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W.L. van Noort, G. de Jong, H.G. van Eijk

Institutions: Erasmus University Rotterdam

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Purification of Isotransferrins by Concanavalin A Sepharose Chromatography and Preparative Isoelectric Focusing

By *W. L. van Noort, G. de Jong and H. G. van Eijk*

Department of Chemical Pathology, Erasmus University Rotterdam, Rotterdam, The Netherlands

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Summary: 1. From pooled serum containing genetically homogeneous transferrin C₁, transferrin was purified and separated in three fractions (tri-tri-, bi-tri- and bi-bi-antennary transferrin C₁), using Concanavalin A-Sepharose.

2. Each of these fractions was separated into its sialic acid-dependent subfractions by preparative isoelectric focusing. Sixteen iso-transferrin C₁ fractions were obtained, which differed in their degree of glycan branching and/or their sialic acid content.

3. Preliminary carbohydrate analyses suggest that in some iso-transferrins the N-acetylglucosamine and the galactose content is lower than expected.

Introduction

The main function of transferrin is the transport of iron between the sites of absorption and storage and the sites of utilization such as myoglobin and haemoglobin synthesis (1–5).

Transferrin is an *M_r* 80 000 monomeric glycoprotein containing two branched oligosaccharide chains, N-glycosidically attached to asparagine residues in the carboxyl-terminal domain (1–5).

A series of papers on the composition analysis of transferrin glycans has been published in the past fifteen years (6–14). Recently we published the sugar analyses of fractions of genetically homogeneous transferrin C₁, obtained by preparative isoelectric focusing in immobilized pH gradients, which produces fractions containing different amounts of sialic acid (15). These methods were optimized for the preparation of subfractions. Separation on Concanavalin A Sepharose (16) resulted in fractions with glycans carrying the same number of antennae.

Using preparative isoelectric focusing in ultra-thin Immobililine gels (15) each tri-tri-, bi-tri- and bi-bi-antennary Fe₂ transferrin fraction was fractionated into 5–6

fractions that differed in their number of sialic acid residues. Thus, sixteen different iso-transferrins were available for study.

Several authors have recently remarked on the desirability of analysing thoroughly purified fractions (17–21), for the investigation of different pathologies, including transferrin analysis in alcoholism (20, 21), and the analysis of other glycoproteins, e.g. in cancer (18, 19).

Materials and Methods

Preparation of transferrin subfractions

Identification of serum containing transferrin C₁

Sera were obtained from adult healthy volunteers and checked for C₁ transferrin genetic homogeneity as follows:

Using Immobililine DryPlate pH 4–7 (Pharmacia) a part of the gel was cut out to the same size as Phast Gel, to obtain a pH-gradient between the electrodes of 5.0–6.0. This piece of DryPlate was rehydrated by overlaying it with 1 ml 170 g/l glycerol for 2–3 hours. Serum (50 µl) was mixed with 2 µl 0.5 mol/l NaHCO₃ and 2 µl 10 mmol/l Fe(III)citrate.

Using PhastSystem (Pharmacia), the Immobililine gel was run with 0.3 µl of twelve different sera, using the program:

Sample application down at			x.2,	0 Vh.
Sample application up at			x.3,	0 Vh.
Extra alarm at			x.1,	350 Vh.
Sep. x.1: 2000 V	2.0 mA	2.0 W	15 °C,	485 Vh.
Sep. x.2: 200 V	0.3 mA	0.3 W	15 °C,	15 Vh.
Sep. x.3: 2000 V	2.0 mA	2.0 W	15 °C,	1300 Vh.

Prefocusing was necessary to obtain sharp bands without smears.

Paraffin oil (50 µl) was applied between the cooling plate and gel backing.

At the end of the run the gel was incubated with 100 µl anti-transferrin solution (Dako-immunoglobulins, Denmark) for 15–30 minutes at room temperature.

The gel was washed overnight in saline (with several changes), incubated with 200 g/l trichloroacetic acid and stained with Coomassie Brilliant Blue R-250 1 g/l at 50 °C for 10 minutes as described in the PhastSystem users Manual (Pharmacia), and destained at 20 °C.

