Survey: **IG2/26**
month: **April 2026**
Participant-No: **9900954**
valid from: **25.04.26**

DAkkS Deutsche Akkreditierungsstelle D-EP-15117-01-00

Nemocnice Pardubickeho Kraje, a.s.
Orlickoustecka nemocnice -1-
Oddeleni klinicke bioch. a diagnostiky
RnDr. Karel Kohoutek
Cs. armady 1076
56218 USTI NAD ORLICI 1
CZECH REPUBLIC

Bonn, 30. April 2026

Certificate

We confirm that you have participated in the survey for immunology/proteins in serum.
This certificate is according to the current version of the RiliBÄK valid until the end of October 2026.
You have met the requirements of the survey for the following analytes:

Albumin	(7)	Transferrin	(4)	IgA	(4)
IgG	(4)	IgM	(4)	Haptoglobin	(4)
C3-complement	(4)	C4-complement	(4)	CRP	(4)
IgE	(5)	sol.Transfe.Receptor	(2)		

Prof. Dr. Dr. K. P. Kohse
EQA scheme director

Dr. Anja Kessler
Head of Reference Institute

Dr. Marika Enders
EQAS-Board

The number in parentheses characterizes the analytical method used.
The assignment of the number to the respective method and/or the respective instrument is to be taken from the total evaluation.

This certificate is valid in conjunction with the final report dated 30.04.26.
This is available for download in the RfB-Online system.

Survey: **IG2/26**
month: **April 2026**
Participant-No: **9900954**

Listing and Evaluation of all your results

Explanations

Certificate

A certificate is issued (given) for an analyte only if the basis for an evaluation of the accuracy is given by the guidelines of the German Medical Association and/or if an evaluation is possible in analogy to these guidelines (see comments on the evaluation) and, if both results for an analyte are within the given acceptance limits. (marked as '+' below C)

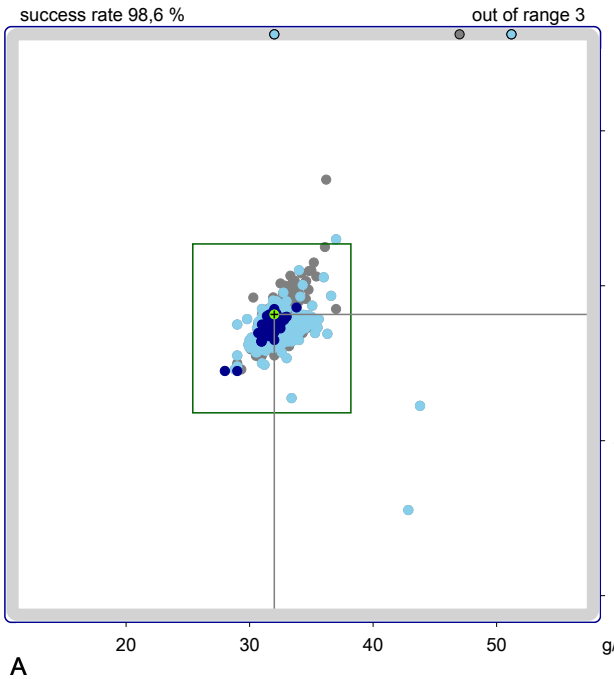
Participation certificate

In the participation certificate all analytes which are included in the list on this page are listed.
If all analytes are listed on the certificate no participation certificate is printed.

Legend:	C=Certification, M=No of method, R=your result, D=difference (R-T) Dmax= maximum allowable amount of difference in measurement, partly given by annax 1 of the guideline of the BÄK. T = target value, either reference method value or assigned value, LL | UL = lower resp. upper limit	Certification:	+ = fulfilled (quotient | D/Dmax | <= 1.0) - = not fulfilled (quotient | D/Dmax | > 1.0 ± = certification cancelled because of technical and/or analytical reasons

	C	M		R	D/Dmax	T	LL	UL
Albumin	+	7	A	32.0	0.03	31.8	25.4	38.2
[g/l]			B	56.3	0.16	54.5	43.6	65.4
Transferrin	+	4	A	1.86	0.27	1.80	1.58	2.02
[g/l]			B	3.76	0.22	3.66	3.22	4.10
IgA	+	4	A	1.55	0.09	1.52	1.21	1.83
[g/l]			B	3.27	-0.03	3.29	2.63	3.95
IgG	+	4	A	9.78	0.03	9.72	7.97	11.5
[g/l]			B	18.3	-0.06	18.5	15.1	21.9
IgM	+	4	A	1.39	-0.08	1.42	1.05	1.79
[g/l]			B	2.81	-0.18	2.95	2.18	3.72
Haptoglobin	+	4	A	1.00	0.00	0.998	0.698	1.30
[g/l]			B	1.61	0.02	1.60	1.12	2.08
C3-complement	+	4	A	1.07	-0.12	1.11	0.777	1.45
[g/l]			B	1.35	-0.05	1.37	0.959	1.79
C4-complement	+	4	A	0.250	0.13	0.240	0.168	0.312
[g/l]			B	0.260	0.13	0.250	0.175	0.325
CRP	+	4	A	50.2	0.02	50.0	40.0	60.0
[mg/l]			B	74.5	-0.03	75.0	60.0	90.0
IgE	+	5	A	60.1	-0.31	66.2	46.3	86.1
[kU/l]			B	121	-0.23	130	91.0	169
sol.Transfe.Receptor	+	2	A	0.930	0.07	0.910	0.637	1.19
[mg/l]			B	1.89	-0.07	1.93	1.35	2.51

