

# **Transferrin FS\***

Diagnostic reagent for quantitative in vitro determination of transferrin (Trf) in serum or plasma on BioMajesty JCA-BM6010/C

# Order Information

Cat. No. 1 7252 99 10 964 R1: 6 x 100 tests R2: 6 x 100 tests

# Method

Immunoturbidimetric test

# Principle

Determination of the transferrin concentration by photometric measurement of antigen-antibody-reaction among antibodies to transferrin and transferrin present in the sample.

# Reagents

#### **Components and Concentrations**

R1:	TRIS	pH 7.5	100 mmol/L
	NaCl		180 mmol/L
R2:	TRIS	pH 8.0	100 mmol/L
	NaCl		300 mmol/L
	Anti-human T	ransferrin antibody (goat)	< 1%

# Storage Instructions and Reagent Stability

The reagents are stable up to the end of the indicated month of expiry, if stored at  $2 - 8^{\circ}$ C, protected from light and contamination is avoided. Do not freeze the reagents!

#### Warnings and Precautions

- The reagents contain sodium azide (0.95 g/L) as preservative. Do not swallow! Avoid contact with skin and mucous membranes!
- Reagent 2 contains animal material. Handle the product as potentially infectious according to universal precautions and good clinical laboratory practices.
- In very rare cases, samples of patients with gammopathy might give falsified results [6].
- 4. Please refer to the safety data sheets and take the necessary precautions for the use of laboratory reagents. For diagnostic purposes, the results should always be assessed with the patient's medical history, clinical examinations and other findings.
- 5. For professional use only!

#### Waste Management

Please refer to local legal requirements.

#### **Reagent Preparation**

The reagents are ready to use. The bottles are placed directly into the reagent rotor.

#### Specimen

Serum, heparin plasma or EDTA plasma Stability [1]:

8 days	at	20 – 25°C
8 days	at	4 – 8°C
6 months	at	–20°C
Freeze only		

Discard contaminated specimens.

# **Calibrators and Controls**

For the calibration the DiaSys TruCal Protein calibrator set is recommended. The assigned values of the calibrators have been made traceable to the reference material ERM<sup>®</sup>-DA470k/IFCC. For internal quality control a DiaSys TruLab Protein control should be assayed. Each laboratory should establish corrective action in case of deviations in control recovery.

	Cat. No.	Kit	size
TruCal Protein (5 levels)	5 9200 99 10 039	5 x	1 mL
TruLab Protein Level 1	5 9500 99 10 046	3 x	1 mL
TruLab Protein Level 2	5 9510 99 10 046	3 x	1 mL

#### Reagent information

# **Performance Characteristics**

Measuring range up to 7.7 g/L (97.0 µmol/L) transferrin, at least up to			
the concentration of the highest calibrator			
(in case of higher concentrations re-measure samples after manual			
dilution with NaCl solution (9 g/L) or use rerun function).			
Limit of detection** 0.01 g/L (0.126 µmol/L) transferrin			
No prozone effect up to 19.9 g/L (251 µmol/L) transferrin			
On-board stability 6 weeks			
Calibration stability 6 weeks			

Interferences < 10% by
Conjugated Bilirubin up to 60 mg/dL
Unconjugated Bilirubin up to 60 mg/dL
Hemoglobin up to 800 mg/dL
Lipemia (triglycerides) up to 2000 mg/dL
For further information on interfering substances refer to Young DS [5].

Precision			
Within run (n=20)	Sample 1	Sample 2	Sample 3
Mean [g/L]	1.65	2.65	4.11
Mean [µmol/L]	20.8	33.4	51.7
Coefficient of variance [%]	1.69	1.50	2.15
Between run (n=20)	Sample 1	Sample 2	Sample 3
Mean [g/L]	1.63	2.46	3.14
Mean [µmol/L]	20.5	31.0	39.6
Coefficient of variance [%]	2.24	3.45	2.31

# Method comparison (n=100

Method comparison (n=100)		
Test x	Competitor Transferrin	
Test y	DiaSys Transferrin FS	
Slope	1.02	
Intercept	-0.012 g/L (-0.151 µmol/L)	
Coefficient of correlation	0.999	

\*\* lowest measurable concentration which can be distinguished from zero mean + 3 SD (n=20) of an analyte free specimen

### **Conversion factor**

Transferrin [mg/dL] x 0.126 = Transferrin [µmol/L]

#### **Reference Range** [2]

200 – 360 mg/dL (25.2 – 45.4 µmol/L)

Each laboratory should check if the reference ranges are transferable to its own patient population and determine own reference ranges if necessary.

