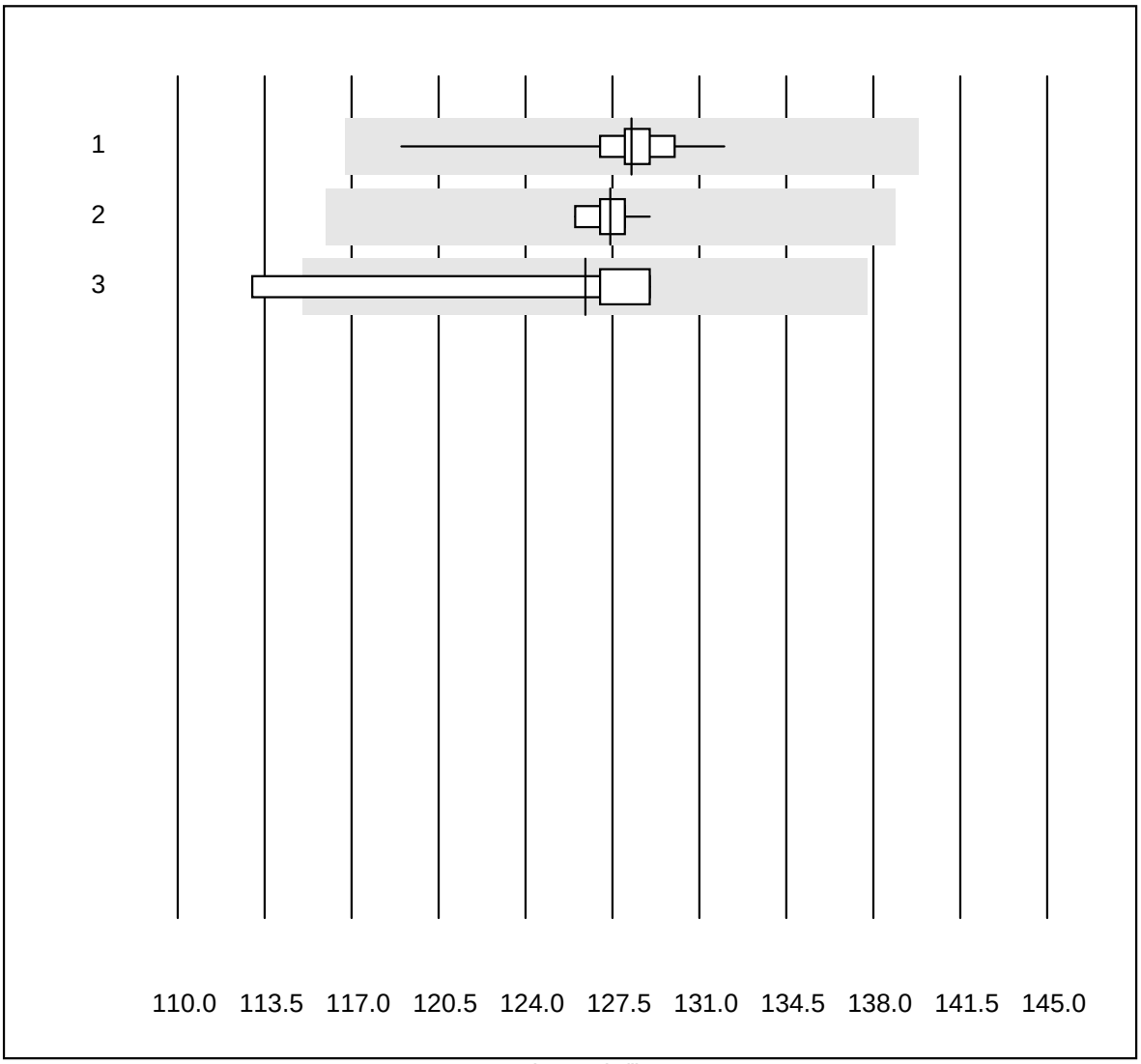
pO2 OR



Deviatione QUALAB : 15 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ABL700/800 Radiomete	79	96.2	1.3	2.5	14.95	4.8	e
2	Radiometer NPT-7	5	100.0	0.0	0.0	13.83	3.8	e
3	ABL 90	27	96.3	3.7	0.0	13.85	5.1	e
4	ABL 80 / Coox	14	92.9	7.1	0.0	14.82	10.5	e*
5	ABL 5	6	100.0	0.0	0.0	14.05	5.0	e*

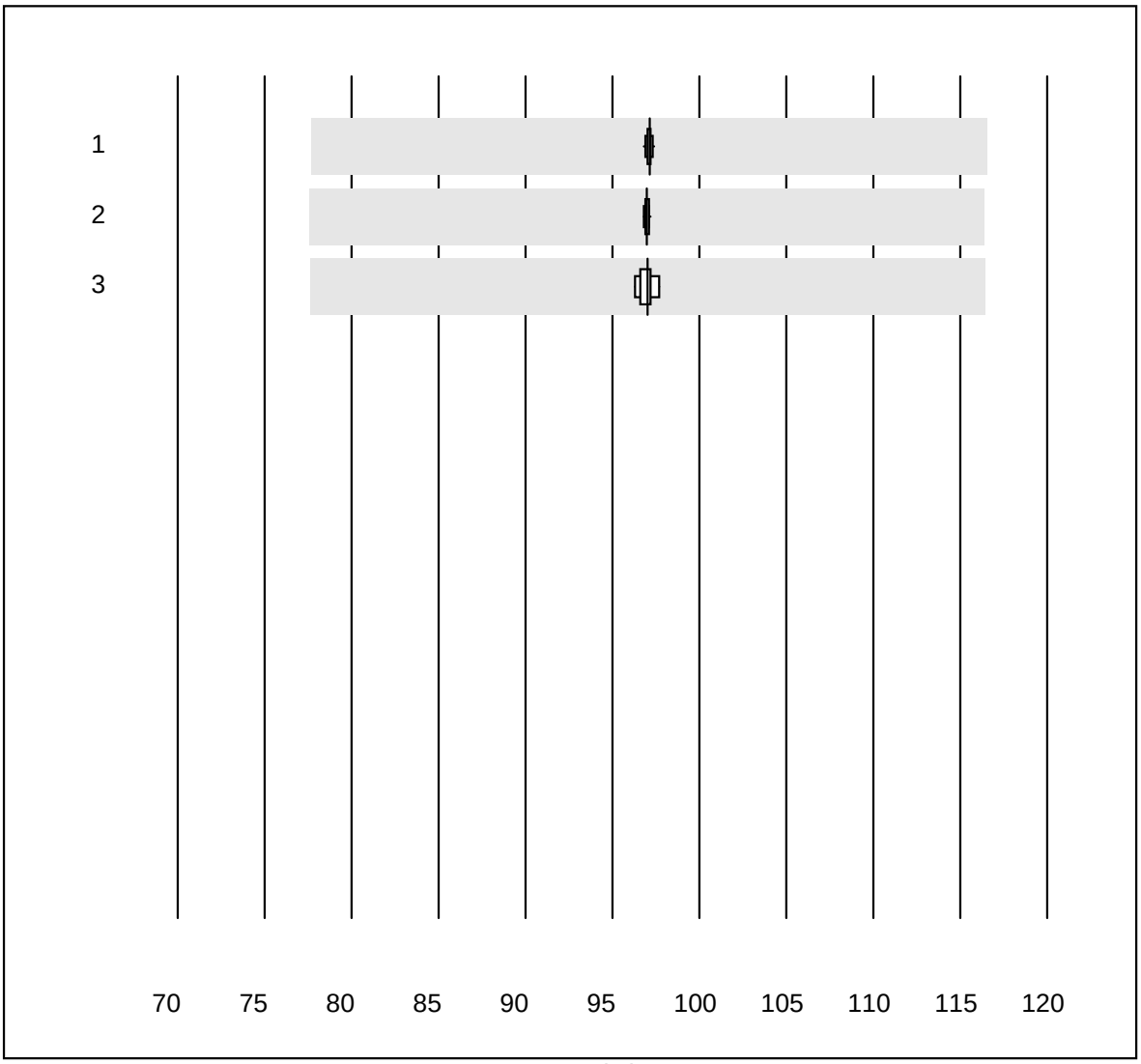
ctHb OR



Deviatione QUALAB : 9 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ABL700/800 Radiomete	70	91.4	0.0	8.6	128.3	1.7	e
2	ABL 90	26	92.3	0.0	7.7	127.4	0.7	e
3	ABL 80 / Coox	11	81.8	9.1	9.1	126.4	3.8	e*

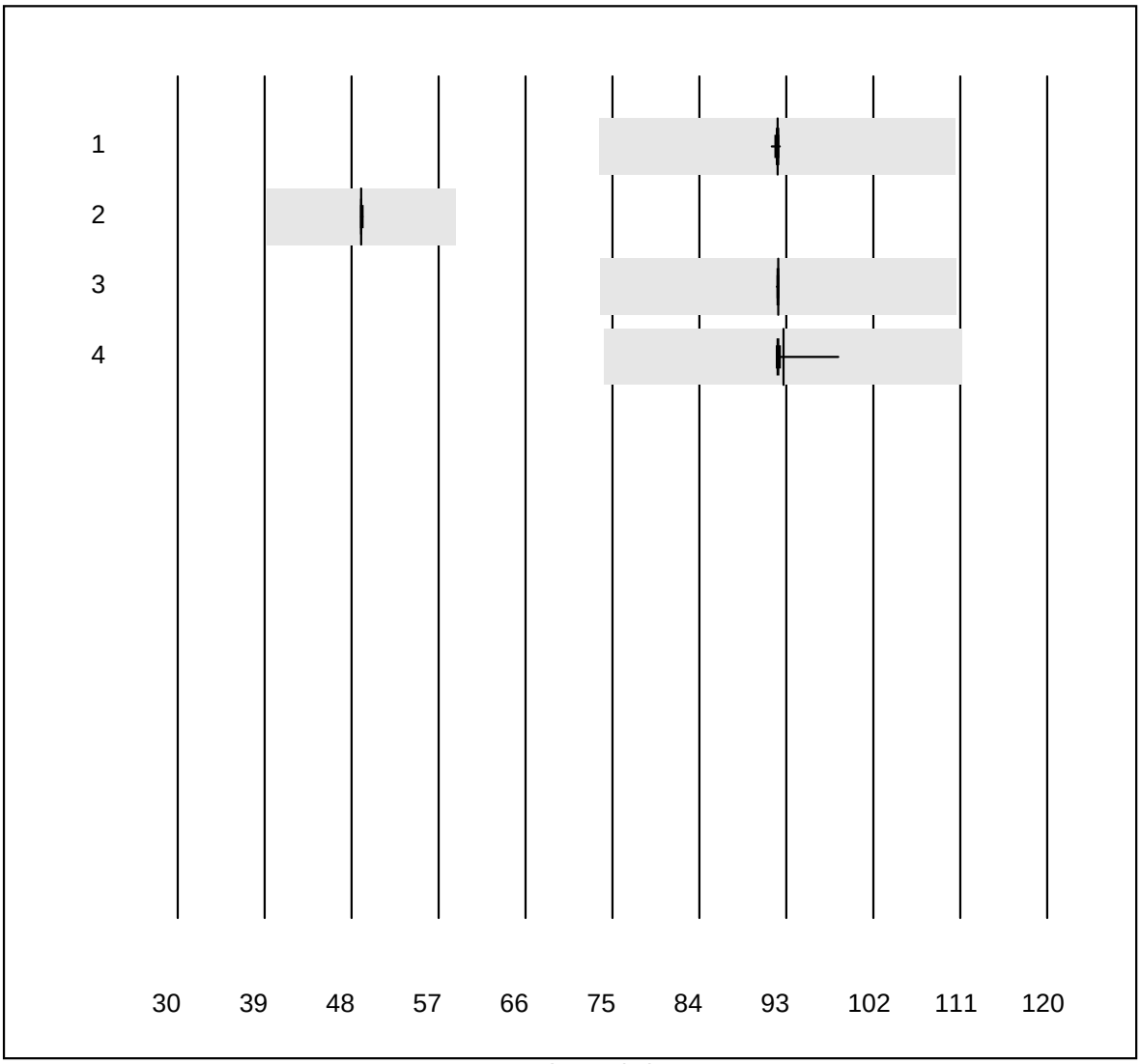
sO2 OR



Deviatione QUALAB : 20 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ABL700/800 Radiomete	54	100.0	0.0	0.0	97.124	0.2	e
2	ABL 90	24	100.0	0.0	0.0	96.983	0.1	e
3	ABL 80 / Coox	10	100.0	0.0	0.0	97.020	0.5	e

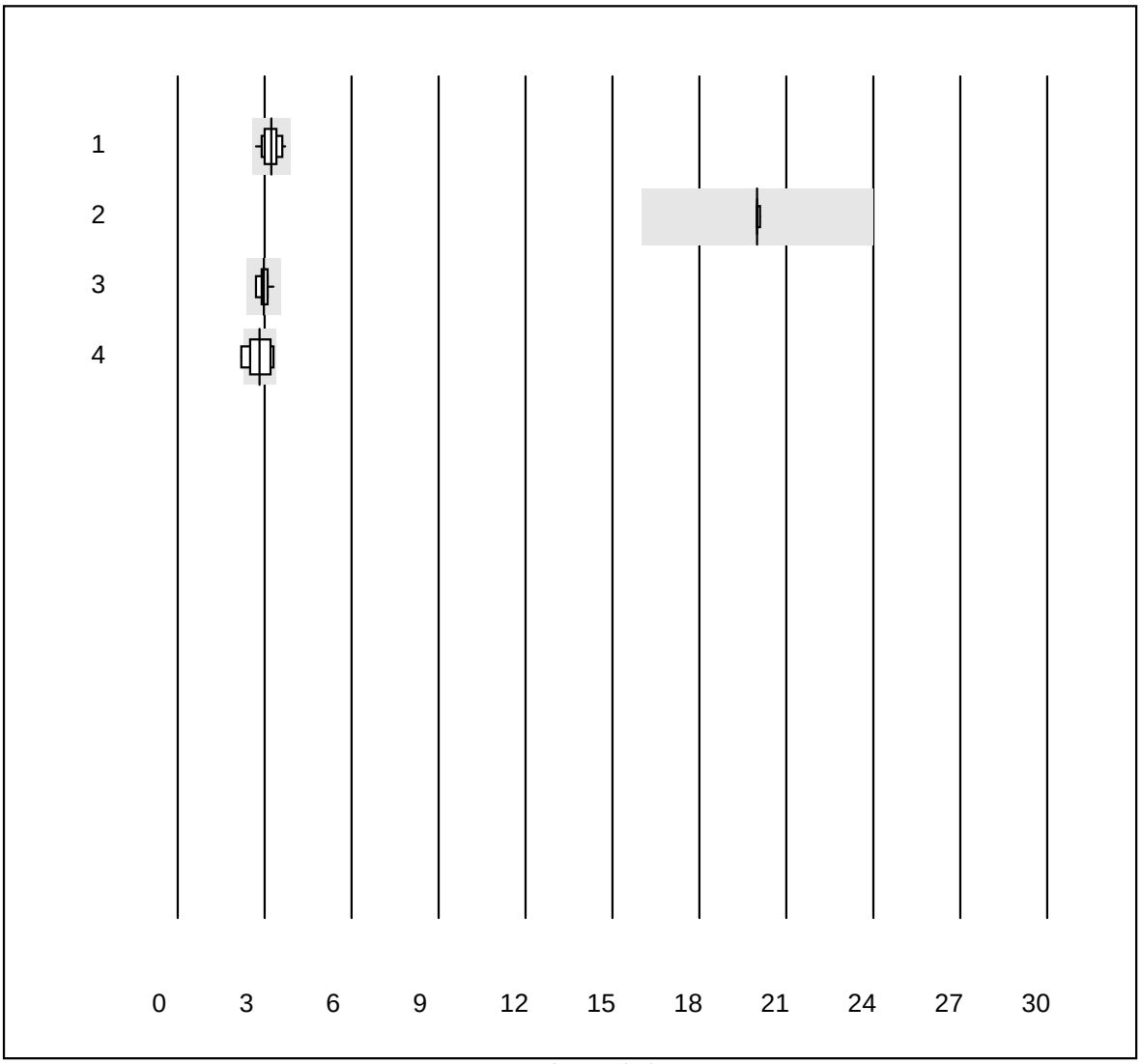
FO2Hb OR



Deviatione QUALAB : 20 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ABL700/800 Radiomete	50	100.0	0.0	0.0	92.056	0.2	e
2	Radiometer NPT-7	4	100.0	0.0	0.0	49.000	0.1	e
3	ABL 90	24	100.0	0.0	0.0	92.163	0.1	e
4	ABL 80 / Coox	11	100.0	0.0	0.0	92.700	2.0	e

FCOHb OR

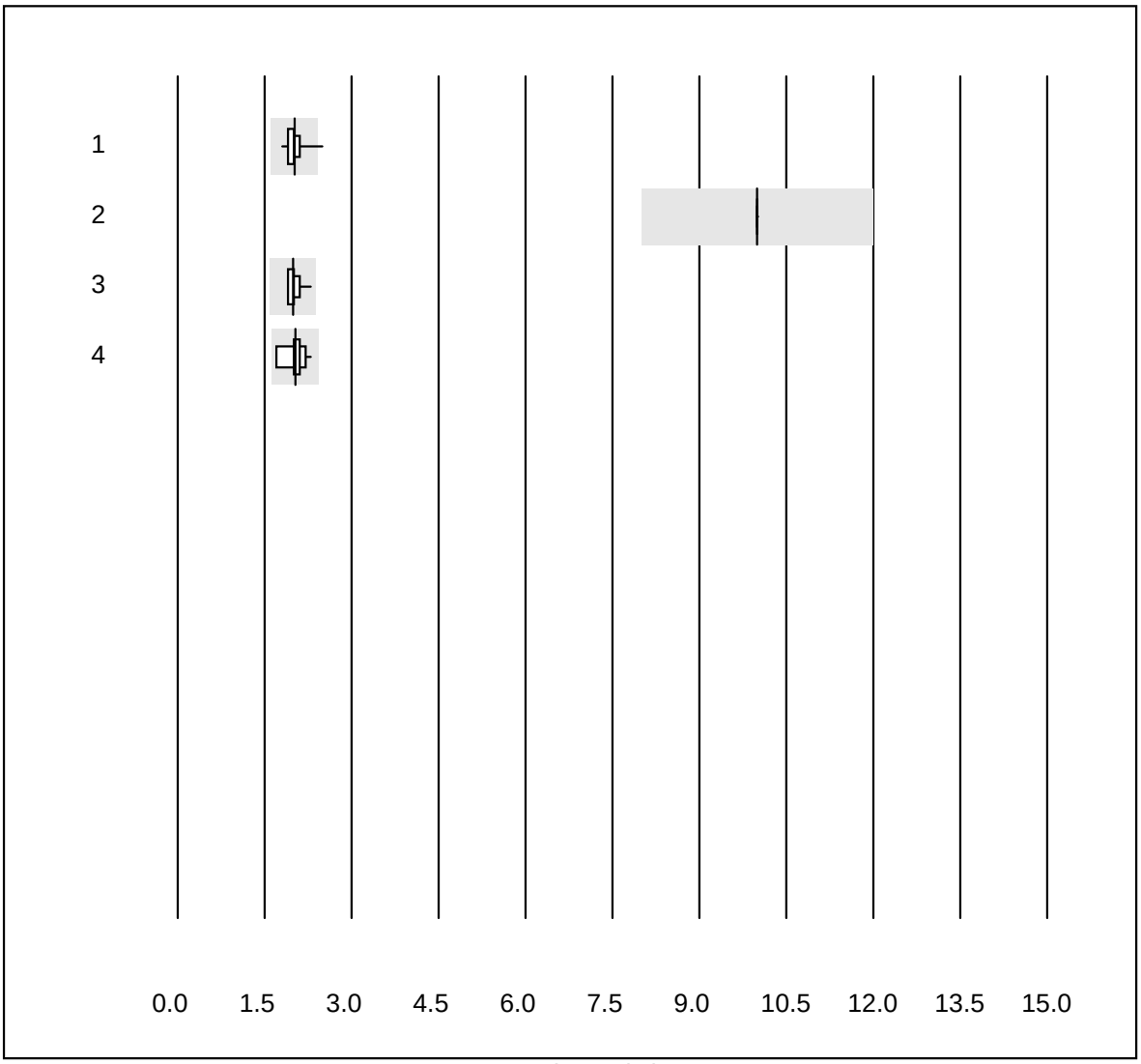


Deviatione QUALAB : 20 %

FCOHb OR (%)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ABL700/800 Radiomete	51	100.0	0.0	0.0	3.228	7.9	e
2	Radiometer NPT-7	4	100.0	0.0	0.0	20.000	0.2	e
3	ABL 90	24	100.0	0.0	0.0	2.971	5.3	e
4	ABL 80 / Coox	10	80.0	10.0	10.0	2.822	14.3	e*

FMetHb OR

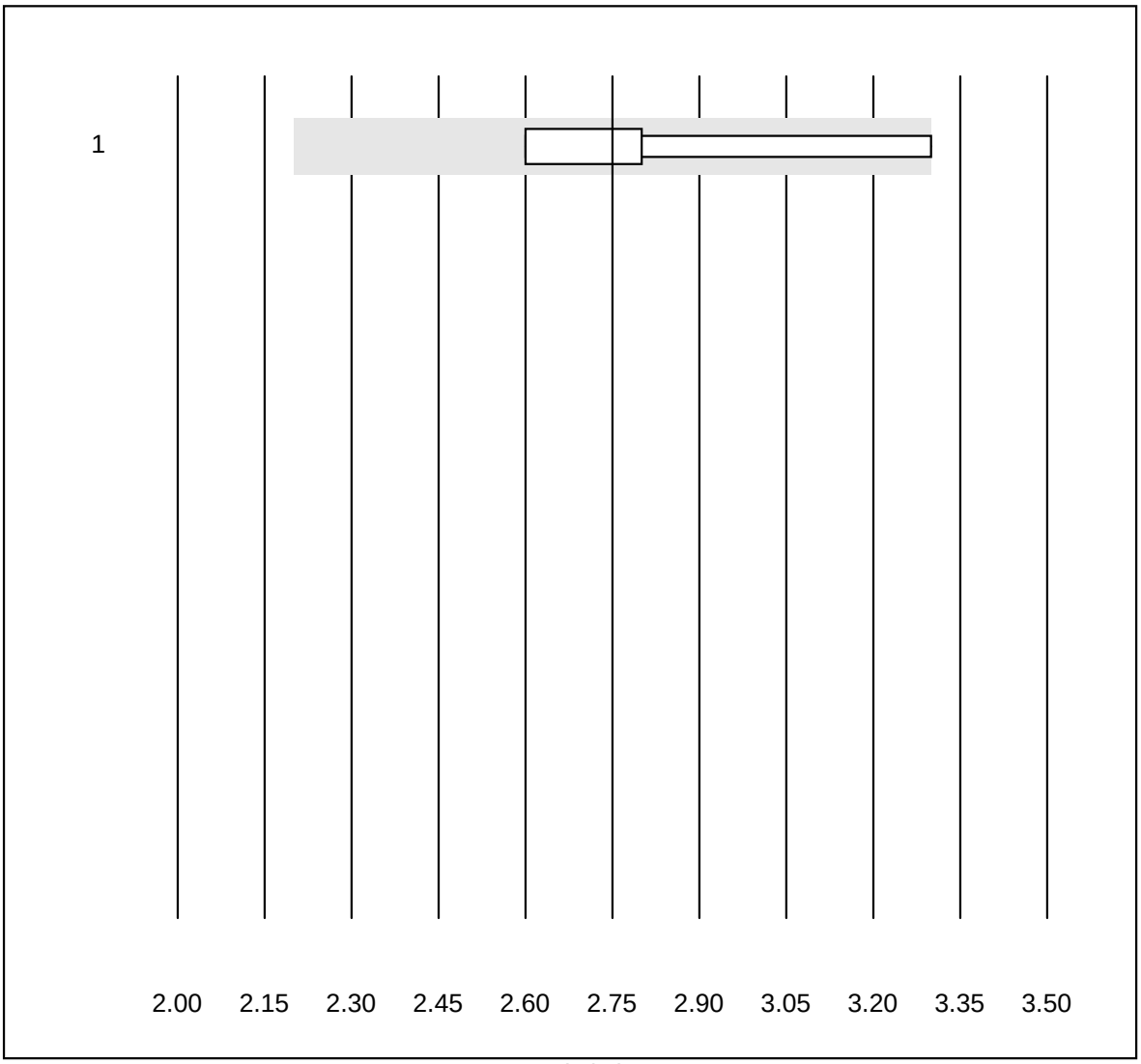


Deviatione QUALAB : 20 %

FMetHb OR (%)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ABL700/800 Radiomete	52	96.2	3.8	0.0	2.012	7.3	e
2	Radiometer NPT-7	4	100.0	0.0	0.0	10.000	0.0	e
3	ABL 90	24	95.8	0.0	4.2	1.987	4.6	e
4	ABL 80 / Coox	11	90.9	0.0	9.1	2.030	8.1	e*

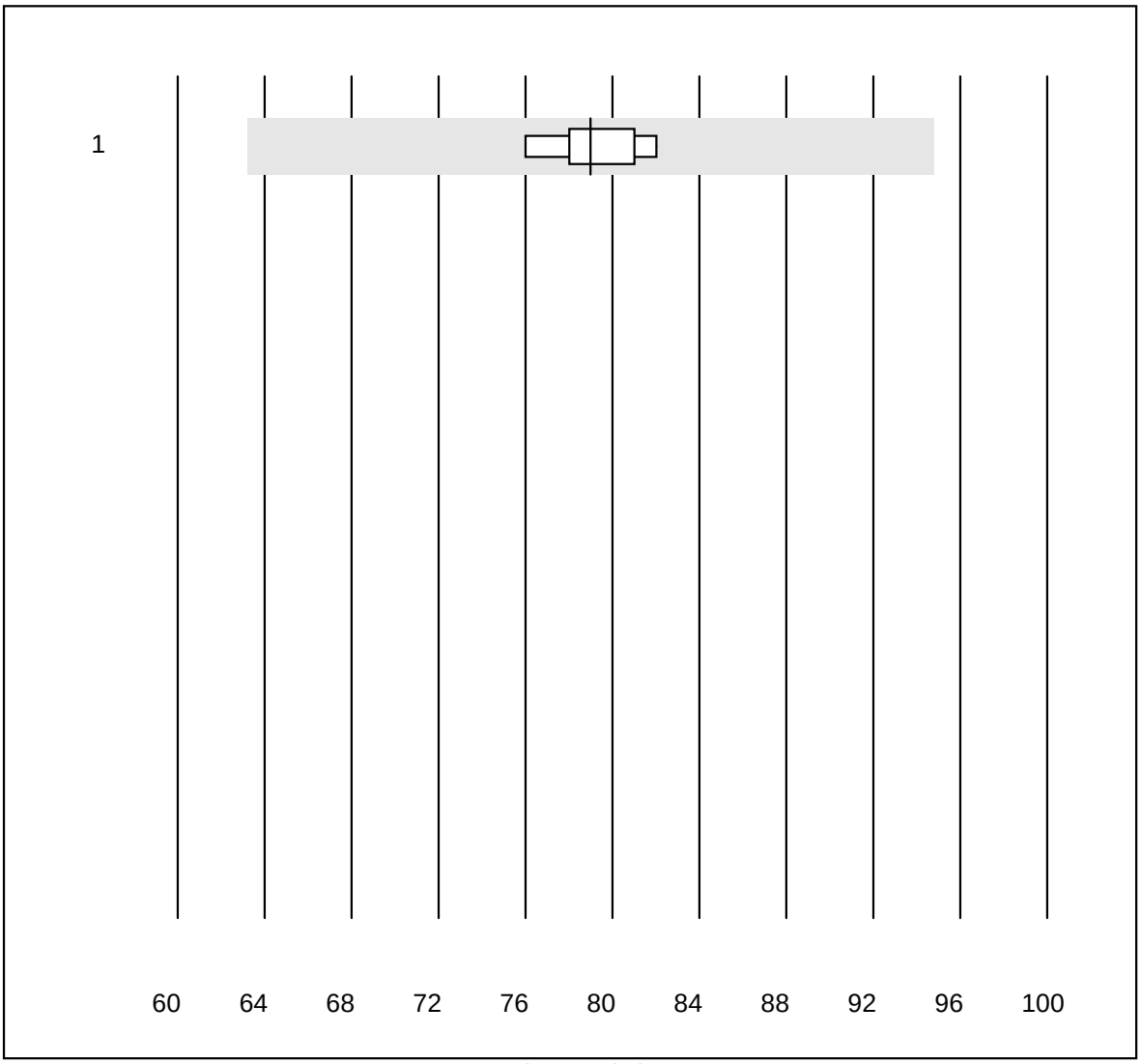
FHHb



Deviatione QUALAB : 20 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ABL 80 / Coox	4	75.0	25.0	0.0	2.750	10.9	e*

