0003015050122c501V8.0 TRSF2 Tina-guant Transferrin ver.2

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Order information

REF	CONTENT		Analyzer(s) on which cobas c pack(s) can be used
03015050 122	Tina-quant Transferrin ver.2/ 100 tests	System-ID 07 6567 8	Roche/Hitachi cobas c 311, cobas c 501/502
11355279 216	Calibrator f.a.s. Proteins (5 x 1 mL)	Code 656	
11355279 160	Calibrator f.a.s. Proteins (5 x 1 mL, for USA)	Code 656	
10557897 122	Precinorm Protein (3 x 1 mL)	Code 302	
10557897 160	Precinorm Protein (3 x 1 mL, for USA)	Code 302	
11333127 122	Precipath Protein (3 x 1 mL)	Code 303	
11333127 160	Precipath Protein (3 x 1 mL, for USA)	Code 303	
10171743 122	Precinorm U (20 x 5 mL)	Code 300	
10171735 122	Precinorm U (4 x 5 mL)	Code 300	
05117003 190	PreciControl ClinChem Multi 1 (20 x 5 mL)	Code 391	
05947626 190	PreciControl ClinChem Multi 1 (4 x 5 mL)	Code 391	
05947626 160	PreciControl ClinChem Multi 1 (4 x 5 mL, for USA)	Code 391	
05117216 190	PreciControl ClinChem Multi 2 (20 x 5 mL)	Code 392	
05947774 190	PreciControl ClinChem Multi 2 (4 x 5 mL)	Code 392	
05947774 160	PreciControl ClinChem Multi 2 (4 x 5 mL, for USA)	Code 392	
04489357 190	Diluent NaCl 9 % (50 mL)	System-ID 07 6869 3	

English

System information For cobas c 311/501 analyzers: TRSF2: ACN 187 For cobas c 502 analyzer: TRSF2: ACN 8187

Intended use

In vitro test for the quantitative determination of transferrin in human serum and plasma on Roche/Hitachi ${\bf cobas}\ {\bf c}$ systems.

Summary^{1,2,3,4,5}

Transferrin is a glycoprotein with a molecular weight of 79570 daltons. It consists of a polypeptide strand with two N-glycosidically linked oligosaccharide chains and exists in numerous isoforms. The rate of synthesis in the liver can be altered in accordance with the body's iron requirements and iron reserves.

Transferrin is the iron transport protein in serum. In cases of iron deficiency, the degree of transferrin saturation appears to be an extremely sensitive indicator of functional iron depletion. The ferritin levels are depressed when there is a deficiency of storage iron. In sideropenia, an iron deficiency can be excluded if the serum transferrin concentration is low, as in inflammations or - less commonly - in cases of ascorbic acid deficiency. In screening for hereditary hemochromatosis, transferrin saturation provides a better indication of the homozygous genotype than does ferritin. The treatment of anemia with erythropoietin in patients with renal failure is only effective when sufficient depot iron is present. The best monitoring procedure is to determine transferrin saturation during therapy. Transferrin saturation in conjunction with ferritin gives a conclusive prediction of the exclusion of iron overloading in patients with chronic liver disease.

A variety of methods are available for determining transferrin including radial immunodiffusion, nephelometry and turbidimetry. The Roche transferrin assay is based on the immunological agglutination principle.

Test principle

Immunoturbidimetric assay.^{6,7,8}

Human transferrin forms a precipitate with a specific antiserum which is determined turbidimetrically.

Reagents - working solutions

- R1 Phosphate buffer: 55 mmol/L, pH 7.2; NaCl: 25 mmol/L; polyethylene glycol: 5 %; preservative
- R2 Anti-human transferrin antibodies (rabbit): dependent on titer; NaCI: 100 mmol/L; preservative

R1 is in position B and R2 is in position C.

Precautions and warnings

For in vitro diagnostic use.