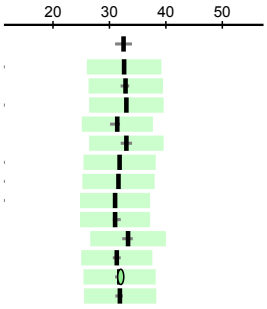
Analyte **Albumin**
Method all methods



No of participants		536	
sample/unit	A	g/l	B
mean	32.6		54.9
standard deviation	1.52		2.71
coefficient of variation	4.67		4.94

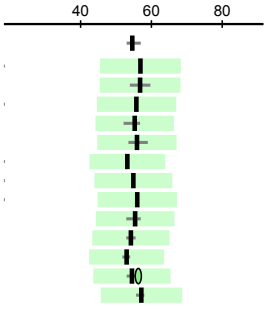
Sample A [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		536	28.0	31.0	32.5	34.0	51.2
2	13	5	30.1		32.6		32.9
2	28	16	31.0	32.0	32.9	33.5	35.0
2	30	4	32.7		33.0		47.0
4	4	12	30.0	30.1	31.4	31.9	32.6
4	30	118	30.3	32.0	33.0	34.0	37.0
4	38	7	30.9		31.8		32.6
4	43	7	31.2		31.6		33.0
4	328	7	29.3		31.0		32.0
7	4	68	29.0	30.6	31.0	32.0	42.9
7	30	191	29.0	32.3	33.3	34.1	43.8
7	38	27	28.8	30.6	31.3	32.0	32.7
7	43	43	28.0	31.0	31.8	32.5	33.8
7	328	8	31.0	31.0	31.9	32.4	32.5



Sample B [g/l]

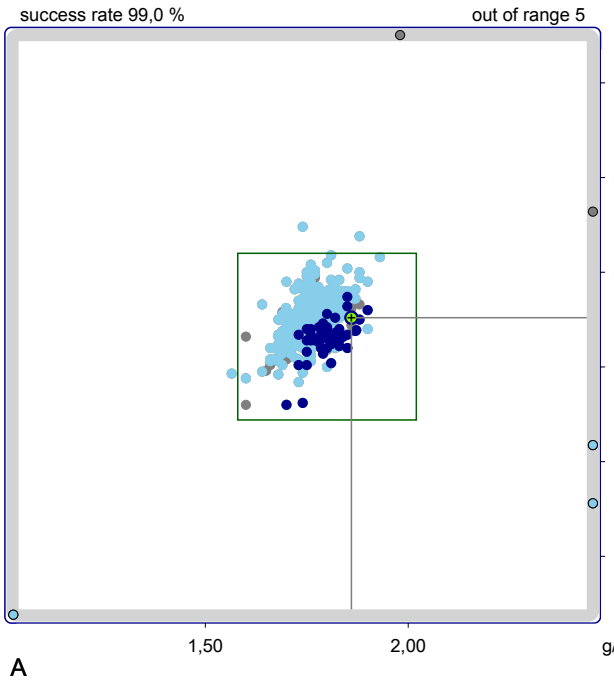
M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		536	31.1	53.0	54.6	57.0	565656
2	13	5	51.8		56.9		60.0
2	28	16	52.7	53.9	56.8	59.7	62.0
2	30	4	53.3		55.8		92.0
4	4	12	52.0	52.1	55.3	56.8	57.0
4	30	118	51.0	53.5	55.9	59.0	65.0
4	38	7	50.9		53.2		54.4
4	43	7	53.6		54.9		56.0
4	328	7	49.2		56.0		58.0
7	4	68	31.1	52.9	55.4	57.0	58.1
7	30	191	44.5	52.9	54.2	55.5	62.0
7	38	27	49.2	51.8	53.0	54.0	54.3
7	43	43	49.0	53.0	54.5	56.3	57.2
7	328	8	55.4	55.7	57.2	58.0	58.0



The deviation of your results from the median of the corresponding sub-collective (kit) is:	A	0.7 %
	B	3.3 %

Other kits (number):
1-30(3), 1-38(1), 2-38(1), 2-43(2), 2-140(1), 4-12(1), 4-13(1), 4-140(3), 5-38(1), 7-02(1), 7-12(1), 7-13(2), 7-40(1), 7-99(1), 7-126(2), 7-166(1),

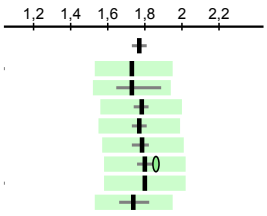
Analyte **Transferrin**
Method all methods



No of participants		653	
sample/unit	A	g/l	B
mean	1.77		3.75
standard deviation	0.045		0.108
coefficient of variation	2.59		2.90

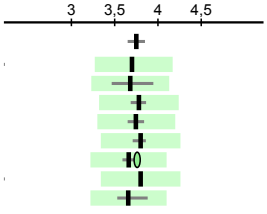
Sample A [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		653	0.001	1.73	1.77	1.81	2.84
1	30	3	1.66		1.73		1.80
2	28	11	1.60	1.65	1.73	1.89	1.98
4	4	96	1.57	1.74	1.78	1.82	2.84
4	30	397	0.001	1.73	1.77	1.81	2.73
4	38	54	1.64	1.73	1.79	1.82	1.90
4	43	54	1.70	1.76	1.80	1.85	1.90
4	126	7	1.68		1.80		1.83
4	328	12	1.60	1.66	1.74	1.82	1.85



Sample B [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		653	0.003	3.65	3.75	3.85	5.29
1	30	3	3.51		3.70		3.85
2	28	11	3.30	3.47	3.68	3.95	5.29
4	4	96	3.09	3.69	3.78	3.86	4.08
4	30	397	0.003	3.65	3.75	3.84	4.19
4	38	54	3.60	3.71	3.80	3.86	4.02
4	43	54	3.30	3.59	3.66	3.73	3.87
4	126	7	3.67		3.80		3.87
4	328	12	3.44	3.53	3.66	3.88	4.24