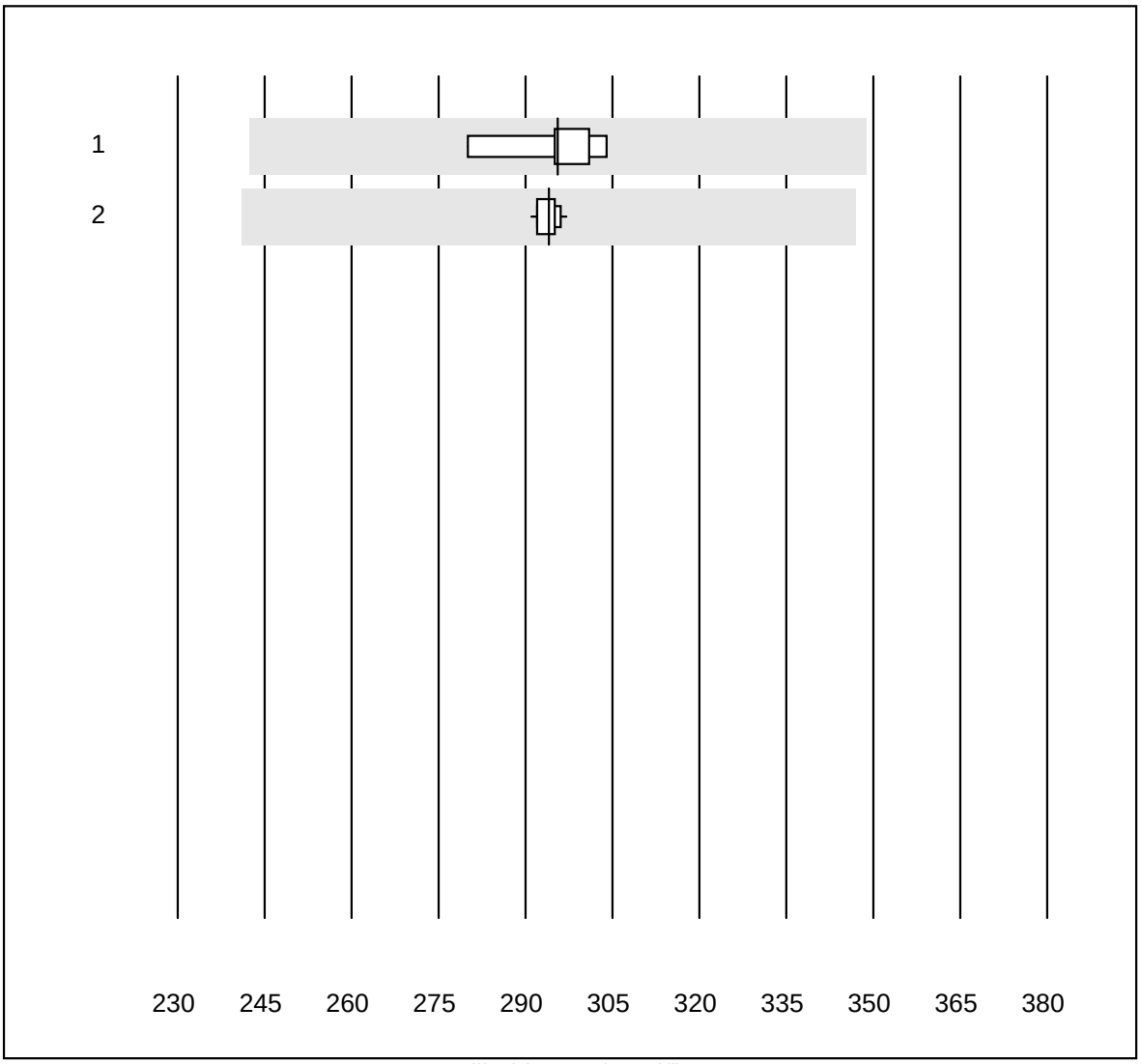
FHbF OR



Deviatione QUALAB : 20 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ABL 90	6	100.0	0.0	0.0	79.000	2.7	e

Bilirubin OR

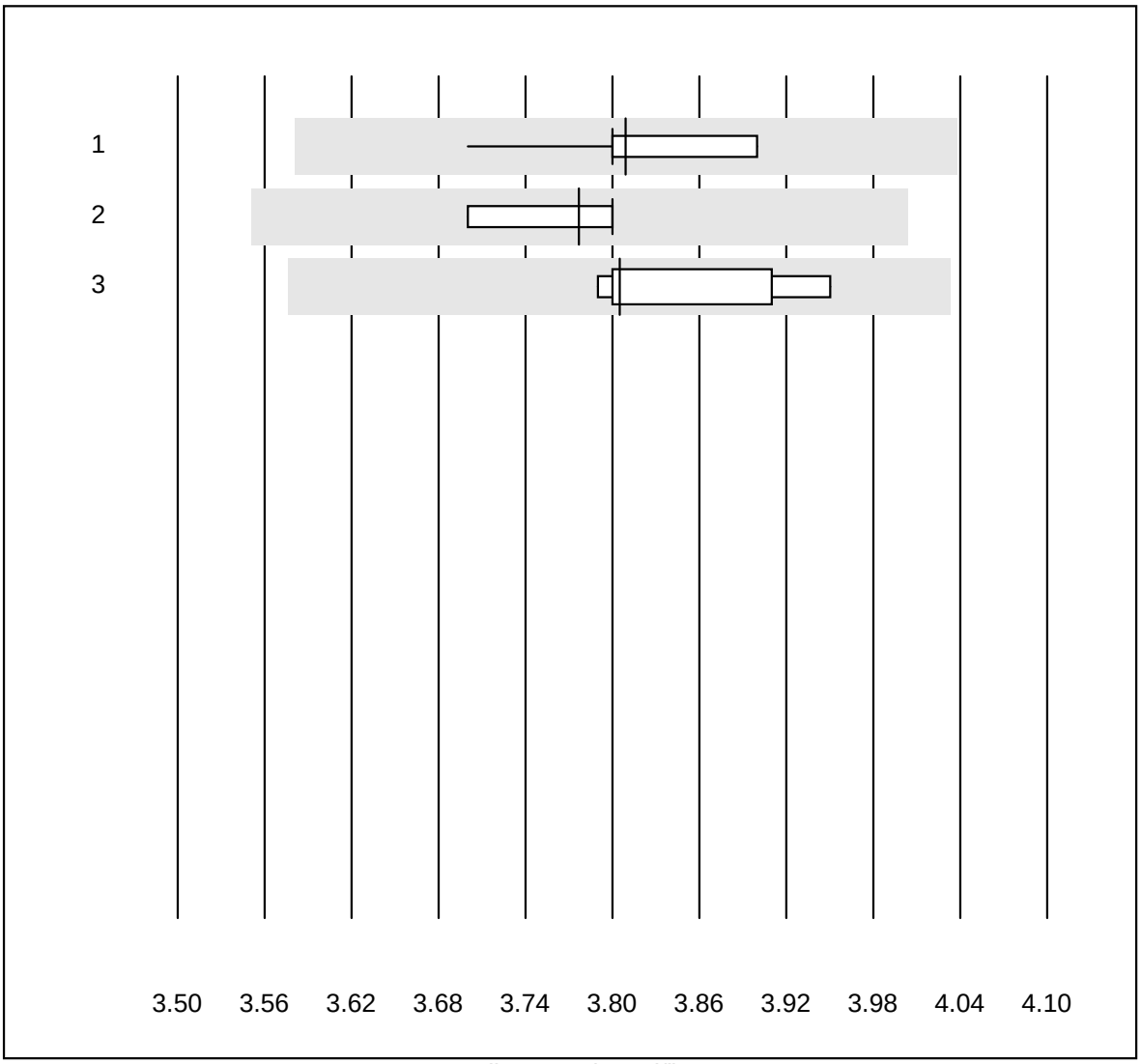


Deviatione QUALAB : 18 %

Bilirubin OR (µmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ABL700/800 Radiomete	8	100.0	0.0	0.0	295.5	2.5	e
2	ABL 90	11	100.0	0.0	0.0	294.0	0.6	e

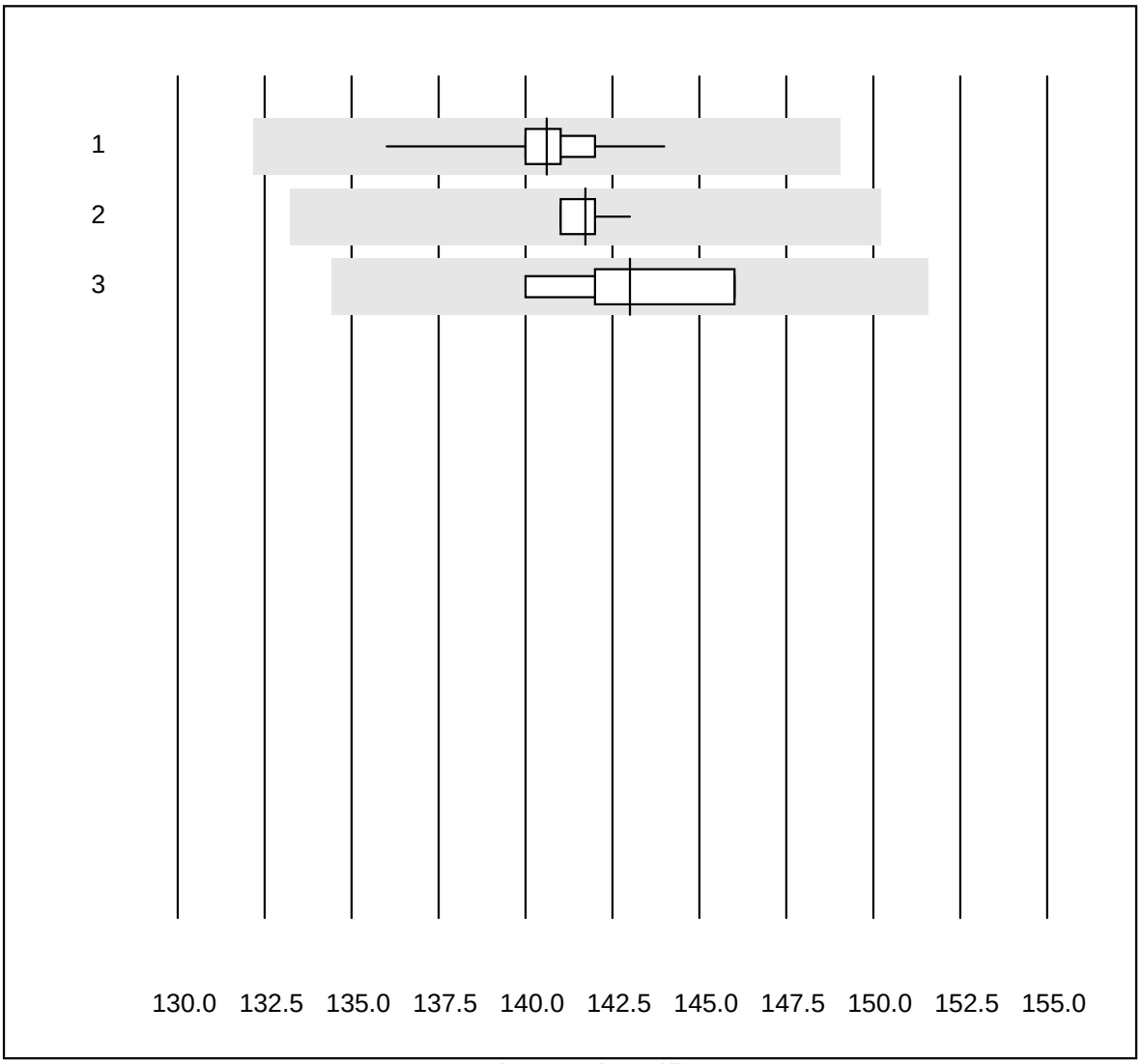
Kalium OR



Deviatione QUALAB : 6 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ABL700/800 Radiomete	66	100.0	0.0	0.0	3.8	1.1	e
2	ABL 90	26	100.0	0.0	0.0	3.8	1.1	e
3	ABL 80 / Coox	6	100.0	0.0	0.0	3.8	1.8	e*

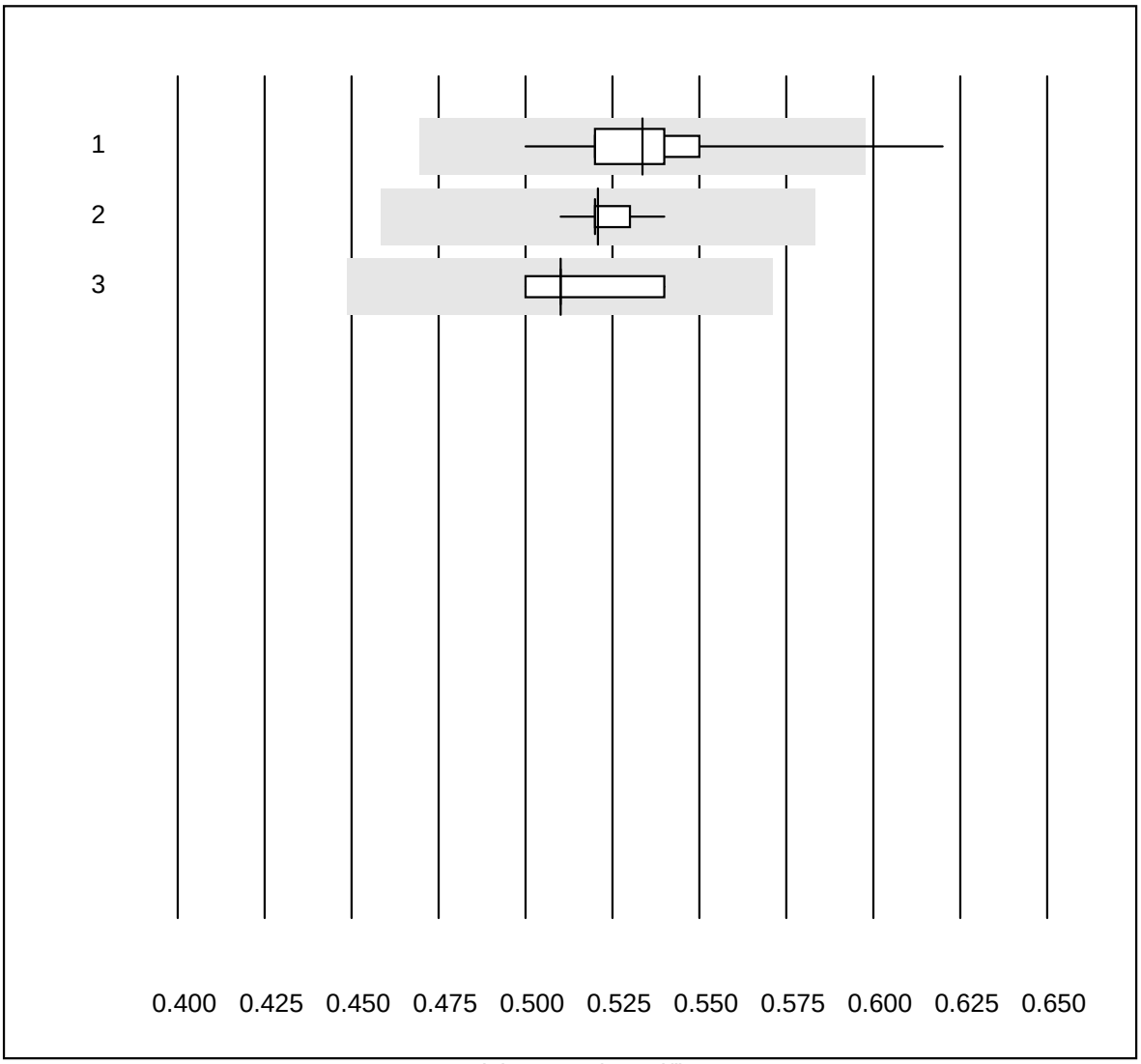
Natrium OR



Deviatione QUALAB : 6 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ABL700/800 Radiomete	64	100.0	0.0	0.0	140.6	0.8	e
2	ABL 90	26	100.0	0.0	0.0	141.7	0.4	e
3	ABL 80 / Coox	5	100.0	0.0	0.0	143.0	1.8	e*

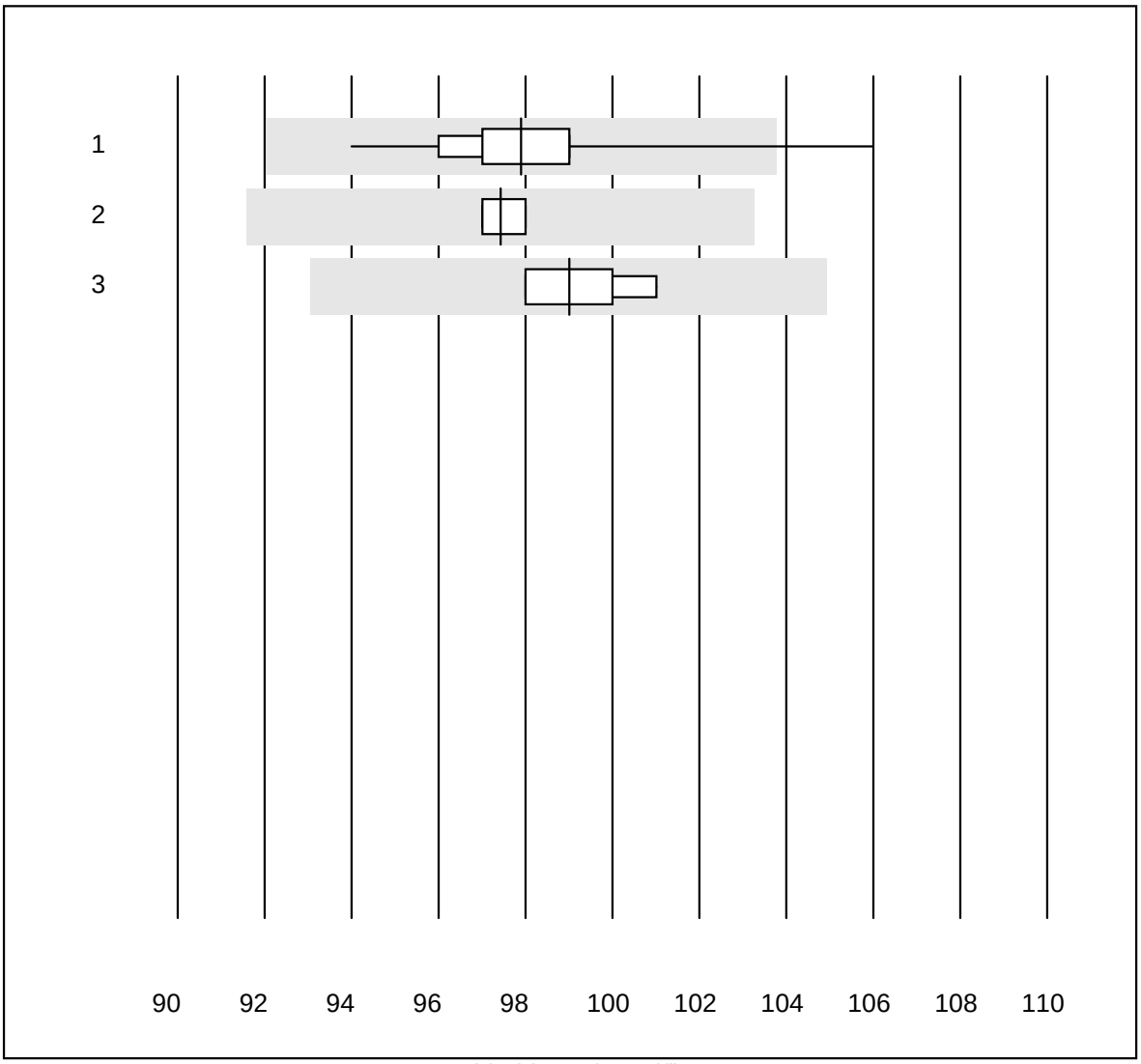
Kalzium OR



Deviatione QUALAB : 12 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ABL700/800 Radiomete	65	96.9	3.1	0.0	0.53	3.8	e
2	ABL 90	26	100.0	0.0	0.0	0.52	1.1	e
3	ABL 80 / Coox	5	100.0	0.0	0.0	0.51	3.0	e

Chlorid OR

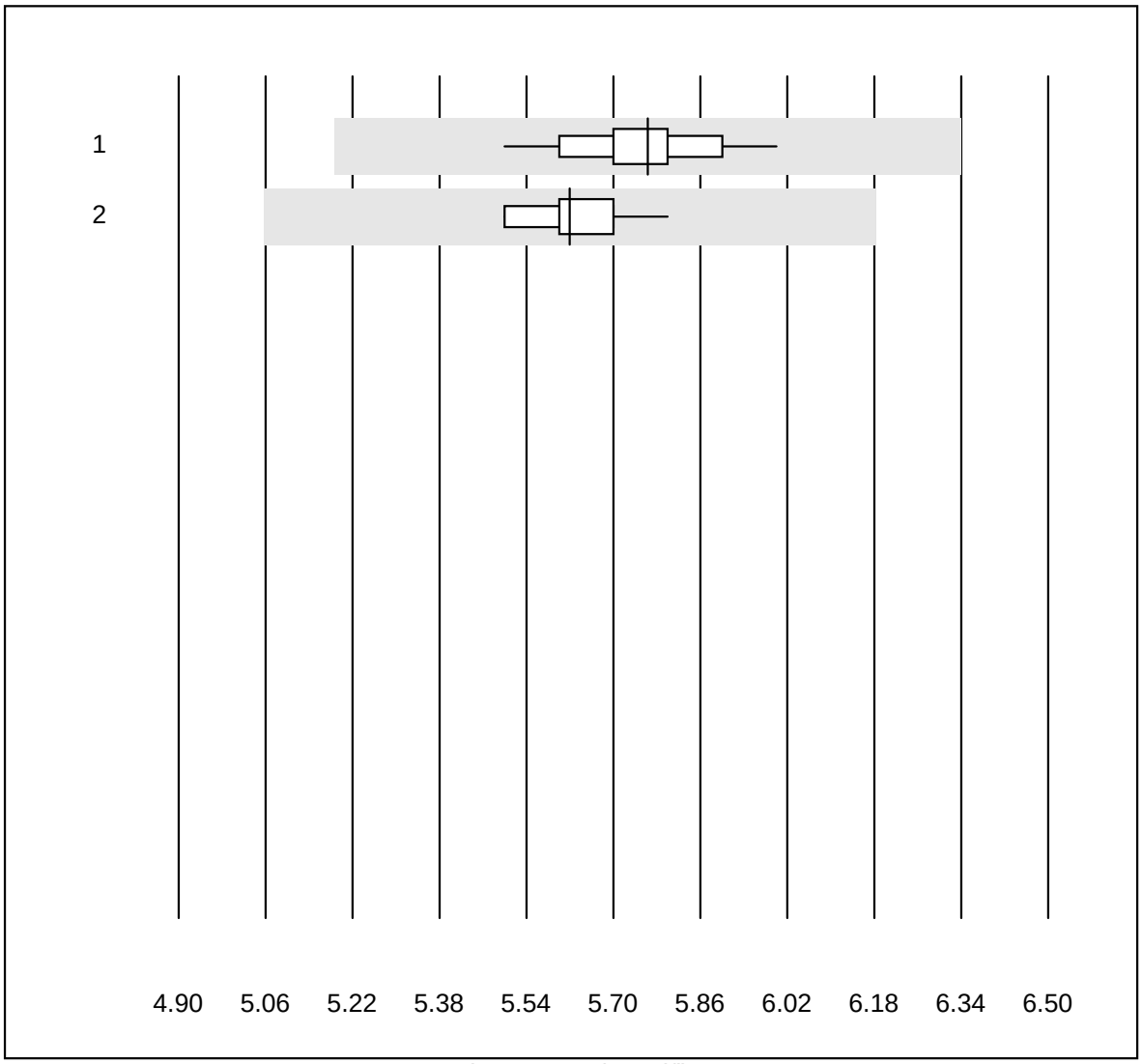


Deviatione QUALAB : 6 %

Chlorid OR (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ABL700/800 Radiomete	53	98.1	1.9	0.0	97.91	1.7	e
2	ABL 90	26	100.0	0.0	0.0	97.42	0.5	e
3	ABL 80 / Coox	5	100.0	0.0	0.0	99.00	1.3	e

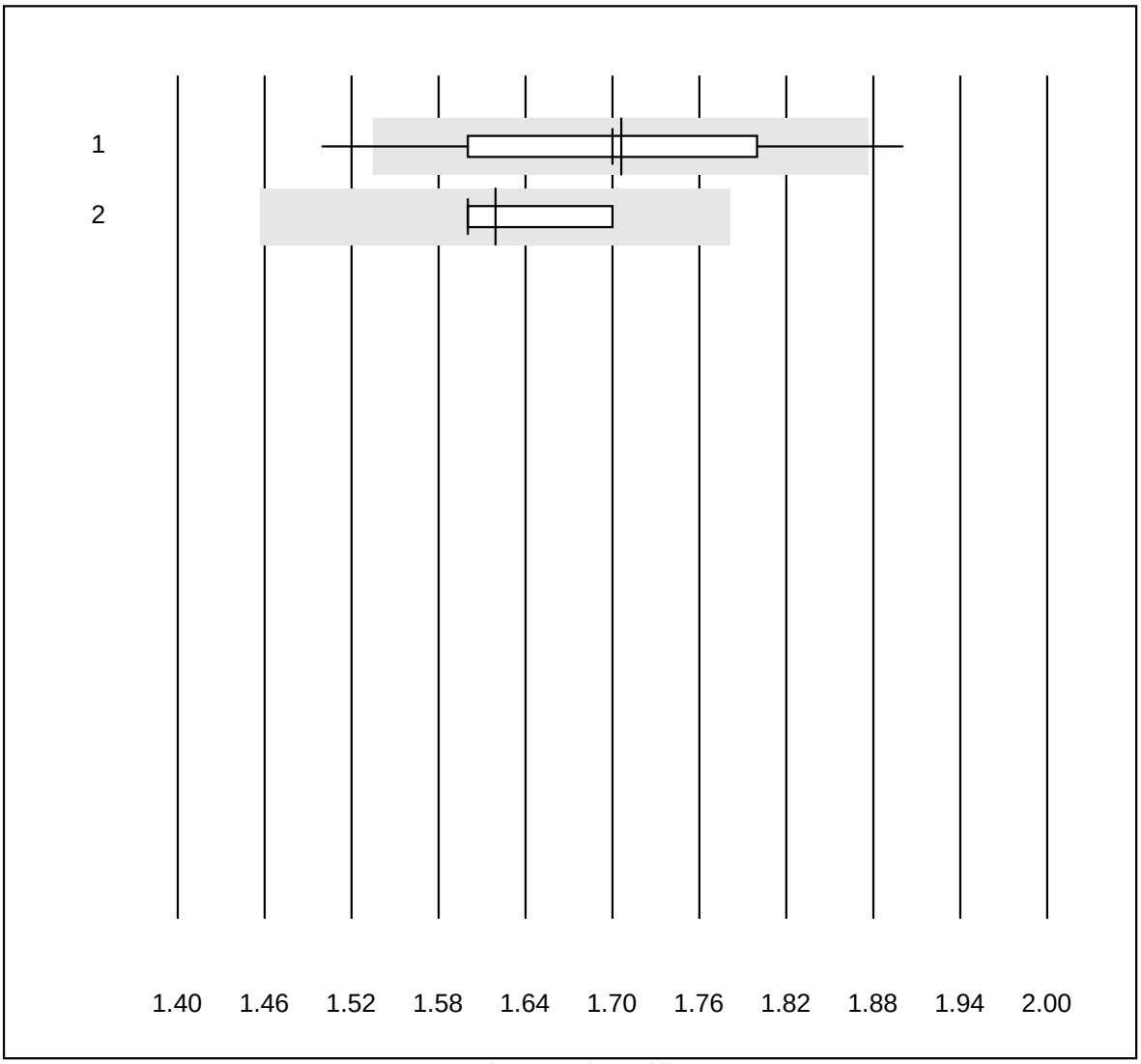
Glucose OR



Deviatione QUALAB : 10 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ABL700/800 Radiomete	65	100.0	0.0	0.0	5.8	1.9	e
2	ABL 90	26	100.0	0.0	0.0	5.6	1.6	e

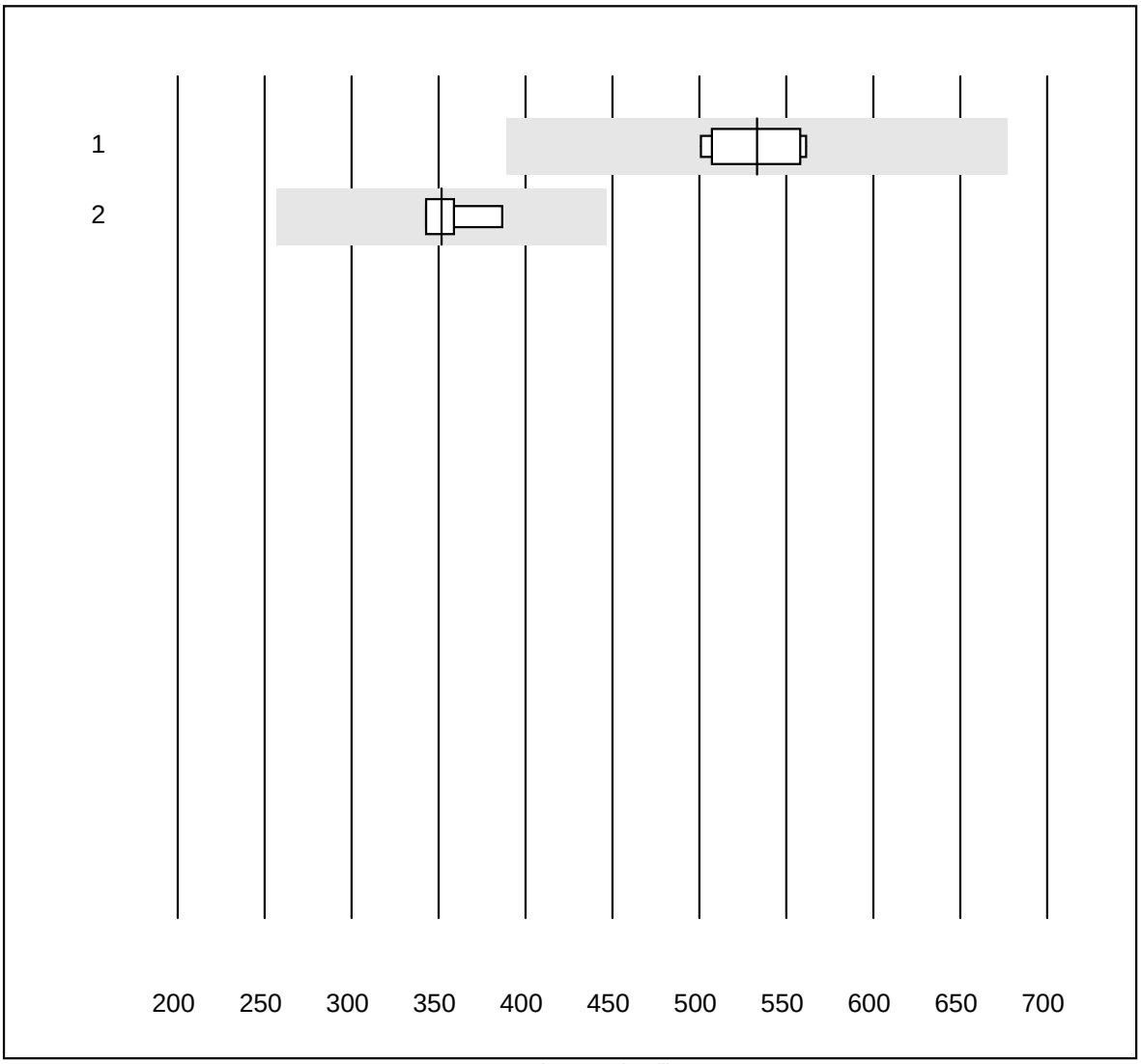
Laktat OR



Deviatione QUALAB : 10 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ABL700/800 Radiomete	68	97.1	2.9	0.0	1.71	3.9	e
2	ABL 90	26	100.0	0.0	0.0	1.62	2.5	e