As transferrin C₁ is the commonly occurring variant (70–80%), other variants are easily recognized by comparing 12 sera on one gel, and can then be excluded from the purification.

Isolation of transferrin from serum

The transferrin C₁ (all transferrin used in this paper is from the same source) was isolated from the serum as described earlier (22) by immuno affinity chromatography.

A specific anti human transferrin Sepharose column was used. Attention must be paid to a complete removal of all other serum proteins that elute with the first buffer overnight.

Separation of transferrin fractions differing in their degree of glycan branching

The transferrin C₁ obtained from step 2 was separated in a fraction with two triantennary glycans (further called tri-tri-transferrin C₁), a fraction with one bi-antennary and one tri-antennary glycan (further called bi-tri-transferrin C₁) and a fraction with two bi-antennary glycans (further called bi-bi-transferrin C₁) by Concanavalin A Sepharose chromatography as described by *Hatton & Berry* (16).

Separation of transferrin subfractions

Each fraction was further separated into fractions with different sialic acid contents, by preparative isoelectric focusing, using Immobiline DryPlate pH 5.0–6.0 on a Multiphor (Pharmacia, Upsala, Sweden) as described previously (15). This method depends on the stable pI-gradient, the high loading capacity and the high resolving power of Immobiline gel. Large amounts of protein become exuded from the gel as concentrated droplets of protein solution. Small amounts of protein can be obtained by diffusion after cutting out pieces of gel containing the appropriate bands with scissors.

Analyses

Densitometry

Densitometric scans of purified iso-transferrin fractions were performed using Ultrosan XL (He/Nelaser, λ = 633 nm, Pharmacia, Sweden).

Check of subfraction homogeneity

The sialic acid-dependent subfractions were checked for homogeneity by isoelectric focusing using PhastGels IEF pH 4–6.5, as recently described (23).

Desialylation

Neuraminidase from *Cl. perfringens*, type V from Sigma was used: 4 Units were dissolved in 1 ml 0.1 mol/l sodium acetate, pH 6. An aliquot (50 µl) of this solution was mixed with 5 µl transferrin preparation (0.5–1 g/l) and incubated for 72 hours at 4 °C.

The pH was then raised to 8 using 2 µl 0.5 mol/l NaHCO₃, and 1 µl 10 mmol/l Fe(III)citrate was added for iron saturation.

After two hours, the samples were analysed using PhastGel IEF 4–6.5 (see above).

Analysis of N-acetylglucosamine and amino acids

a) Hydrolysis and sample preparation

An adaptation of the previously described method employing 3 mol/l *p*-toluene sulphonic acid for hydrolysis was used (24).

Protein solution (50 µl containing 0.5–1 g/l protein and 10 mmol/l NaHCO₃) was mixed with 150 µl of a 4 mol/l *p*-toluene sulphonic acid solution. The mixture was degassed at < 1 mm Hg, sealed and incubated at 110 °C for 24 hours.

The hydrolysate was titrated with 1 mol/l NaOH to pH 2.00 ± 0.02.

After filtration over a 0.45 µm disk filter (Millipore, Japan) 25 or 50 µl were analysed using an ALPHA PLUS 4151 amino acid analyser (Pharmacia/LKB/Biochrom-Cambridge, England).

b) Analysis

The column eluate (21 ml/h) was mixed with *o*-phthalaldehyde reagent (8 ml/h), prepared according to the apparatus Handbook, except that the volume of β-mercaptoethanol was doubled (all chemicals from Merck, Germany).

The T-piece in which eluate and reagent are mixed, was connected directly with a fluorimeter (F1000, Hitachi). The excitation wavelength was set at 340 nm and the emission wavelength at 455 nm. The fluorimeter outlet was connected with a standard photometer (coil temperature: 50 °C) to generate some back-pressure but the back-pressure valve spring was removed to avoid the back-pressure becoming too high. The fluorimeter signal was received and stored by a peak-integrator (Chromato-Integrator D2000, Hitachi).