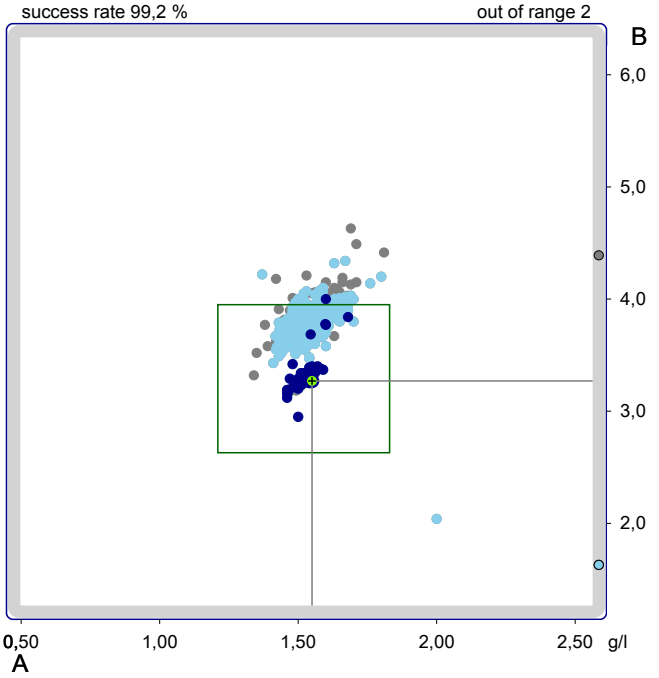


The deviation of your results from the median of the corresponding sub-collective (kit) is:	A	3.3 %
	B	2.6 %

Other kits (number):
1-04(1), 2-13(2), 2-30(2), 2-38(1), 2-43(2), 4-08(1), 4-13(2), 5-04(1), 5-30(2), 6-30(1), 7-30(2), 7-43(2),

Analyte **IgA**
Method all methods

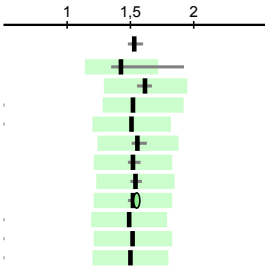
success rate 99,2 % out of range 2



No of participants		525	
sample/unit	A	g/l	B
mean	1.54		3.77
standard deviation	0.067		0.217
coefficient of variation	4.39		5.75

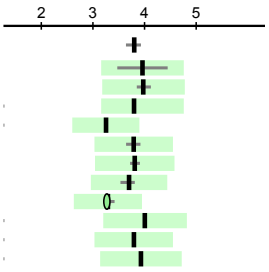
Sample A [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		525	1.34	1.48	1.53	1.60	2.67
2	13	10	1.34	1.35	1.43	1.92	2.66
2	28	38	1.45	1.55	1.62	1.67	1.81
2	30	3	1.51		1.52		1.56
2	43	4	1.48		1.51		1.63
4	4	80	1.45	1.51	1.56	1.63	2.67
4	30	291	1.42	1.48	1.52	1.58	2.00
4	38	33	1.41	1.50	1.54	1.59	1.65
4	43	43	1.46	1.48	1.52	1.57	1.68
4	126	4	1.37		1.49		1.57
4	140	6	1.43		1.52		1.60
4	328	5	1.47		1.50		1.66



Sample B [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		525	1.63	3.64	3.80	3.93	4.63
2	13	10	3.32	3.47	3.96	4.45	4.63
2	28	38	3.60	3.85	3.98	4.12	4.49
2	30	3	3.77		3.80		3.82
2	43	4	3.18		3.25		3.67
4	4	80	1.63	3.64	3.79	3.92	4.05
4	30	291	2.04	3.72	3.81	3.91	4.34
4	38	33	3.43	3.53	3.70	3.81	3.90
4	43	43	2.95	3.21	3.29	3.42	4.00
4	126	4	3.79		4.01		4.22
4	140	6	3.49		3.79		3.91
4	328	5	3.81		3.93		4.07

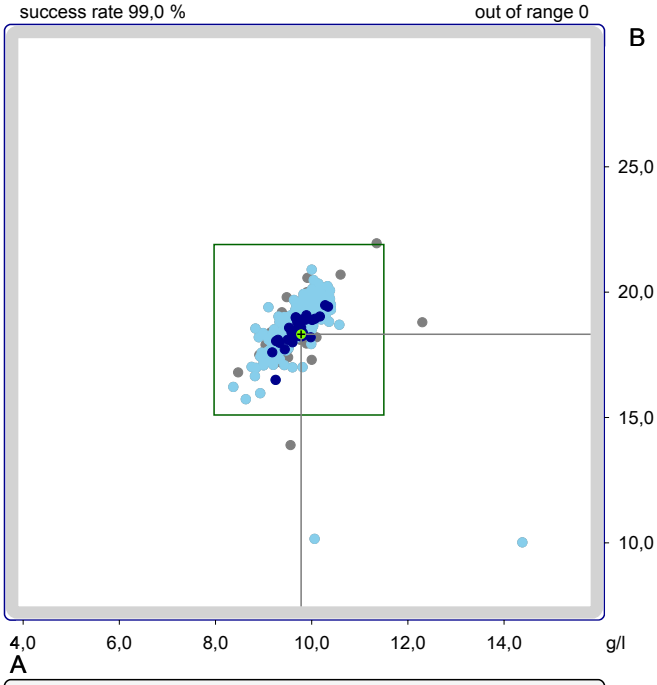


The deviation of your results from the median of the corresponding sub-collective (kit) is:
A 1.9 %
B -0.6 %

