Survey: **IG2/25**
month: **April 2025**
Participant-No: **9900954**
valid from: **26.04.25**

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

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

Bonn, 16. May 2025

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 2025.
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 14.05.25.
This is available for download in the RfB-Online system.

Survey: **IG2/25**
month: **April 2025**
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)

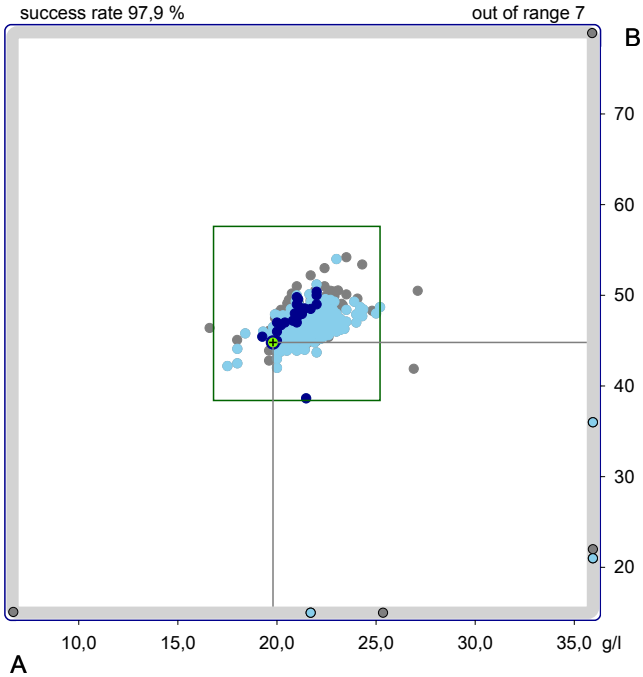
Certificate of participation

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	19.8	-0.29	21.0	16.8	25.2
[g/l]			B	44.8	-0.33	48.0	38.4	57.6
Transferrin	+	4	A	1.43	0.17	1.40	1.23	1.57
[g/l]			B	3.88	0.40	3.70	3.25	4.15
IgA	+	4	A	1.01	0	1.01	0.808	1.22
[g/l]			B	3.08	0	3.08	2.46	3.70
IgG	+	4	A	4.92	0.28	4.68	3.83	5.53
[g/l]			B	16.7	-0.25	17.5	14.3	20.7
IgM	+	4	A	0.470	0.08	0.460	0.340	0.580
[g/l]			B	2.95	0.09	2.88	2.13	3.63
Haptoglobin	+	4	A	0.640	-0.20	0.680	0.476	0.884
[g/l]			B	1.61	-0.25	1.74	1.21	2.27
C3-complement	+	4	A	0.400	0	0.400	0.280	0.520
[g/l]			B	0.940	-0.10	0.970	0.679	1.27
C4-complement	+	4	A	0.030	0	0.030	0	0.080
[g/l]			B	0.080	-0.37	0.090	0.063	0.117
CRP	+	4	A	12.5	0.25	11.9	9.52	14.3
[mg/l]			B	57.4	-0.09	58.5	46.8	70.2
IgE	+	5	A	94.1	0.15	90.0	63.0	117
[kU/l]			B	220	0.22	206	144	268
sol.Transfe.Receptor	+	2	A	0.630	0.01	0.627	0.438	0.816
[mg/l]			B	1.76	0.11	1.70	1.19	2.21

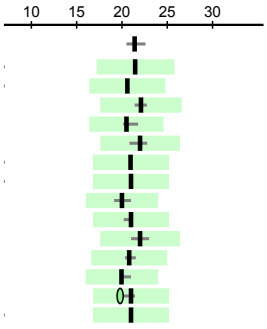
Analyte **Albumin**
Method all methods



No of participants		533	
sample/unit	A	g/l	B
mean	21.5		46.9
standard deviation	1.15		1.80
coefficient of variation	5.34		3.84

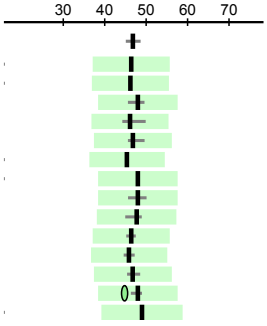
Sample A [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		533	0.021	20.5	21.4	22.6	21800
1	30	6	20.8		21.5		23.0
2	13	5	19.8		20.6		21.3
2	28	21	21.0	21.4	22.1	22.7	47.0
4	4	12	20.0	20.1	20.5	21.8	21.9
4	30	106	0.021	20.8	22.0	22.8	21800
4	38	6	19.8		21.0		21.6
4	43	7	19.6		21.0		26.9
4	328	9	18.0	19.1	20.0	21.0	21.0
7	4	65	19.9	20.2	21.0	21.1	46.9
7	30	186	17.5	21.0	22.0	23.0	25.2
7	38	38	19.8	20.3	20.8	21.5	22.4
7	40	8	19.8	19.8	20.0	21.0	21.0
7	43	33	19.3	20.0	21.0	21.4	22.0
7	328	7	19.9		21.0		22.1



Sample B [g/l]