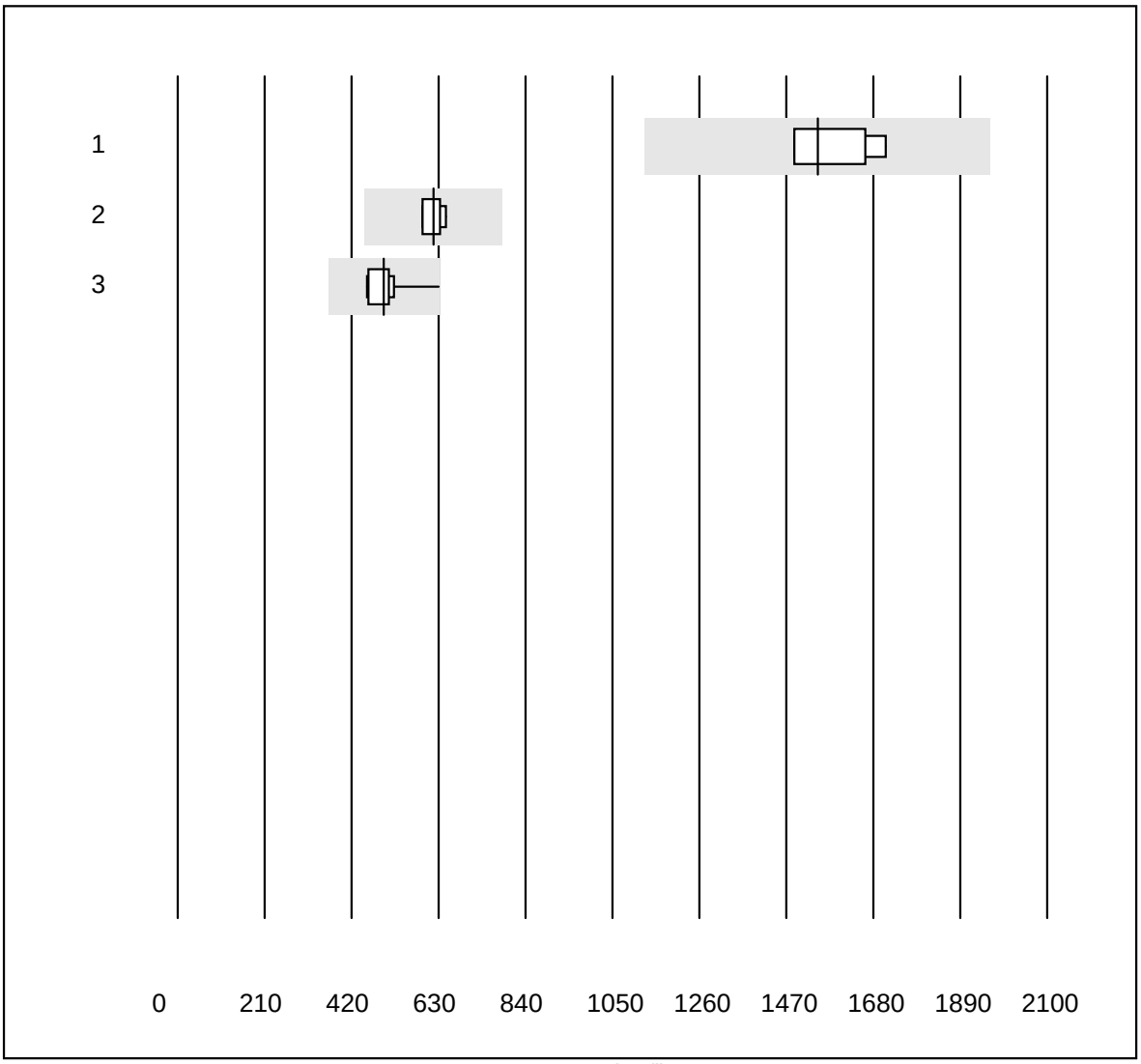
BNP Plasma



Deviatione QUALAB : 27 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ADVIA Centaur XP/CP	5	100.0	0.0	0.0	533.0	5.3	e
2	Architect	4	100.0	0.0	0.0	351.5	5.7	e

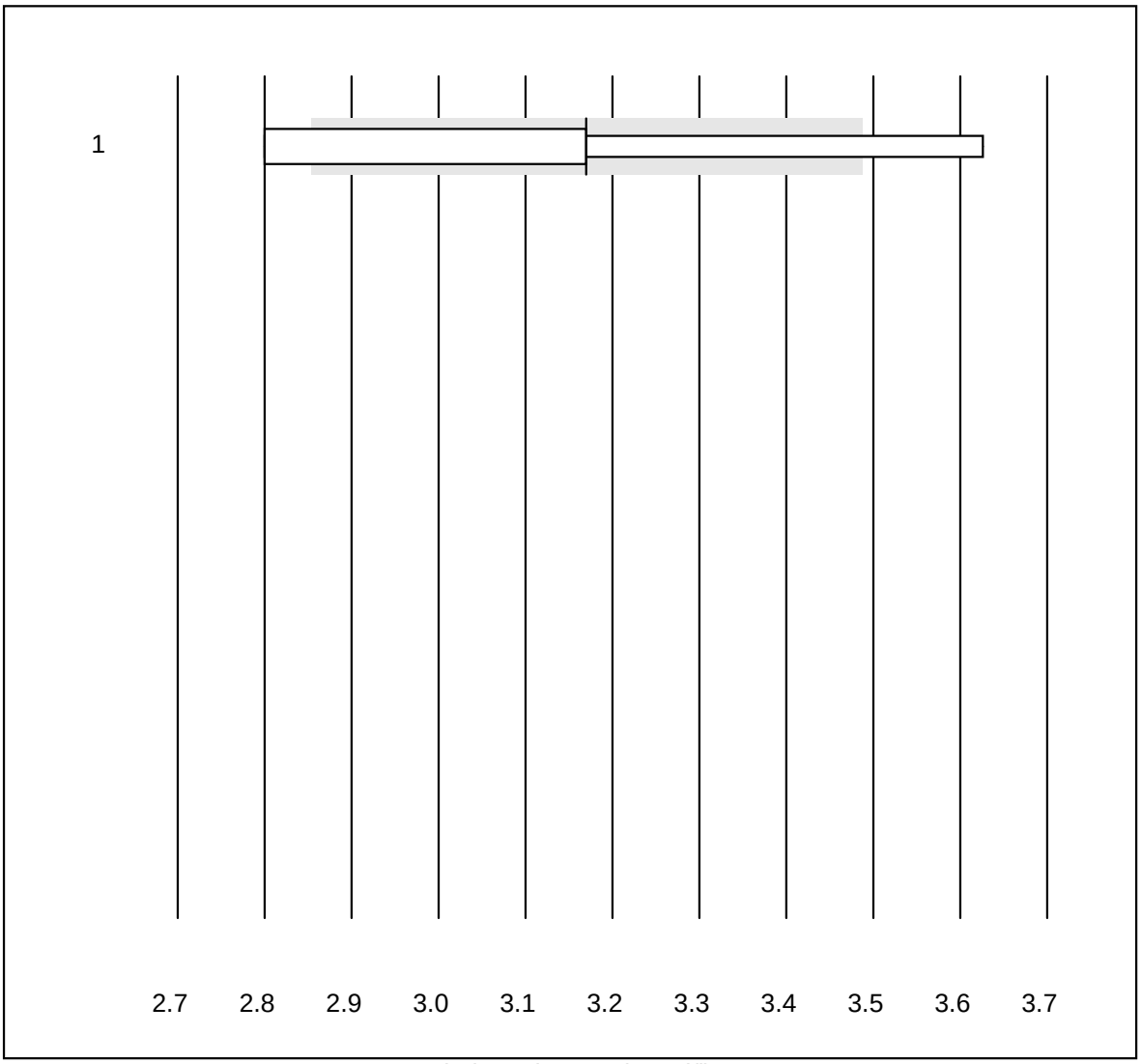
NT-proBNP



Deviatione QUALAB : 27 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	AQT 90 FLEX	5	100.0	0.0	0.0	1545.0	6.2	e
2	Vidas	4	100.0	0.0	0.0	617.0	4.4	e
3	Cobas E / Elecsys	11	100.0	0.0	0.0	496.8	9.9	e

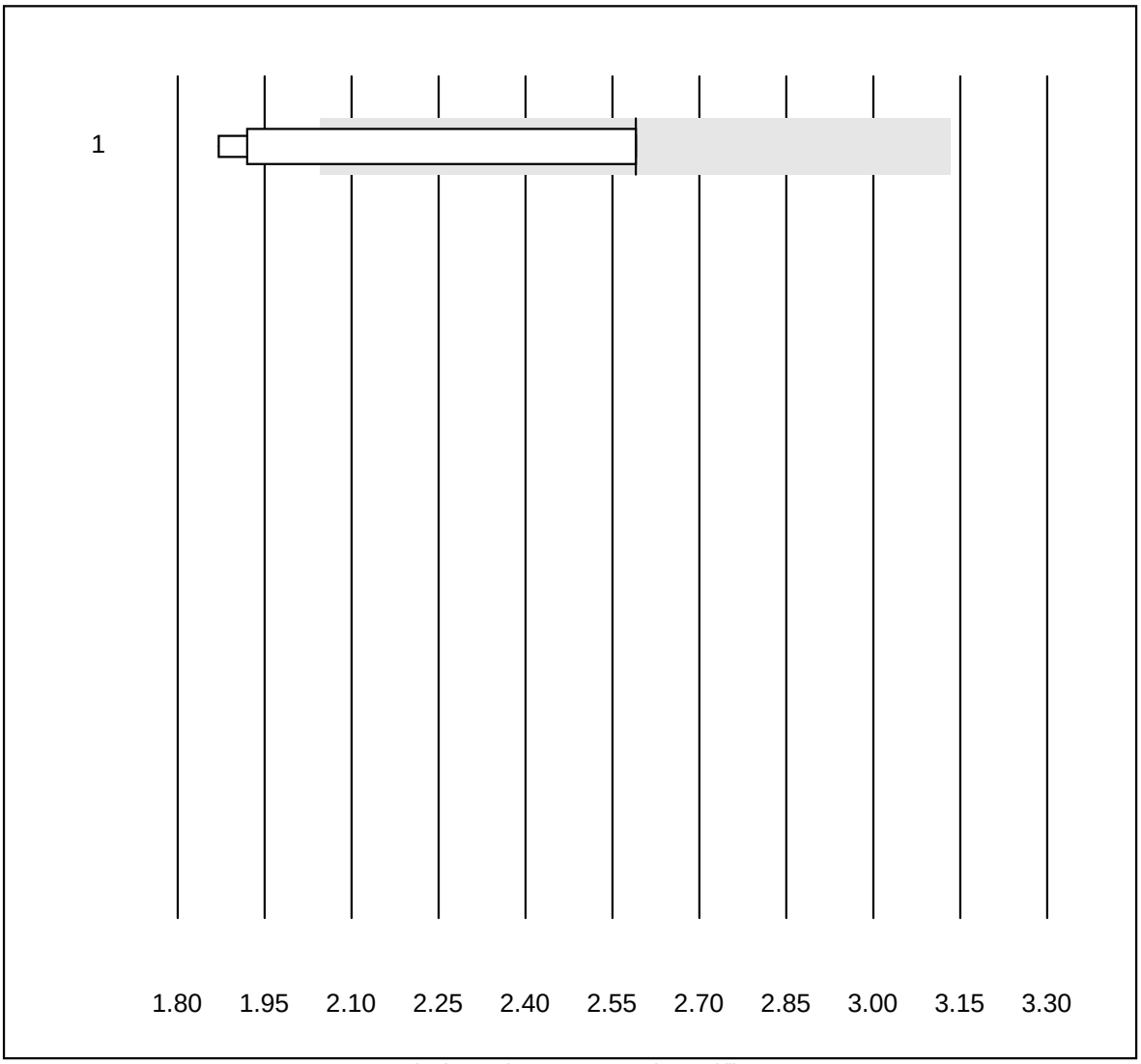
Cholesterin PTS



Deviatione QUALAB : 10 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	CardioChek	5	40.0	40.0	20.0	3.2	10.7	e*

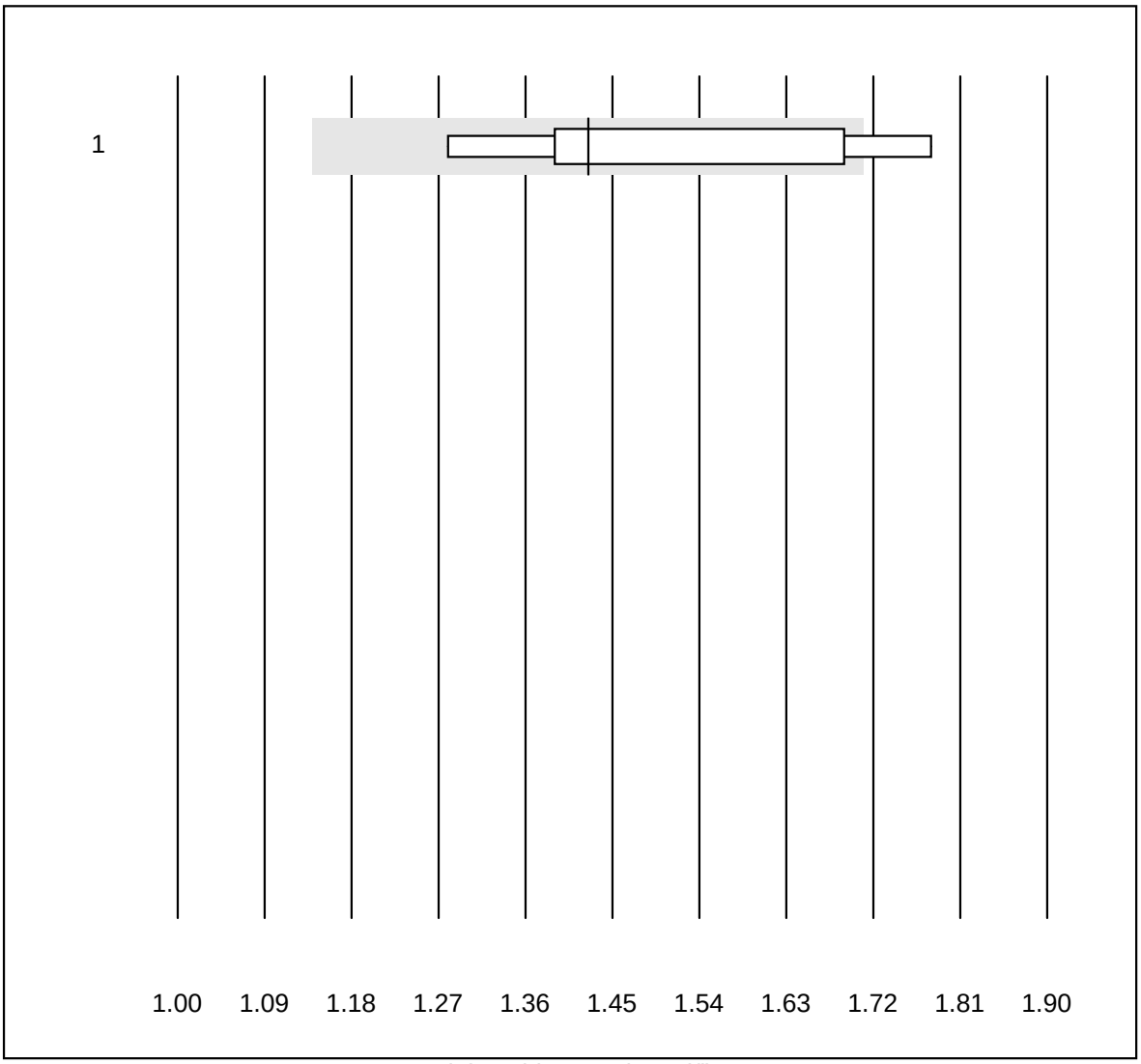
Cholesterin HDL PTS



Deviatione QUALAB : 21 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	CardioChek	5	60.0	40.0	0.0	2.6	16.5	e*

Triglyceride PTS

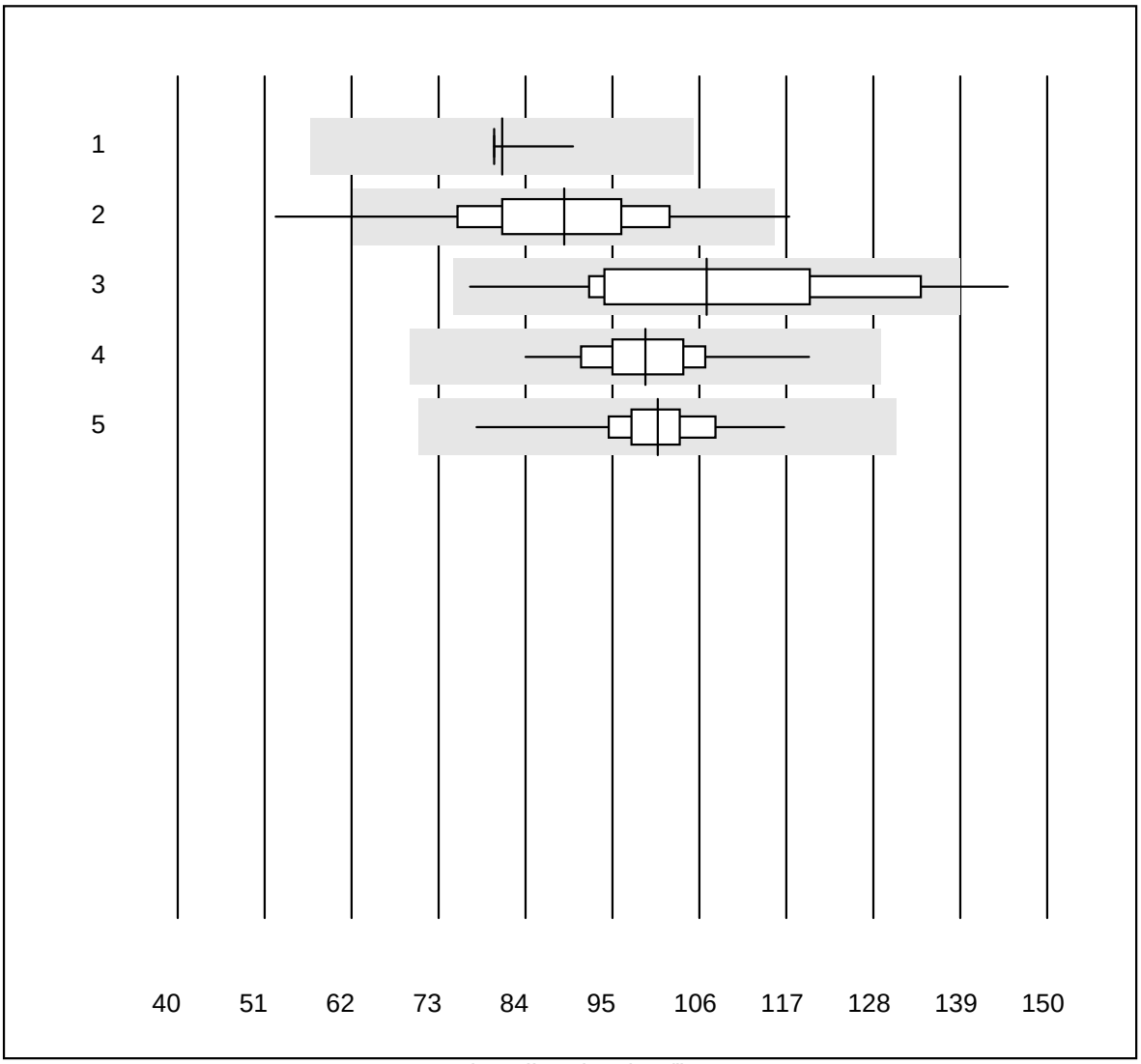


Deviatione QUALAB : 20 %

Triglyceride PTS (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	CardioChek	5	80.0	20.0	0.0	1.43	14.0	e*

Microalbumina

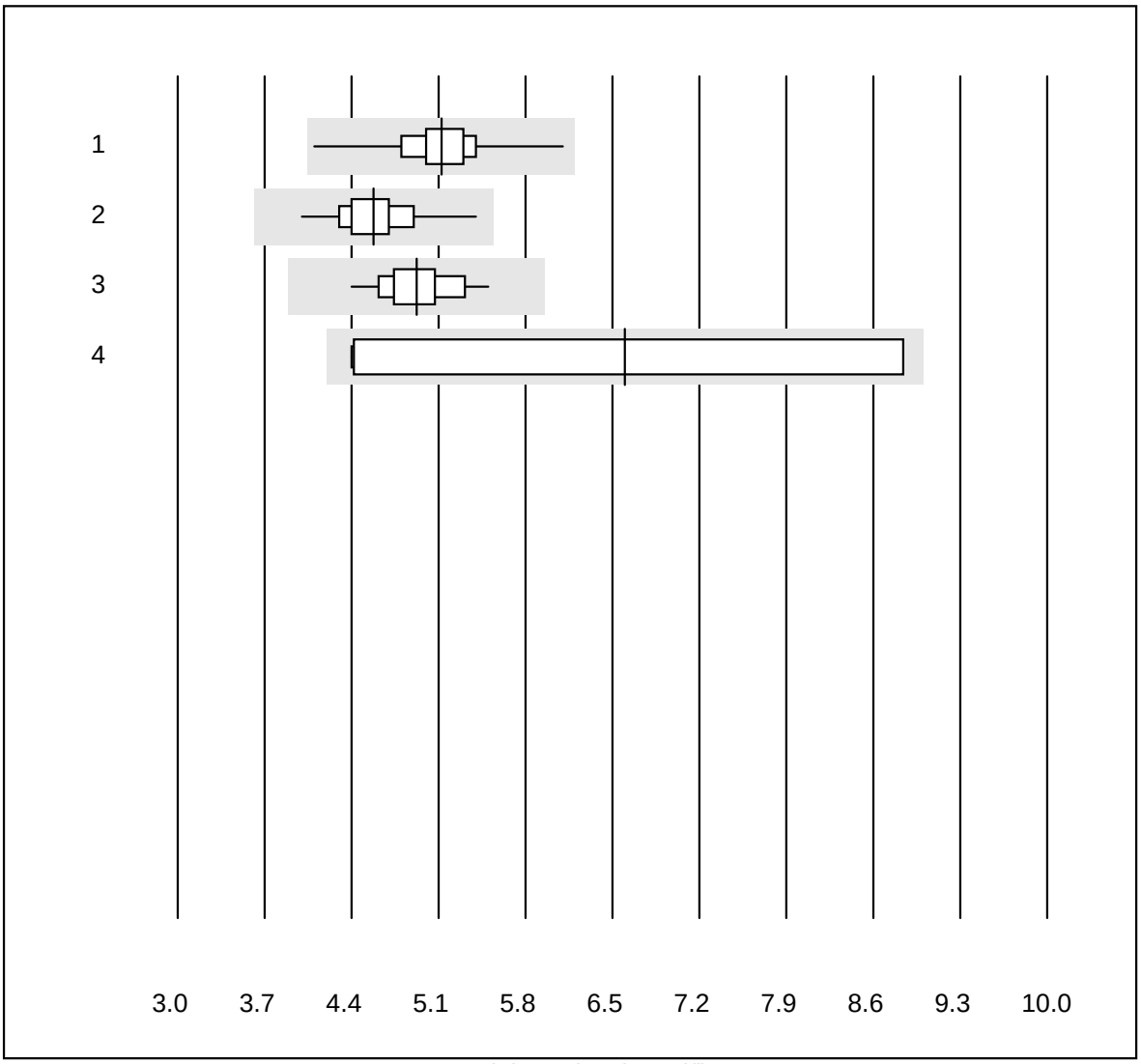


Deviatione QUALAB : 30 %

Microalbumina (mg/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Siemens Clinitek	11	90.9	0.0	9.1	81.0	3.9	e
2	Afinion	326	96.1	1.8	2.1	88.9	12.1	e
3	NycoCard	16	87.4	6.3	6.3	106.9	16.5	e*
4	Turbidimetrie	18	100.0	0.0	0.0	99.2	7.9	e
5	DCA2000/Vantage	113	95.6	0.0	4.4	100.8	5.6	e

Creatinina urina

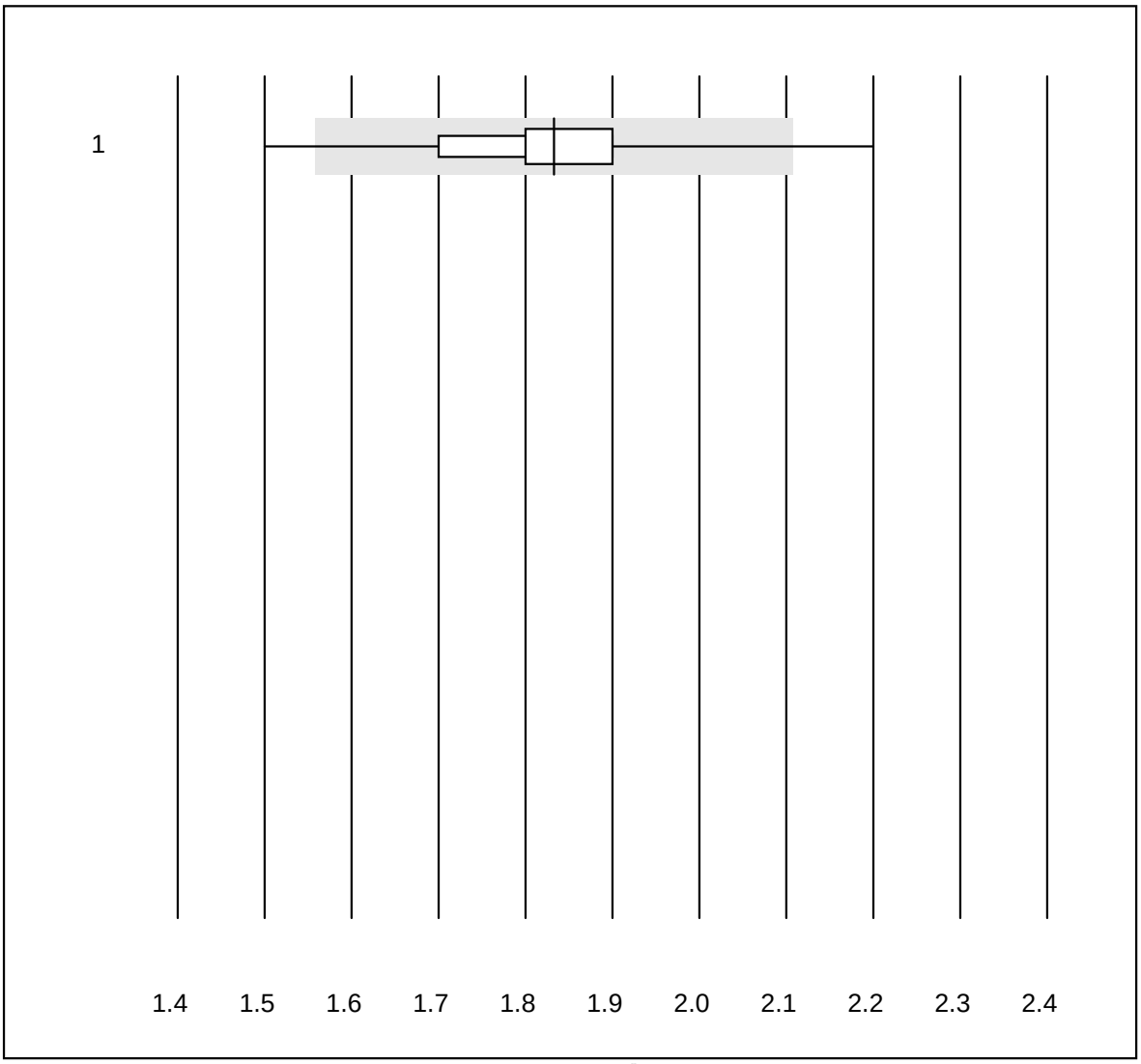


Deviatione QUALAB : 21 %

Creatinina urina (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	DCA2000/Vantage	112	96.4	0.0	3.6	5.1	5.2	e
2	Afinion	327	98.5	0.0	1.5	4.6	5.6	e
3	Chimica umida	27	100.0	0.0	0.0	4.9	5.3	e
4	Siemens Clinitek	10	80.0	0.0	20.0	6.6	31.8	a