The analysis of hexoses

We have already described this method extensively (25). After hydrolysis with 2 mol/l trifluoroacetic acid for 16 hours at 100 °C and < 1 mm Hg, analyses are performed in the subnanomole range using an adapted amino acid analyser.

Partition chromatography was used, and the eluate was mixed with tetrazolium blue chloride (Fluka Chemie, Belgium – cat. No. 88190) in 0.18 mol/l NaOH. The reaction product was detected at 570 nm and the signal was stored by the same integrator as used for the amino acid analyses.

Results

Purification of sixteen different transferrin C₁ fractions

Transferrin subfractions were obtained according to the scheme of figure 1.

Figure 2 shows a typical example of the genetic homogeneity of twelve normal sera of healthy adult volun-

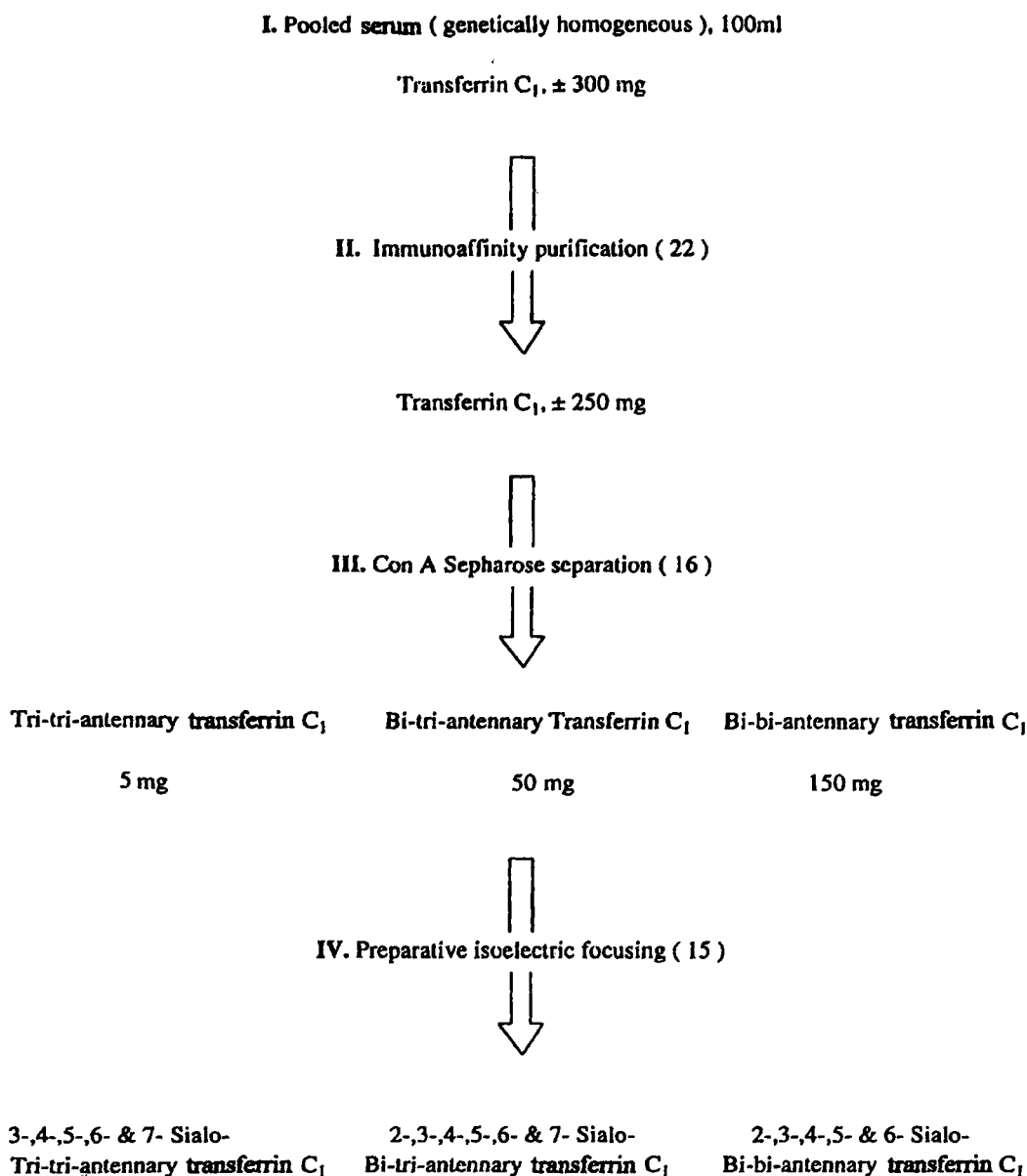


Fig. 1 Outline of the main techniques and the essential results of the sialo-transferrin preparations (22, 16, 15).

teers. While eight lanes show the same pattern of microheterogeneity, sera in lanes a, c, f and k show an apparently abnormal pattern. Due to diallelic expression of the transferrin gene, the normal pattern and the pattern

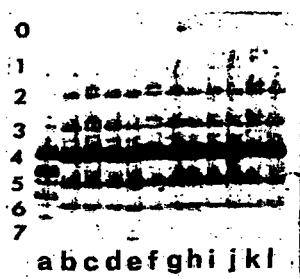


Fig. 2 On high resolution Immobililine gels, twelve different sera show eight patterns with the common occurring transferrin C₁ variant (70%). The four deviating patterns (in the lanes a, c, f, and k) are easy to recognize.