Exercise the normal precautions required for handling all laboratory reagents. Disposal of all waste material should be in accordance with local guidelines. Safety data sheet available for professional user on request. For USA: Caution: Federal law restricts this device to sale by or on the order of a physician. Reagent handling

Ready for use

Storage and stability

TRSF2

See expiration date on cobas c pack label.
8 weeks
See expiration date on cobas c pack label.

On-board in use and refrigerated on the analyzer: 12 weeks

Specimen collection and preparation

For specimen collection and preparation only use suitable tubes or collection containers.

Only the specimens listed below were tested and found acceptable. Serum.

Plasma: Li-heparin plasma. Do not use EDTA or citrate plasma.

The sample types listed were tested with a selection of sample collection tubes that were commercially available at the time of testing, i.e. not all available tubes of all manufacturers were tested. Sample collection systems from various manufacturers may contain differing materials which could affect the test results in some cases. When processing samples in primary tubes (sample collection systems), follow the instructions of the tube manufacturer.

Centrifuge samples containing precipitates before performing the assay.

Stability:9

8 days at 15-25 °C 8 days at 2-8 °C 6 months at (-15)-(-25) °C



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Materials provided						Sample	e	Diluent (NaCl)
See "Reagents – working solutions" section for reagents.			Normal	12.5 µL	9 µL		180 µL	
Materials required (but not	provided)			Decreased	12.5 µL	5 µL		152 µL
 See "Order information" section 			Increased	12.5 µL	18 µL		180 µL	
 General laboratory equipment 			Calibration					
Assay	the energy falles	u the alive	ations sives in this	Calibrators	S1- H O			
document for the analyzer co manual for analyzer-specific a	For optimum performance of the assay follow the directions given in this document for the analyzer concerned. Refer to the appropriate operator's manual for analyzer-specific assay instructions				S2-S6: C.f.a	a.s. Proteir	าร	
The performance of application and must be defined by the u	ons not validate ser.	ed by Rocl	he is not warranted		Multiply the calibrator va	lot-specific alue by the	c C.f.a.s. factors	. Proteins below to
Application for serum and p	olasma				determine t	he standar calibration	d concei curve:	ntrations for
cobas c 311 test definition					S2·0 120	callorallori	S5: 1.0	0
Assay type	2-Point End				S3: 0.239		S6: 1.9	1
Reaction time / Assay points	10 / 6-25			S4: 0.478				
Wavelength (sub/main)	700 / 505 nm			Calibration mode	BCM2			
Reaction direction	Increase			Calibration frequency	Full calibrat	ion		
Units	g/L (µmol/L, n	ng/dL)		Calibration nequency	after reade	ent lot char	nae	
Reagent pipetting		Diluent ((H ₂ O)		 as require 	d following	quality	control
R1	140 µL	_			procedures			