INR CCXS

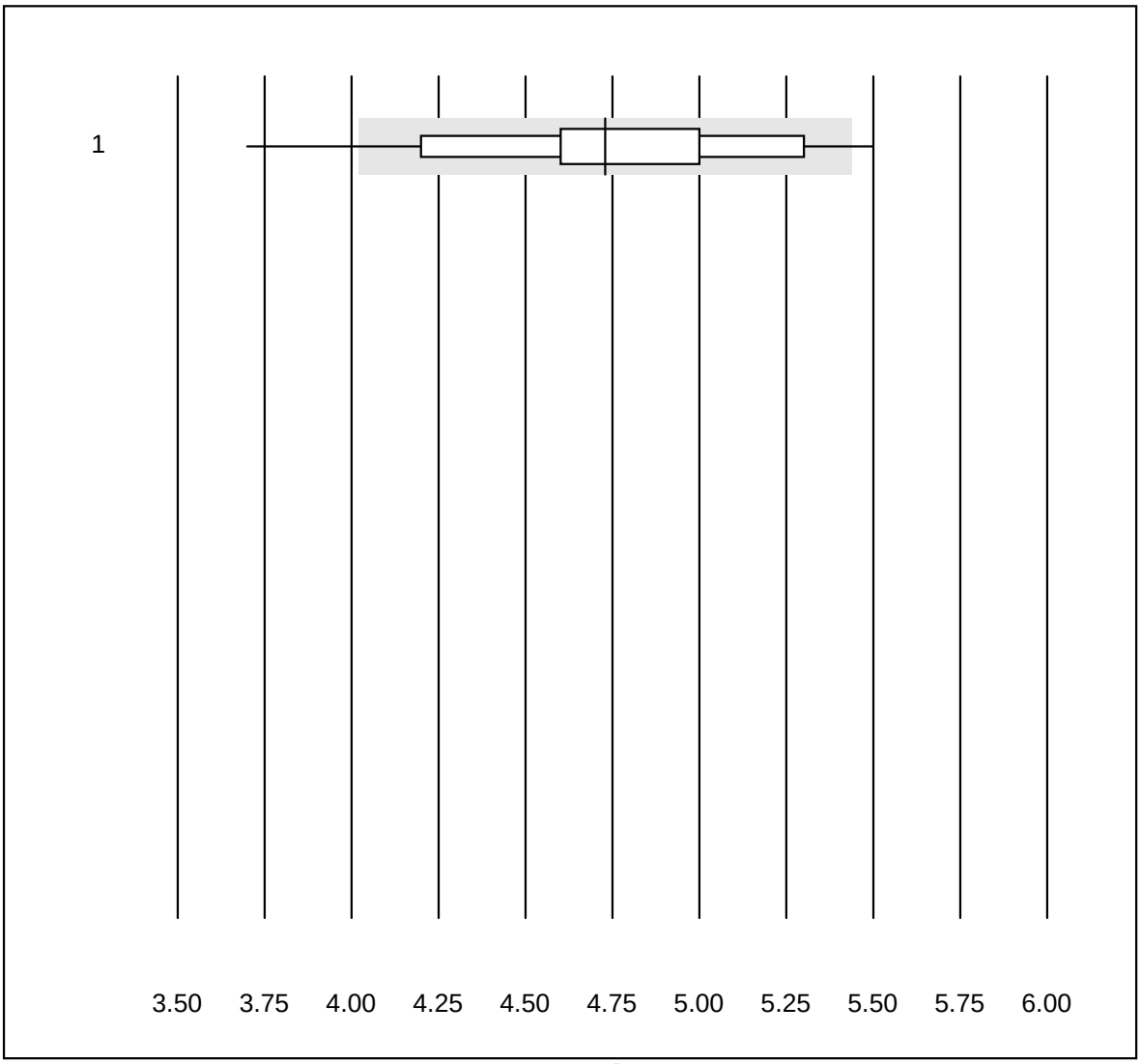


Deviatione QUALAB : 15 %

INR CCXS ()

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	CoaguChek XS	2335	99.1	0.3	0.6	1.8	4.3	e

INR HC

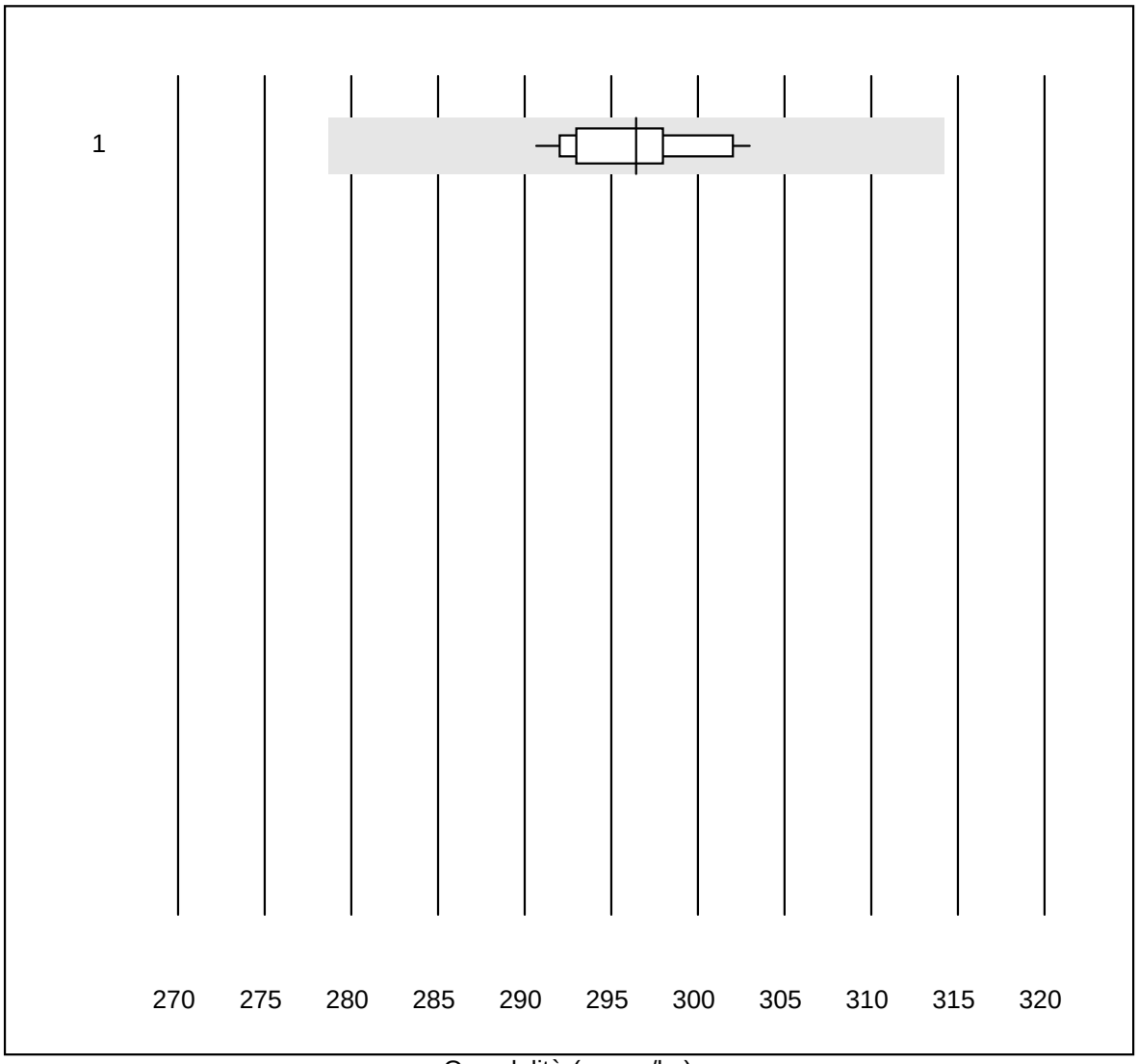


Deviatione QUALAB : 15 %

INR HC ()

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Hemochron j.	20	60.0	10.0	30.0	4.7	10.1	e*

Osmolalità

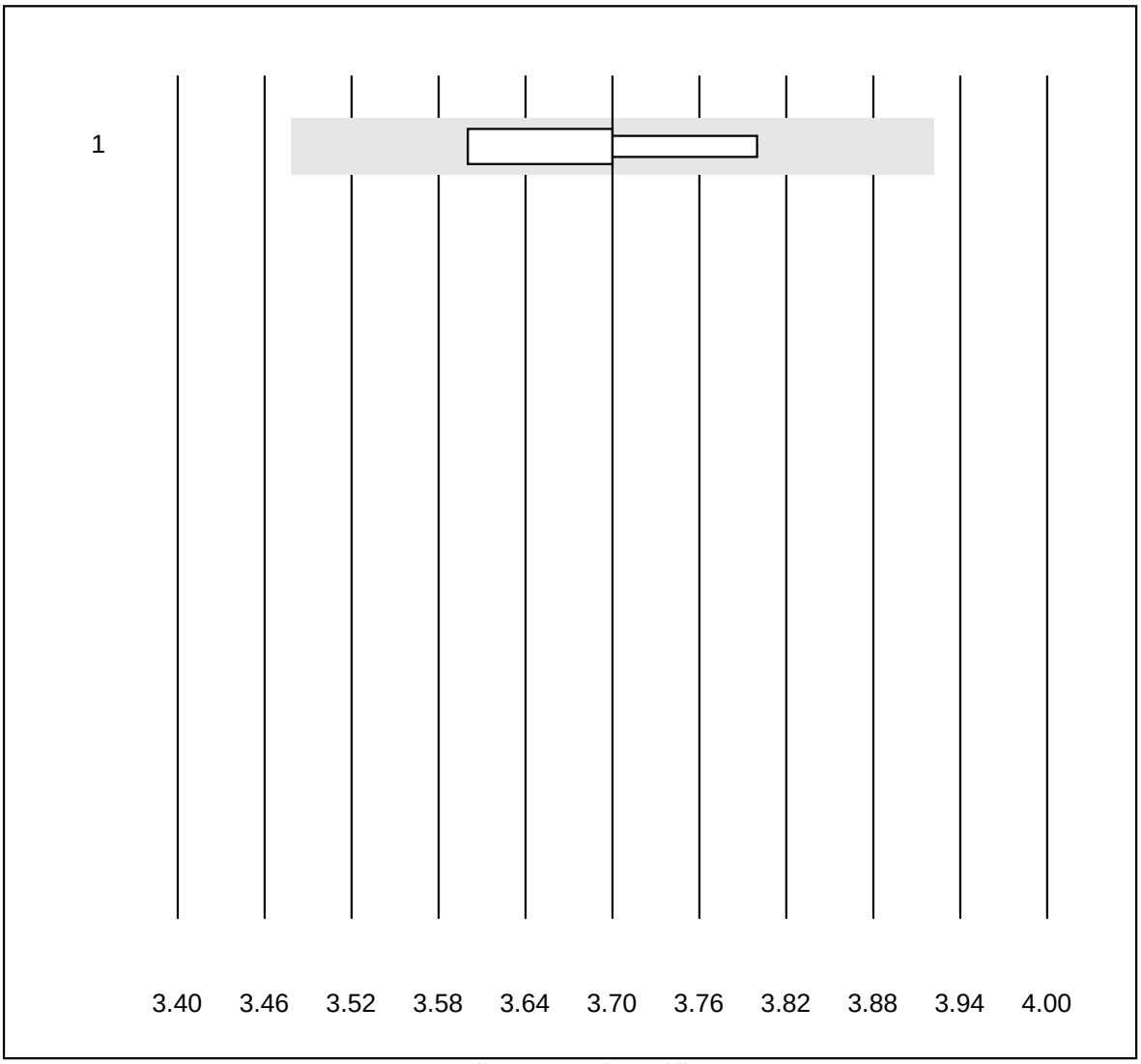


Deviatione QUALAB : 6 %

Osmolalità (mosm/kg)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cryoscopie	11	100.0	0.0	0.0	296	1.3	e

Kalium - K22

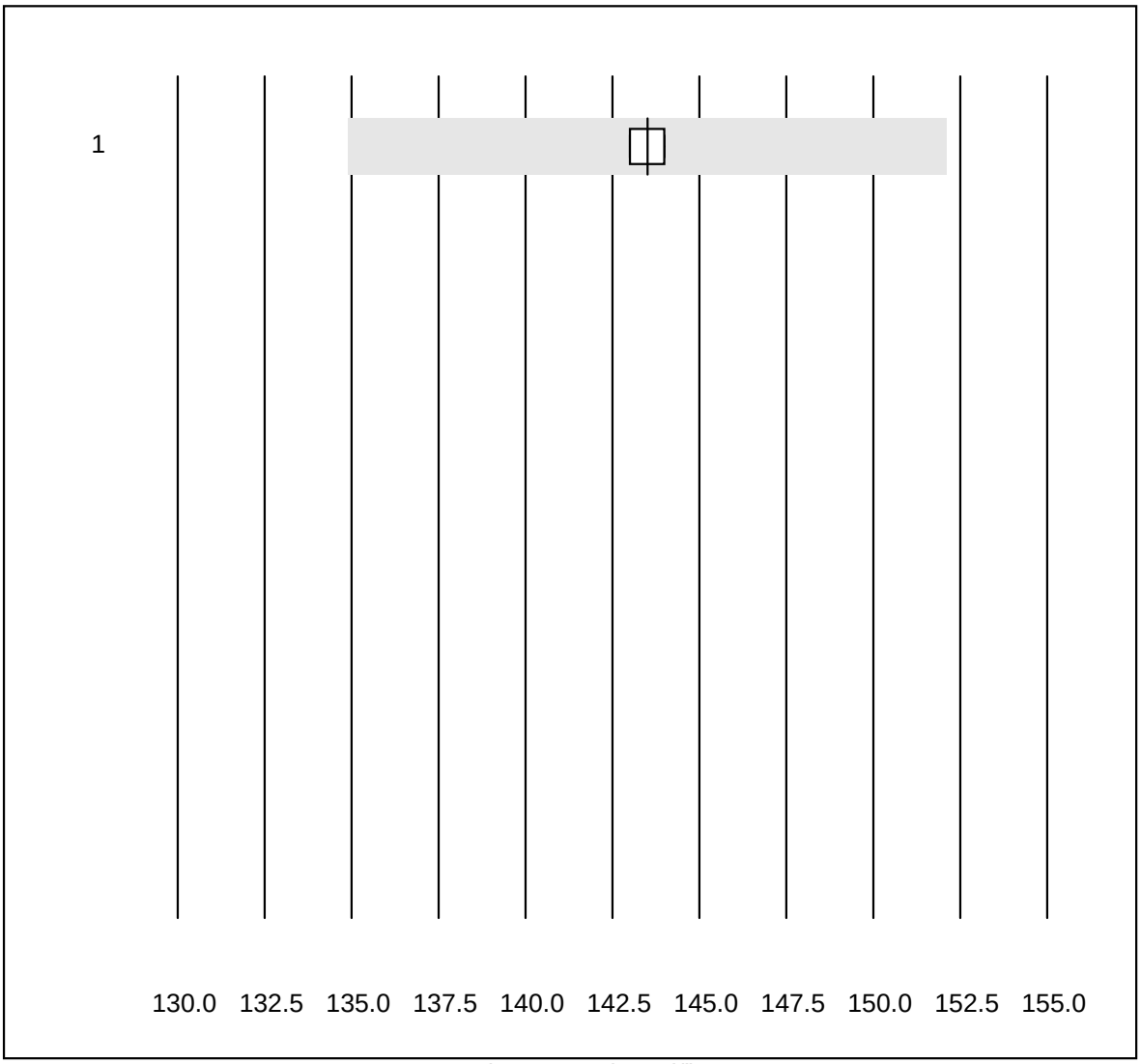


Deviatione QUALAB : 6 %

Kalium - K22 (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ISE	8	100.0	0.0	0.0	3.7	1.8	e

Natrium - K22

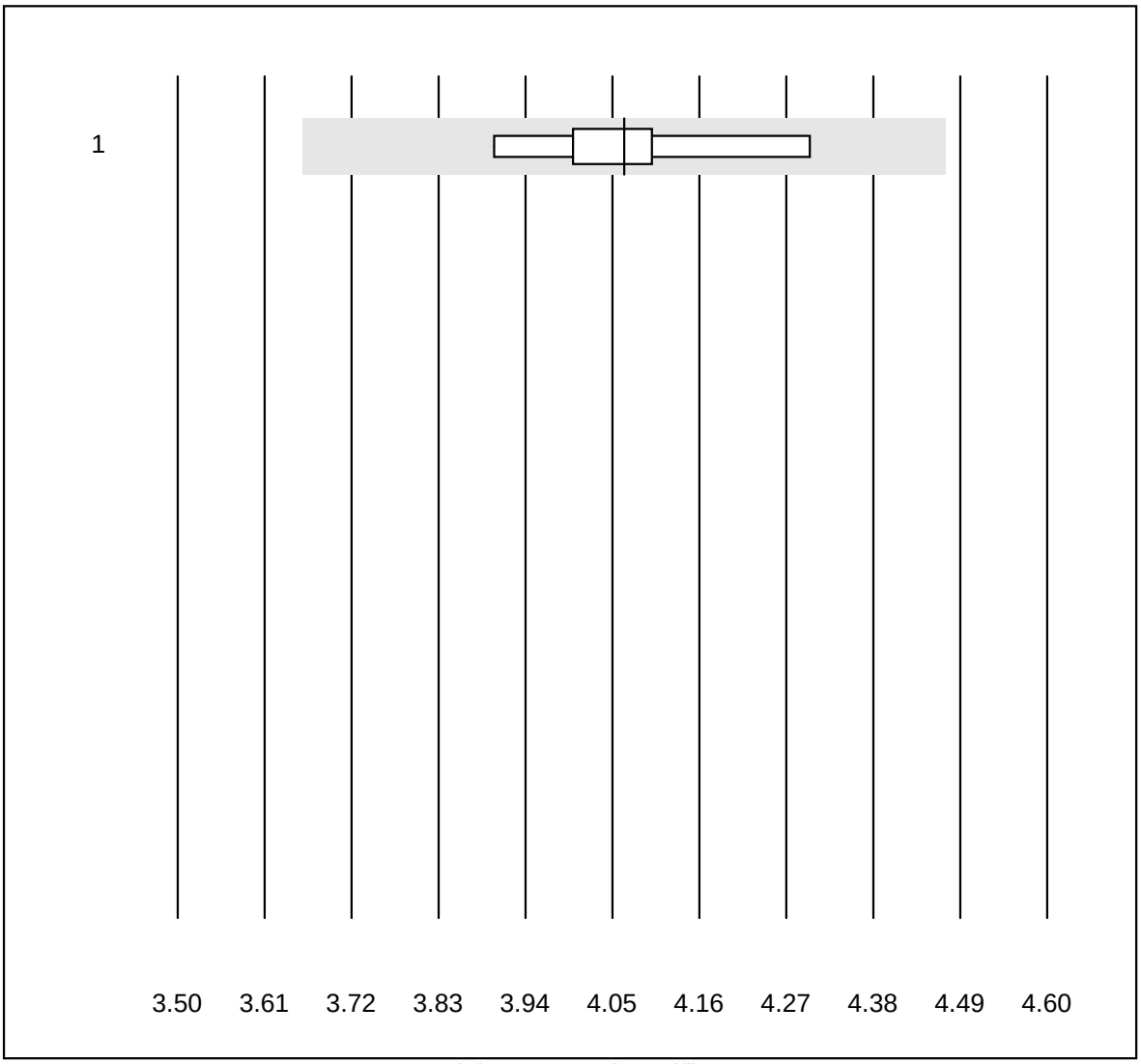


Deviatione QUALAB : 6 %

Natrium - K22 (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	ISE	8	100.0	0.0	0.0	144	0.4	e

Glukose - K22

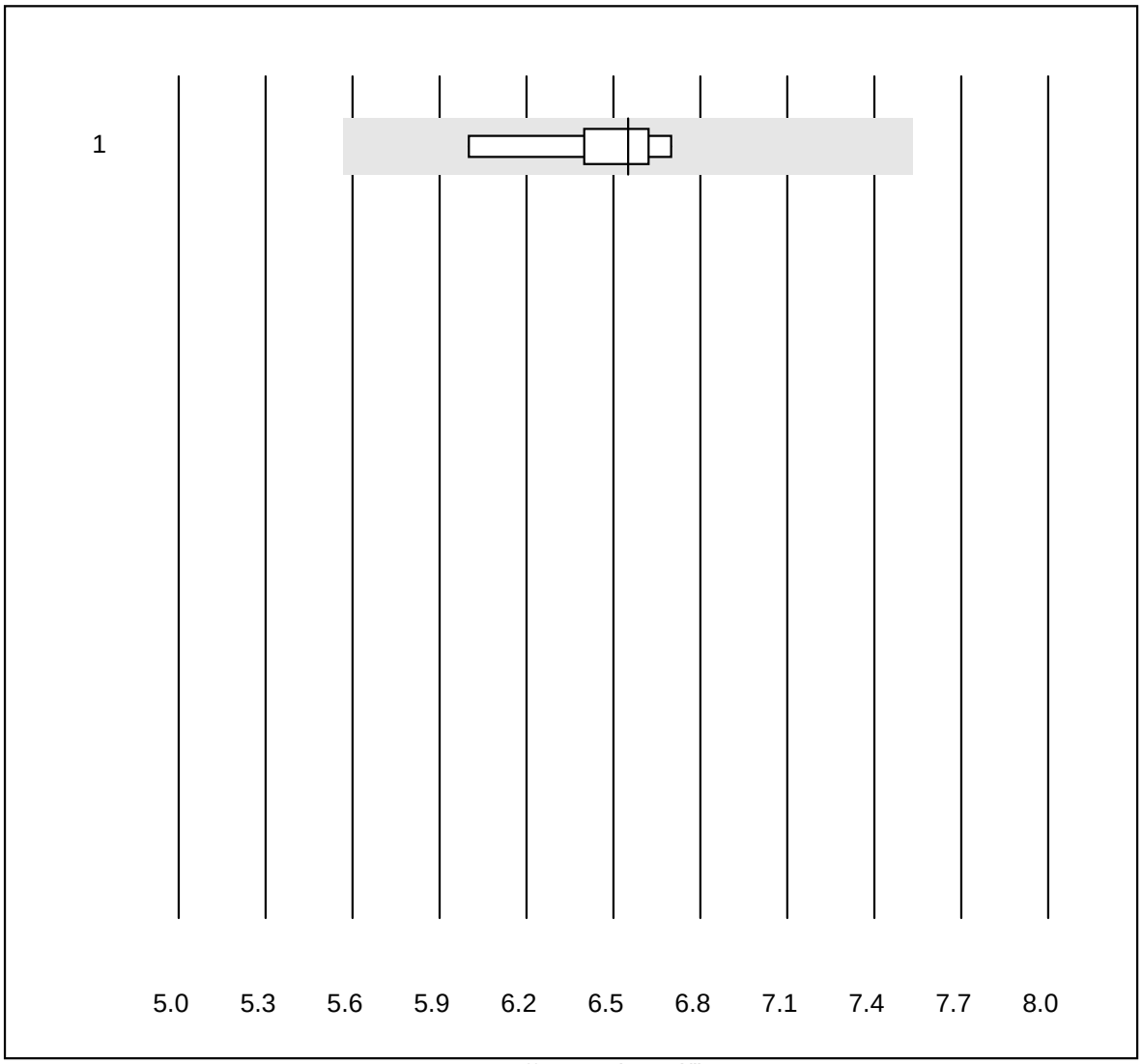


Deviatione QUALAB : 10 %

Glukose - K22 (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	8	100.0	0.0	0.0	4.1	2.9	e

Harnstoff - K22

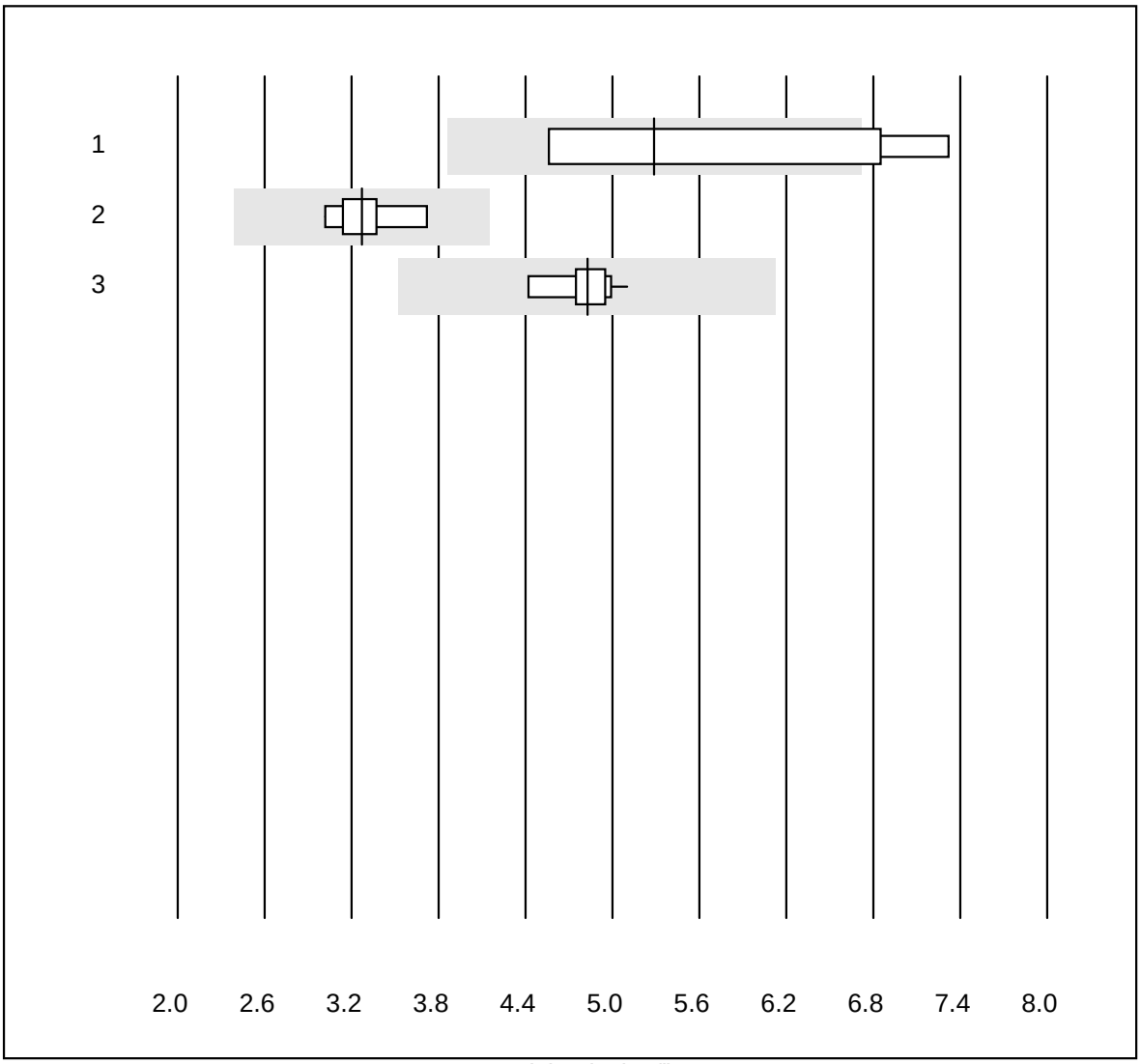


Deviatione QUALAB : 15 %

Harnstoff - K22 (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Chimica umida	8	87.5	0.0	12.5	6.6	3.6	e

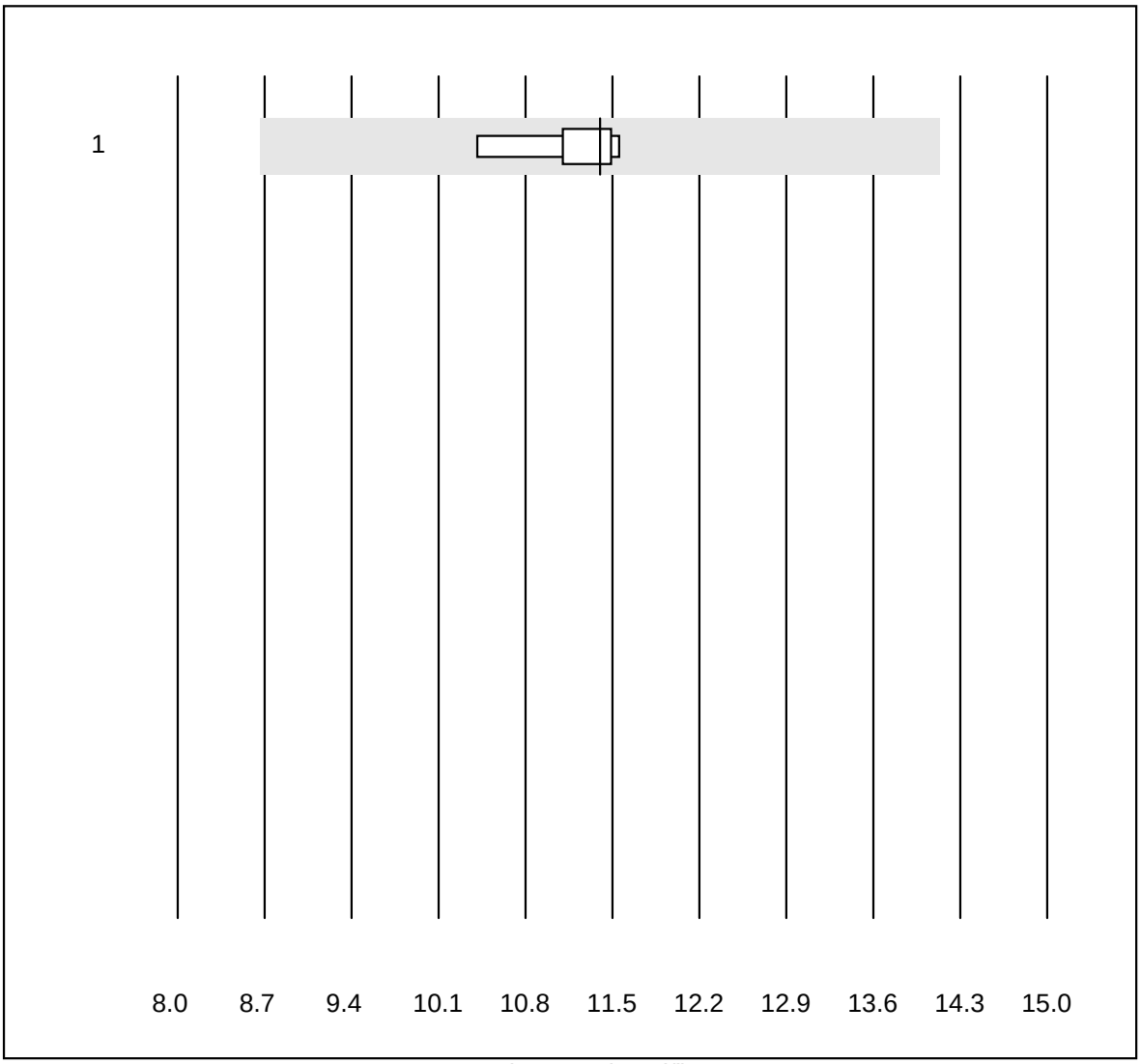
Procalcitonin



Deviatione QUALAB : 27 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	altro	5	40.0	40.0	20.0	5.29	19.6	e*
2	Cobas	9	100.0	0.0	0.0	3.27	6.3	e
3	Mini Vidas	10	100.0	0.0	0.0	4.83	3.9	e