Other kits (number):
1-04(1), 1-30(2), 2-140(1), 4-13(2), 5-04(1), 6-30(1),

Analyte **IgG**
Method all methods

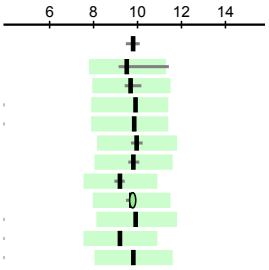
success rate 99,0 % out of range 0



No of participants		527	
sample/unit	A	g/l	B
mean	9.78		18.7
standard deviation	0.411		0.925
coefficient of variation	4.20		4.93

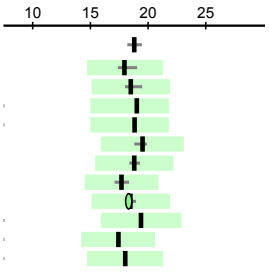
Sample A [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		527	8.37	9.47	9.80	10.1	14.4
2	13	11	9.04	9.14	9.51	11.4	12.3
2	28	38	8.47	9.42	9.69	10.2	10.6
2	30	3	9.74		9.91		10.2
2	43	3	8.91		9.85		9.96
4	4	83	9.22	9.70	9.96	10.2	14.4
4	30	293	8.82	9.58	9.81	10.1	10.6
4	38	34	8.37	8.95	9.20	9.42	9.62
4	43	40	9.18	9.47	9.72	9.93	10.3
4	126	4	8.75		9.92		10.1
4	140	5	8.93		9.20		9.99
4	328	5	9.15		9.81		10.2



Sample B [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		527	10.0	18.2	18.8	19.5	22.0
2	13	11	17.2	17.4	18.0	19.1	22.0
2	28	38	13.9	18.0	18.5	19.5	20.7
2	30	3	19.0		19.0		19.1
2	43	3	17.5		18.8		19.2
4	4	83	10.0	18.8	19.5	19.9	20.9
4	30	293	10.2	18.4	18.8	19.3	20.1
4	38	34	16.0	17.1	17.7	18.4	19.0
4	43	40	16.5	18.1	18.5	19.0	19.5
4	126	4	17.0		19.4		19.9
4	140	5	17.3		17.4		19.1
4	328	5	17.0		18.0		20.2

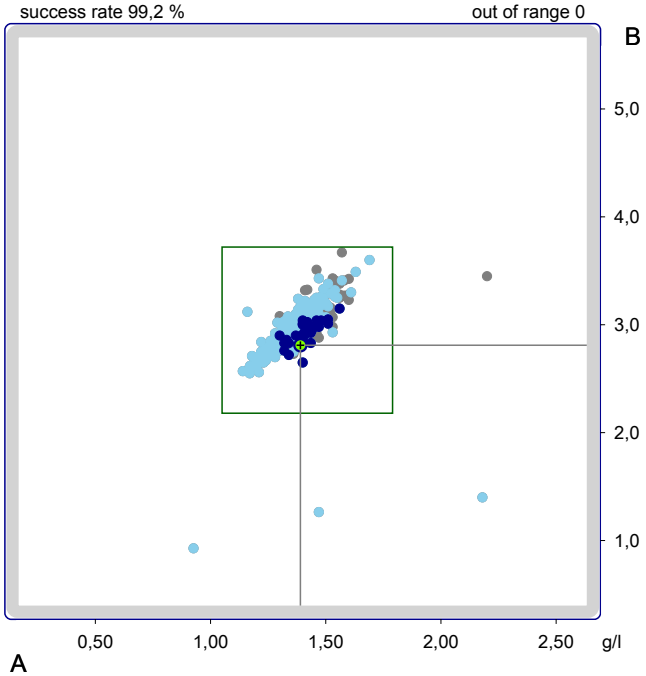


The deviation of your results from the median of the corresponding sub-collective (kit) is:
A 0.6 %
B -0.7 %

Other kits (number):
1-04(1), 1-30(2), 2-140(1), 4-13(2), 5-04(1), 6-30(1),

Analyte **IgM**
Method all methods

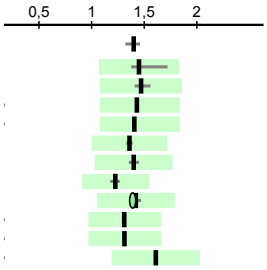
success rate 99,2 % out of range 0



No of participants		521	
sample/unit	A	g/l	B
mean	1.39		3.01
standard deviation	0.095		0.212
coefficient of variation	6.84		7.04

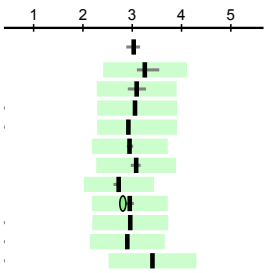
Sample A [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		521	0.926	1.32	1.40	1.46	2.20
2	13	10	1.30	1.38	1.45	1.72	2.20
2	28	37	1.32	1.41	1.47	1.56	1.60
2	30	3	1.40		1.43		1.47
2	43	3	1.30		1.41		1.42
4	4	82	1.26	1.33	1.36	1.39	2.18
4	30	288	0.926	1.35	1.40	1.45	1.55
4	38	34	1.14	1.18	1.23	1.27	1.32
4	43	41	1.30	1.36	1.42	1.47	1.56
4	126	4	1.22		1.31		1.42
4	140	5	1.16		1.31		1.39
4	328	5	1.54		1.61		1.69



Sample B [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		521	0.928	2.88	3.03	3.16	3.67
2	13	10	3.08	3.10	3.26	3.55	3.67
2	28	37	2.73	2.91	3.09	3.28	3.43
2	30	3	3.05		3.06		3.09
2	43	3	2.80		2.92		2.94
4	4	82	1.40	2.89	2.95	3.02	3.25
4	30	288	0.928	2.97	3.08	3.17	3.43
4	38	34	2.55	2.62	2.73	2.77	2.84
4	43	41	2.65	2.83	2.95	3.03	3.15
4	126	4	2.84		2.96		3.08
4	140	5	2.83		2.90		3.12
4	328	5	3.26		3.41		3.60