of the variant gene product co-exist, thus duplicating the number of bands without an actual change in the microheterogeneity pattern.

Figure 3 shows the densitometric scans of the transferrin C₁ obtained directly after separation using an anti-transferrin column (22), and the three fractions that were obtained after separation of the purified transferrin by Concanavalin A Sepharose: i.e. tri-tri-antennary transferrin C₁, bi-tri-antennary transferrin C₁, and bi-bi-antennary transferrin C₁ (16). Although each fraction is homogeneous with respect to their branching, they all contain at least five sialo-transferrin fractions (see the figures 5–7 and table 1).

This sialic acid-dependent heterogeneity is confirmed in figure 4, in which similar fractions are seen in lanes g, e and a. Lanes b and c show a partial desialylation of

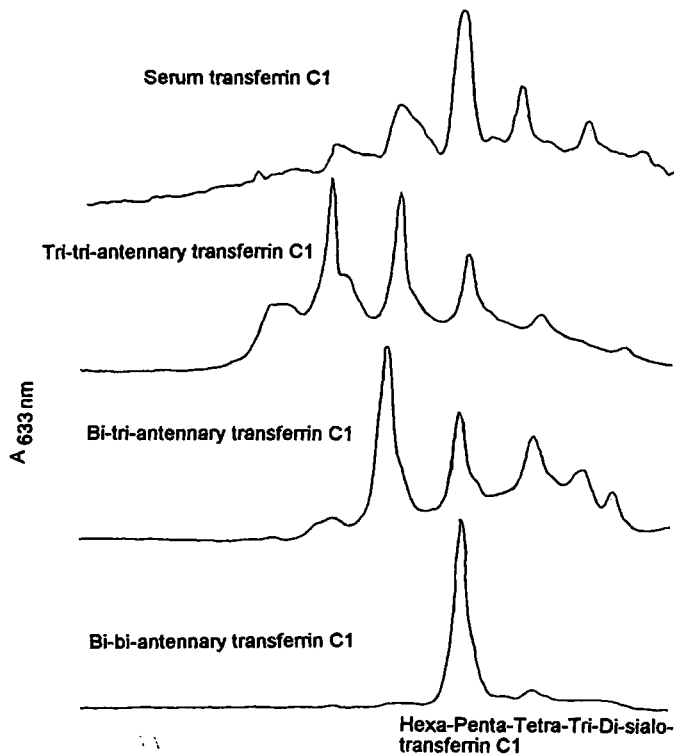


Fig. 3 Densitometric scans of the unfractionated purified transferrin (top) and the three transferrin fractions obtained after Concanavalin A Sepharose separation (10, 16, 23).

the bi-bi-transferrin C_1 , while lanes d, f and h show the asialo variants of bi-bi-transferrin C_1 , bi-tri-transferrin C_1 , and tri-tri-transferrin C_1 , respectively. Prolonged exposure to neuraminidase abolished all variability in each fraction, leaving behind only the asialo-transferrin in all fractions, confirming that microheterogeneity is the cause of the electrophoretic diversity in all fractions.