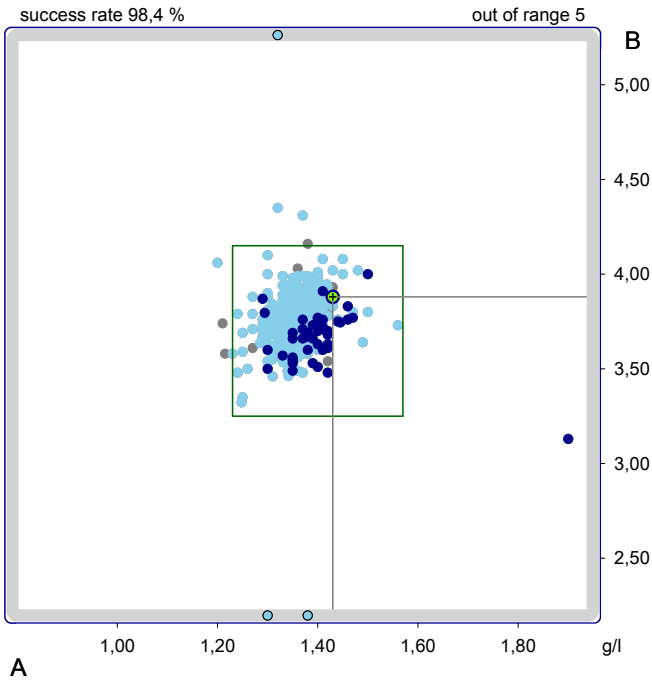
M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		533	0.045	45.1	46.8	48.7	44800
1	30	6	44.7		46.4		48.1
2	13	5	45.4		46.2		48.9
2	28	21	22.0	45.6	48.0	49.6	50.5
4	4	12	43.0	44.3	46.1	49.9	50.5
4	30	106	0.045	45.6	46.8	49.6	44800
4	38	6	44.0		45.4		45.9
4	43	7	41.9		48.0		49.1
4	328	9	45.1	45.6	48.0	50.1	51.0
7	4	65	21.0	45.0	47.7	48.9	50.1
7	30	186	42.0	45.2	46.4	47.5	54.0
7	38	38	43.0	44.6	45.9	47.2	48.8
7	40	8	45.4	45.4	46.8	48.6	49.0
7	43	33	38.6	46.3	48.0	49.0	50.4
7	328	7	47.8		49.0		49.5



The deviation of your results from the median of the corresponding sub-collective (kit) is:	A	-5.7 %
	B	-6.6 %

Other kits (number):
1-04(1), 2-30(3), 2-38(1), 2-43(2), 2-140(1), 2-228(1), 4-12(1), 4-13(1), 4-40(2), 4-140(2), 6-08(1), 7-12(1), 7-13(2), 7-32(1), 7-126(4),

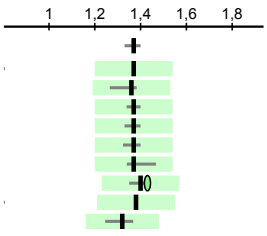
Analyte **Transferrin**
Method all methods



No of participants		647	
sample/unit	A	g/l	B
mean	1.37		3.76
standard deviation	0.045		0.118
coefficient of variation	3.34		3.14

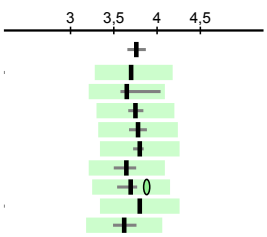
Sample A [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		647	1.20	1.33	1.37	1.40	3.90
1	30	3	1.31		1.37		1.40
2	28	11	1.22	1.27	1.36	1.38	1.42
4	4	98	1.30	1.34	1.37	1.40	3.90
4	30	381	1.23	1.33	1.37	1.40	1.45
4	38	51	1.20	1.32	1.37	1.40	1.48
4	40	12	1.31	1.34	1.37	1.47	1.49
4	43	48	1.29	1.35	1.40	1.43	1.90
4	126	5	1.32		1.38		1.50
4	328	15	1.24	1.25	1.32	1.37	1.56



Sample B [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		647	0.392	3.66	3.76	3.87	43.3
1	30	3	3.61		3.70		3.75
2	28	11	3.54	3.58	3.65	4.04	4.16
4	4	98	1.44	3.67	3.75	3.84	4.02
4	30	381	0.392	3.68	3.78	3.88	4.31
4	38	51	0.430	3.72	3.80	3.84	4.06
4	40	12	3.46	3.50	3.65	3.76	3.80
4	43	48	3.13	3.54	3.70	3.77	4.00
4	126	5	3.73		3.80		43.3
4	328	15	3.35	3.49	3.62	3.76	4.35

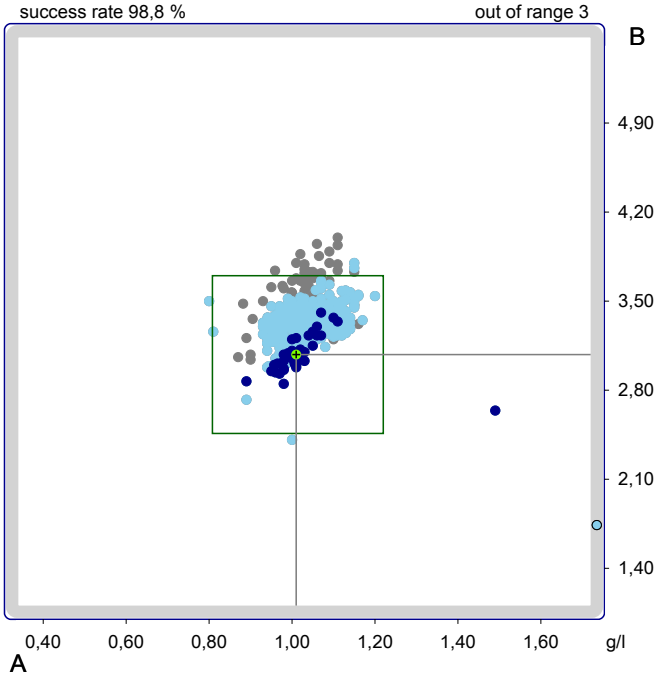


The deviation of your results from the median of the corresponding sub-collective (kit) is:	A	2.1 %
	B	5.0 %