Parathormon

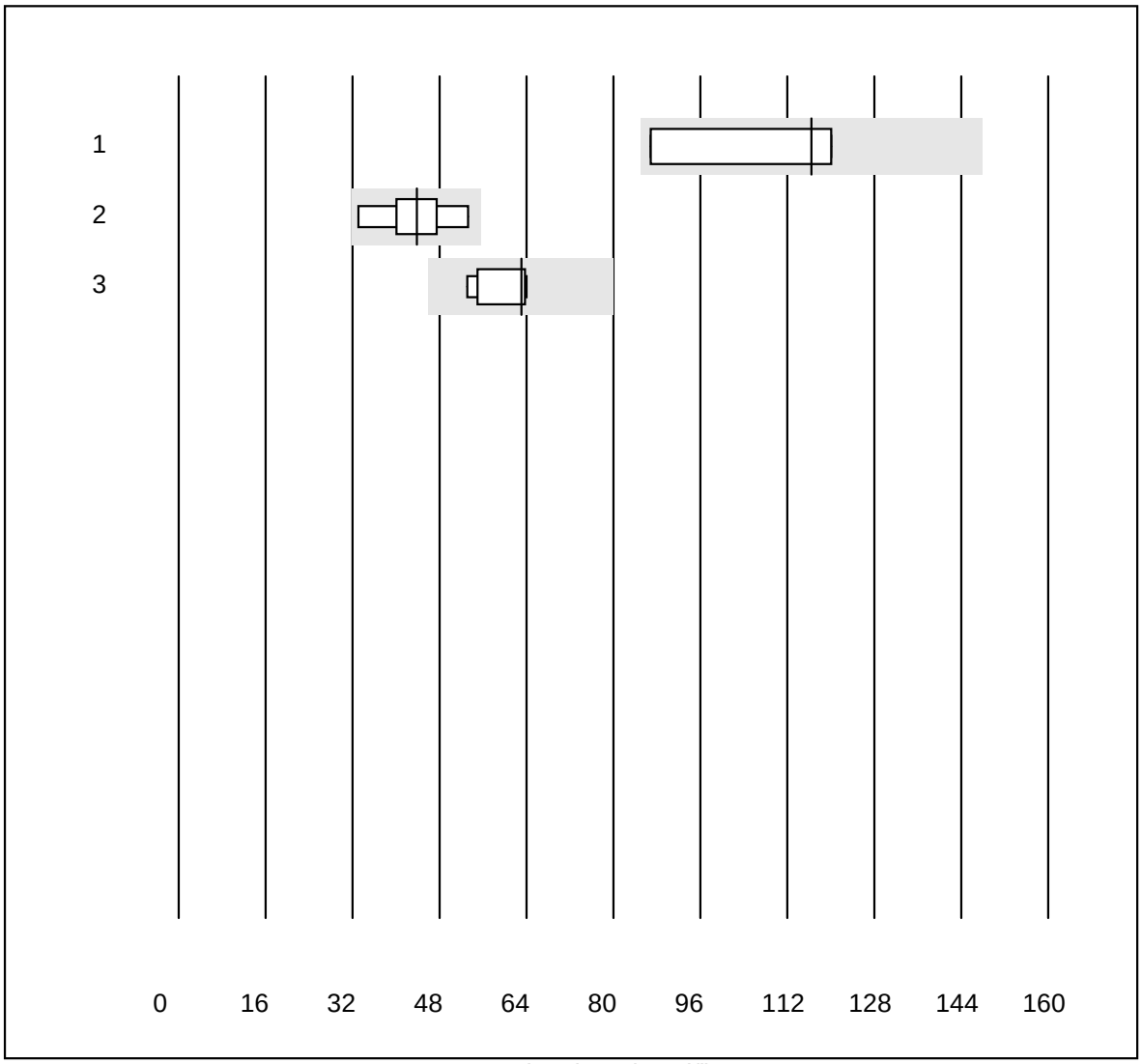


Deviatione QUALAB : 24 %

Parathormon (pmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas PTH STAT	5	100.0	0.0	0.0	11.4	4.2	e

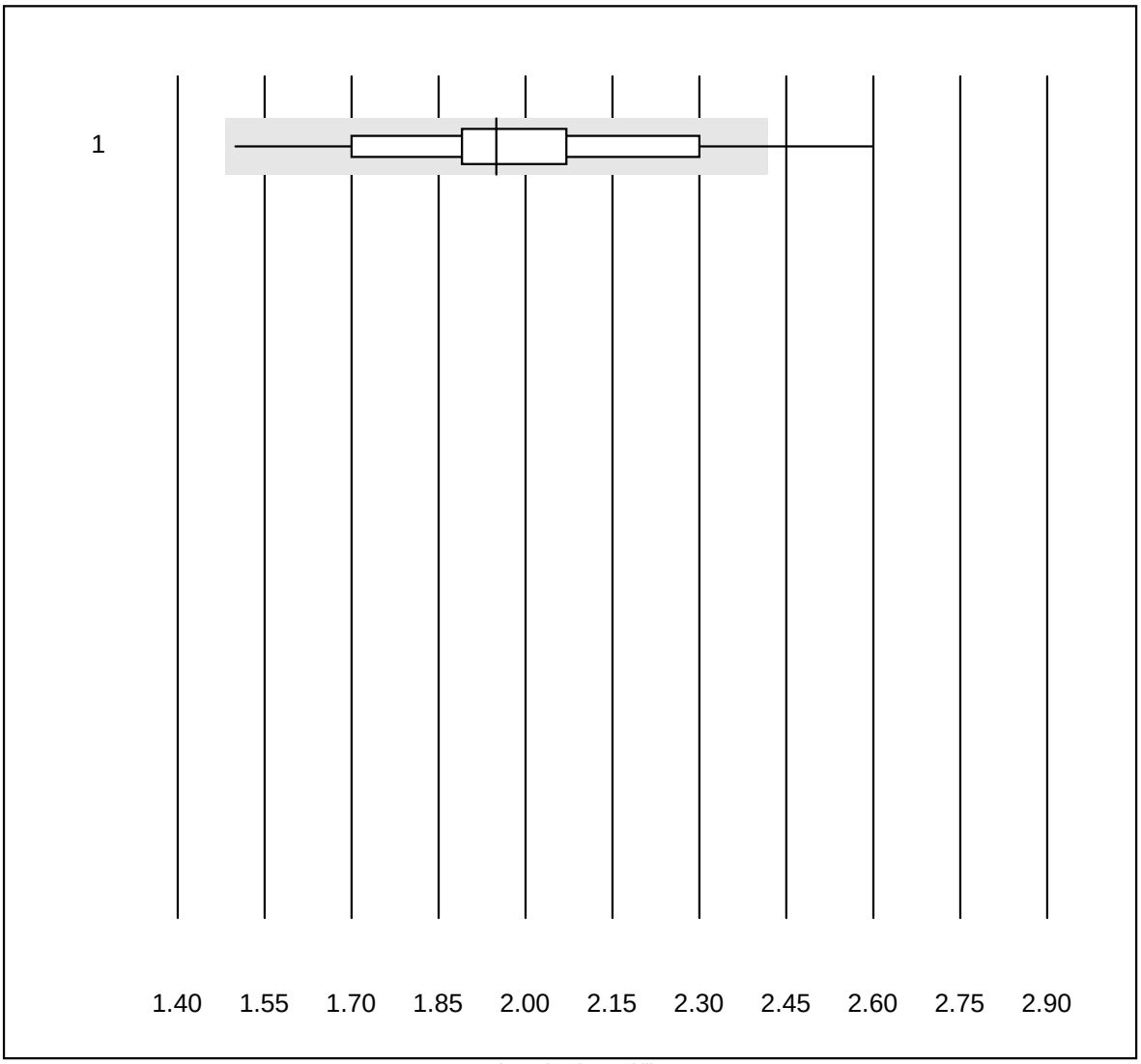
25-OH Vitamin D



Deviatione QUALAB : 27 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Qualigen	4	75.0	0.0	25.0	116.5	16.4	e*
2	Cobas	7	100.0	0.0	0.0	43.8	14.5	e*
3	Architect	5	100.0	0.0	0.0	63.0	8.8	e*

Digoxin

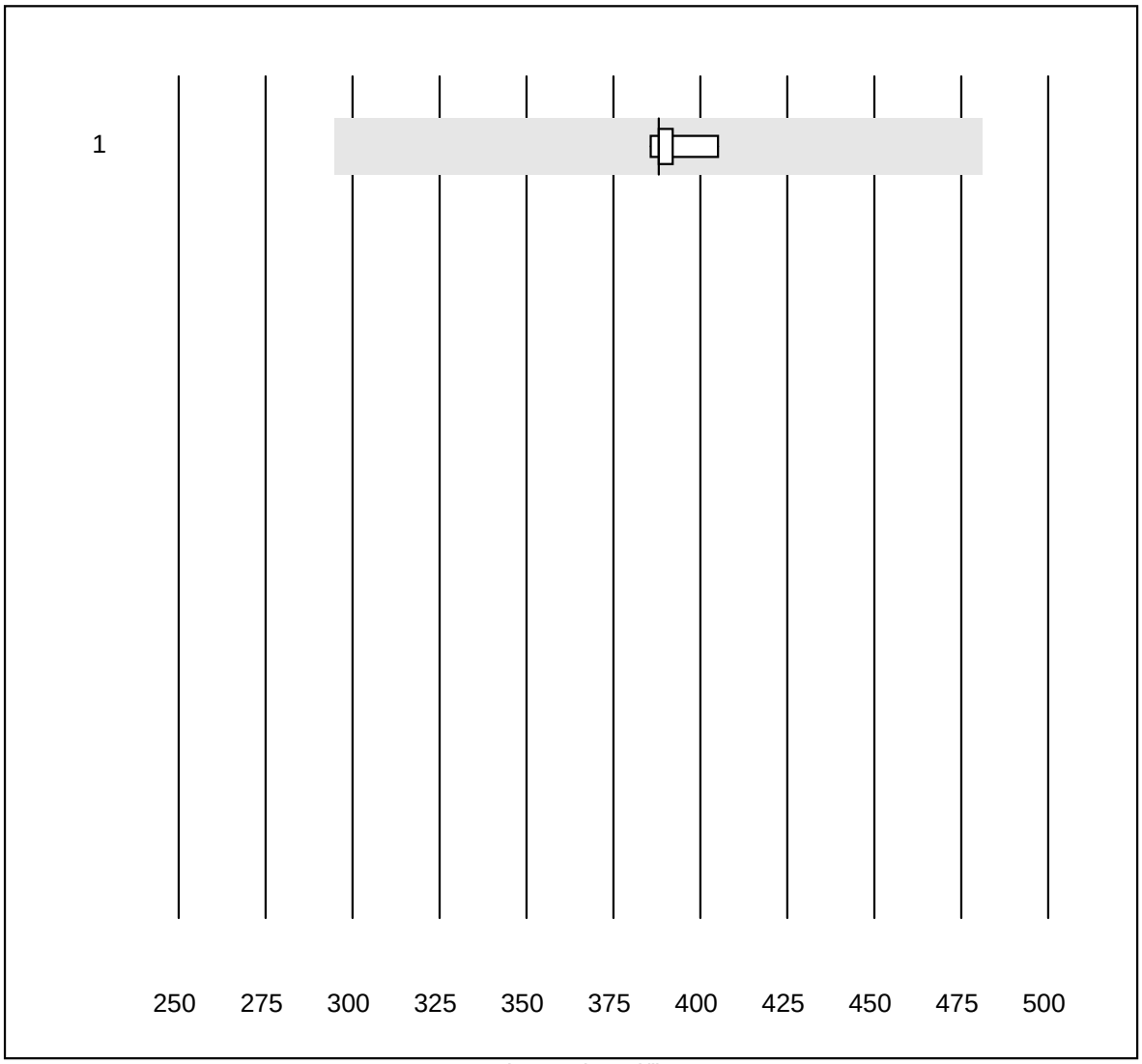


Deviatione QUALAB : 24 %

Digoxin (nmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	altro	11	90.9	9.1	0.0	1.95	14.5	e*

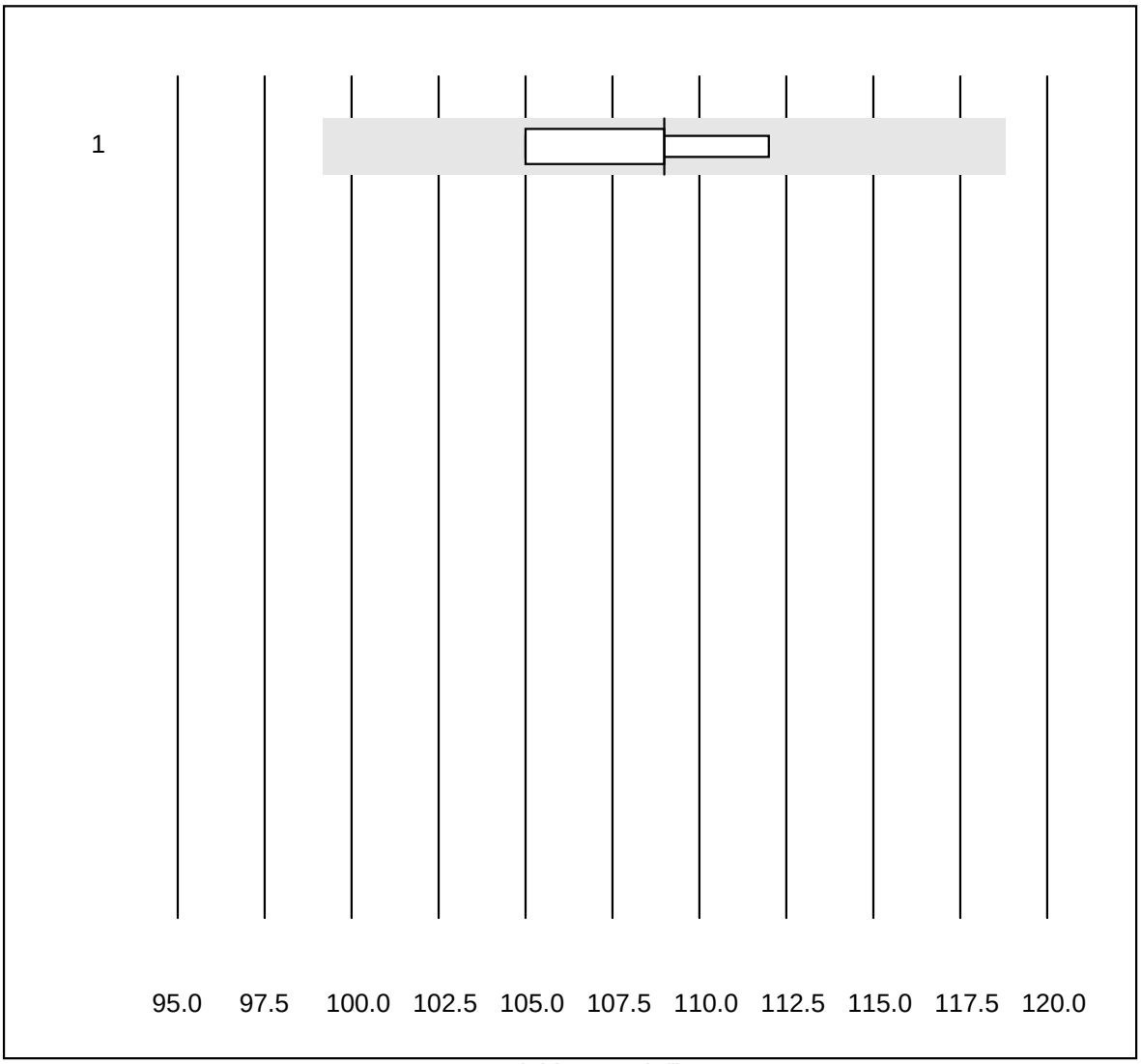
Valproat



Deviatione QUALAB : 24 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	tutti	5	100.0	0.0	0.0	388.0	2.0	e

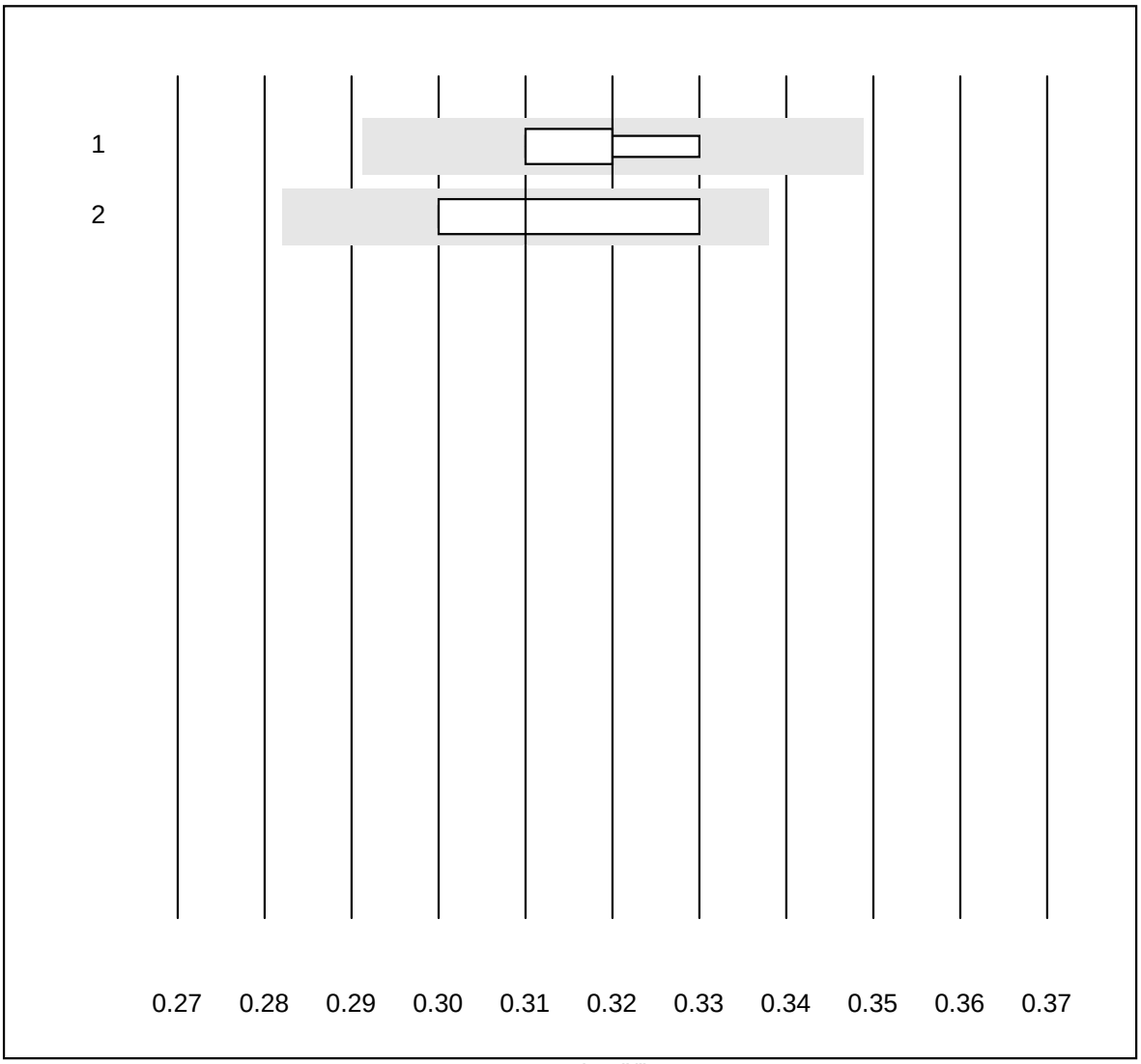
Emoglobina BG



Deviatione QUALAB : 9 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	iStat	4	100.0	0.0	0.0	109.0	2.6	e*

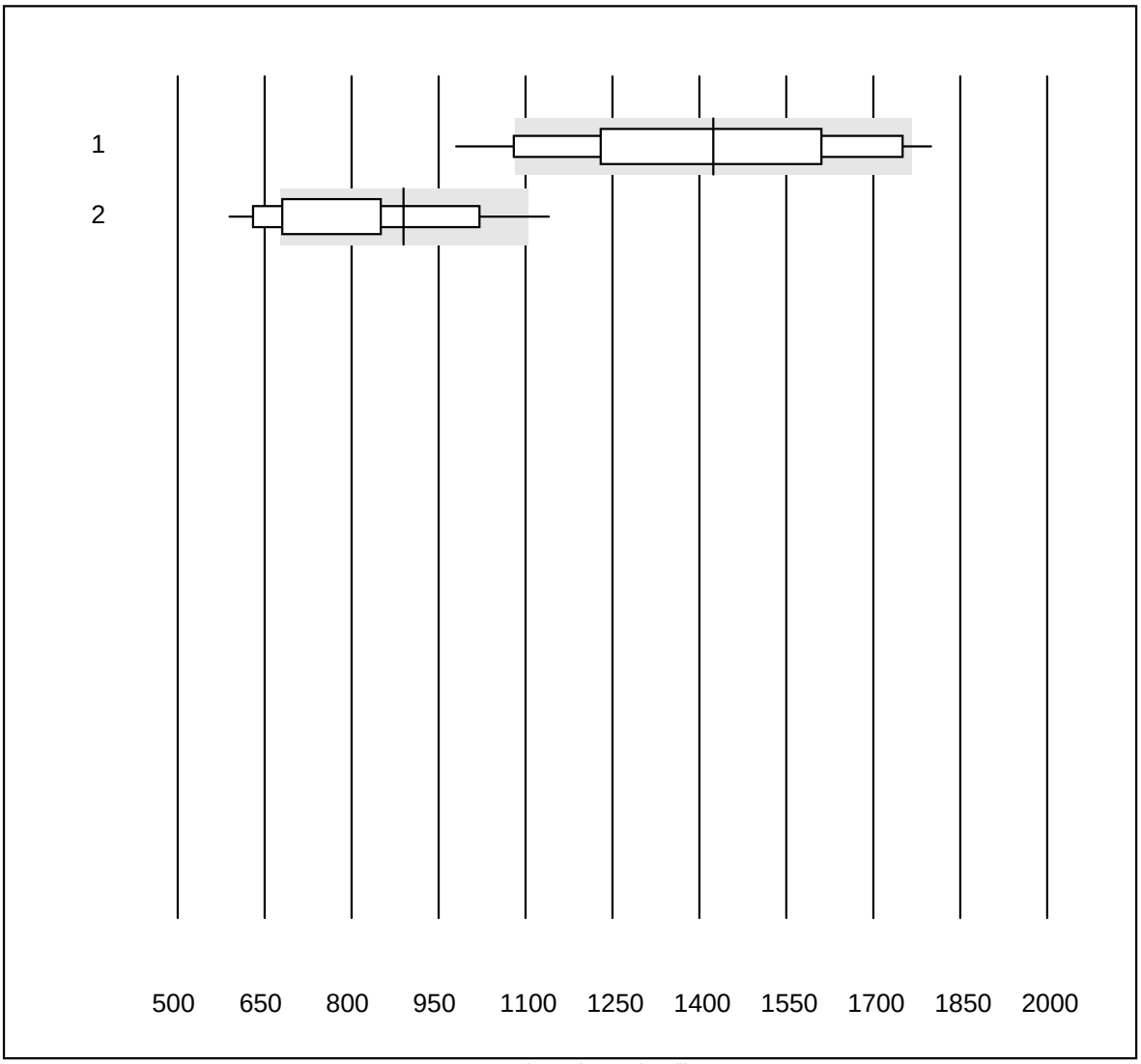
Ematocrito



Deviatione QUALAB : 9 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	iStat	4	100.0	0.0	0.0	0.32	2.6	e*
2	EPOC	4	75.0	0.0	25.0	0.31	4.8	e*

Troponin Triage

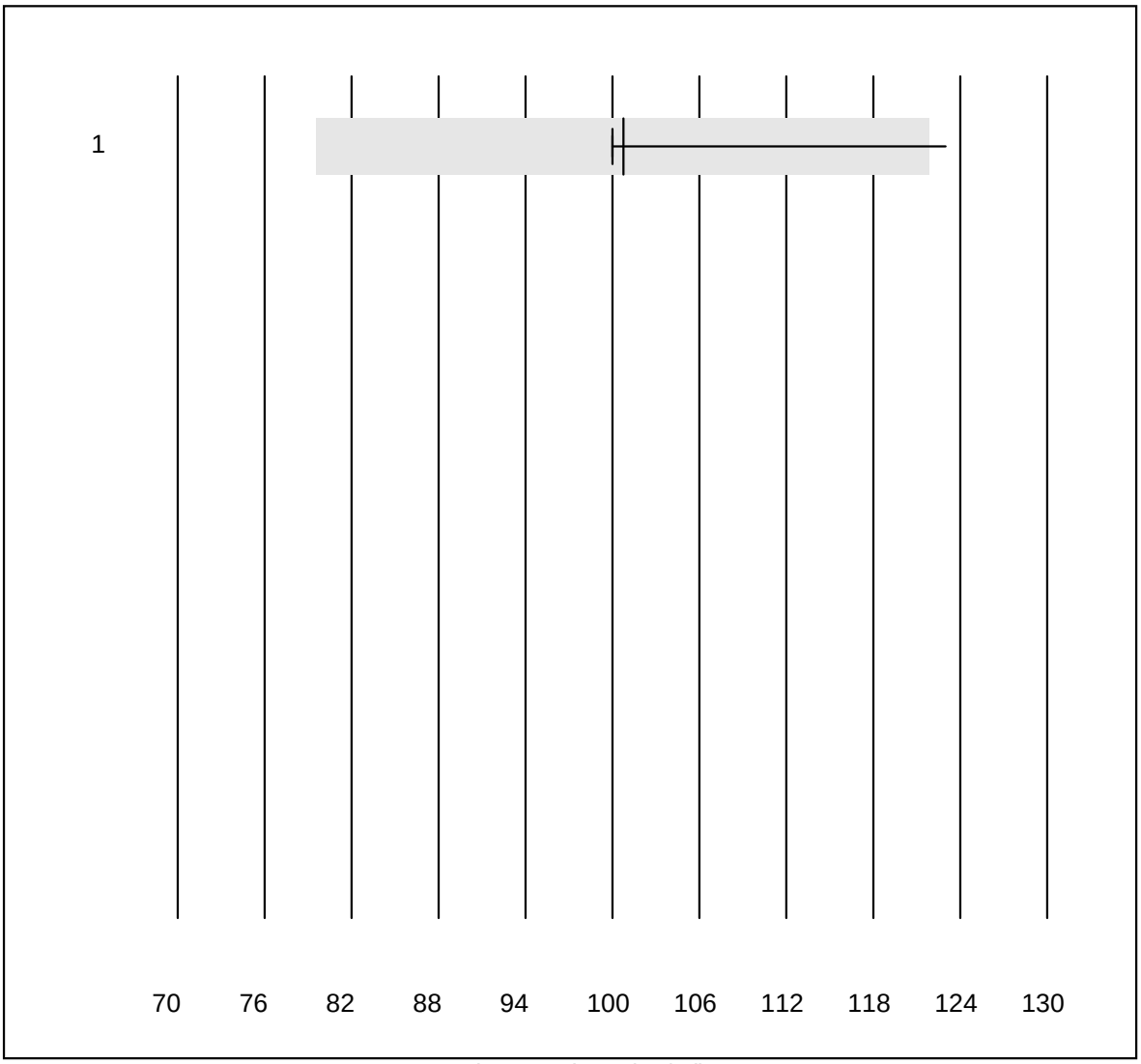


Deviatione QUALAB : 24 %

Troponin Triage (ng/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Triage Next Gen	18	61.1	16.7	22.2	1424.40	18.5	a
2	Triage SOB/Cardiac	24	62.5	20.8	16.7	890.00	18.5	a

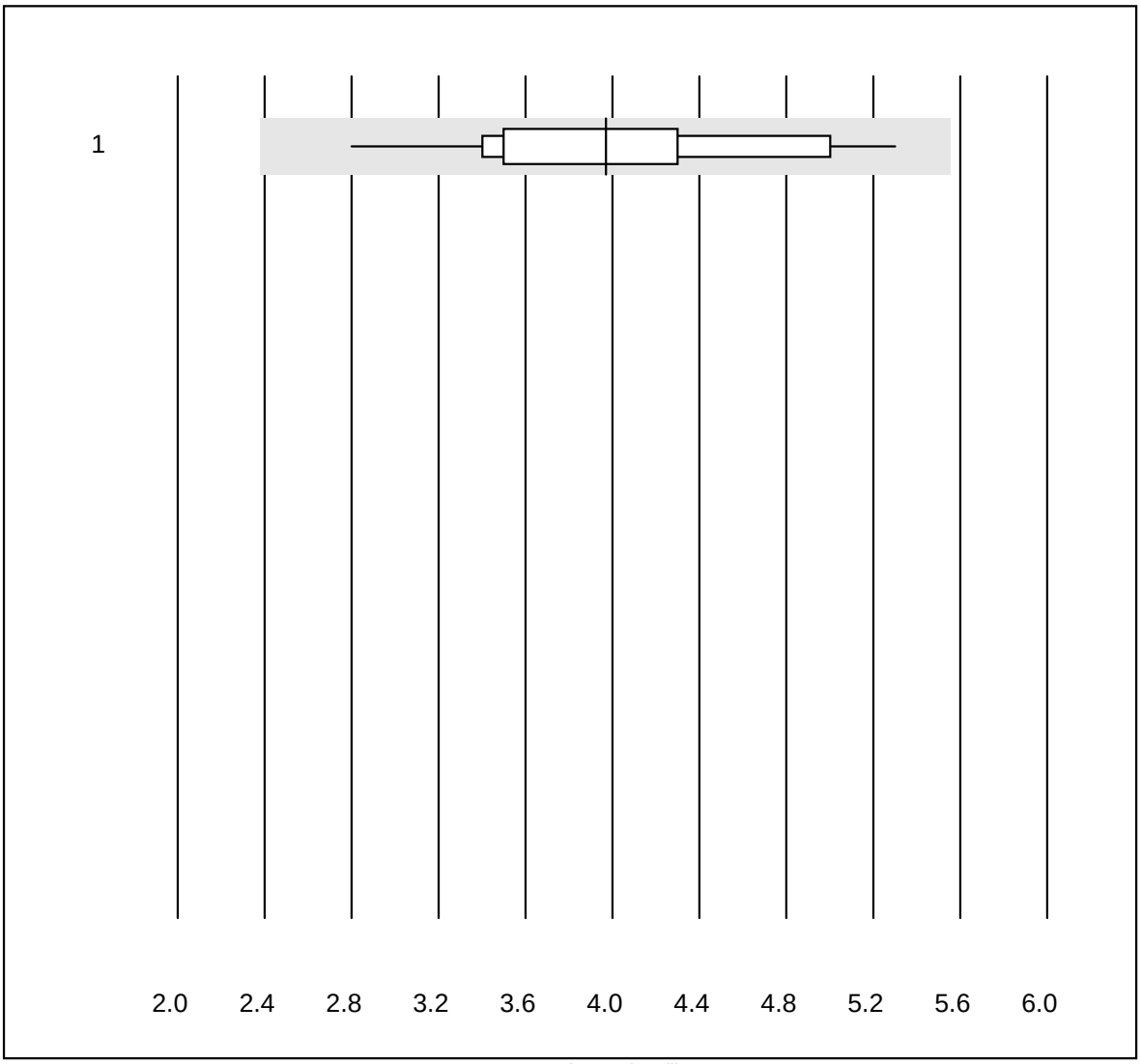
D-Dimere Triage



Deviatione QUALAB : 21 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Triage Meter	40	92.5	2.5	5.0	100.74	3.8	e