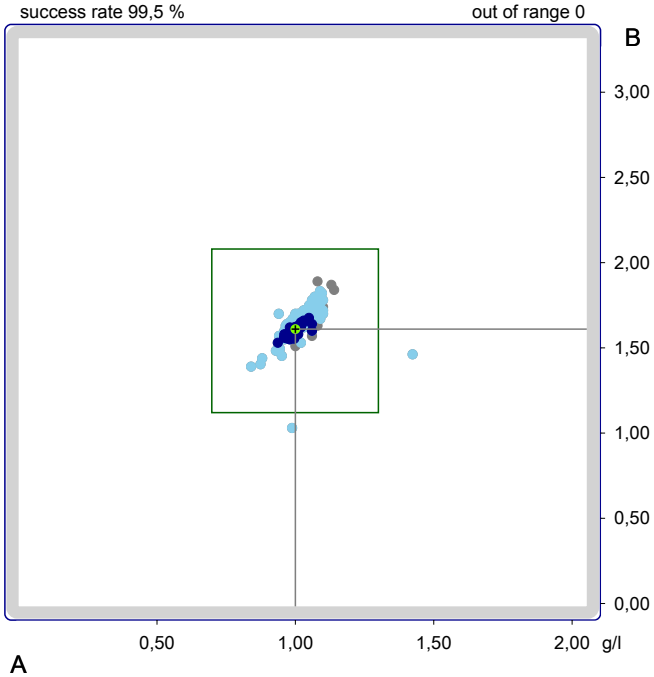


The deviation of your results from the median of the corresponding sub-collective (kit) is:
A -2.1 %
B -4.7 %

Other kits (number):
1-04(1), 1-30(2), 2-140(1), 4-13(2), 5-04(1), 5-30(1), 6-30(1),

Analyte **Haptoglobin**
Method all methods

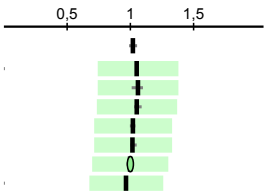
success rate 99,5 % out of range 0



No of participants		401	
sample/unit	A	g/l	B
mean	1.02		1.65
standard deviation	0.041		0.069
coefficient of variation	4.02		4.20

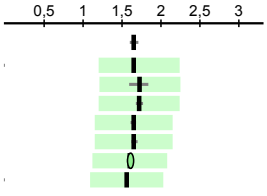
Sample A [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		401	0.840	0.990	1.02	1.05	1.42
2	13	3	1.00		1.05		1.07
2	28	18	0.980	1.01	1.06	1.10	1.14
4	4	60	0.990	1.03	1.05	1.09	1.42
4	30	234	0.930	0.996	1.02	1.04	1.10
4	38	24	0.944	1.00	1.02	1.05	1.07
4	43	46	0.937	0.970	0.998	1.01	1.06
4	140	6	0.874		0.966		0.994



Sample B [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		401	1.03	1.60	1.65	1.71	1.89
2	13	3	1.51		1.65		1.74
2	28	18	1.57	1.59	1.73	1.84	1.89
4	4	60	1.46	1.68	1.72	1.77	1.83
4	30	234	1.03	1.61	1.65	1.68	1.76
4	38	24	1.50	1.62	1.65	1.70	1.72
4	43	46	1.53	1.56	1.60	1.63	1.68
4	140	6	1.40		1.56		1.60

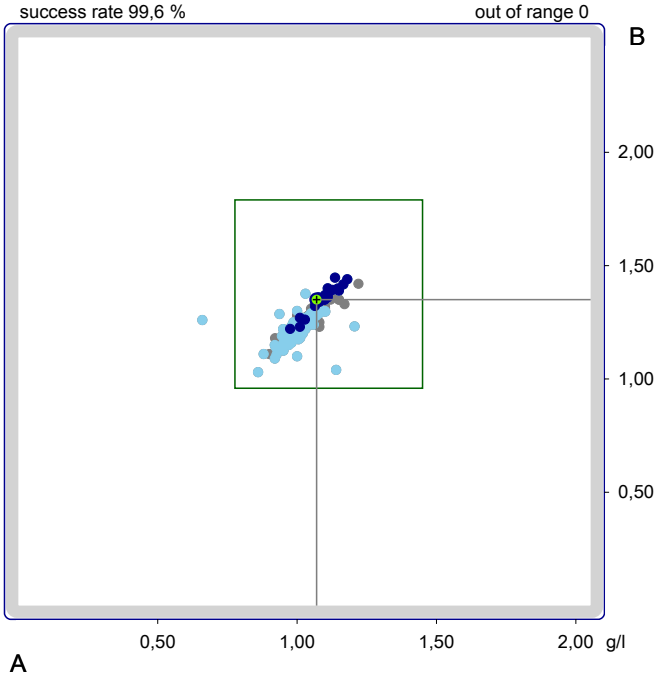


The deviation of your results from the median of the corresponding sub-collective (kit) is:
A 0.2 %
B 0.6 %

Other kits (number):
1-30(1), 2-43(1), 4-12(1), 4-13(1), 4-45(1), 4-328(2), 5-30(1), 6-30(1), 7-43(1),

Analyte **C3-complement**
Method all methods

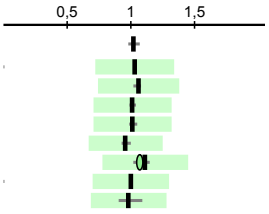
success rate 99,6 % out of range 0



No of participants		289	
sample/unit	A	g/l	B
mean	1.02		1.24
standard deviation	0.056		0.063
coefficient of variation	5.51		5.11