Other kits (number):
1-04(2), 2-13(2), 2-30(2), 2-38(1), 2-43(2), 2-228(1), 4-13(2), 4-16(1), 4-128(1), 4-154(1), 5-04(1), 5-30(1), 6-30(1), 7-30(1), 7-38(2), 7-43(2),

Analyte **IgA**
Method all methods

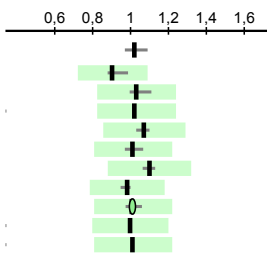
success rate 98,8 % out of range 3



No of participants		**538**	
sample/unit	A	g/l	B
mean	1.03		3.35
standard deviation	0.060		0.179
coefficient of variation	5.91		5.35

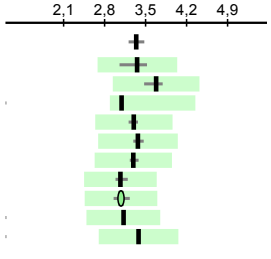
Sample A [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		538	0.800	0.970	1.02	1.09	3.63
2	13	10	0.870	0.879	0.903	0.987	0.999
2	28	48	0.950	0.995	1.03	1.11	3.63
2	43	3	0.969		1.02		1.09
4	4	81	0.890	1.03	1.07	1.10	1.17
4	30	276	0.930	0.970	1.01	1.07	1.20
4	38	39	0.800	1.06	1.10	1.13	1.15
4	40	13	0.940	0.948	0.982	1.00	1.03
4	43	39	0.890	0.974	1.01	1.06	1.49
4	140	6	0.970		0.998		1.02
4	328	5	0.810		1.01		1.09



Sample B [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		538	1.02	3.21	3.34	3.48	4.00
2	13	10	3.04	3.06	3.36	3.52	3.60
2	28	48	1.02	3.48	3.68	3.79	4.00
2	43	3	2.98		3.09		3.66
4	4	81	2.73	3.21	3.30	3.37	3.53
4	30	276	3.11	3.29	3.37	3.47	3.80
4	38	39	3.07	3.23	3.29	3.38	3.50
4	40	13	2.98	2.99	3.07	3.20	3.20
4	43	39	2.64	2.95	3.08	3.23	3.41
4	140	6	3.01		3.12		3.23
4	328	5	3.26		3.38		3.63

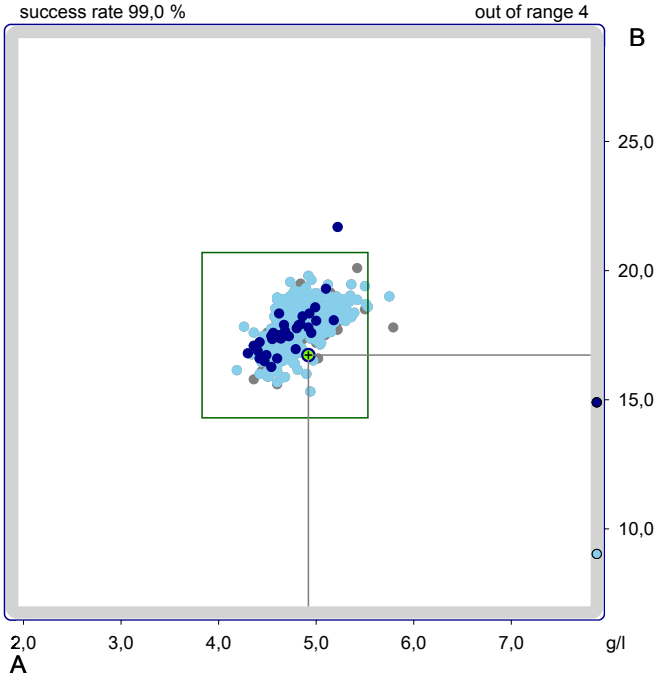


The deviation of your results from the median of the corresponding sub-collective (kit) is: A 0.2 %, B 0 %

Other kits (number):
1-04(1), 1-30(2), 2-30(2), 2-126(1), 2-140(1), 2-228(1), 4-13(2), 4-126(2), 4-128(1), 4-154(1), 5-04(1), 6-30(1), 7-30(1), 7-38(1),

Analyte **IgG**
Method all methods

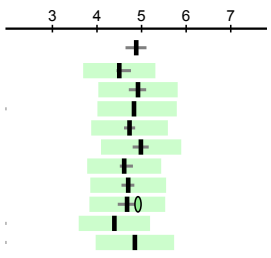
success rate 99,0 % out of range 4



No of participants		**536**	
sample/unit	A	g/l	B
mean	4.87		17.9
standard deviation	0.234		0.773
coefficient of variation	4.81		4.32

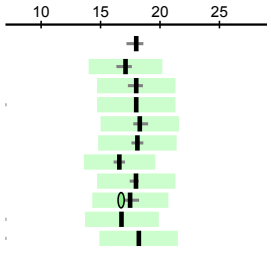
Sample A [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		536	4.19	4.64	4.88	5.11	18.4
2	13	11	4.41	4.44	4.50	4.76	4.90
2	28	47	4.51	4.71	4.92	5.10	18.4
2	43	3	4.71		4.83		5.16
4	4	83	4.26	4.61	4.73	4.86	5.16
4	30	276	4.45	4.80	4.99	5.16	5.75
4	38	39	4.43	4.51	4.61	4.81	4.97
4	40	13	4.46	4.55	4.70	4.84	4.86
4	43	36	4.30	4.47	4.68	4.99	7.83
4	140	5	4.19		4.39		4.42
4	328	5	4.60		4.85		5.50