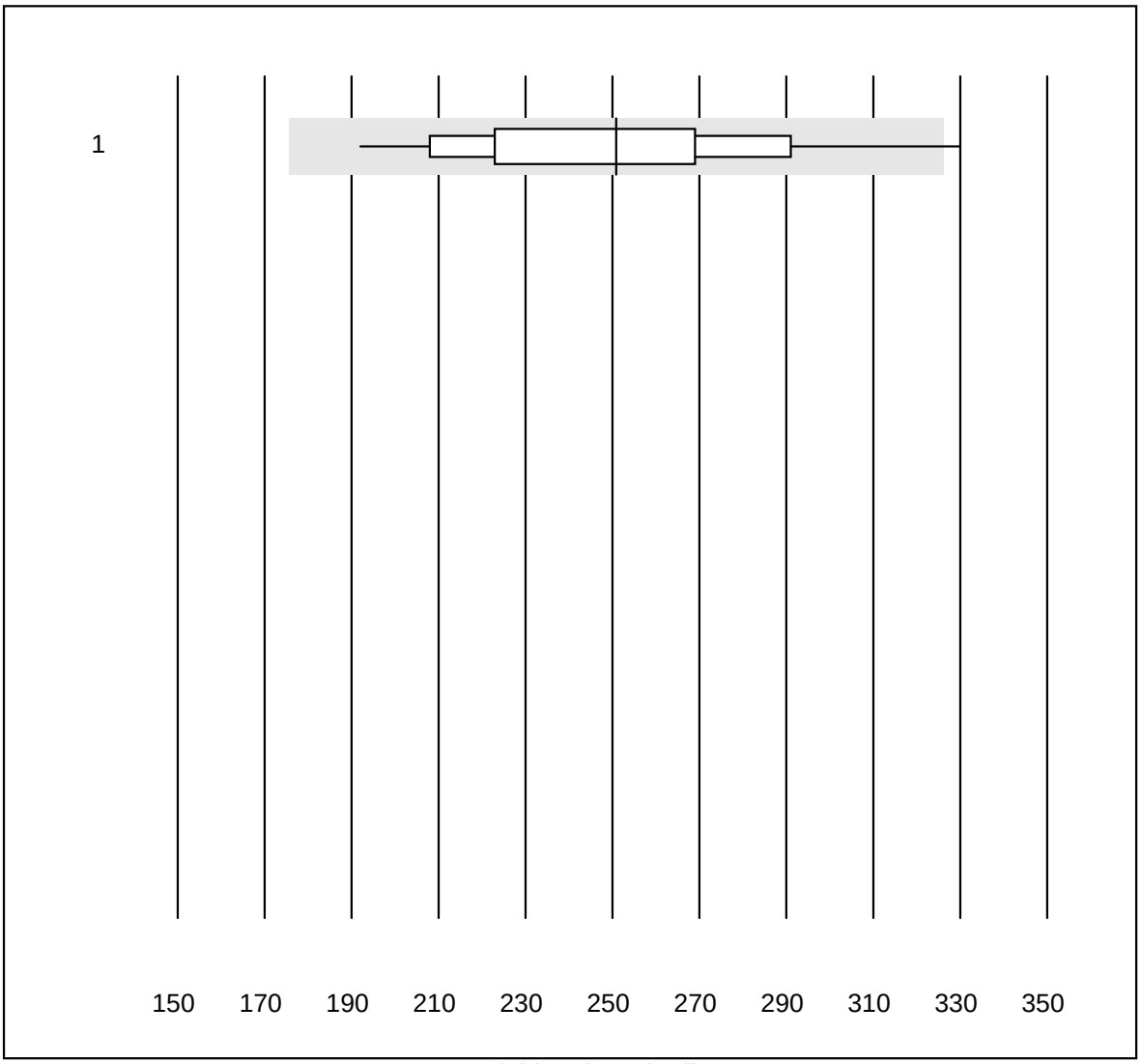
CK-MB Triage



Deviatione QUALAB : 40 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Triage Meter	23	95.7	0.0	4.3	4.0	15.5	e

Myoglobin Triage

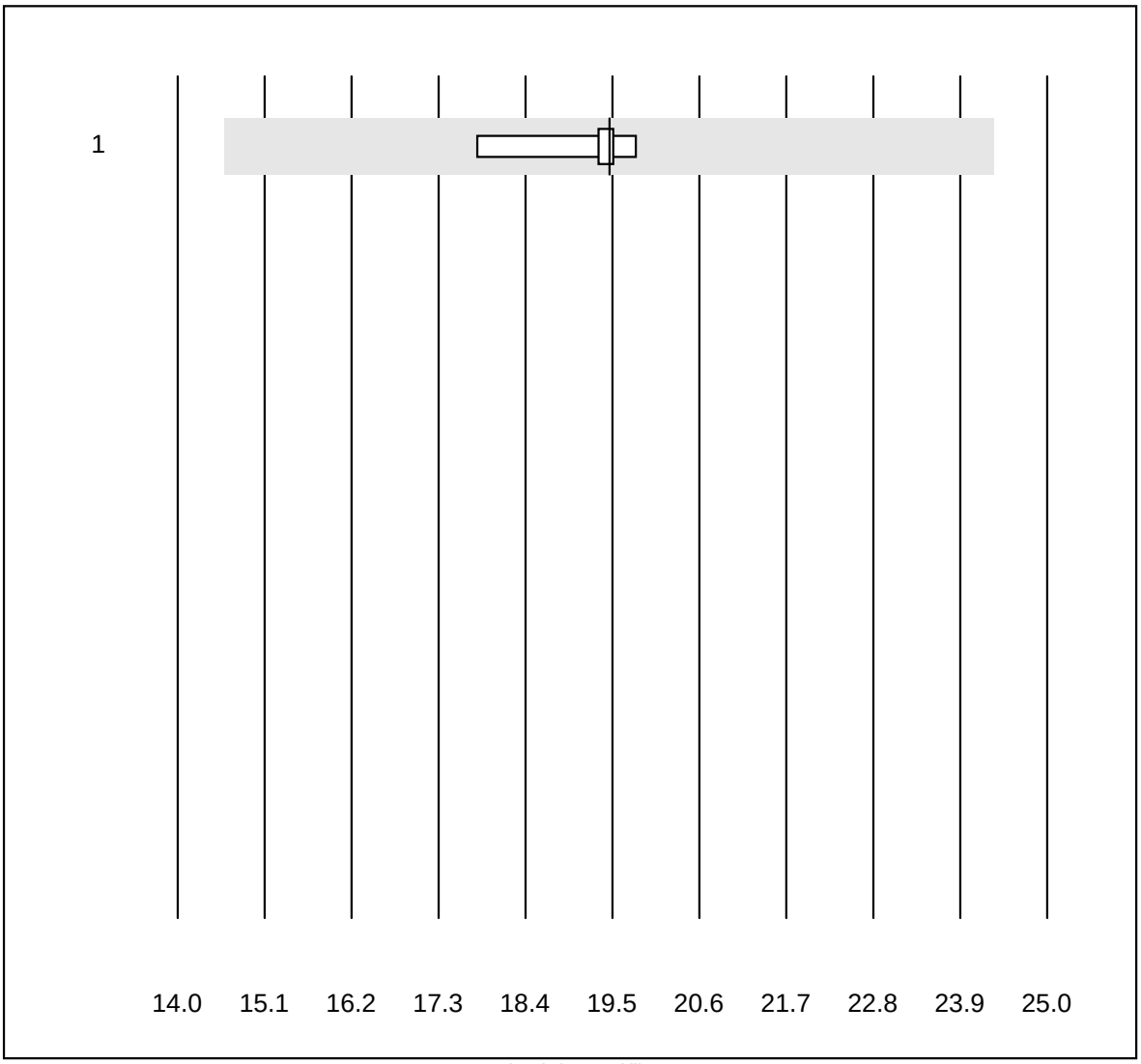


Deviatione QUALAB : 30 %

Myoglobin Triage (µg/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Triage Meter	21	90.4	4.8	4.8	250.9	14.0	e

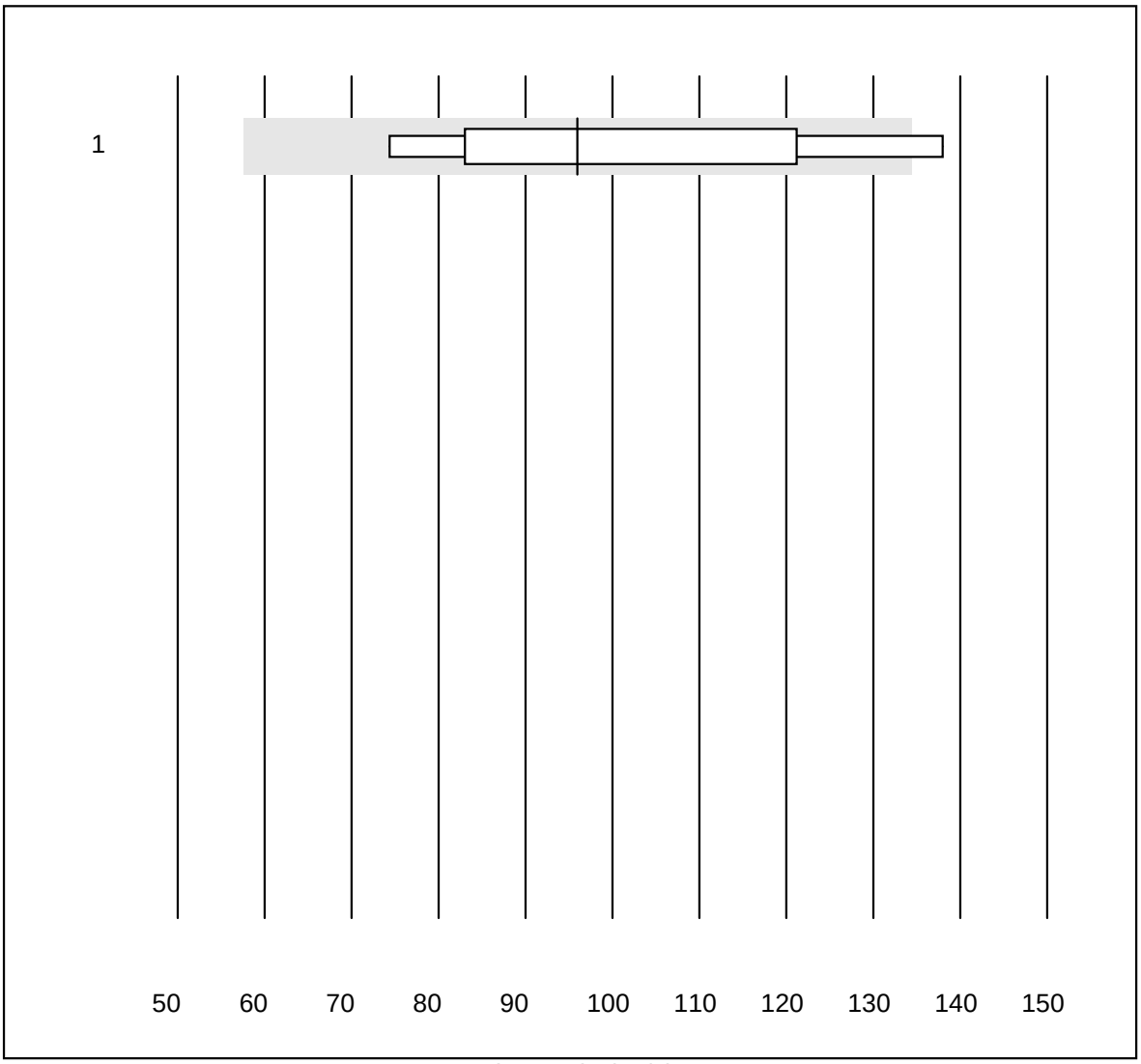
Alcol



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	tutti	5	100.0	0.0	0.0	19.5	4.1	e

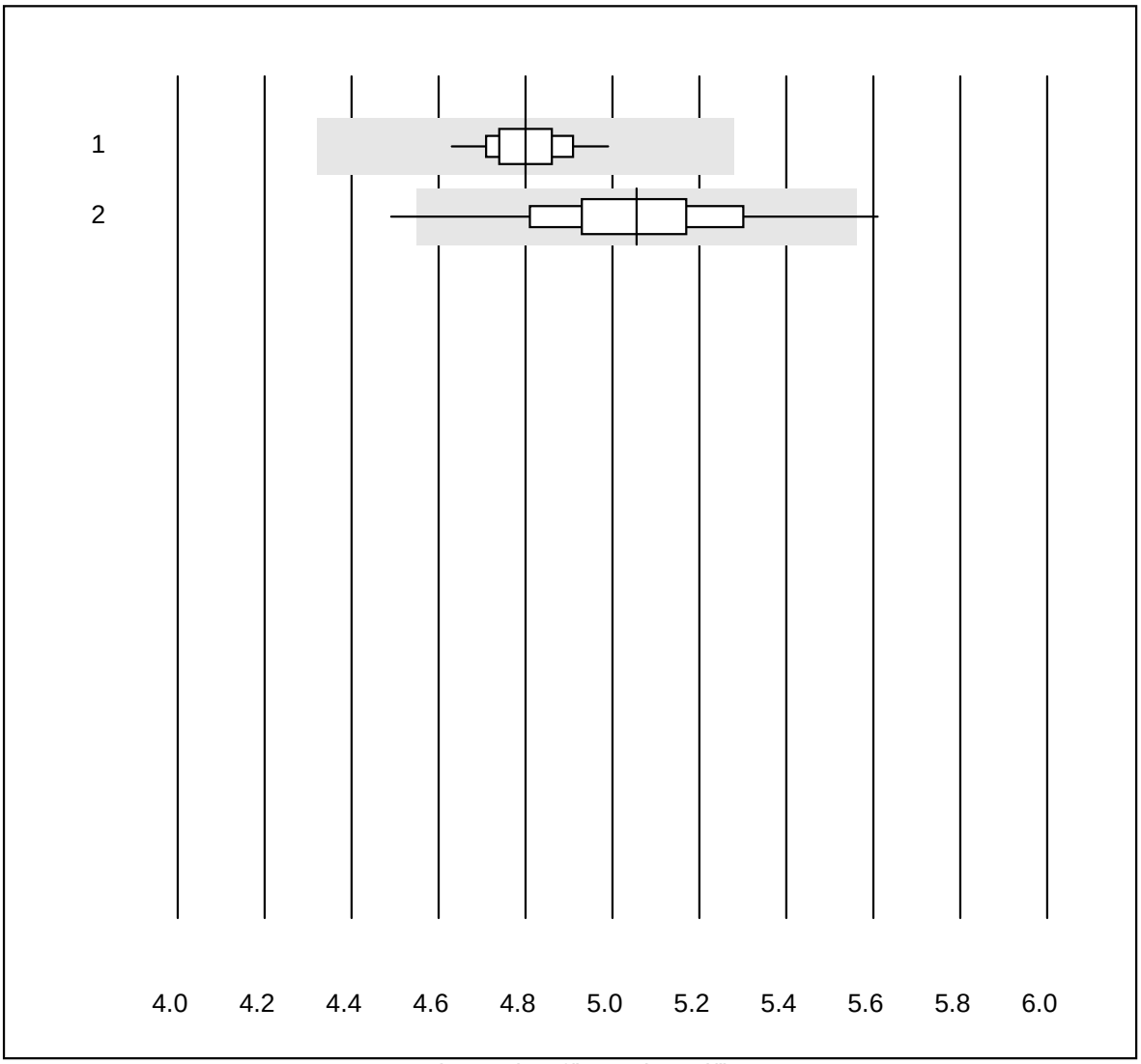
Calprotectin



Deviatione QUALAB : 40 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Bühlmann	7	71.4	14.3	14.3	96	23.1	e*

Colesterolo Af/b101

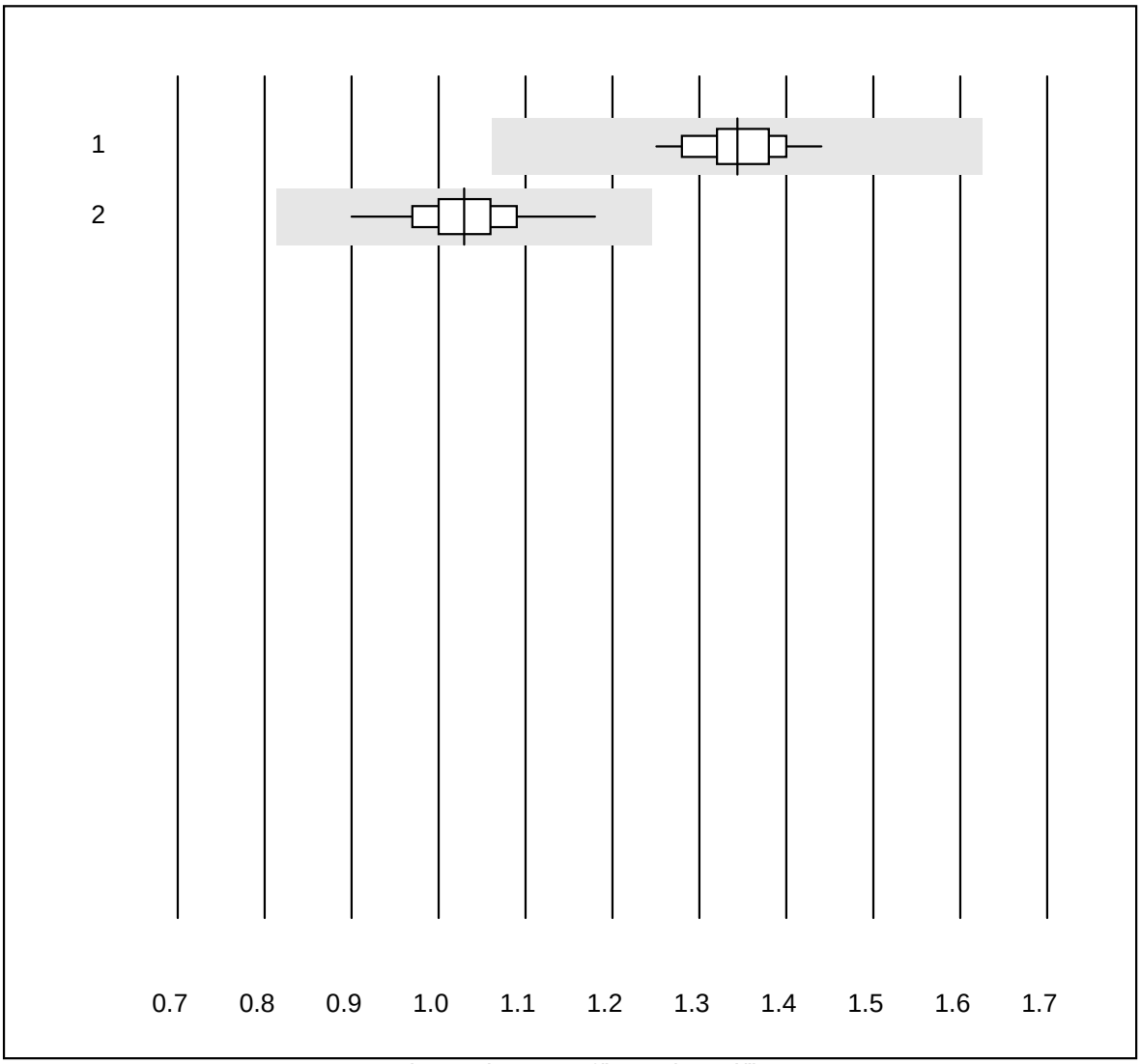


Deviatione QUALAB : 10 %

Colesterolo Af/b101 (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas b101	33	100.0	0.0	0.0	4.8	1.7	e
2	Afinion	244	98.4	1.6	0.0	5.1	3.8	e

Colesterolo HDL Af/b101

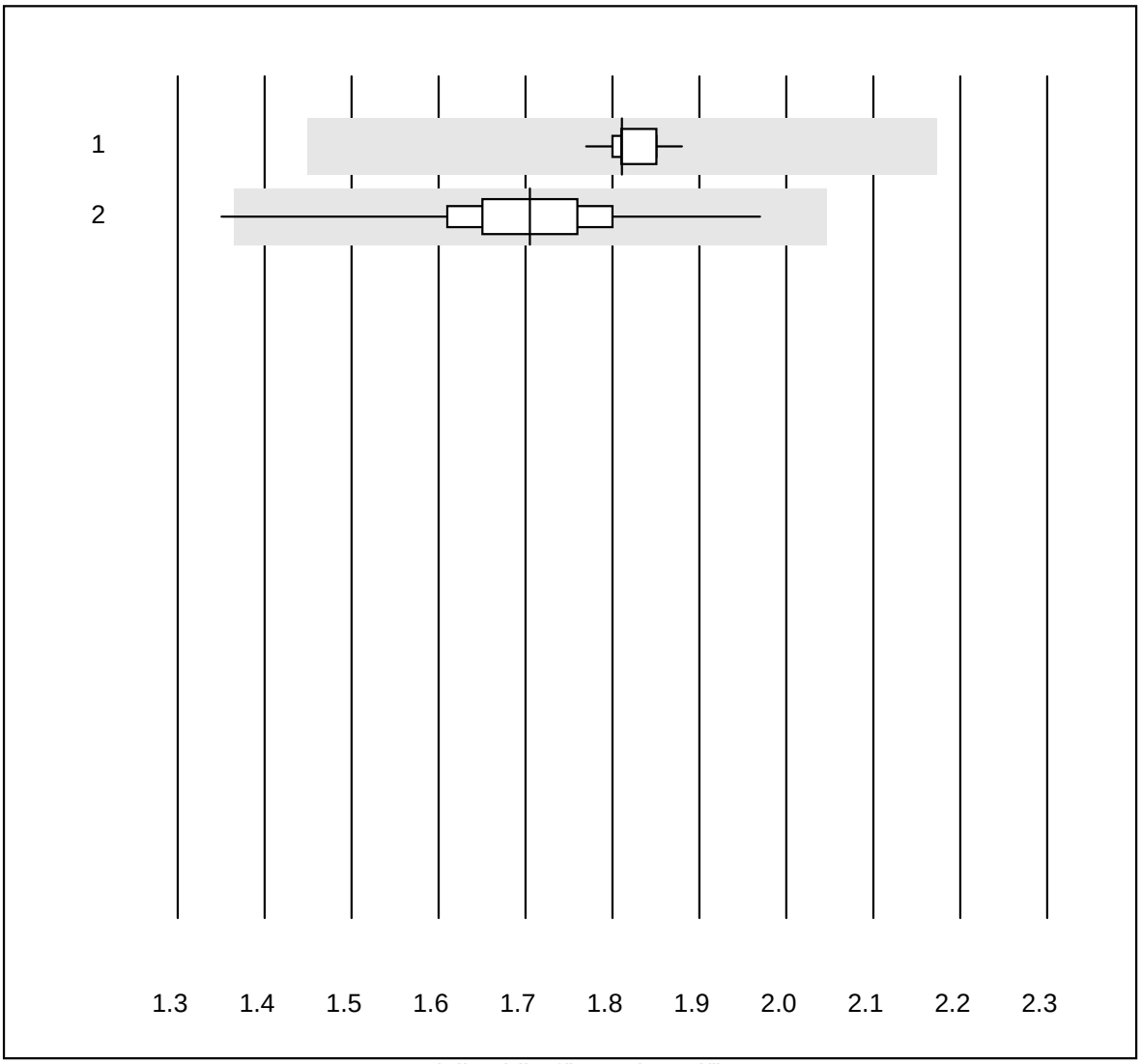


Deviatione QUALAB : 21 %

Colesterolo HDL Af/b101 (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas b101	33	100.0	0.0	0.0	1.3	3.5	e
2	Afinion	244	94.7	0.0	5.3	1.0	4.4	e

Trigliceridi Af/b101

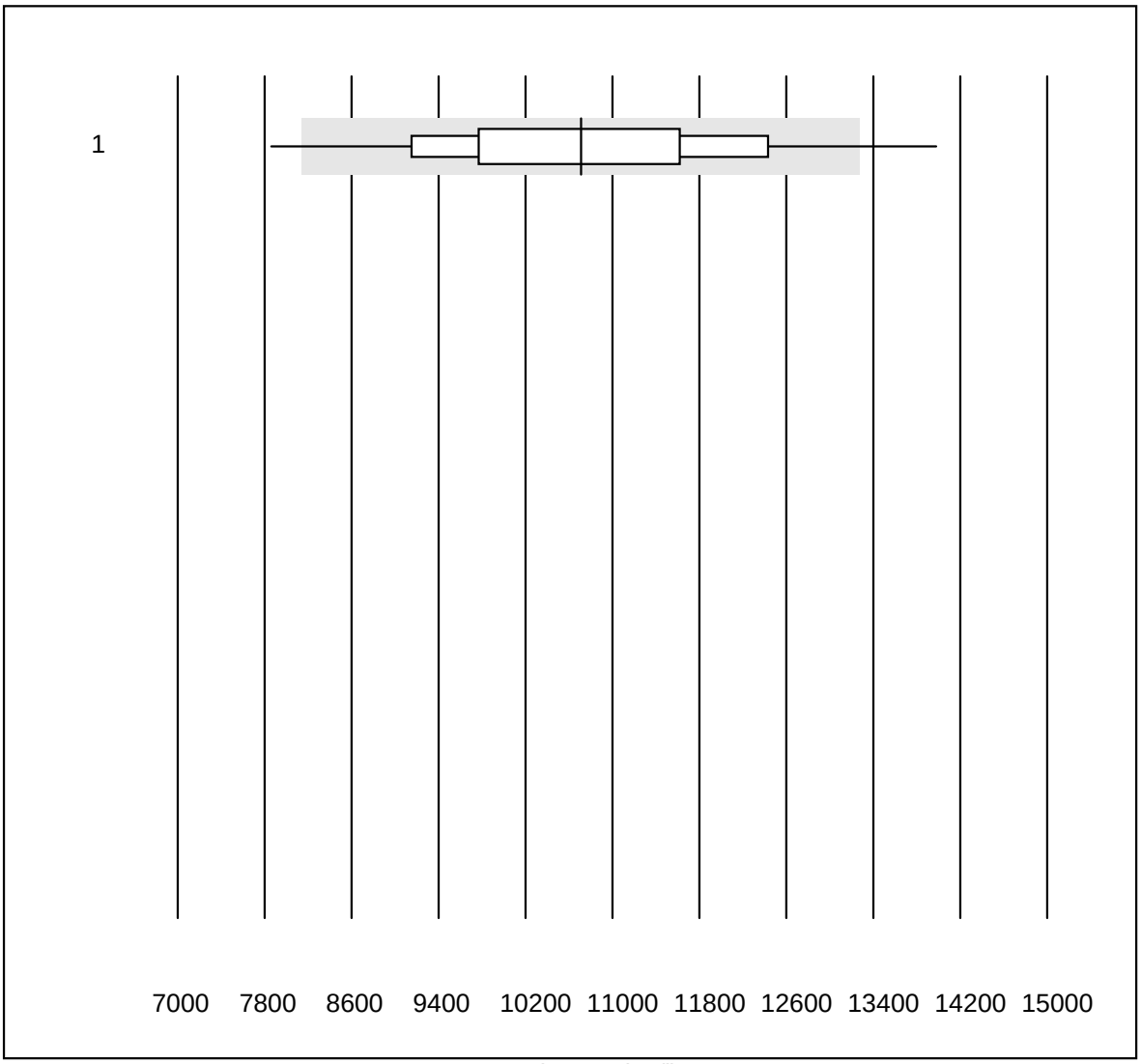


Deviatione QUALAB : 20 %

Trigliceridi Af/b101 (mmol/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Cobas b101	33	100.0	0.0	0.0	1.81	1.4	e
2	Afinion	244	99.2	0.4	0.4	1.71	4.7	e

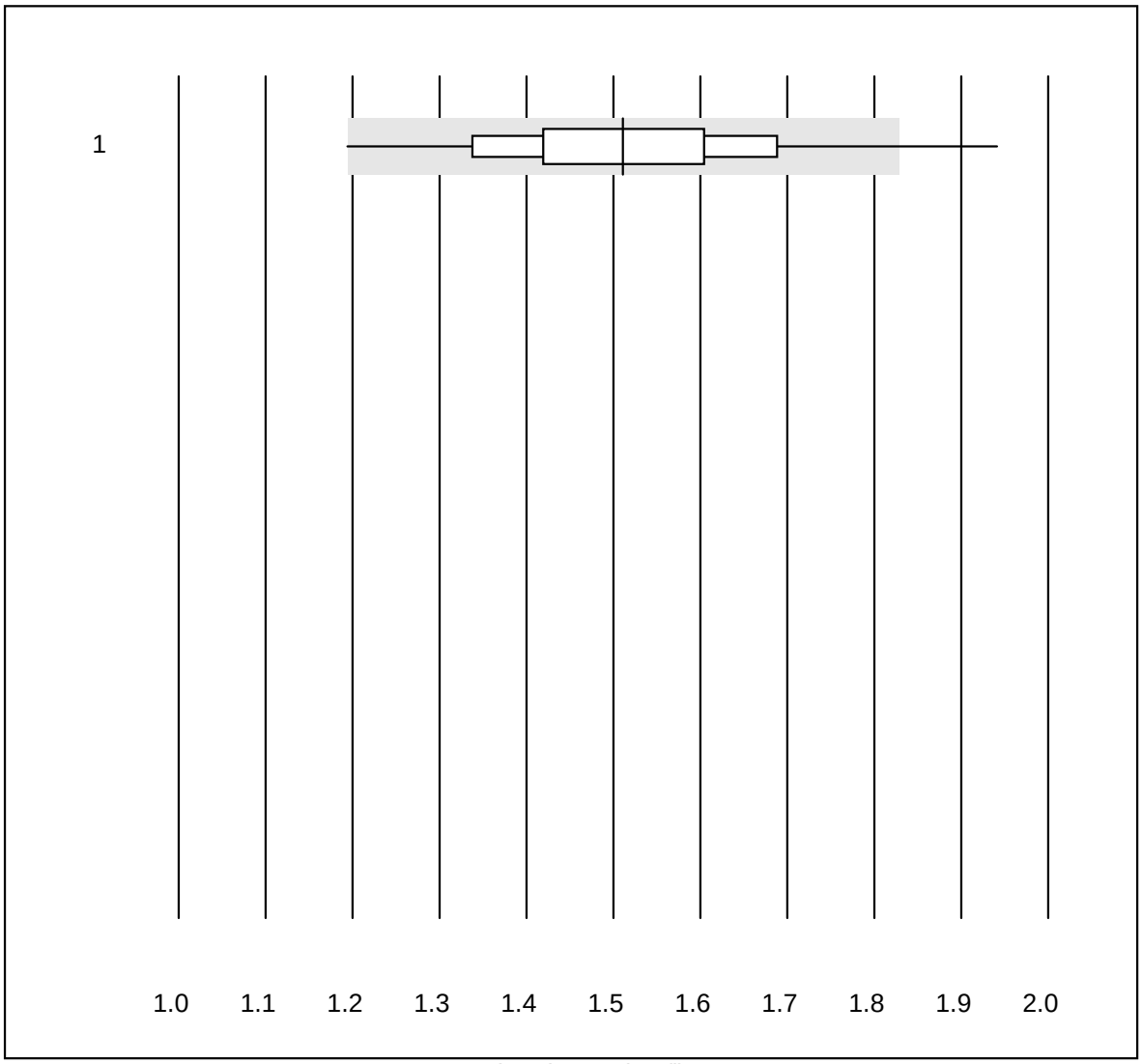
Troponina I S



Deviatione QUALAB : 24 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Samsung LABGEO IB10	72	86.2	6.9	6.9	10710.45	12.2	e