The three fractions from the Concanavalin A-Sepharose separation were each separated into their sialic acid-dependent fractions and isolated by preparative isoelectric focusing on ultra thin Immobiline gels (Pharmacia) as described recently (15).

Figure 5 shows the fractions obtained from tri-tri-antennary transferrin C_1 , ranging from 3-sialo-tri-tri-antennary transferrin C_1 to the 7-sialo-tri-tri-antennary transferrin C_1 in lanes a–e, while in lanes f–h contain 4-sialo-bi-bi-antennary preparations for comparison.

Although we only obtained less than one milligram of most of the subfractions from 100 ml of serum, we were able to analyse both their amino acid and their carbohydrate composition.

Figure 6 shows the purification products of the bi-tri-antennary transferrin C_1 , ranging from 2-sialo-bi-tri-antennary transferrin C_1 to 7-sialo-bi-tri-antennary transferrin C_1 in lanes a–f, while lanes g and h contain a 4-sialo-bi-bi-transferrin C_1 as reference. A similar display of bi-bi-antennary transferrin C_1 sialo-variants is

found in figure 7, lanes a–e, respectively, with 4-sialo-transferrin C_1 references in lanes f–h.

As mentioned already by *Hatton & Berry* (16), serum transferrin contains about 4% tri-tri-antennary transferrin and about 18% bi-tri-antennary transferrin. Consequently less starting material could be applied in the purification of tri-tri-transferrin C_1 sialo-variants. As a result, minor contamination of subfractions are seen as vicinal bands in bi-bi- and bi-tri-antennary transferrin preparations, albeit at negligible amounts.

Results of the sugar analyses of the transferrin fractions

Table 1 summarizes the results for N-acetylglucosamine, galactose and fucose.

According to *l.c.* (29–31) a glycan can bear an additional sialic acid and/or an additional N-acetylglucosamine as visualized in figure 8. It is therefore possible for 5- and 6-sialo transferrin to occur, although there are only two antennae on each glycan. The galactose values

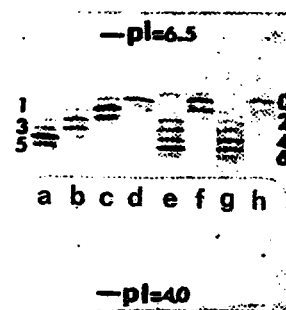


Fig. 4



Fig. 6

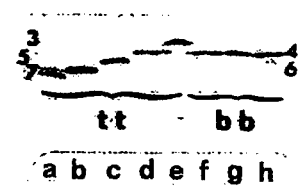


Fig. 5

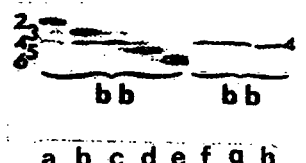


Fig. 7

Fig. 4 PhastGel electrophoresis of the three transferrin fractions. Lanes a, e and g contain a bi-bi-transferrin C_1 , a bi-tri-transferrin C_1 , and a tri-tri-transferrin C_1 preparation, respectively. Lanes b and c show partially desialylated bi-bi-transferrin C_1 . Lanes d, f and h show the completely desialylated products of the preparations in lanes a, e and g, respectively, all resulting in the same asialo-transferrin C_1 band (23) (asialo band at the right hand top-side of the gel).

Fig. 5 PhastGel electrophoresis of tri-tri-transferrins. Lanes a–e contain the 7-, 6-, 5-, 4- and 3-sialo-tri-tri-transferrin C_1 fractions, respectively, as obtained by preparative isoelectric focusing (15). Lanes f–h contain 4-sialo-bi-bi-transferrin C_1 as a reference.