Sample B [g/l]

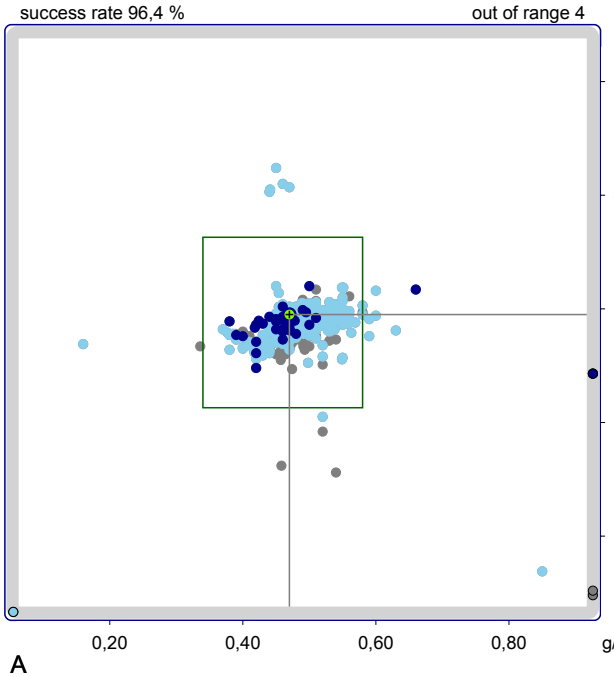
M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		536	4.88	17.2	18.0	18.6	21.7
2	13	11	16.1	16.3	17.1	17.6	18.0
2	28	47	4.88	17.3	18.0	18.6	20.1
2	43	3	17.9		18.0		18.2
4	4	83	17.1	17.8	18.3	19.0	19.8
4	30	276	16.6	17.6	18.1	18.6	19.5
4	38	39	15.7	16.1	16.6	17.1	17.5
4	40	13	17.3	17.4	18.0	18.2	18.2
4	43	36	14.9	16.7	17.5	18.2	21.7
4	140	5	16.0		16.8		17.0
4	328	5	16.9		18.2		19.4



The deviation of your results from the median of the corresponding sub-collective (kit) is: A 5.2 %, B -4.3 %

Other kits (number):
1-04(1), 1-30(2), 2-30(2), 2-126(1), 2-140(1), 2-228(1), 4-13(2), 4-126(2), 4-128(1), 4-154(1), 5-04(1), 6-30(1), 7-30(1), 7-38(1),

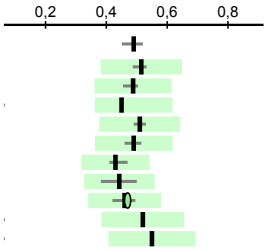
Analyte **IgM**
Method all methods



No of participants		532	
sample/unit	A	g/l	B
mean	0.487		2.86
standard deviation	0.044		0.213
coefficient of variation	9.14		7.44

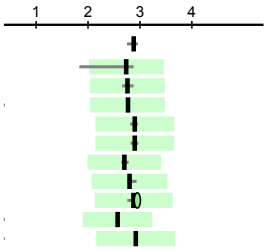
Sample A [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		532	0.047	0.451	0.490	0.520	2.83
2	13	10	0.467	0.487	0.515	0.532	0.540
2	28	47	0.336	0.455	0.488	0.504	2.83
2	43	3	0.402		0.450		0.490
4	4	83	0.441	0.490	0.510	0.530	0.560
4	30	272	0.047	0.460	0.490	0.515	0.600
4	38	39	0.160	0.410	0.430	0.470	0.500
4	40	14	0.377	0.382	0.443	0.500	0.530
4	43	36	0.380	0.420	0.460	0.495	1.06
4	140	5	0.498		0.520		0.550
4	328	5	0.370		0.550		0.630



Sample B [g/l]

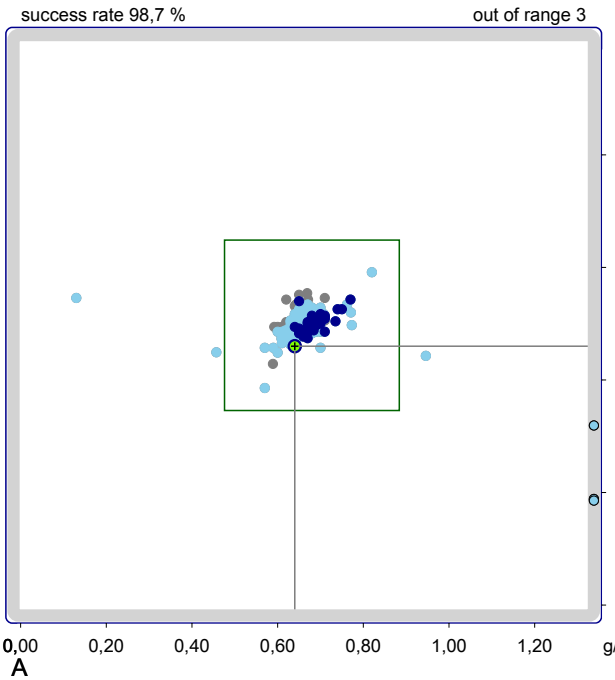
M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		532	0.283	2.75	2.88	2.96	4.24
2	13	10	1.56	1.83	2.74	2.88	2.90
2	28	47	0.480	2.66	2.76	2.88	3.07
2	43	3	2.75		2.77		2.83
4	4	83	2.64	2.81	2.90	2.96	3.19
4	30	272	0.283	2.82	2.90	2.97	4.24
4	38	39	2.56	2.64	2.70	2.78	2.90
4	40	14	2.61	2.76	2.80	2.94	2.97
4	43	36	2.43	2.76	2.88	2.96	3.20
4	140	5	2.53		2.57		2.68
4	328	5	2.81		2.92		3.16