D Dimeri qn S

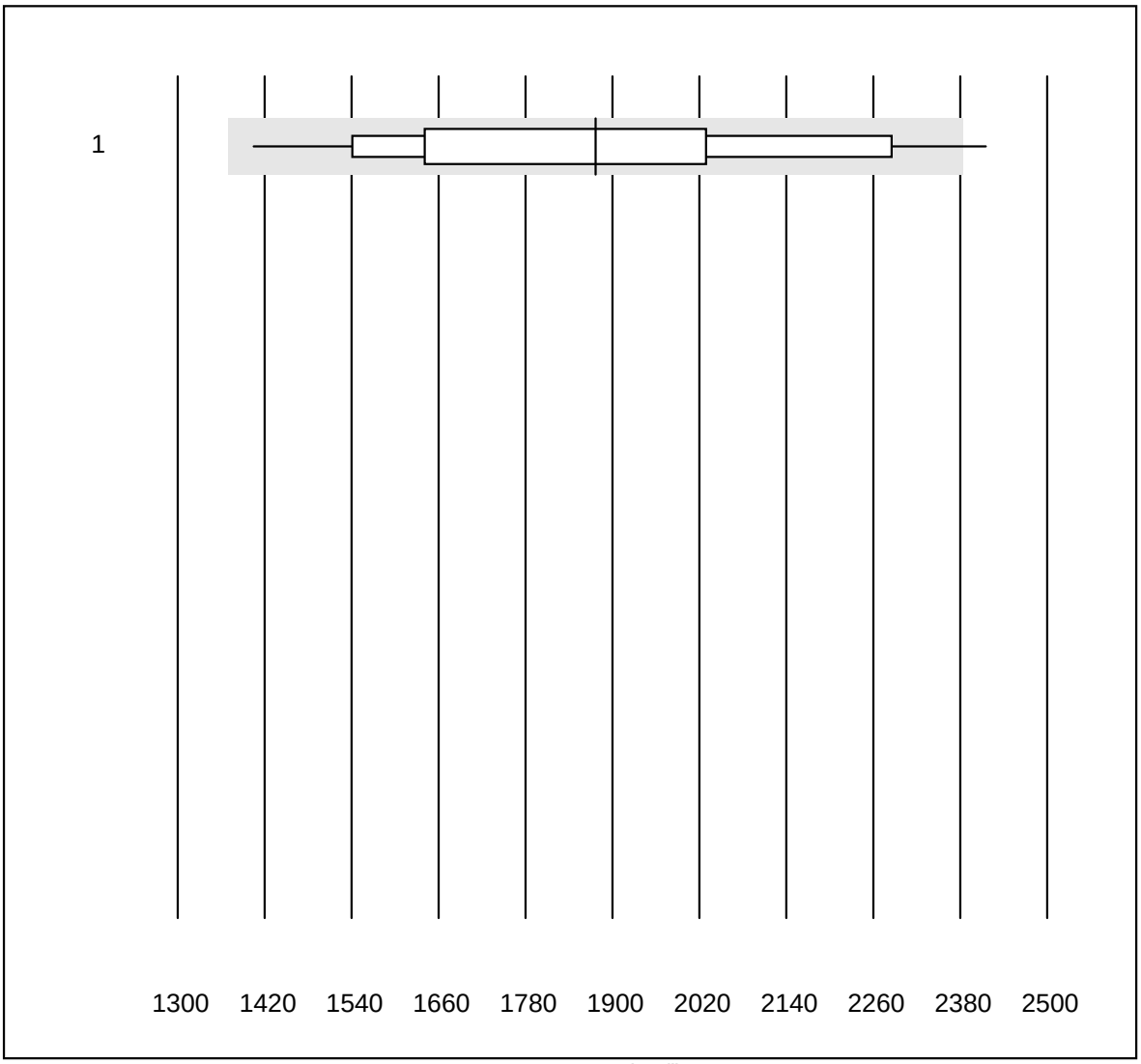


Deviatione QUALAB : 21 %

D Dimeri qn S (mg/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Samsung LABGEO IB10	88	96.6	2.3	1.1	1.51	9.4	e

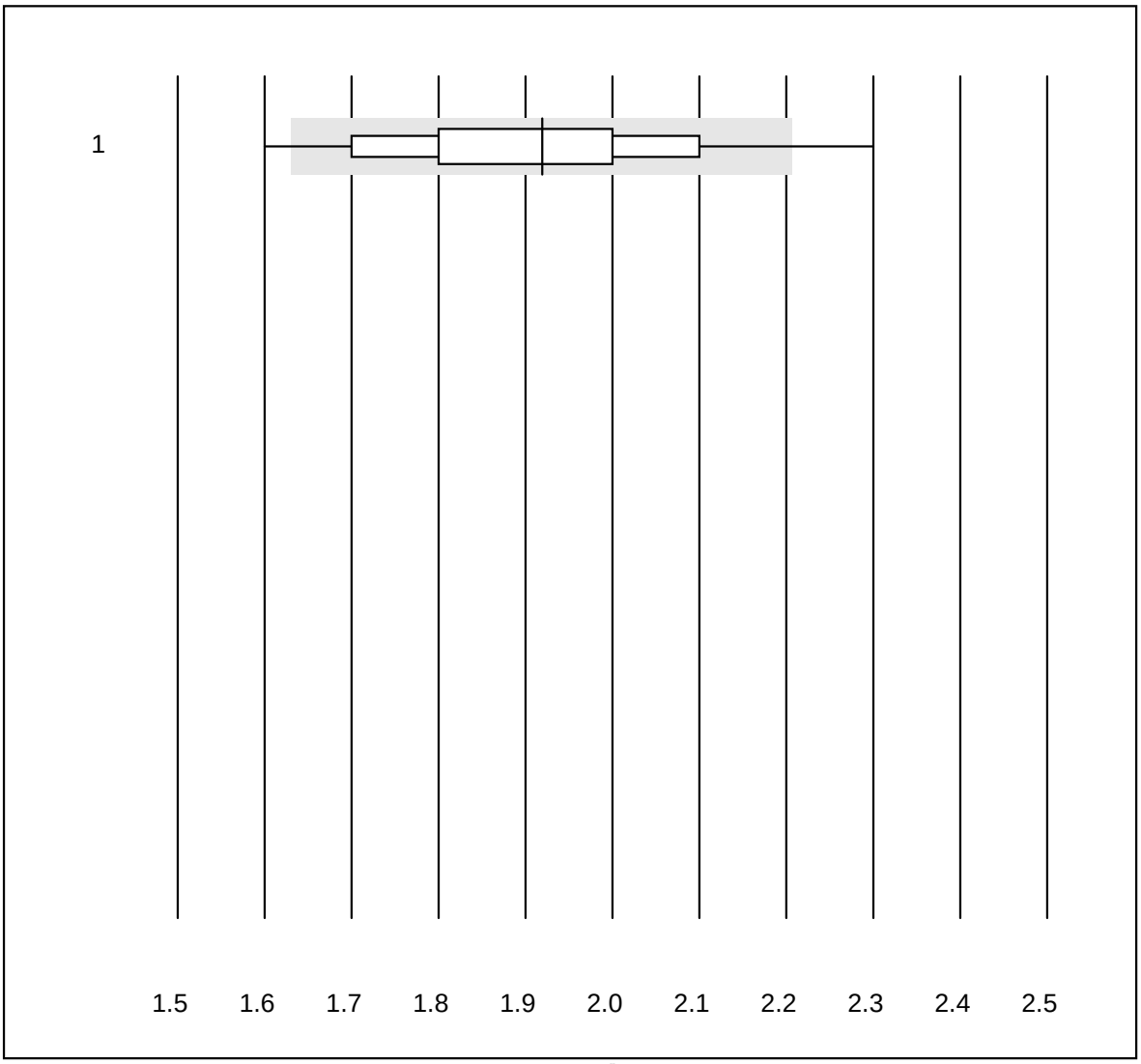
NT-pro BNP S



Deviatione QUALAB : 27 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Samsung LABGEO IB10	57	94.7	1.8	3.5	1877.1	14.1	e

INR MI

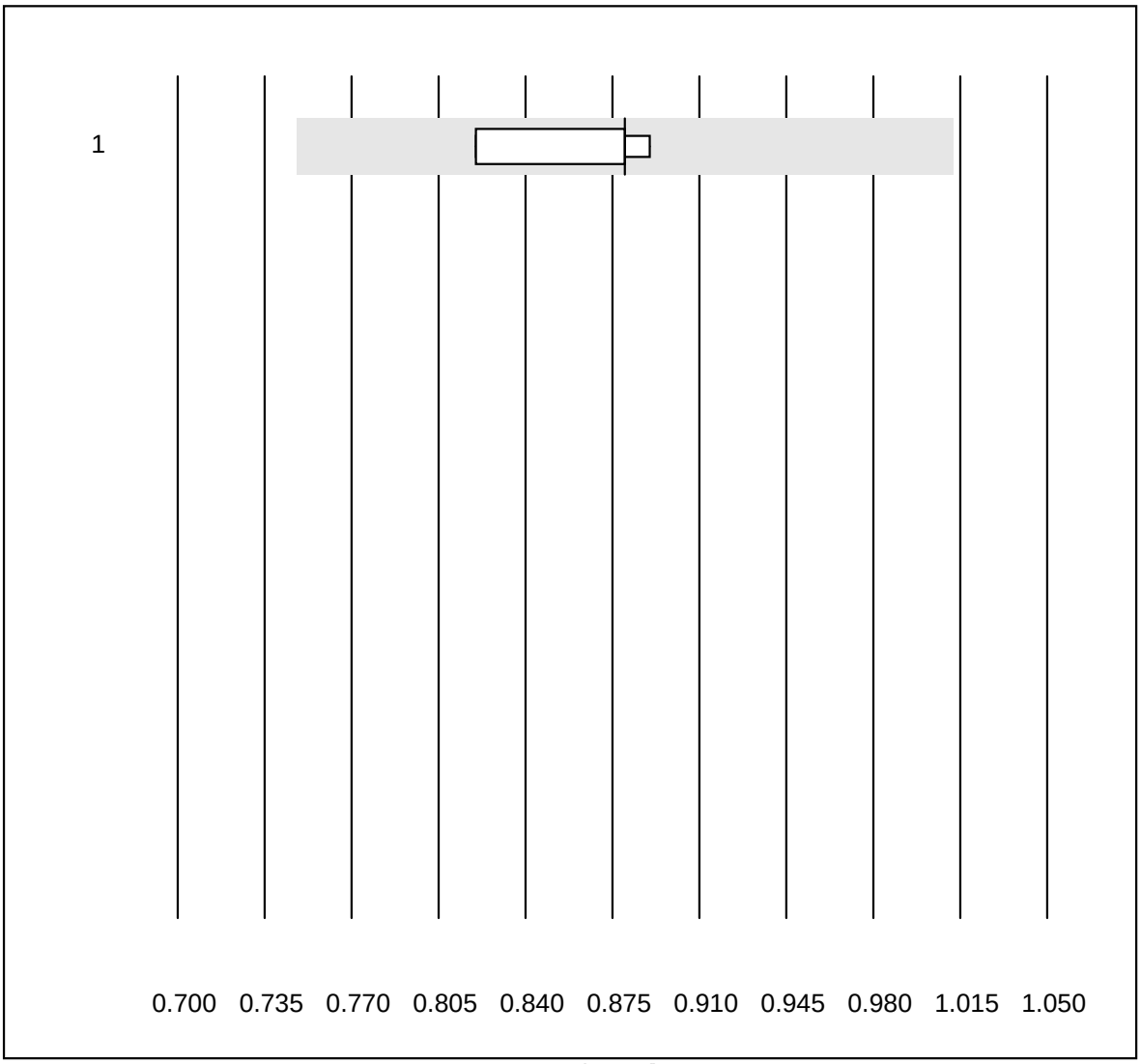


Deviatione QUALAB : 15 %

INR MI ()

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	microINR	64	84.4	7.8	7.8	1.9	8.0	e

INR Eurolyser

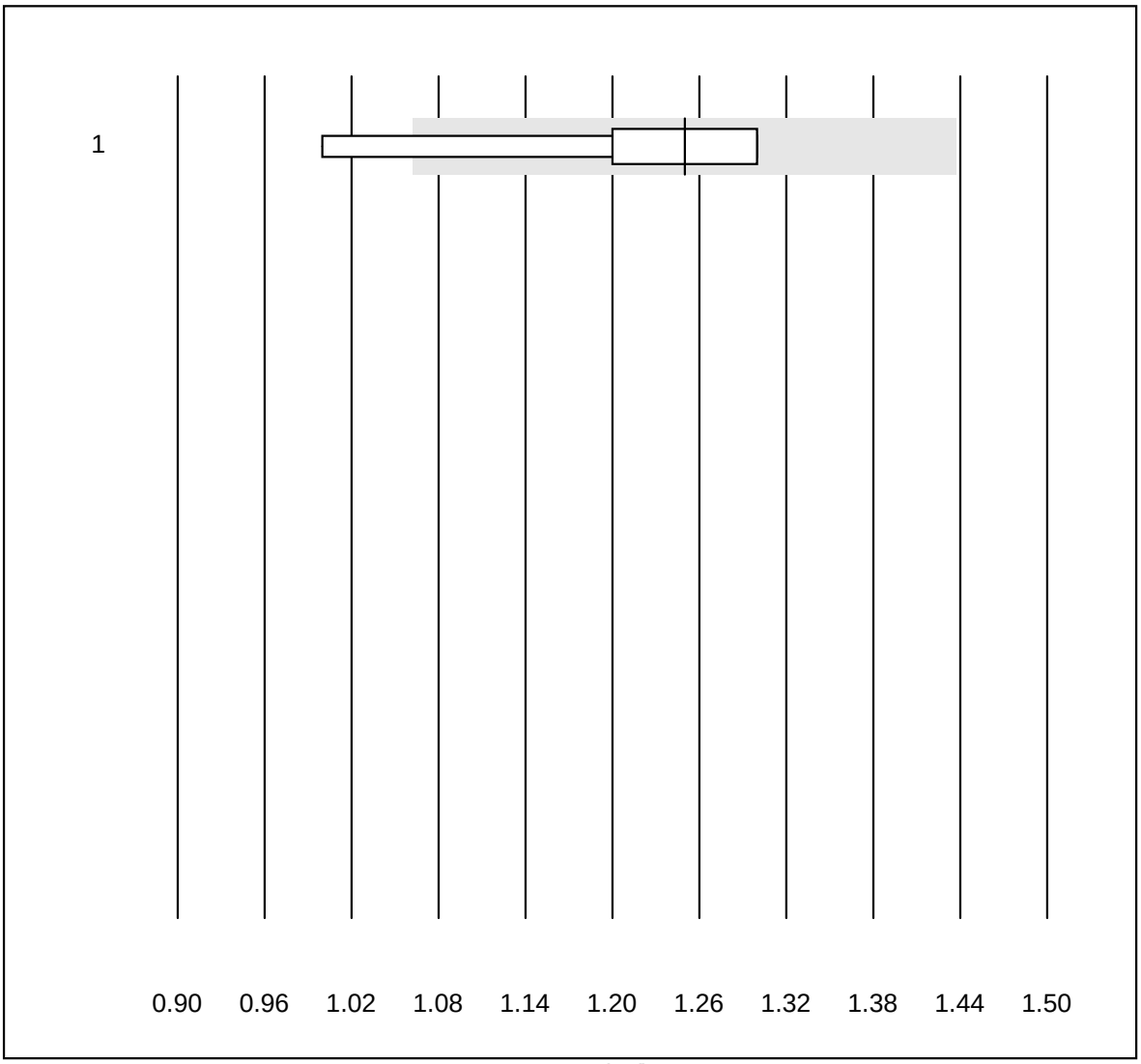


Deviatione QUALAB : 15 %

INR Eurolyser ()

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Eurolyser	5	80.0	0.0	20.0	0.9	3.6	e

INR Xprecia

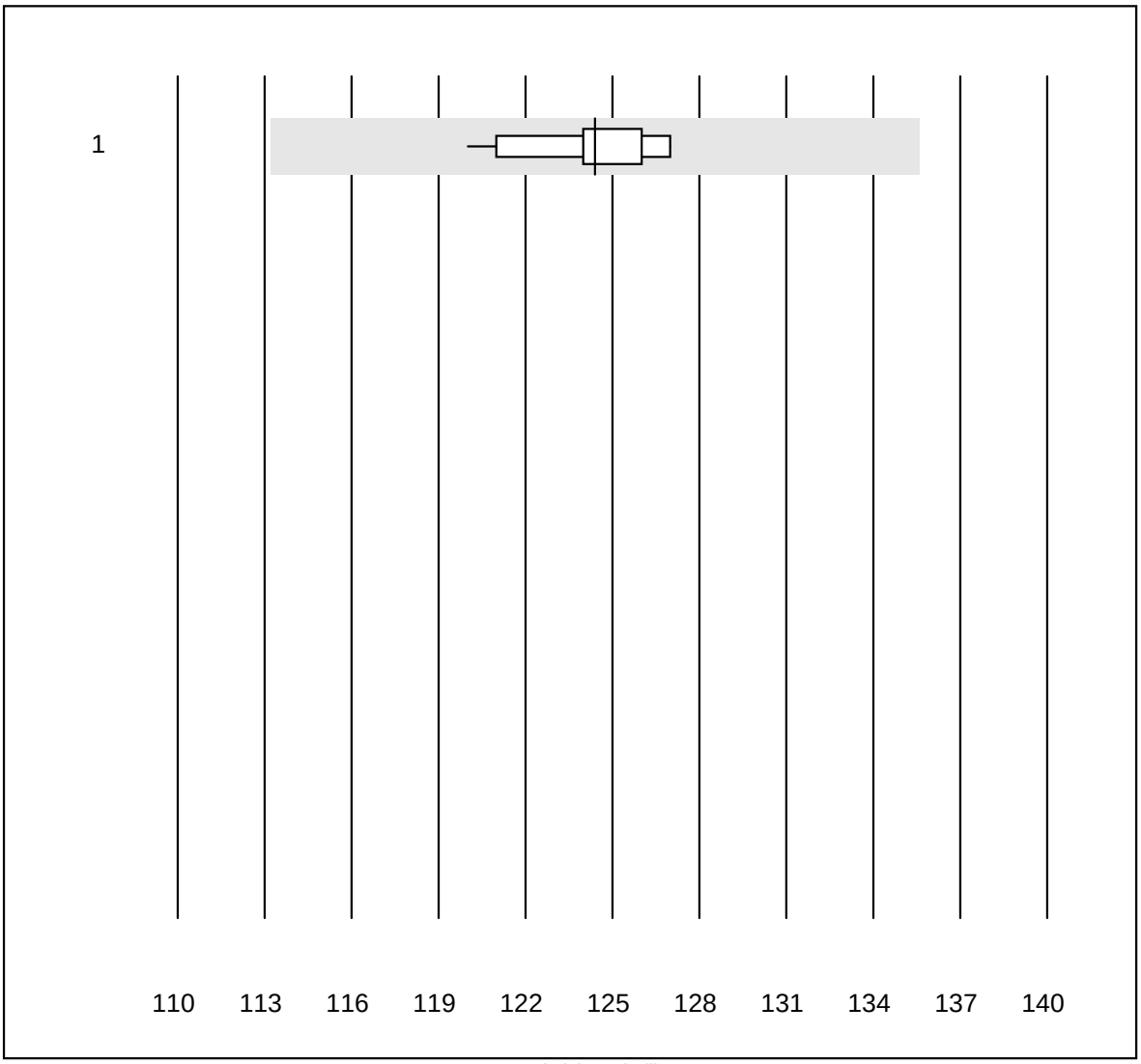


Deviatione QUALAB : 15 %

INR Xprecia ()

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Xprecia	6	83.3	16.7	0.0	1.3	9.6	e*

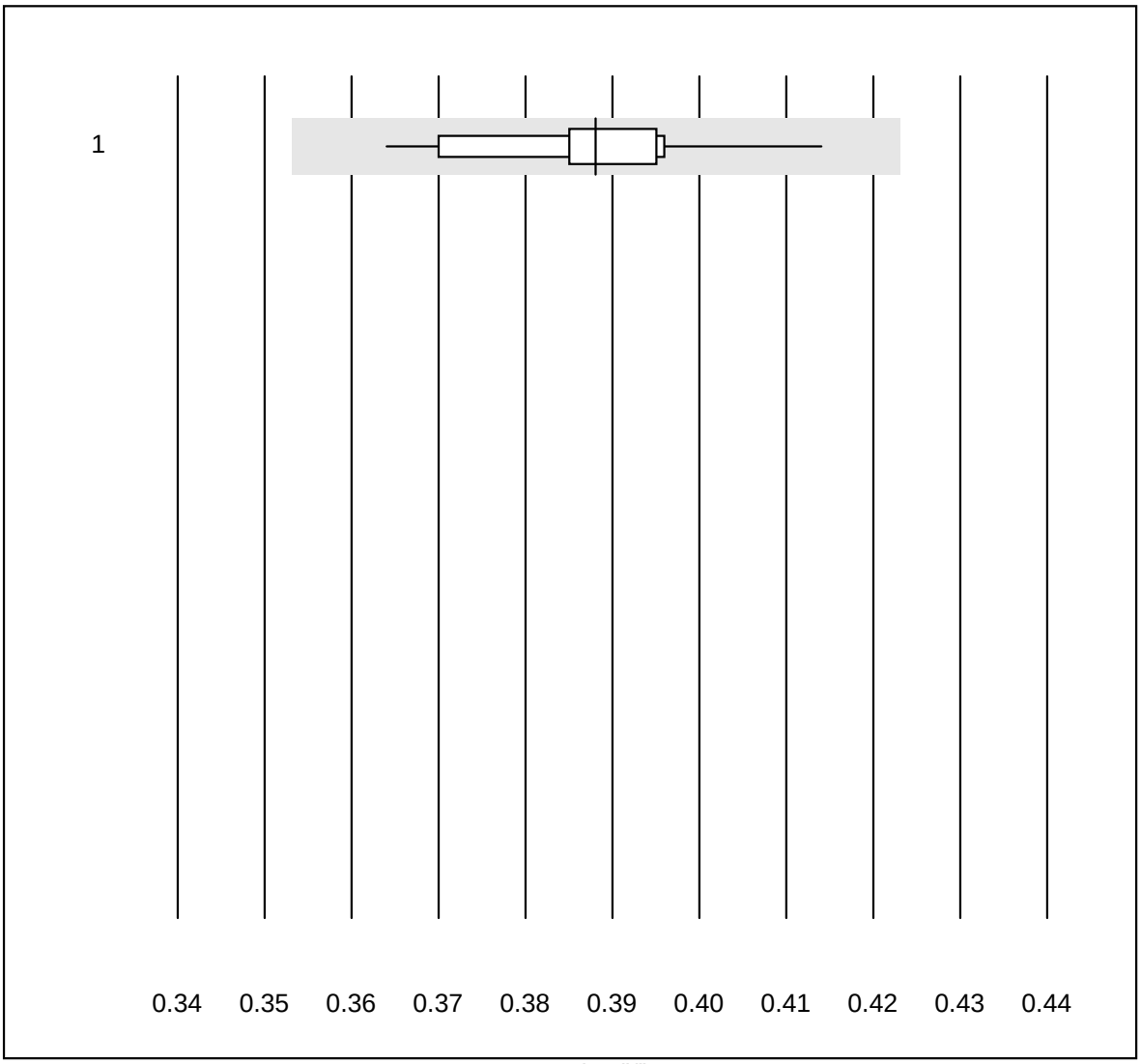
Emoglobina



Deviatione QUALAB : 9 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Sysmex	13	100.0	0.0	0.0	124.4	1.7	e

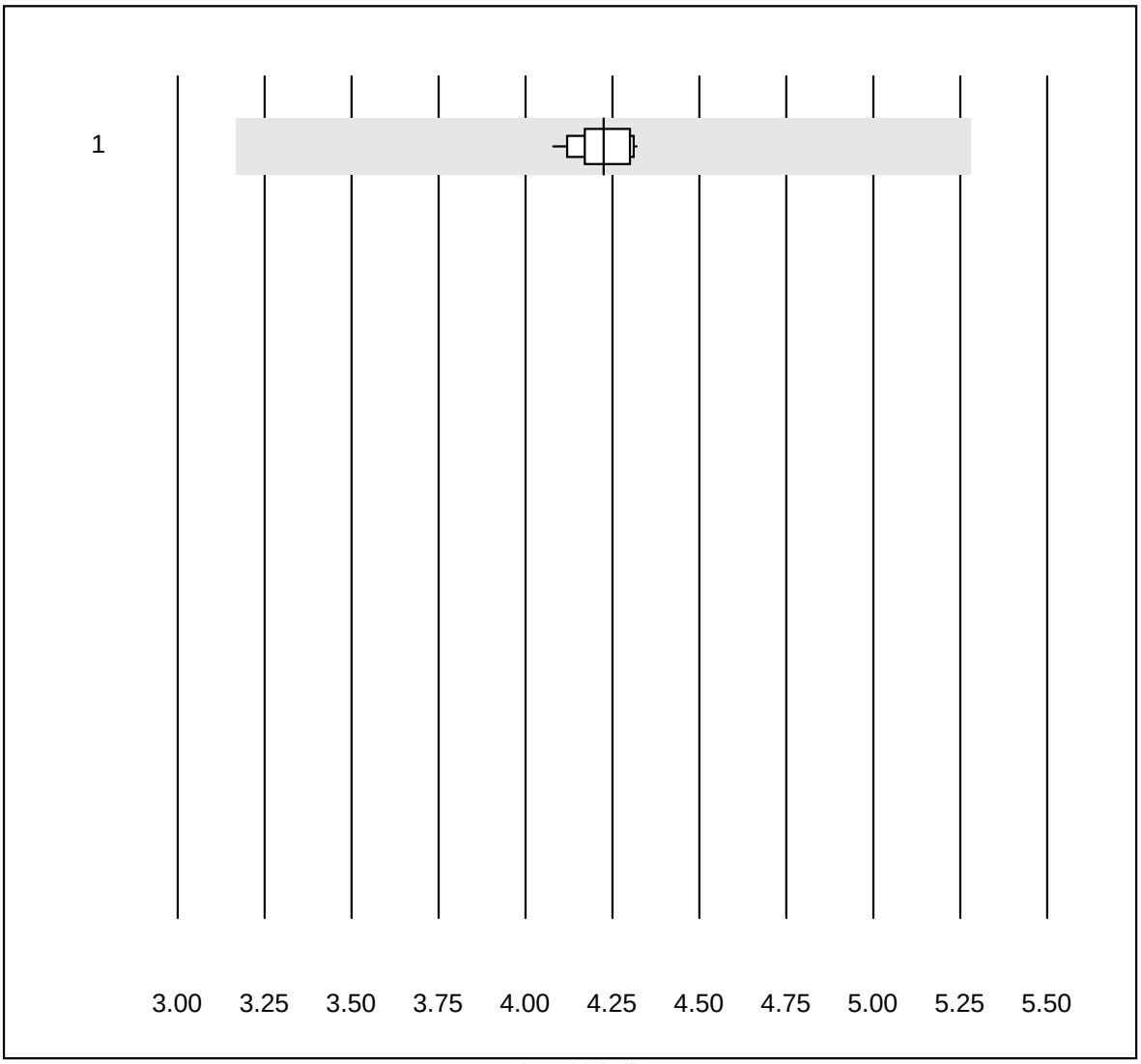
Ematocrito



Deviatione QUALAB : 9 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Sysmex	14	100.0	0.0	0.0	0.39	3.1	e