Fig. 6 PhastGel electrophoresis of bi-tri-transferrins. Lanes a–f contain the 2-, 3-, 4-, 5-, 6- and 7-sialo-transferrin C_1 . Lanes g and h contain two 4-sialo-bi-bi-transferrin C_1 fractions for reference (15, 23).

Fig. 7 PhastGel electrophoresis of bi-bi-transferrins. Five fractions were analysed (lanes a–e). Standards of 4-sialo-bi-bi-transferrin C_1 (lanes f–g) (15, 23).

Tab. 1 Values of galactose and N-acetylglucosamine expressed in moles sugar/mole protein for the sixteen purified fractions of human serum transferrin.

Calculated percentage occurrences of each fraction in normal human serum are also given. These expected values are based on the

types of branching: tri-tri-, bi-tri- and bi-bi-transferrin C₁ according to l. c. (16) and on the recovery of sialo-transferrins after preparative isoelectric focusing purification (15). Of the three fractions of the Concanavalin A separation the fucose content is noted. The expected values agree with the literature (16).

Sialylation degree of transferrin C ₁	tri-tri-transferrin C ₁			bi-tri-transferrin C ₁			bi-bi-transferrin C ₁		
	GlcNH ₂	Gal	%	GlcNH ₂	Gal	%	GlcNH ₂	Gal	%
disialylation of degree transferrin C ₁	—	—	<1	5.3	4.4	2	7.4	4.1	<1
trisialylation degree of transferrin C ₁	8.1	5.6	<1	7.5	4.8	3	7.4	4.0	6
tetrasialylation degree of transferrin C ₁	8.0	5.7	<1	8.2	5.1	4	8.0	4.1	65
pentasialylation degree of transferrin C ₁	8.7	5.7	<1	9.0	5.0	5	7.7	4.2	6
hexasialylation degree of transferrin C ₁	x	5.5	3	8.8	5.2	1	8.8	4.1	1
heptasialylation degree of transferrin C ₁	7.4	x	<1	7.8	4.3	<1	—	<1	
Expected (16)	10	6	4	9	5	18	8	4	78
Fucose		0.6			0.4			0.1	

% the calculated percentage occurrences of this variant in normal serum is given.
 — not detectable

x not available for this analysis
 GlcNH₂ = glucosamine
 Gal = galactose

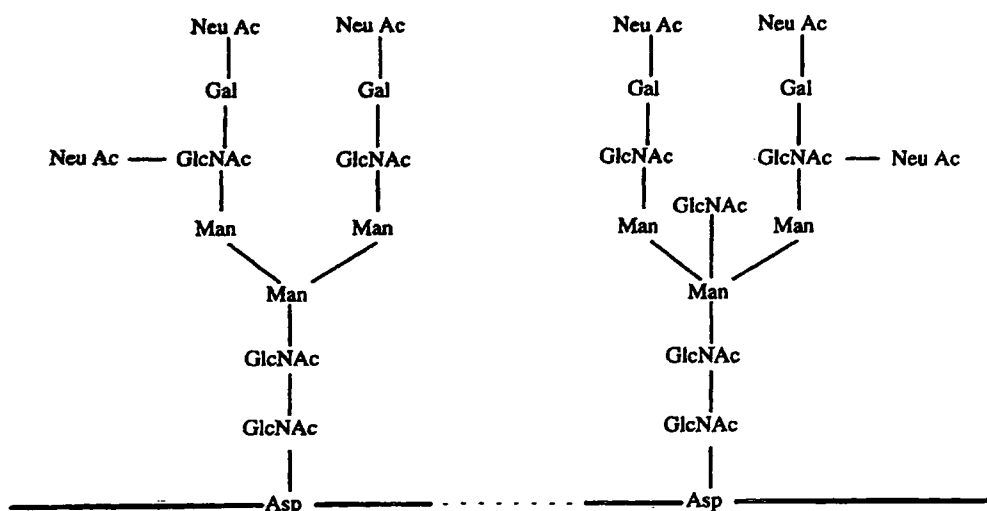


Fig. 8 The two presumed glycan structures of the 6-sialo-bi-bi-transferrin C₁ isotransferrin containing two additional sialic acids

and an additional N-acetylglucosamine (see tab. 1), derived according to l. c. (29–31).

in most fractions are near to the expected values (see tab. 1) although in particular the 2- and 2-sialo-bi-tri-transferrin C₁ fractions seem to contain less galactose than expected.