The deviation of your results from the median of the corresponding sub-collective (kit) is:	A	2.1 %
	B	2.6 %

Other kits (number):
1-04(1), 1-30(2), 2-30(2), 2-126(1), 2-140(1), 2-228(1), 4-13(2), 4-126(1), 4-128(1), 4-154(1), 5-04(1), 5-30(1), 6-30(1), 7-30(1), 7-38(1),

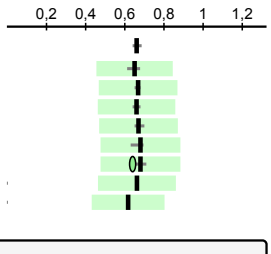
Analyte **Haptoglobin**
Method all methods



No of participants		396	
sample/unit	A	g/l	B
mean	0.662		1.75
standard deviation	0.042		0.062
coefficient of variation	6.44		3.58

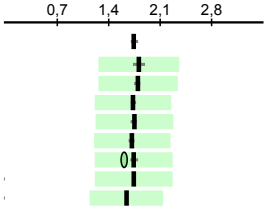
Sample A [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		396	0.130	0.640	0.661	0.686	1.83
2	28	19	0.589	0.612	0.650	0.678	0.700
4	4	59	0.620	0.650	0.668	0.680	1.83
4	30	223	0.590	0.640	0.660	0.680	0.946
4	38	23	0.640	0.650	0.670	0.700	0.708
4	40	8	0.630	0.630	0.680	0.696	0.700
4	43	39	0.640	0.650	0.680	0.710	0.770
4	140	4	0.610		0.662		0.686
4	328	5	0.457		0.617		0.773



Sample B [g/l]

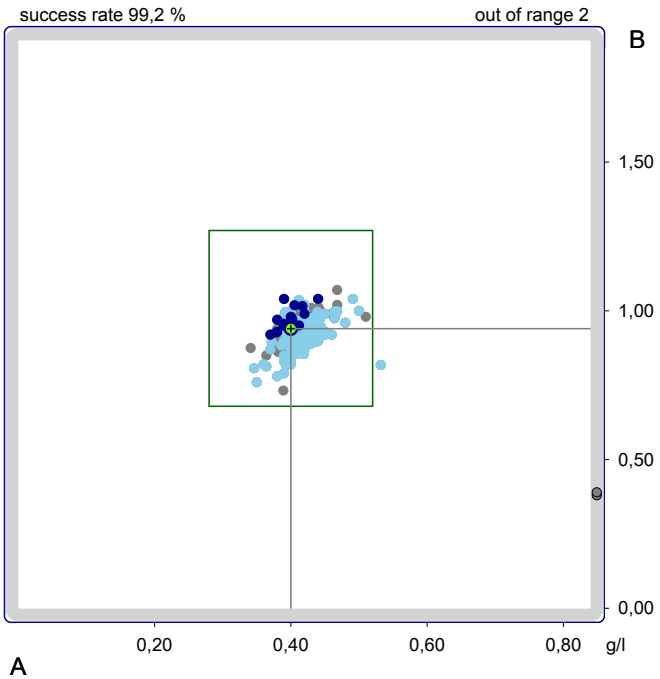
M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		396	0.650	1.70	1.74	1.80	2.07
2	28	19	1.50	1.73	1.81	1.89	1.93
4	4	59	0.650	1.75	1.80	1.83	1.87
4	30	223	1.55	1.70	1.73	1.77	2.07
4	38	23	1.60	1.70	1.75	1.78	1.81
4	40	8	1.67	1.67	1.72	1.75	1.76
4	43	39	1.61	1.69	1.74	1.80	1.90
4	140	4	1.70		1.74		1.76
4	328	5	1.57		1.64		1.74



The deviation of your results from the median of the corresponding sub-collective (kit) is:	A	-5.8 %
	B	-7.4 %

Other kits (number):
1-04(1), 1-30(1), 2-13(2), 2-43(2), 2-140(1), 4-12(1), 4-13(1), 4-45(1), 4-126(1), 4-154(1), 5-04(1), 6-30(1), 7-38(1), 7-43(1),

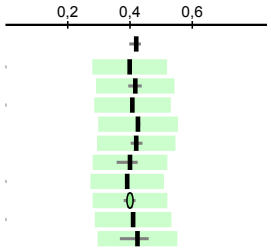
Analyte **C3-complement**
Method all methods



No of participants		285	
sample/unit	A	g/l	B
mean	0.417		0.935
standard deviation	0.023		0.049
coefficient of variation	5.73		5.31

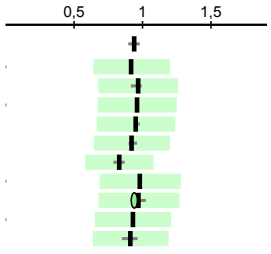
Sample A [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		285	0.341	0.398	0.420	0.435	0.940
2	13	6	0.380		0.399		0.420
2	28	63	0.341	0.394	0.417	0.438	0.940
2	43	4	0.380		0.408		0.415
4	4	26	0.410	0.420	0.426	0.433	0.450
4	30	118	0.370	0.402	0.420	0.440	0.532
4	38	16	0.346	0.357	0.400	0.425	0.440
4	40	5	0.387		0.391		0.412
4	43	16	0.370	0.380	0.400	0.418	0.440
4	126	7	0.386		0.410		0.440
4	140	8	0.364	0.368	0.424	0.460	0.468