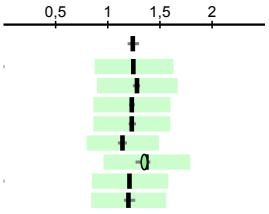
Sample A [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		289	0.660	0.980	1.02	1.07	1.22
2	13	6	0.965		1.03		1.17
2	28	57	0.921	1.02	1.06	1.08	1.22
4	4	27	0.950	0.989	1.01	1.04	1.14
4	30	130	0.920	0.987	1.01	1.05	1.10
4	38	16	0.860	0.927	0.956	1.00	1.01
4	43	21	0.975	1.02	1.11	1.15	1.18
4	126	7	0.923		1.00		1.05
4	140	11	0.660	0.904	0.980	1.09	1.21



Sample B [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		289	1.03	1.19	1.24	1.30	1.45
2	13	6	1.18		1.25		1.35
2	28	57	1.16	1.24	1.28	1.31	1.42
4	4	27	1.04	1.20	1.23	1.26	1.29
4	30	130	1.12	1.20	1.23	1.27	1.31
4	38	16	1.03	1.10	1.14	1.18	1.19
4	43	21	1.22	1.27	1.37	1.41	1.45
4	126	7	1.14		1.21		1.29
4	140	11	1.11	1.16	1.20	1.26	1.30

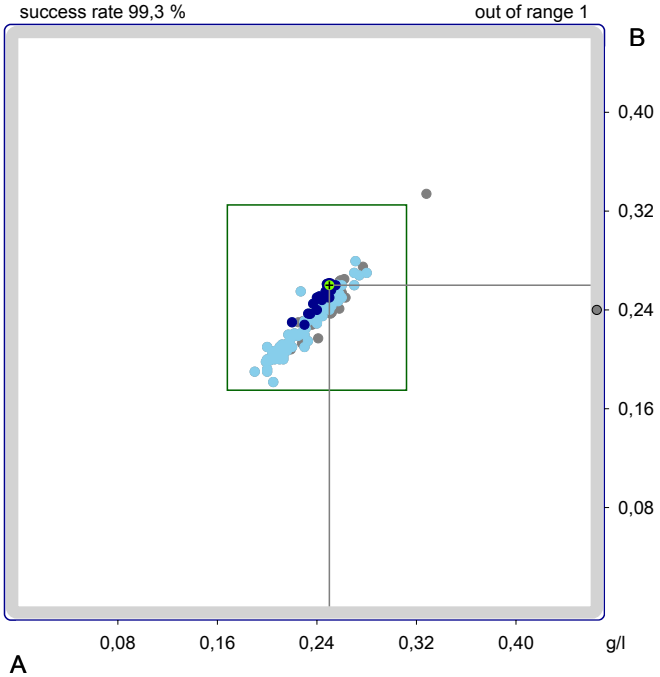


The deviation of your results from the median of the corresponding sub-collective (kit) is:
A -3.6 %
B -1.4 %

Other kits (number):
1-04(1), 1-30(1), 2-30(1), 2-43(1), 2-140(1), 2-156(1), 4-12(2), 4-28(1), 4-328(2), 5-04(1), 6-30(1), 7-02(1),

Analyte **C4-complement**
Method all methods

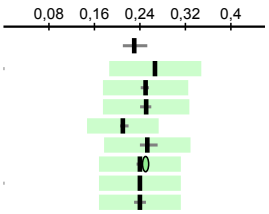
success rate 99,3 % out of range 1



No of participants		290	
sample/unit	A	g/l	B
mean	0.231		0.228
standard deviation	0.020		0.022
coefficient of variation	9.04		9.70

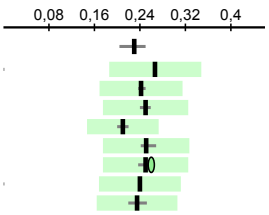
Sample A [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		290	0.190	0.210	0.230	0.253	0.760
2	13	6	0.229		0.267		0.328
2	28	58	0.219	0.241	0.250	0.256	0.760
4	4	26	0.236	0.240	0.251	0.260	0.270
4	30	130	0.190	0.205	0.210	0.220	0.271
4	38	16	0.200	0.240	0.253	0.271	0.280
4	43	22	0.220	0.234	0.240	0.250	0.255
4	126	7	0.230		0.240		0.250
4	140	11	0.227	0.230	0.240	0.251	0.258



Sample B [g/l]

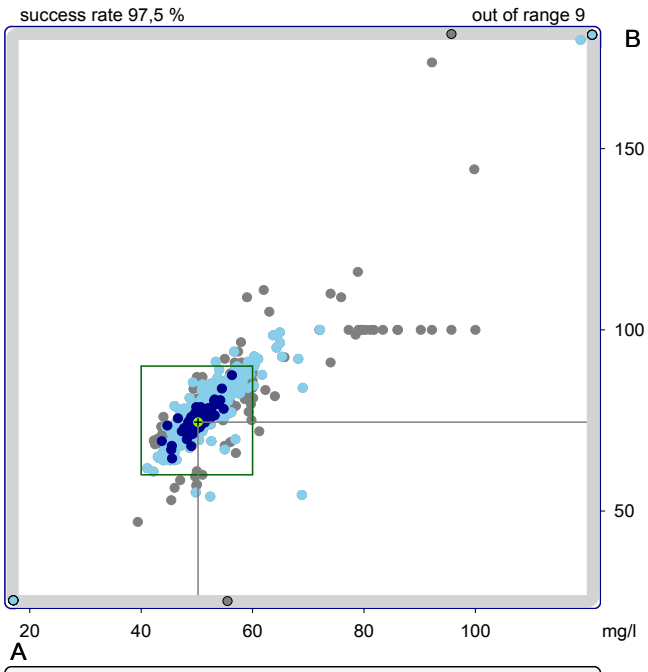
M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		290	0.182	0.204	0.230	0.250	0.334
2	13	6	0.231		0.267		0.334
2	28	58	0.208	0.237	0.242	0.250	0.264
4	4	26	0.235	0.240	0.250	0.259	0.270
4	30	130	0.190	0.200	0.210	0.220	0.280
4	38	16	0.200	0.241	0.251	0.269	0.270
4	43	22	0.228	0.237	0.250	0.260	0.261
4	126	7	0.210		0.240		0.250
4	140	11	0.215	0.220	0.235	0.252	0.255