# Literature

- Guder WG, Zawta B et al. The Quality of Diagnostic Samples. 1<sup>st</sup> ed. Darmstadt: GIT Verlag; 2001; p. 22-3.
- Dati F, Schumann G, Thomas L, Aguzzi F, Baudner S, Bienvenu J et al. Consensus of a group of professional societies and diagnostic companies on guidelines for interim reference ranges for 14 proteins in serum based on the standardization against the IFCC/BCR/CAP reference material (CRM 470). Eur J Clin Chem Clin Biochem 1996; 34: 517-20.
- Wick M, Pingerra W, Lehmann P. Iron metabolism: diagnosis and therapy of anemias. 3<sup>rd</sup> ed. Vienna, New York: Springer Verlag, 1996.
- Fairbanks VF, Klee GG. Biochemical aspects of hematology. In: Burtis CA, Ashwood ER, editors. Tietz Textbook of Clinical Chemistry. 3<sup>rd</sup> ed. Philadelphia: W.B Saunders Company; 1999. p. 1642-1710.
- Young DS. Effects of Drugs on Clinical Laboratory Tests. 5th ed. Volume 1 and 2. Washington, DC: The American Association for Clinical Chemistry Press 2000.
- Bakker AJ, Mücke M. Gammopathy interference in clinical chemistry assays: mechanisms, detection and prevention. ClinChemLabMed 2007;45(9):1240-1243.

# Manufacturer



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# **Transferrin FS**

# Chemistry code 10 725

# Application for serum and plasma samples

This application was set up and evaluated by DiaSys. It is based on the standard equipment at that time and does not apply to any equipment modifications undertaken by unqualified personnel.

Analytical Conditions		
R1 volume	125	
R2e volume	0	
R2 volume	25	
R1 diluent vol	0	
R2e diluent vol	0	
R2 diluent vol	0	
Sample vol (S)	1	
Sample vol (U)	1	
Reagent 1 mix	weak	
Reagent 2e mix	weak	
Reagent 2 mix	weak	
Reaction time	10	

Sub-analy. Conditions		
Name	TRF	
Digits	2	
M-wave L.	571	
S-wave.L	****	
Analy.mthd.	EPA	
Calc.mthd.	MSTD	
Qualit. judge	No	

Analysis Test Condition Setting (M)			
Sample Type	Serum	Urine	
Reac. sample vol.	1	1	
Diluent method	No dil	No dil	
Undil. sample vol.	0	0	
Diluent volume	0	0	
Diluent position	0	0	

Endpoint Method		
Re.absorb (u)	9.999	
Re.absorb (d)	-9.999	

Calculation Method Setting		
M-DET.P.I	0	
M-DET.P.m	41	
M-DET.P.n	42	
S-DET.P.p	17	
S-DET.P.r	18	
Check D.P.I.	0	
Limit value	0.003	
Variance	10	
Reac.type	Inc	

Reaction Rate Method					
Cycle	2				
Factor	2				
E2 corre	Not do				
Blank (u)	9.999				
Blank (d)	-9.999				
Sample (u)	9.999				
Sample (d)	-9.999				

Prozone					
Prozone form	No				
Prozone limit	9.999				
Prozone judge	Upper limit				
Judge limit	9.999				
M-DET.P.m	0				
M-DET.P.n	0				
S-DET.P.p	0				
S-DET.P.r	0				

MULTI-STD Setting											
Formula	Logit Log	2 Axis Co	onv No	conv							
Blank	Blank is	0 Points	ints 6								
	FV	Reac.	Dil.	Dil. smp.	Diluent	Diluent	STD H	STD L			
		smp. vol.	method	vol.	vol.	pos.					
BLK	#	1	No dil	0	0	0	9.999	-9.999			
1	#	1	No dil	0	0	0	9.999	-9.999			
2	#	1	No dil	0	0	0	9.999	-9.999			
3	#	1	No dil	0	0	0	9.999	-9.999			
4	#	1	No dil	0	0	0	9.999	-9.999			
5	#	1	No dil	0	0	0	9.999	-9.999			

# entered by user

MUL TLSTD Setting