The N-acetylglucosamine values are lower than expected, in particular in bi-tri- and tri-tri-transferrin C₁ fractions.

However, when the calculated percentage occurrences of each fraction in serum are taken into account, it ap-

pears that about 75% of all transferrin variants contain the expected number of N-acetylglucosamines. The content of fucose was determined in each of the sixteen fractions. In table 1 they are only specified according to the antennae-dependent fractions. Yet, it is very clear that transferrin fractions that are more branched apparently contain more fucose. This may be related to the fact that in transferrins isolated from diseased persons both the branching of glycoproteins and the fucose content (predominantly in cancer) are much higher than

in normals (26), but at the moment we cannot confirm this finding.

Amino acid determination

Table 2 shows the results of the amino acid analyses, in which the N-acetylglucosamine values were also obtained (see tab. 1). Within the margins of error, the results indicate that all fractions show the same amino acid composition (24).

Discussion

Electrophoretic mobility of transferrin is known to be influenced by genetically determined differences in amino acid content, as well as differences in iron content, sialic acid content and the number of glycan antennae. The interpretation of the patterns has been extensively reviewed (1–5).

Theoretically, if partially Fe-saturated, genetically pure transferrin C₁C₂ is focused on an Immobiline gel, this would result in the separation of 72 variants (fig. 9), illustrating the wide heterogeneity of transferrin variants that must exist in one individual serum (4). For our investigation we halved this number by selecting only genetically homogeneous transferrin, transferrin C₁. To further simplify the pattern, we limited the number of fractions by a factor 4 using only fully iron-saturated transferrin (15, 23, 27) (fig. 9b). Applying Concanavalin A Sepharose we distinguished not only the various sialo-transferrins, but also the three transferrin forms which differ in their glycan branching, tri-tri-, bi-tri- and bi-bi-

transferrin C₁ (16). Figure 3 shows that each of these three forms from fresh serum can be further fractionated according to the types of sialo-transferrins. The number of terminal sialic acids contributes to the differences in isoelectric point, irrespective of the number of antennae, which is confirmed in figure 4.

The conclusions that can be drawn from the results of table 1 are that all of the sixteen analysed fractions differ either in sialic acid content (see figures 3–7), in the number of antennae (with consequent structural differences in both N-acetylglucosamine, galactose and sialic acid content), or in fucose content.

In some variants the content of N-acetylglucosamine or galactose or both is less than that calculated for model glycans that would be expected to migrate to these positions, based on our present knowledge of glycan structure. As sialic acid can also be bound to the N-acetylglucosamine, instead of galactose, 5- and 6-sialo-bi-bi-transferrin C₁ may occur. An additional N-acetylglucosamine of the 6-sialo-bi-bi-transferrin C₁ can be bound to the central mannose (29–31), which may explain these differences.

Figure 8 shows the presumed glycan structures of the latter iso-transferrin.

Several authors have reported changes in the (percentual) sialo-transferrin distribution in various diseases as well as during pregnancy (4, 5, 13, 23, 28).

Despite their analytical accuracies, these results leave room for interpretation. One might suppose that in haemochromatosis, 5- and 6-sialo-bi-tri-transferrin variants are elevated, while in rheumatoid arthritis the 5-, 6- and 7-sialo-tri-tri-transferrin fractions could be increased. Results presented in this paper demonstrate that it is possible to find subtle differences in the various transferrin subfractions, which may provide additional valuable information concerning glycan structures.

This is important, because the terminal sugar of even only one of the four to six antennae might influence the properties of the glycoprotein (7, 17, 18, 32). By combining the data of our sugar analyses (tab. 1) and the published structures of glycans (29–31), it is possible to derive interesting abnormal structures. These differences can be determined by analysis of the sialo-transferrins distribution in conjunction with crossed affinity isoelectric focusing, a method which we are developing for transferrin.