The deviation of your results from the median of the corresponding sub-collective (kit) is:
A 4.1 %
B 4.0 %

Other kits (number):
1-04(1), 1-30(1), 2-04(1), 2-30(1), 2-43(1), 2-140(1), 4-12(2), 4-28(1), 4-328(2), 5-04(1), 6-30(1), 7-02(1),

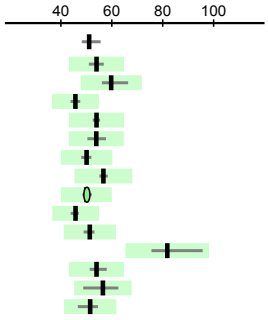
Analyte **CRP**
Method all methods



No of participants		1280	
sample/unit	A	mg/l	B
mean	52.2		77.9
standard deviation	6.14		7.96
coefficient of variation	11.8		10.2

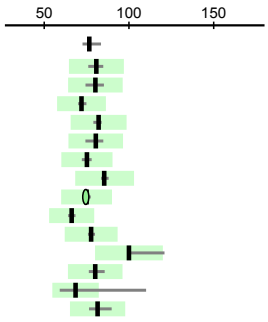
Sample A [mg/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		1280	1.73	48.2	51.1	55.6	521
1	61	50	50.0	51.0	54.0	56.8	60.0
2	13	10	48.2	56.1	59.8	66.4	74.0
2	28	43	42.3	43.8	45.7	47.6	49.9
4	4	149	44.5	52.5	54.0	55.4	68.9
4	7	50	5.60	50.4	53.9	57.7	64.9
4	30	623	1.73	48.0	50.1	52.0	521
4	38	83	52.5	55.1	56.7	58.4	64.9
4	43	80	43.7	48.4	50.0	52.1	56.3
4	126	20	42.2	43.8	45.8	47.0	49.6
4	328	32	48.2	48.9	51.4	53.2	55.0
5	44	23	9.66	75.6	81.8	95.7	100
5	61	26	49.0	51.3	54.0	58.0	60.0
6	89	8	47.0	48.8	56.5	62.6	63.0
6	130	8	44.0	46.6	51.5	54.6	55.0



Sample B [mg/l]

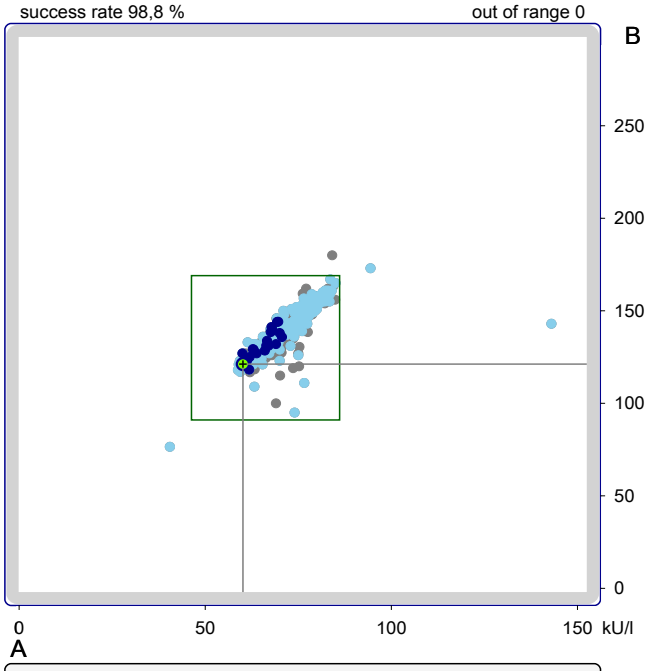
M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		1280	0.500	72.6	76.6	83.5	775
1	61	50	71.0	76.0	80.8	84.8	91.0
2	13	10	72.0	74.4	80.2	85.2	91.0
2	28	43	68.4	70.0	72.0	74.9	77.5
4	4	149	54.4	79.0	82.1	83.9	98.5
4	7	50	8.12	74.3	80.4	84.9	99.3
4	30	623	1.33	72.2	75.2	78.0	775
4	38	83	80.0	83.5	85.4	88.0	96.4
4	43	80	64.5	72.8	75.0	77.5	87.5
4	126	20	60.9	64.1	66.2	68.4	74.3
4	328	32	69.3	75.8	77.7	80.0	84.4
5	44	23	14.4	100	100	121	324
5	61	26	75.0	76.3	80.1	85.7	91.0
6	89	8	58.5	59.2	68.5	110	111
6	130	8	76.0	76.4	81.5	89.8	92.0



The deviation of your results from the median of the corresponding sub-collective (kit) is:
A 0.4 %
B -0.6 %

Other kits (number):
1-04(1), 1-30(1), 1-99(1), 1-159(1), 2-99(1), 3-159(1), 4-507(6), 4-08(3), 4-12(5), 4-13(3), 4-32(1), 4-33(6), 4-34(2), 4-40(5), 4-89(1), 4-99(3), 4-128(1), 4-140(2), 4-166(1), 4-168(1), 4-364(6), 5-38(1), 5-89(1), 5-159(5), 5-335(1), 6-08(6), 6-30(1), 6-99(1), 6-159(1), 7-30(2), 7-99(3), 7-151(1),