R2	30 µL	_		Calibration interval may be	e extended base	ed on acce	ptable ve	erification of
Sample volumes	Sample	S	ample dilution	Traceability: This method	ry. has been stand:	ardized ana	ainst the	reference
		Sample	Diluent (NaCl)	preparation of the IRMM (I	Institute for Refe	erence Mat	erials an	nd
Normal	12.5 uL	9 uL	180 uL	Measurements) BCR470/0 Proteins in Human Serum	CRM470 (RPPH	IS - Refere	nce Prep	paration for
Decreased	12.5 uL	5 uL	152 uL					
Increased	12.5 µL	9 µL	180 µL	For quality control, use consection.	ntrol materials a	s listed in t	the "Orde	er information"
cobas c 501 test definition				In addition, other suitable	control material	can be use	ed.	
Assay type	2-Point End			The control intervals and li	imits should be	adapted to	each lat	boratory's
Reaction time / Assay points	10 / 10-36			individual requirements. V	alues obtained sould establish co	should fall v prrective me	within the	e defined to be taken if
Wavelength (sub/main)	700/505 nm			values fall outside the defined limits.				
Reaction direction	Increase			Follow the applicable gove	ernment regulati	ons and lo	cal guide	elines for
Units	g/L (µmol/L, mg/dL)		quality control.					
Reagent pipetting	0 (1)	Diluent ((H ₂ O)	Calculation Boche/Hitachi cohas c sv	stems automatic	cally calcul	ate the a	analyte
R1	140 uL	_	(2-)	concentration of each sample.		inaryto		
R2	30 ul	_		Conversion factors:	ma/dL x 0 0	l/n = 1(a/L x 12	$P 6 = \mu mol/l$
Sample volumes	Sample	S	ample dilution		n/l x 100 −	ma/dl		$x = 0.0796 - \alpha/l$
Campie Volamee	Campio	Samnla	Diluent (NaCl)		9/EX 100 -	ilig/u⊑	µmoi/ E	x 0.0750 – g/L
Normal	12.5 ul		180 ul	Criterion: Becovery within	e ⊥10 % of initial	valuo at a	tranefor	rin
Decreased	12.5 µ⊑ 12.5 µl	5 μ⊑ 5 μl	150 µL	concentration of 2 g/L (25.	$\frac{1}{2} \mu mol/L, 200 m$	ng/dL).	liansien	
Increased	12.5 μL 12.5 μL	5 μL 9 μL	132 μL 180 μL	Icterus: ¹¹ No significant interference up to an I index of 60 for conjugated and unconjugated bilirubin (approximate conjugated and unconjugated				
cobas c 502 test definition				Hemolysis: ¹¹ No significan	∠o µmoi/∟ or 60 it interference ui	rng/ɑ∟). o to an H ir	ndex of 1	000
Assay type	2-Point End			(approximate hemoglobin	concentration: 6	621 µmol/L	or 1000	mg/dL).
Reaction time / Assay points	10 / 10-36			Lipemia (Intralipid): ¹¹ No s	ignificant interfe	rence up to	o an L in	dex of 500.
Wavelength (sub/main)	700/505 nm			triglycerides concentration			sponus u	o turbiuity) anu
Reaction direction	Increase			Rheumatoid factors up to	1200 IU/mL do r	not interfer	e.	
Units	g/L (µmol/L, n	ng/dL)		High dose hook-effect: No	false result occ	urs up to a	transfer	rrin
Reagent pipetting		Diluent ((H ₂ O)	Drugs: No interference wa	s found at thera	nuy/uL).	contratio	ne usina
R1	140 µL	-		common drug panels. ^{12,13}				na uality
R2	30 µL	_		In very rare cases, gammopathy, in particular type IgM (Waldenström's		denström's		
Sample volumes	Sample	S	ample dilution	macroglobulinemia), may cause unreliable results. ¹⁴				