Sample B [g/l]

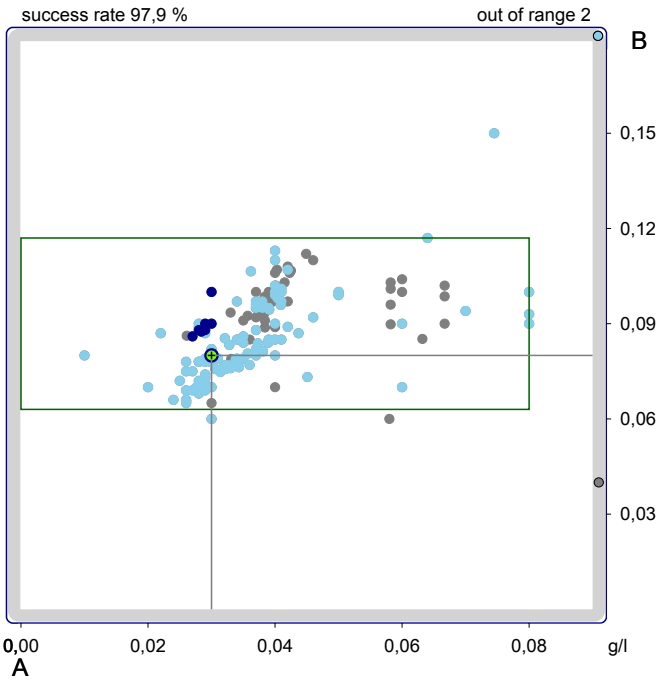
M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		285	0.380	0.896	0.939	0.980	1.07
2	13	6	0.862		0.917		0.970
2	28	63	0.380	0.915	0.968	0.994	1.07
2	43	4	0.940		0.960		1.00
4	4	26	0.930	0.940	0.950	0.980	1.02
4	30	118	0.818	0.898	0.920	0.960	1.04
4	38	16	0.760	0.787	0.830	0.870	0.898
4	40	5	0.927		0.980		1.04
4	43	16	0.920	0.929	0.970	1.02	1.04
4	126	7	0.884		0.930		1.04
4	140	8	0.813	0.849	0.911	0.964	0.999



The deviation of your results from the median of the corresponding sub-collective (kit) is:
A 0 %
B -3.0 %

Other kits (number):
1-28(1), 1-30(1), 2-30(1), 2-126(1), 2-140(1), 2-228(2), 3-40(1), 4-12(1), 4-28(1), 4-328(2), 5-04(1), 5-43(1), 6-30(1), 7-38(1),

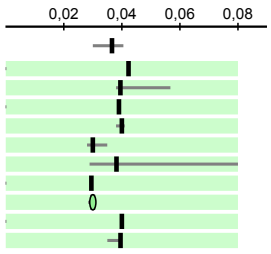
Analyte **C4-complement**
Method all methods



No of participants		286	
sample/unit	A	g/l	B
mean	0.036		0.087
standard deviation	0.010		0.012
coefficient of variation	28.0		14.7

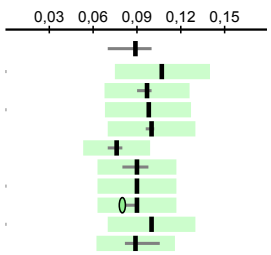
Sample A [g/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		286	0.010	0.030	0.036	0.040	0.274
2	13	7	0.040		0.042		0.046
2	28	62	0.030	0.038	0.039	0.056	0.100
2	43	3	0.026		0.039		0.060
4	4	25	0.037	0.038	0.040	0.041	0.070
4	30	118	0.010	0.028	0.030	0.035	0.074
4	38	16	0.020	0.028	0.038	0.080	0.080
4	40	6	0.028		0.029		0.050
4	43	17	0.027	0.028	0.030	0.030	0.030
4	126	7	0.036		0.040		0.060
4	140	8	0.035	0.035	0.039	0.040	0.040



Sample B [g/l]

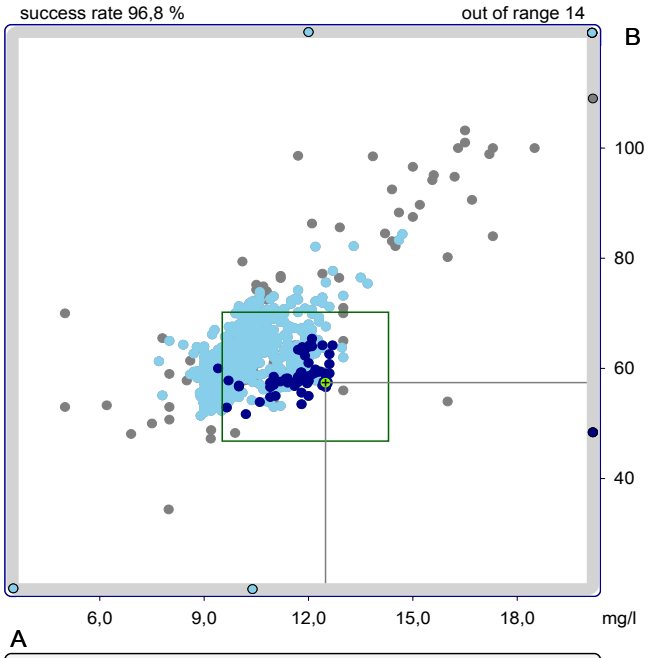
M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		286	0.040	0.070	0.089	0.100	0.208
2	13	7	0.103		0.107		0.112
2	28	62	0.040	0.090	0.096	0.100	0.108
2	43	3	0.086		0.098		0.100
4	4	25	0.094	0.096	0.100	0.102	0.113
4	30	118	0.060	0.070	0.076	0.080	0.150
4	38	16	0.070	0.080	0.090	0.097	0.100
4	40	6	0.087		0.090		0.099
4	43	17	0.080	0.080	0.090	0.090	0.100
4	126	7	0.090		0.100		0.110
4	140	8	0.080	0.081	0.089	0.106	0.110