Conclusion

By applying the set of techniques presented in this paper it is possible to obtain a sufficient quantity of each of

Tab. 2 Average values from the amino acid analyses of the five or six sialo-transferrin fractions of each Concanavalin A fraction. In the far right column (seq. anal. 33) the values of the sequence analyses obtained from l. c. (33) are given.

Amino acids	Concanavalin A transferrin fractions			Reference values Sequence analyses (33)
	bi-bi- x̄	bi-tri- x̄	tri-tri- x	
Asx	79	78	78	79
Thr	31	29	29	30
Ser	41	43	44	41
Glx	59	58	60	59
Gly	49	50	54	50
Ala	57	59	58	57
Val	46	45	45	45
Met	9	9	8	9
Ile	15	17	18	15
Leu	58	58	56	58
Tyr	27	26	24	26
Phe	28	28	27	28
Lys	56	58	56	58
His	19	19	18	19
Arg	26	26	26	26

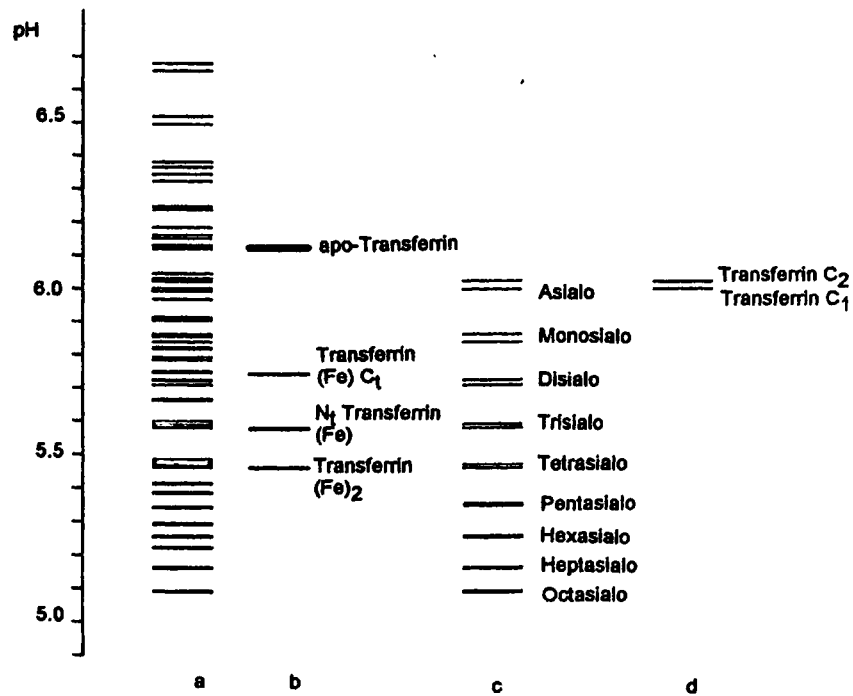


Fig. 9 Diagrammatic representation of transferrin microheterogeneity of an individual with the C_1C_2 genotype detectable on IEF. a) Total number of bands that can theoretically be distinguished on an Immobiline gel (4).

b) Major iron-dependent bands as revealed by IEF in carrier ampholyte (Ampholine) gels (23).

c) Microheterogeneous forms of transferrin that can be separated from iron-saturated transferrin (15).

d) Reduction of the number of bands from 72 (a) to 2 by iron-saturation and neuraminidase treatment of the transferrin, a procedure that can be useful in the assessment of genotypic variations (see fig. 4).

(Modified after l. c. (4) with permission.)

the sixteen purified isotransferrins to quantify their specific set of carbohydrates, even though the purification started from only 100 ml serum. Serum of individuals

with genetically homogeneous transferrin is required for optimal results.

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Prof. Dr. H. G. van Eijk
Erasmus University Rotterdam
Department of Chemical Pathology
Room EE 600
P.O. Box 1738
NL-3000 DR Rotterdam
The Netherlands