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For diagnostic purposes, the results should always be assessed in conjunction with the patient's medical history, clinical examination and other findings

ACTION REQUIRED

Special Wash Programming: The use of special wash steps is mandatory when certain test combinations are run together on Roche/Hitachi cobas c systems. The latest version of the carry-over evasion list can be found with the NaOHD-SMS-SmpCln1+2-SCCS Method Sheets. For further instructions refer to the operator's manual. cobas c 502 analyzer: All special wash programming necessary for avoiding carry-over is available via the cobas link, manual input is not required.

Where required, special wash/carry-over evasion programming must be implemented prior to reporting results with this test.

Limits and ranges

Measuring range

0.1-5.2 g/L (1.26-65.5 µmol/L, 10-520 mg/dL)

Determine samples having higher concentrations via the rerun function. Dilution of samples via the rerun function is a 1:1.5 dilution. Results from samples diluted using the rerun function are automatically multiplied by a factor of 1.5.

Lower limits of measurement

Lower detection limit of the test

0.1 g/L (1.26 µmol/L, 10 mg/dL)

The lower detection limit represents the lowest measurable analyte level that can be distinguished from zero. It is calculated as the value lying 3 standard deviations above that of the lowest standard (standard 1 + 3 SD, repeatability, n = 21).

Expected values¹⁵ I

2.0-3.6 g/L (25.2-45.4 µmol/L; 200-360 mg/dL)

Each laboratory should investigate the transferability of the expected values to its own patient population and if necessary determine its own reference ranges

Specific performance data

Representative performance data on the analyzers are given below. Results obtained in individual laboratories may differ.

Precision

Precision was determined using human samples and controls in an internal protocol with repeatability (n = 21) and intermediate precision (3 aliquots per run, 1 run per day, 21 days). The following results were obtained:

Repeatability	Mean	SD	CV
	g/L (µmol/L, mg/dL)	g/L (µmol/L, mg/dL)	%
Precinorm Protein	2.62 (33.0, 262)	0.03 (0.4, 3)	1.2
Precipath Protein	4.01 (50.5, 401)	0.07 (0.9, 7)	1.7
Human serum 1	1.27 (16.0, 127)	0.02 (0.3, 2)	1.2
Human serum 2	2.63 (33.1, 263)	0.04 (0.5, 4)	1.5
Intermediate precision	Mean	SD	CV
Intermediate precision	Mean g/L (µmol/L, mg/dL)	SD g/L (µmol/L, mg/dL)	CV %
Intermediate precision Precinorm Protein	Mean g/L (μmol/L, mg/dL) 2.55 (32.1, 255)	SD g/L (µmol/L, mg/dL) 0.07 (0.9, 7)	CV % 2.9
Intermediate precision Precinorm Protein Precipath Protein	Mean g/L (μmol/L, mg/dL) 2.55 (32.1, 255) 3.95 (49.8, 395)	SD g/L (μmol/L, mg/dL) 0.07 (0.9, 7) 0.13 (1.6, 13)	CV % 2.9 3.2
Intermediate precision Precinorm Protein Precipath Protein Human serum 3	Mean g/L (µmol/L, mg/dL) 2.55 (32.1, 255) 3.95 (49.8, 395) 2.14 (27.0, 214)	SD g/L (µmol/L, mg/dL) 0.07 (0.9, 7) 0.13 (1.6, 13) 0.06 (0.8, 6)	CV % 2.9 3.2 2.6
Intermediate precision Precinorm Protein Precipath Protein Human serum 3 Human serum 4	Mean g/L (μmol/L, mg/dL) 2.55 (32.1, 255) 3.95 (49.8, 395) 2.14 (27.0, 214) 2.96 (37.3, 296)	SD g/L (μmol/L, mg/dL) 0.07 (0.9, 7) 0.13 (1.6, 13) 0.06 (0.8, 6) 0.08 (1.0, 8)	CV % 2.9 3.2 2.6 2.6

Method comparison

Transferrin values for human serum and plasma samples obtained on a Roche/Hitachi cobas c 501 analyzer (y) were compared with those determined using the corresponding reagent on a Roche/Hitachi 917 analyzer (x)

Sample size (n) = 117

Passing/Bablok¹⁶

Linear regression

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y = 1.030x - 0.068 g/L
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The sample concentrations were between 1.05 and 4.51 g/L (13.2 and 56.7 µmol/L, 105 and 450 mg/dL).

References

T = 0.964

1 Wick M, Pinggera W, Lehmann P, eds. Iron Metabolism, Diagnosis and Therapy of Anemias. 5th ed. Vienna/New York: Springer-Verlag 1999.

y = 1.018x - 0.044 g/L

r = 0.998

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- 16 Bablok W, Passing H, Bender R, et al. A general regression procedure for method transformation. Application of linear regression procedures for method comparison studies in clinical chemistry, Part III. J Clin Chem Clin Biochem 1988 Nov;26(11):783-790.

A point (period/stop) is always used in this Method Sheet as the decimal separator to mark the border between the integral and the fractional parts of a decimal numeral. Separators for thousands are not used.

Symbols

Roche Diagnostics uses the following symbols and signs in addition to those listed in the ISO 15223-1 standard (for USA: see https://usdiagnostics.roche.com for definition of symbols used):

CONTENT	
GTIN	

Contents of kit Volume after reconstitution or mixing Global Trade Item Number





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