The deviation of your results from the median of the corresponding sub-collective (kit) is:
A 0 %
B -11.11 %

Other kits (number):
1-28(1), 1-30(1), 2-04(1), 2-30(1), 2-126(1), 2-140(1), 2-228(2), 4-12(1), 4-28(1), 4-154(1), 4-328(2), 5-04(1), 5-43(1), 6-30(1), 7-38(1),

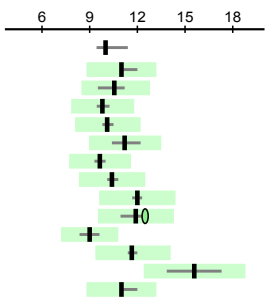
Analyte **CRP**
Method all methods



No of participants		1288	
sample/unit	A	mg/l	B
mean	10.3		61.9
standard deviation	1.22		6.37
coefficient of variation	11.8		10.3

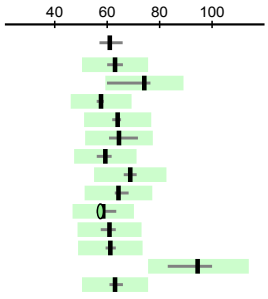
Sample A [mg/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		1288	2.20	9.43	10.0	11.4	1000
1	61	50	5.00	11.0	11.0	12.0	13.0
2	13	10	7.98	9.52	10.6	11.2	11.2
2	28	46	9.19	9.45	9.80	10.2	10.8
4	4	151	9.30	9.80	10.1	10.5	66.0
4	7	55	9.80	10.4	11.2	12.2	70.5
4	30	599	2.20	9.30	9.64	10.0	1000
4	38	94	9.70	10.1	10.4	10.8	13.0
4	40	20	10.0	11.7	12.0	12.3	14.6
4	43	64	9.40	10.9	11.9	12.5	57.6
4	126	21	7.70	8.36	9.00	9.61	13.0
4	328	43	10.4	11.4	11.7	12.0	12.4
5	44	24	10.1	13.9	15.6	17.3	20.7
5	61	17	10.0	11.0	11.0	12.0	12.0



Sample B [mg/l]

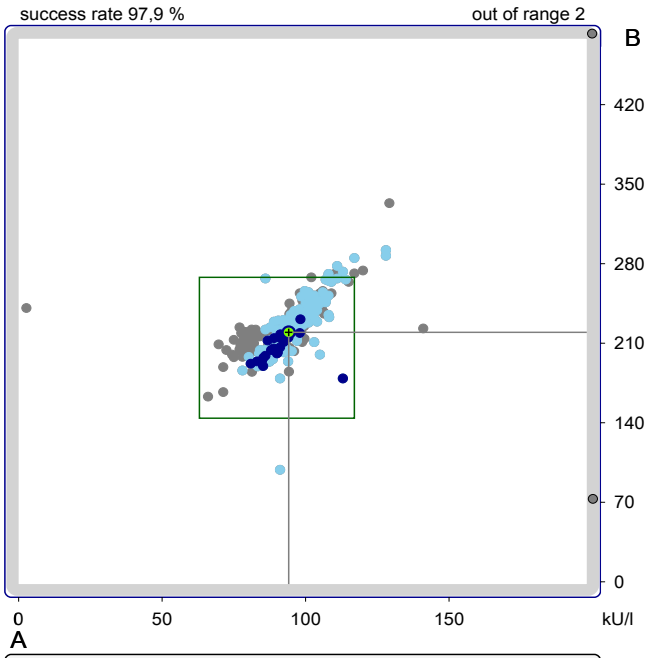
M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		1288	0.270	57.1	61.0	65.9	6210
1	61	50	56.0	60.0	63.0	66.0	70.0
2	13	10	34.4	59.9	74.2	76.5	76.8
2	28	46	47.3	56.0	57.7	58.7	63.0
4	4	151	10.0	61.9	64.0	65.3	74.2
4	7	55	13.0	60.7	64.5	71.7	84.4
4	30	599	1.70	56.0	59.3	61.7	6210
4	38	94	0.270	66.3	68.8	71.2	73.9
4	40	20	61.7	62.9	64.3	68.2	6060
4	43	64	12.4	56.0	58.5	63.5	65.4
4	126	21	55.1	57.6	60.9	63.3	65.0
4	328	43	56.8	59.4	61.2	63.3	65.7
5	44	24	79.4	83.1	94.5	100	109
5	61	17	60.0	60.9	63.0	66.0	68.0



The deviation of your results from the median of the corresponding sub-collective (kit) is:
A 4.9 %
B -1.8 %

Other kits (number):
1-30(3), 1-99(1), 1-159(1), 2-04(1), 2-30(1), 2-43(2), 2-228(1), 4-03(5), 4-507(7), 4-08(1), 4-12(6), 4-13(2), 4-32(1), 4-33(5), 4-36(2), 4-61(1), 4-99(3), 4-128(2), 4-140(3), 4-364(2), 5-34(1), 5-89(1), 5-99(1), 5-159(3), 6-04(1), 6-08(8), 6-30(1), 6-33(1), 6-61(1), 6-89(8), 6-99(1), 6-126(1), 6-130(8), 7-30(2), 7-33(1), 7-38(1), 7-43(1), 7-99(2), 7-151(1),