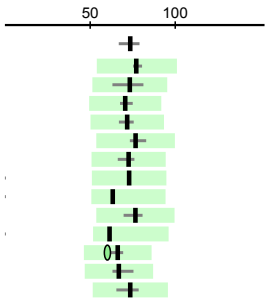
Analyte **IgE**
Method all methods



No of participants		352	
sample/unit	A	kU/l	B
mean	73.2		142
standard deviation	7.26		12.5
coefficient of variation	9.92		8.81

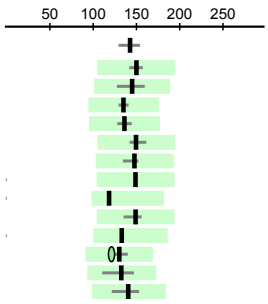
Sample A [kU/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		352	40.5	66.8	73.6	79.0	143
1	30	9	73.6	75.2	77.1	80.5	80.8
1	72	14	60.8	63.1	73.3	81.3	84.9
2	28	15	66.0	67.5	70.6	75.0	77.6
4	4	34	65.0	66.9	71.8	75.6	77.8
4	30	12	70.2	73.3	76.7	82.9	84.3
4	38	9	61.0	66.2	72.6	76.0	79.0
BeckCou2		7	70.6		72.9		76.2
4	140	3	62.0		63.3		66.8
5	30	156	61.8	69.7	76.6	80.8	85.0
5	40	3	59.3		61.4		68.1
5	43	22	60.0	61.1	66.2	69.4	70.6
5	44	16	40.5	63.1	66.9	75.4	94.4
5	72	38	58.8	65.4	73.6	78.5	143



Sample B [kU/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		352	76.5	129	143	154	180
1	30	9	139	142	150	157	158
1	72	14	126	127	145	160	180
2	28	15	115	129	135	141	144
4	4	34	120	128	136	145	151
4	30	12	139	142	150	161	163
4	38	9	120	134	148	152	154
BeckCou2		7	119		149		159
4	140	3	117		118		126
5	30	156	122	135	149	156	165
5	40	3	123		133		135
5	43	22	118	123	130	140	144
5	44	16	76.5	110	133	147	173
5	72	38	95.0	121	141	153	167

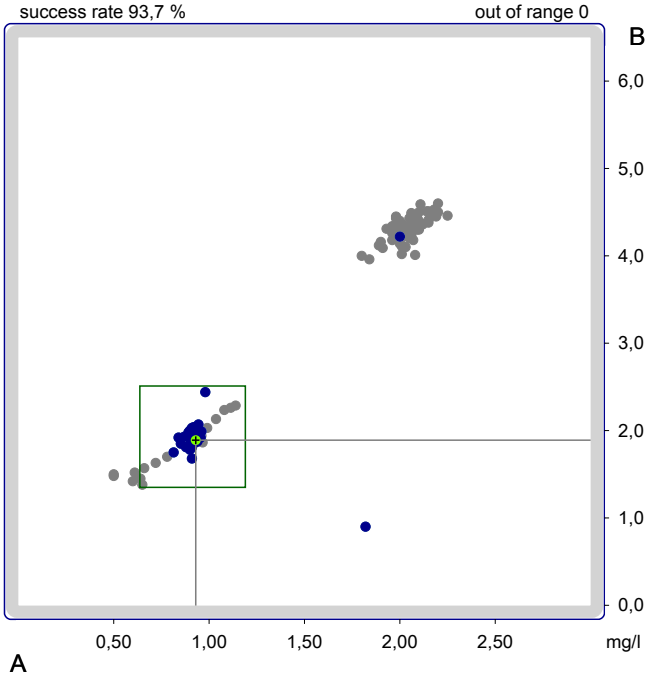


The deviation of your results from the median of the corresponding sub-collective (kit) is:
A -9.1 %
B -6.9 %

Other kits (number):
1-04(1), 1-43(1), 2-13(1), 2-99(1), 3-30(1), 4-12(2), 4-13(2), 4-16(1), 4-43(2), 5-04(1), 5-23(1),

Analyte **sol.Transfe.Receptor**
Method all methods

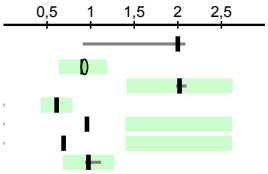
success rate 93,7 % out of range 0



No of participants		145	
sample/unit	A	mg/l	B
mean	1.65		3.48
standard deviation	0.556		1.17
coefficient of variation	33.6		33.6

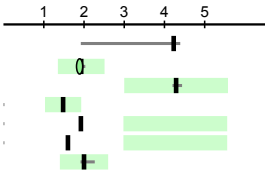
Sample A [mg/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		145	0.500	0.910	2.00	2.09	2.25
2	28	29	0.814	0.877	0.910	0.957	2.00
4	30	94	1.80	1.98	2.02	2.10	2.25
4	43	5	0.500		0.610		0.650
4	140	2	0.950		0.958		0.966
4	154	2	0.660		0.690		0.720
5	13	10	0.920	0.935	0.975	1.12	1.14



Sample B [mg/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		145	0.900	1.92	4.23	4.39	4.60
2	28	29	0.900	1.80	1.93	2.03	4.22
4	30	94	3.96	4.20	4.29	4.44	4.60
4	43	5	1.38		1.48		1.52
4	140	2	1.86		1.92		1.98
4	154	2	1.57		1.60		1.63
5	13	10	1.89	1.91	2.00	2.27	2.29



The deviation of your results from the median of the corresponding sub-collective (kit) is:
A 2.2 %
B -2.0 %

Other kits (number):
1-154(1), 4-105(1), 5-30(1),