Eritrociti

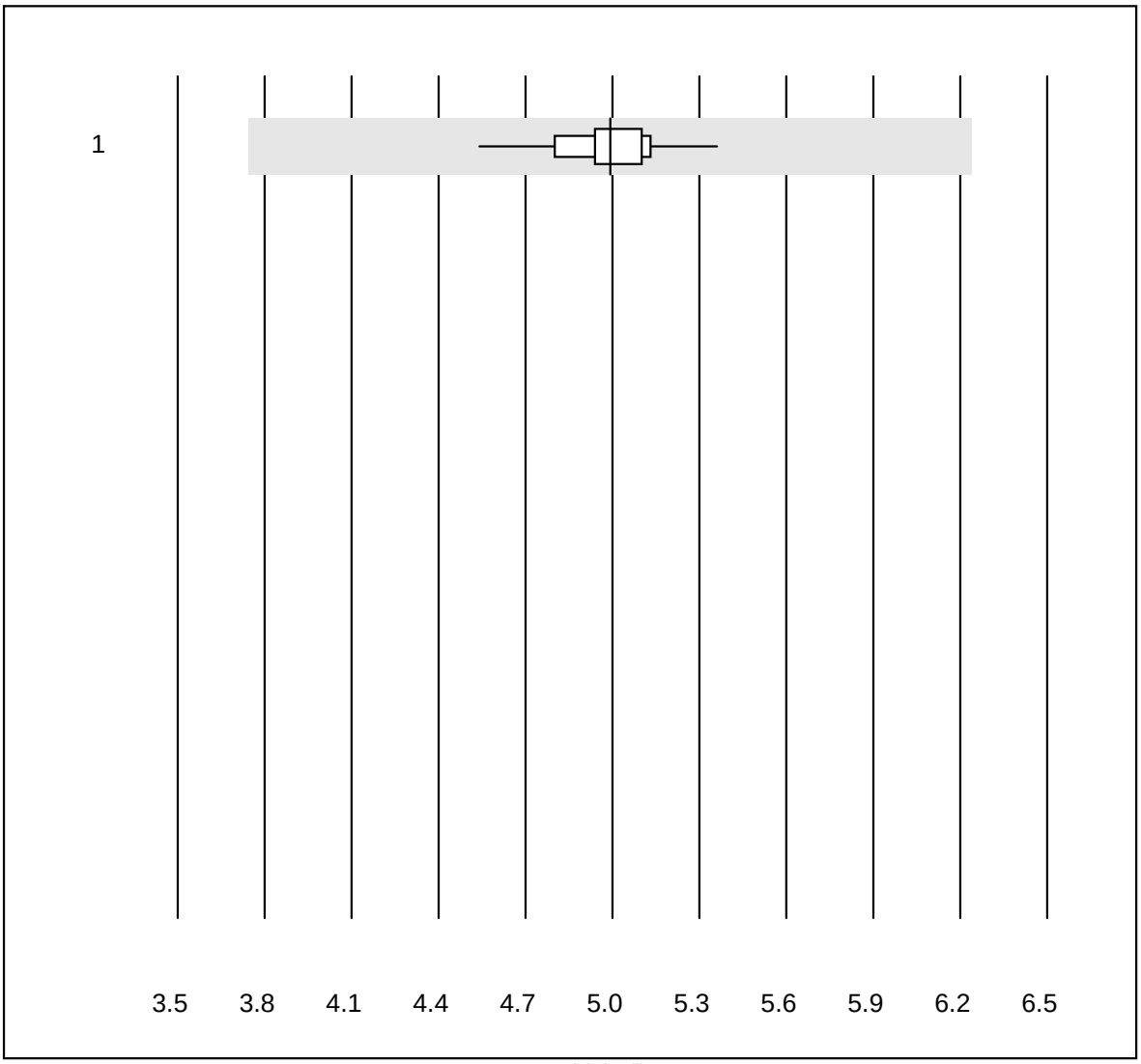


Deviatione QUALAB : 25 %

Eritrociti (T/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Sysmex	13	100.0	0.0	0.0	4.22	1.8	e

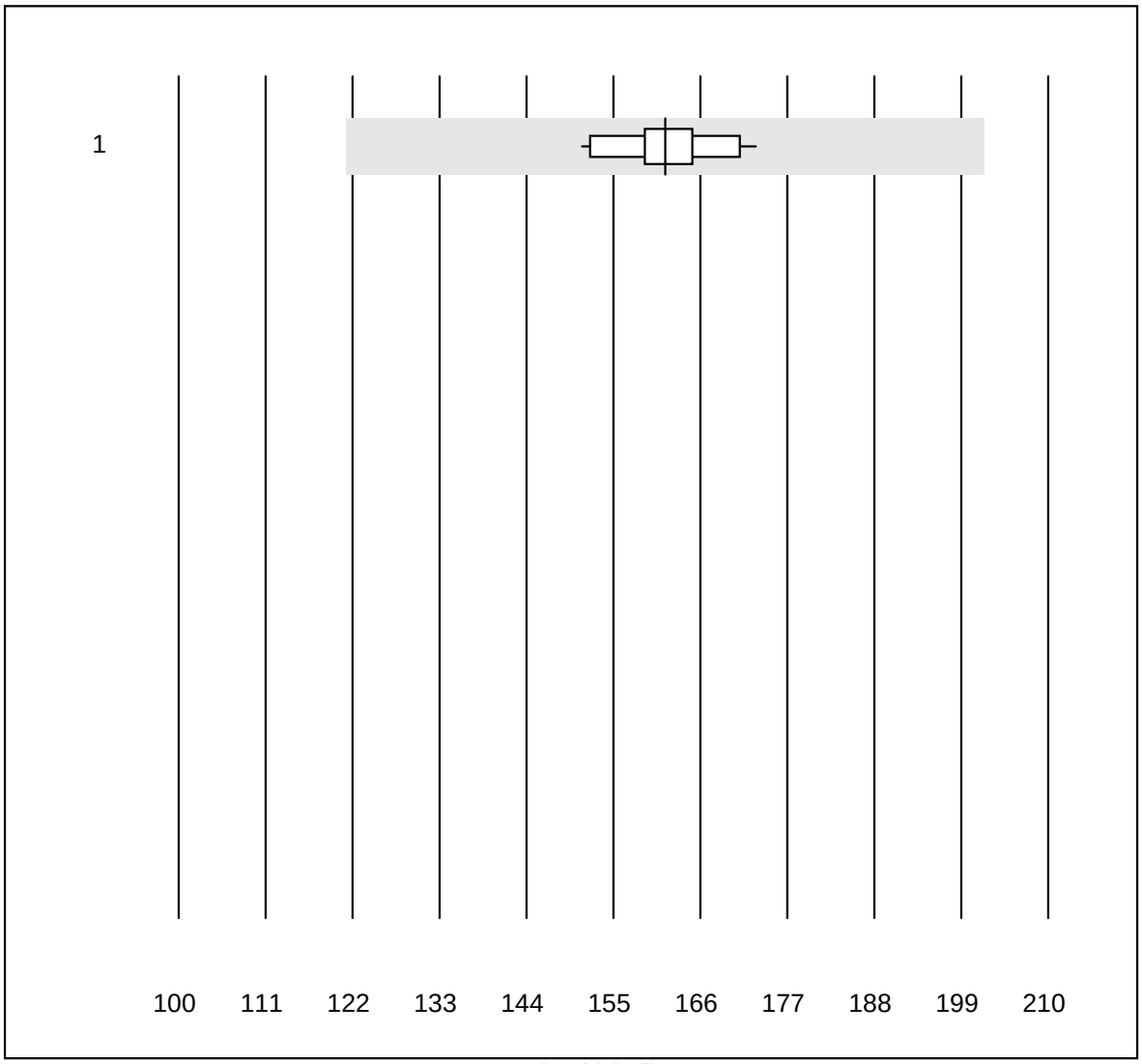
Leucociti



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Sysmex	13	100.0	0.0	0.0	4.99	3.9	e

Trombociti

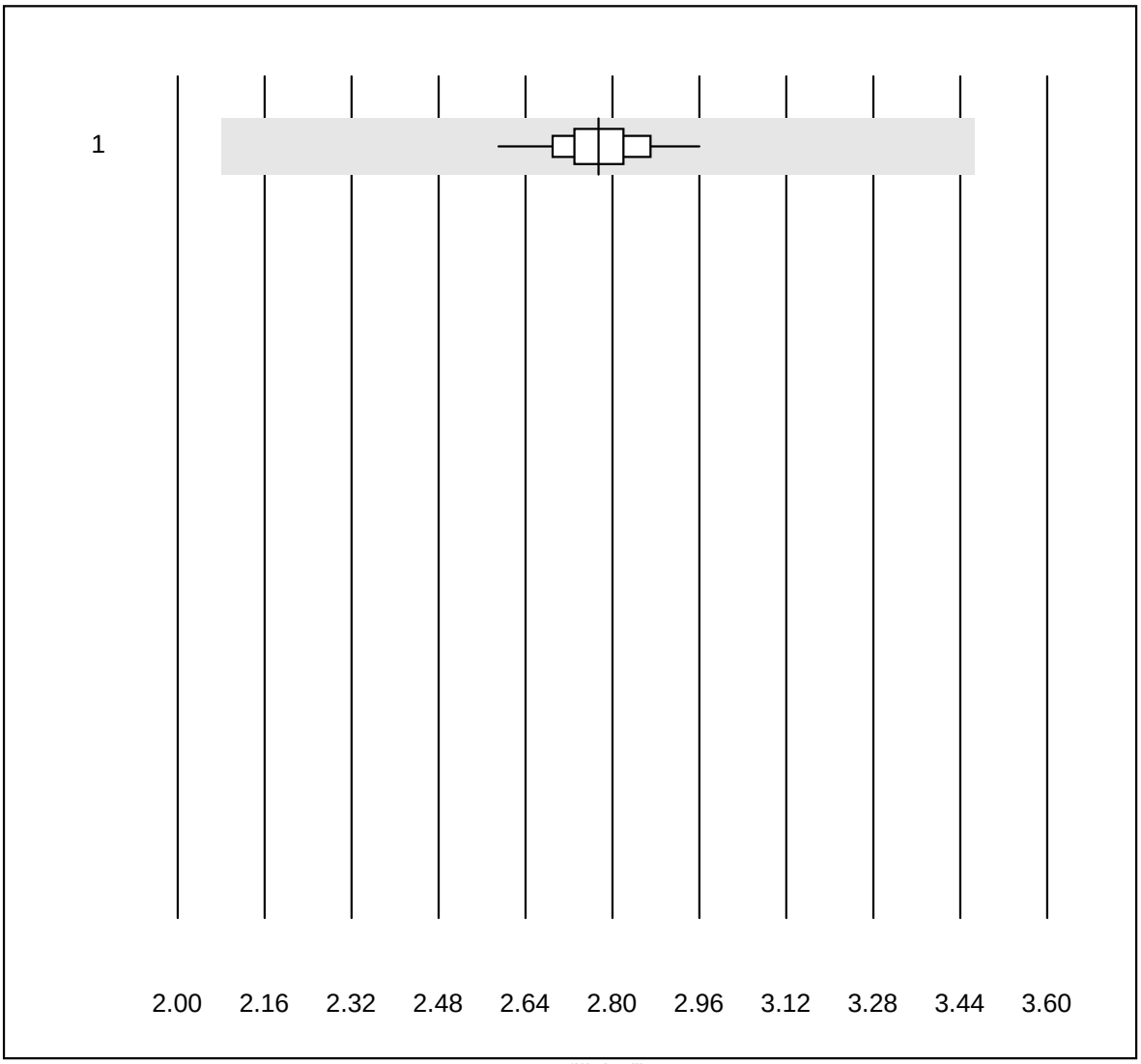


Deviatione QUALAB : 25 %

Trombociti (G/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Sysmex	13	100.0	0.0	0.0	161.5	4.2	e

Neutrofili

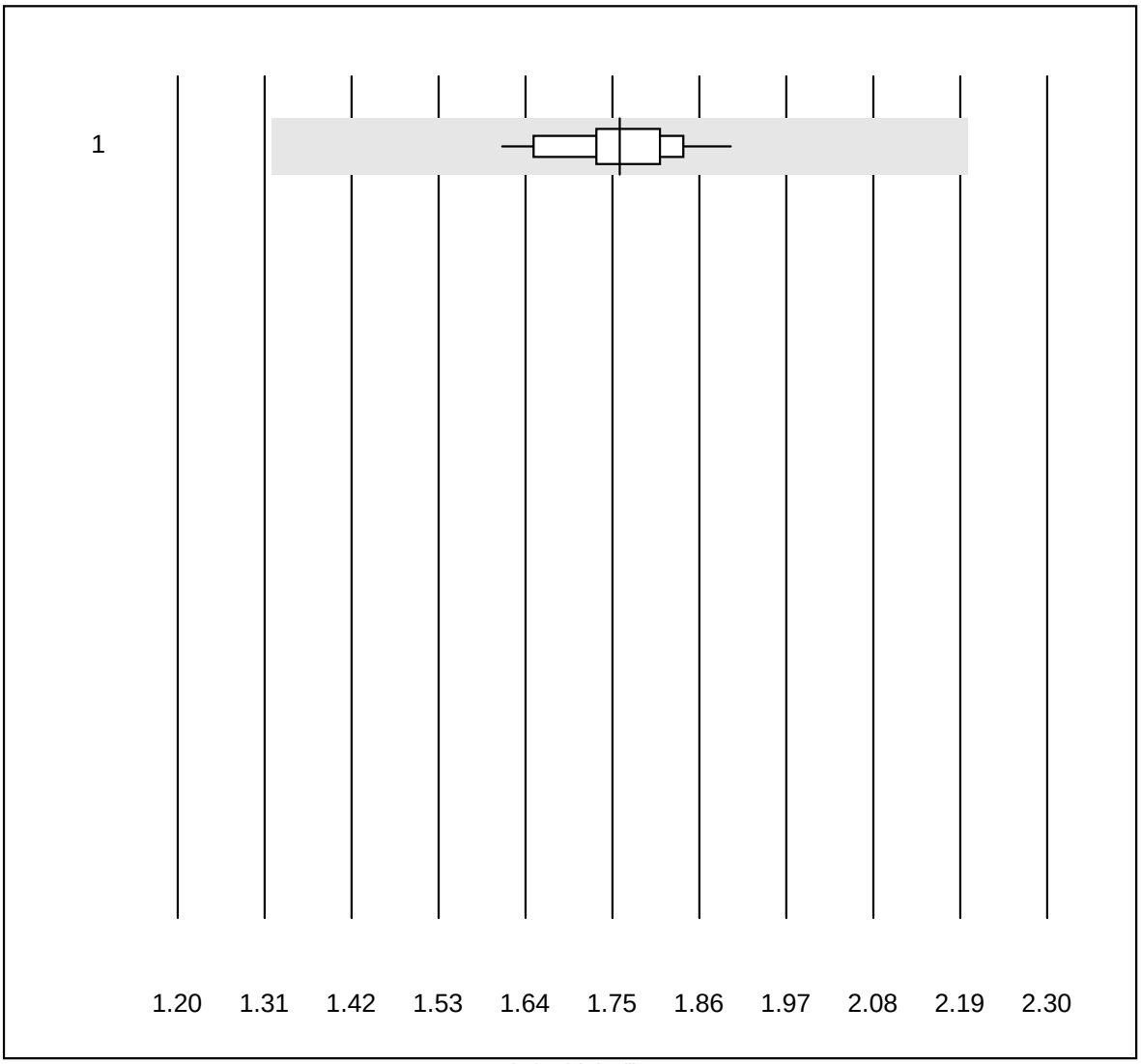


Deviatione QUALAB : 25 %

Neutrofili (G/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Sysmex	13	100.0	0.0	0.0	2.77	3.4	e

Linfociti

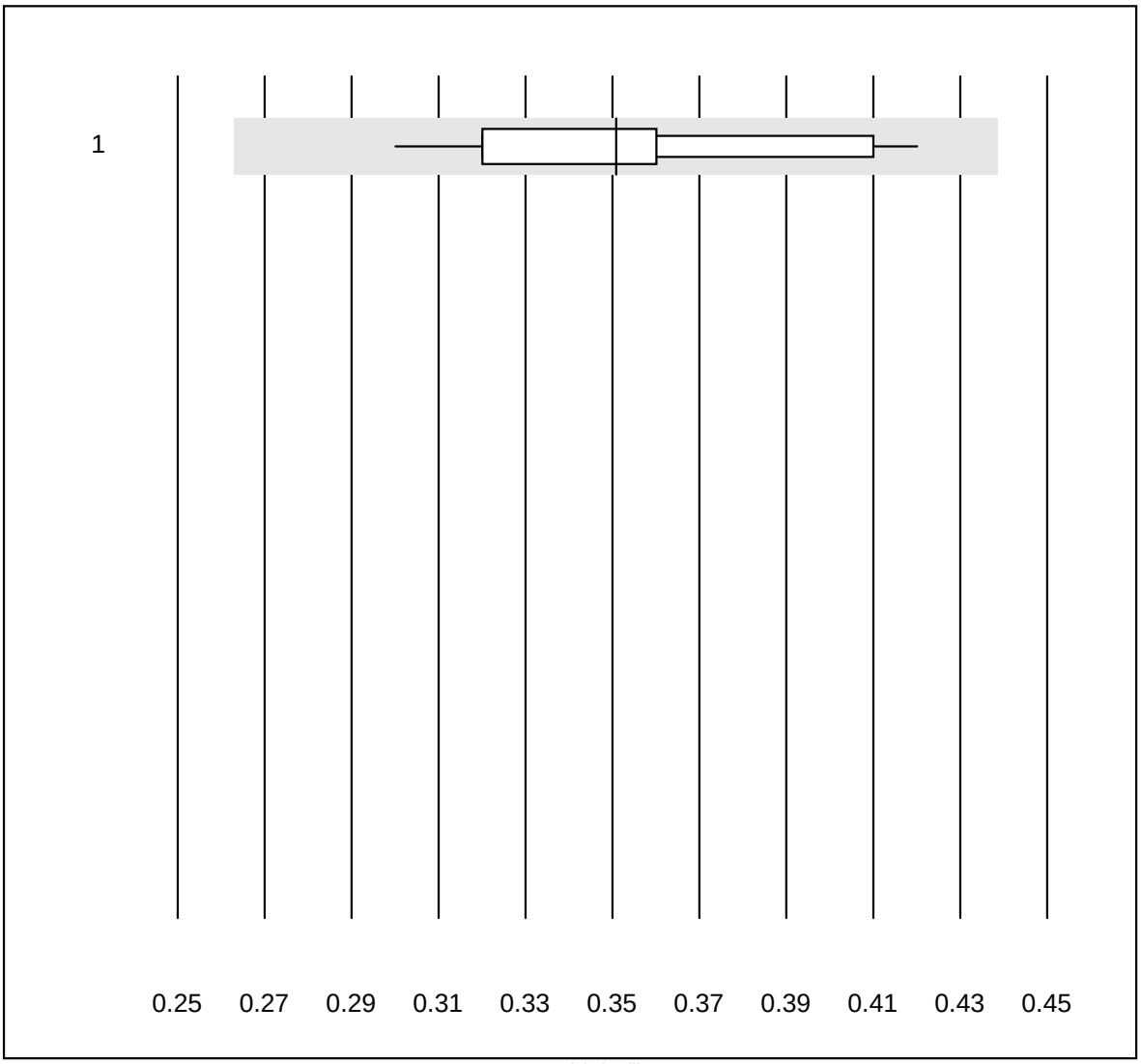


Deviatione QUALAB : 25 %

Linfociti (G/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Sysmex	13	100.0	0.0	0.0	1.76	4.5	e

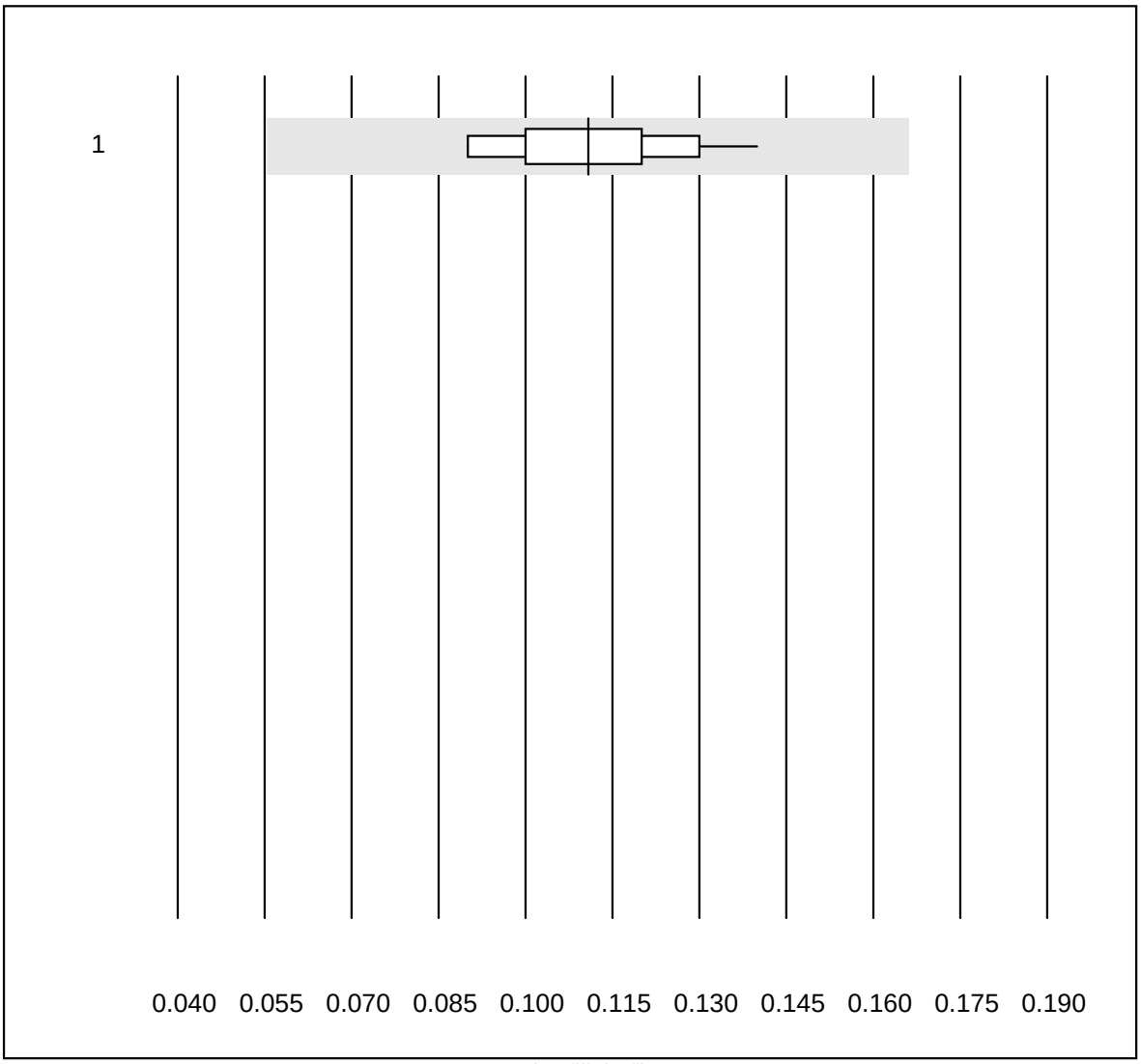
Monociti



Deviatione QUALAB : 25 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Sysmex	13	92.3	0.0	7.7	0.35	10.4	e

Eosinofili

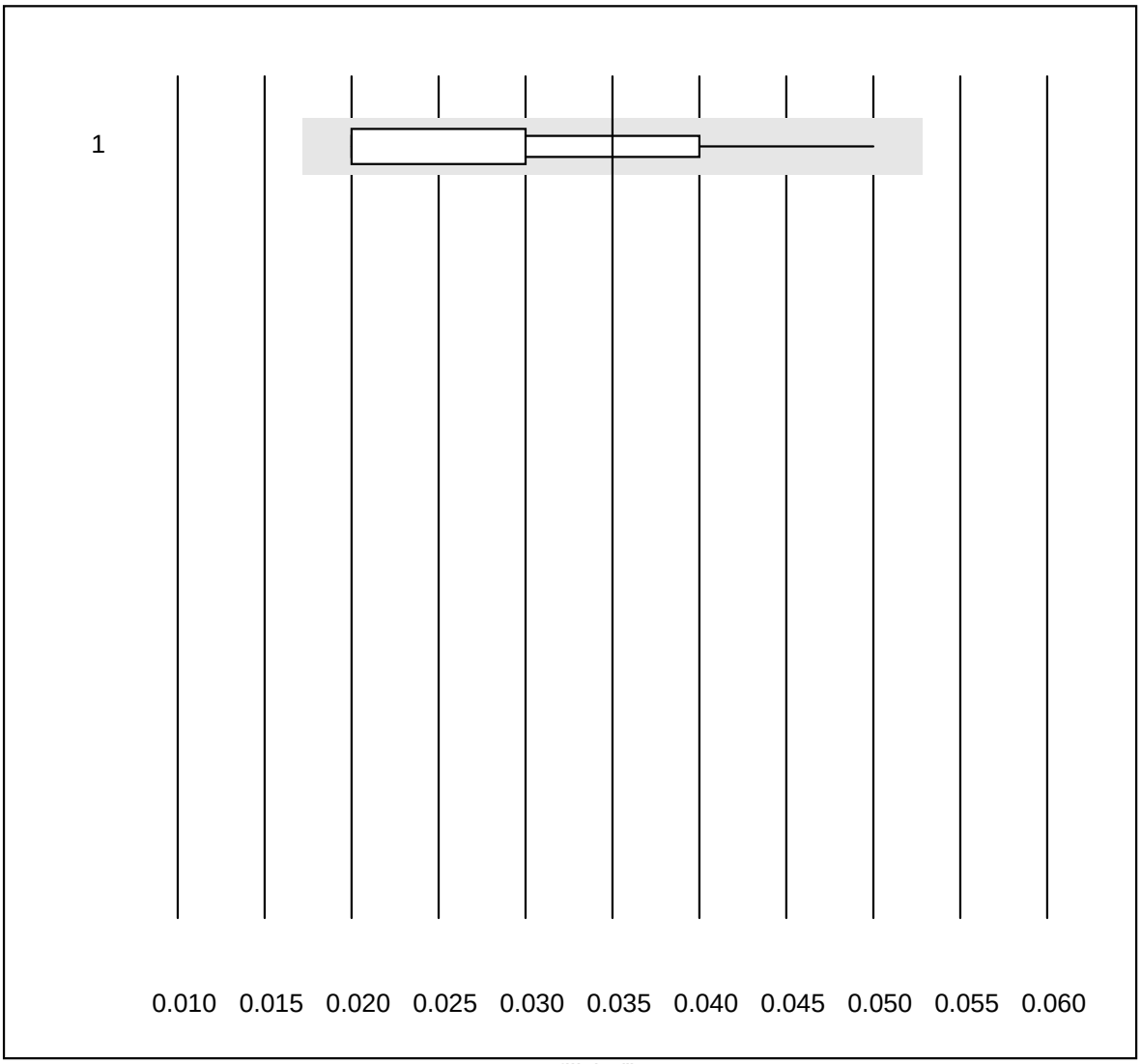


Deviatione QUALAB : 50 %

Eosinofili (G/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Sysmex	13	100.0	0.0	0.0	0.11	14.5	e

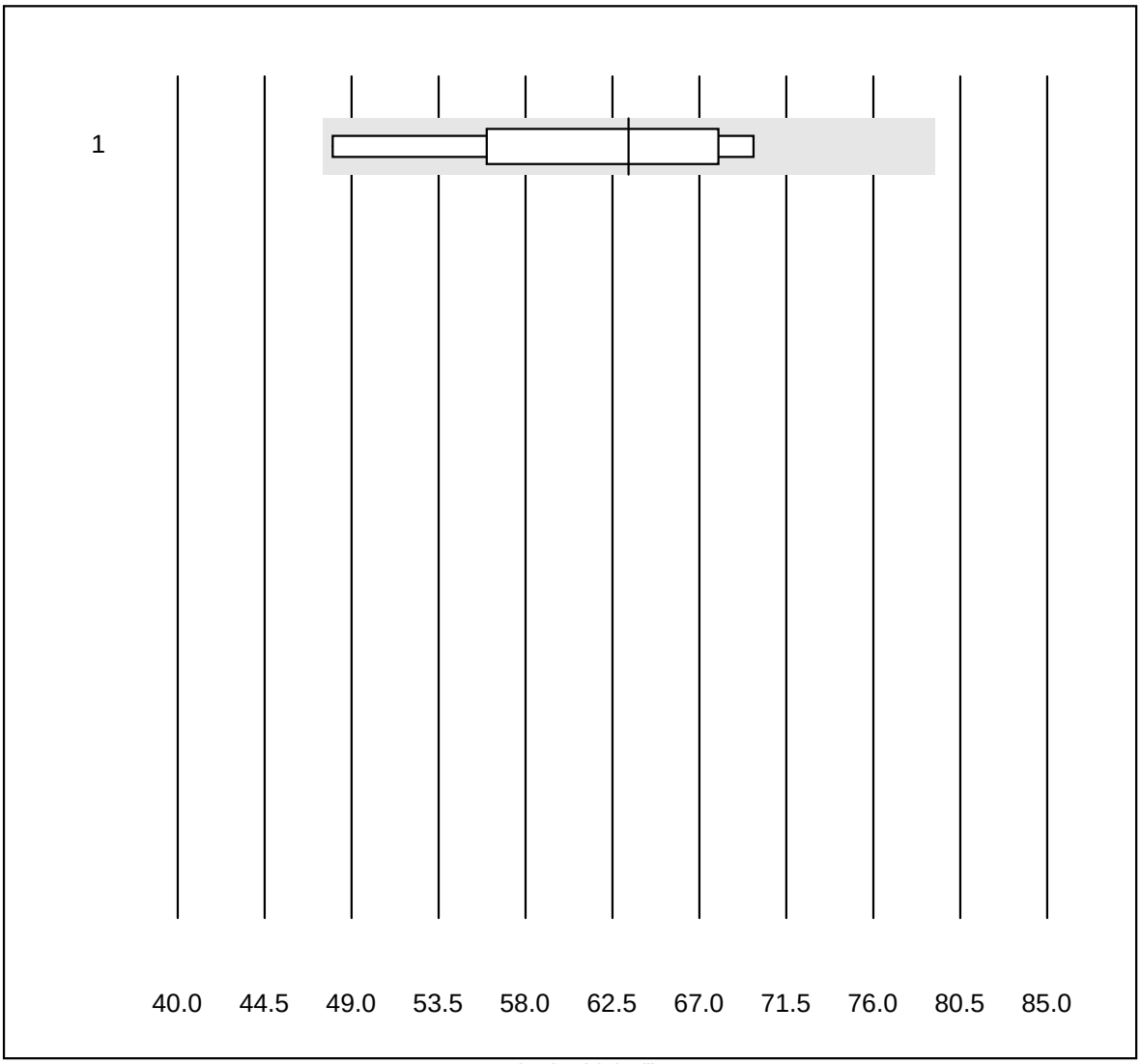
Basofili



Deviatione QUALAB : 80 %

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Sysmex	13	100.0	0.0	0.0	0.04	34.7	a

Reticolociti



Deviatione QUALAB : 25 %

Reticolociti (G/l)

No.	Metodo	Totale	% conforme	% insuff.	% outlier	Valore ideale	CV%	Typ
1	Sysmex	9	100.0	0.0	0.0	63.3	12.6	e*