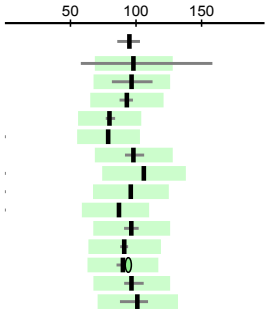
Analyte **IgE**
Method all methods



No of participants		342	
sample/unit	A	kU/l	B
mean	94.3		224
standard deviation	10.8		21.6
coefficient of variation	11.5		9.60

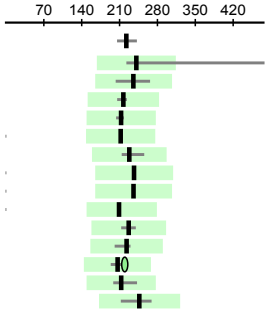
Sample A [kU/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		342	2.70	85.7	95.0	103	237
1	30	9	2.70	57.9	98.0	158	237
1	72	14	72.4	81.6	96.6	113	117
2	28	15	86.0	87.4	93.0	97.7	98.8
4	4	30	74.4	76.9	79.8	84.1	129
4	12	6	69.7		78.8		200
4	30	11	90.0	91.6	98.0	106	120
4	38	6	78.0		106		141
BeckCou2		7	94.4		96.0		100
4	43	3	86.8		87.0		88.2
5	30	147	78.0	90.9	96.4	102	114
5	40	9	86.0	87.9	91.0	94.0	95.4
5	43	23	80.9	85.1	90.0	94.7	113
5	44	16	84.1	91.0	96.5	106	111
5	72	36	80.2	87.7	101	109	128



Sample B [kU/l]

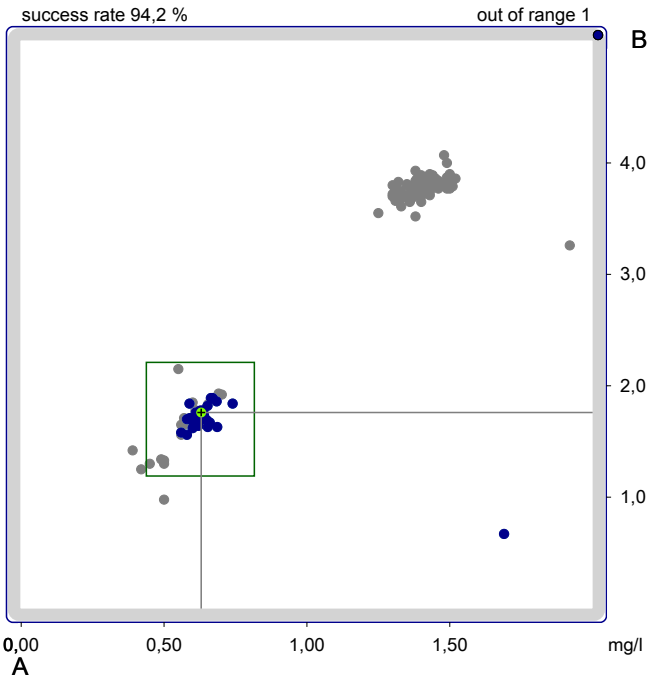
M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		342	73.0	205	223	242	847
1	30	9	219	223	241	492	847
1	72	14	185	203	236	266	272
2	28	15	196	206	217	223	230
4	4	30	198	204	213	218	333
4	12	6	73.0		212		220
4	30	11	209	214	228	256	274
4	38	6	198		237		245
BeckCou2		7	203		236		248
4	43	3	198		209		209
5	30	147	186	213	227	240	266
5	40	9	199	201	223	230	231
5	43	23	179	194	206	219	231
5	44	16	179	198	213	242	278
5	72	36	98.6	212	247	269	292



The deviation of your results from the median of the corresponding sub-collective (kit) is:
A 4.5 %
B 6.3 %

Other kits (number):
1-04(1), 1-44(1), 2-13(1), 3-30(1), 4-13(2), 4-16(1), 4-140(1), 5-23(1), 7-99(1),

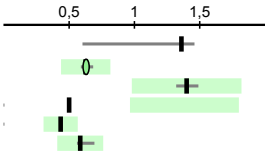
Analyte **sol.Transfe.Receptor**
Method all methods



No of participants		140	
sample/unit	A	mg/l	B
mean	1.13		3.04
standard deviation	0.394		1.03
coefficient of variation	34.8		34.0

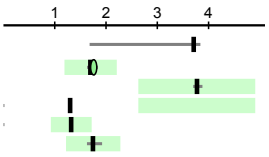
Sample A [mg/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		140	0.390	0.602	1.36	1.46	610
2	28	30	0.560	0.590	0.627	0.684	610
4	30	91	1.25	1.32	1.40	1.49	1.92
4	43	5	0.500		0.500		0.550
4	154	4	0.390		0.435		0.490
5	13	10	0.560	0.560	0.585	0.694	0.702



Sample B [mg/l]

M	Kit	N	Min	16.P	50.P	84.P	Max
Alle		140	0.670	1.68	3.71	3.84	1680
2	28	30	0.670	1.63	1.70	1.84	1680
4	30	91	3.26	3.70	3.77	3.88	4.07
4	43	5	0.978		1.30		2.15
4	154	4	1.25		1.32		1.42
5	13	10	1.56	1.63	1.75	1.92	1.93



The deviation of your results from the median of the corresponding sub-collective (kit) is:
A 0.5 %
B 3.8 %

Other kits (number):