Apolipoprotein H: a two-step isolation method

Roberto Gambino, Gianluca Ruiu,* Maurizio Cassader,¹ and Gianfranco Pagano

Istituto di Medicina Interna, Universitá di Torino, *and Labmatorio Gentrale,* * *'Baldi e Riberi': Azienda Ospeduliera S. Giovanni Battista, Torino, Italy*

Summary A new method for the purification of apolipopro- tein H by **affinity** chromatography followed by continuouselution electrophoresis is described. It is both simpler and less complicated than the chromatographic and electrophoretic methods usually used. In addition, apolipoprotein H is isolated in a pure, structurally uncleaved form. This is of importance, **as** impairment has been detected in commercial preparations. The separation and purification of apolipoprotein H is a necessary prelude to its quantitative determination and phenotyping, and hence the clarification of its physiopathological mechanisms in lipid metabolism.-Gambino, R, **C. Ruiu, M. Cassader, and G. Pagano.** Apolipoprotein H: a two-step isolation method. *J. Lipid Res.* 1996. 37: 902-904.

Supplementary key words β 2-glycoprotein I • affinity chromatography \bullet continuous-elution electrophoresis.

Apolipoprotein H (apoH), also known as β 2-glycoprotein I, is a 50 kDa plasma protein that has recently attracted considerable interest on account of its many functions, especially those associated with lipoprotein metabolism (1, **2).**

Investigated because of its involvement in coagulation (3), apoH has recently been identified as an obligate cofactor for the formation of anti-phospholipid antibodies **(4).** It is present in plasma and is found in the two major lipoprotein fractions (VLDL and HDL): about 35% is lipoprotein-associated, and the rest is free plasma protein (5). Its ability to activate lipoprotein lipase suggests that it plays a part in triglyceride (Tg) metabolism (6). The rationale for this role seems to lie in apoH's structural polymorphism: three alleles at a single locus code for three isoforms whose phenotypes appear to be correlated with Tg and Chol-HDL levels (1, 2). It has long been known, in fact, that the phenotype of polymorphous apolipoproteins and their plasma level influence lipoprotein levels in various ways (7).

Separation and purification of apoH is a necessary prelude to its quantitative determination and the investigation of its phenotype, and hence the clarification of its physiopathological mechanisms in lipoprotein metabolism. *As* an alternative to the somewhat complicated chromatographic and electrophoretic methods usually used **(4,** 8), this report describes a two-step procedure for the isolation of high-purity apoH from normal sera by means of affinity chromatography followed by continuous-elution electrophoresis.

METHODS AND RESULTS

Rabbit anti-apoH antibodies (22 mg) (Behring, Scop pito, Italy) were immobilized on cyanogen-bromide-activated Sepharose 6B (Pharmacia Fine Chemicals, Piscataway, NJ) in 5 ml total volume and packed into a 0.7 **x** 13.5 cm column equilibrated with 0.1 M phosphate buffer, 0.3 M NaCl, 0.05% NaNs, pH 7.6 (buffer A) at a flow rate of 0.3 ml/min. Absorbance was monitored at **280** nm. Normal serum samples (5 ml per run) were diluted into **40** ml buffer A and applied to the column at 0.17 ml/min. The column was washed with buffer A at 0.3 ml/min until the absorbance reached the baseline, then equilibrated at 0.3 ml/min for 1 h with a lower ionic strength buffer composed of 0.01 M phosphate buffer, 0.15 M NaCl, 0.05% NaN₃, pH 6.8. The apoH bound to the column was eluted with 0.1 **M** glycine, **0.05%** NaNs, pH 2.5. Sample fractions were collected every minute and the pH was immediately adjusted with 60μ I 0.5 M phosphate buffer, pH 7.6.

Eluates from four column loadings were pooled and concentrated with Centriprep-10 concentrators (Amicon Inc., Beverly, MA) at 3,000 rpm and 20°C in a Beckman-JGB centrifuge (Beckman Instruments, Palo Alto, CA) to a final volume of about 1 ml.

The purity of the sample thus obtained was assessed by electrophoresis in **12%** SDS-PAGE. The gel was then silver-stained. As can be seen in Fig. **1** (lane l), the approximately **50** kDa band corresponding to apoH was accompanied by higher-weight bands. The presence of these contaminating proteins showed that further purification was required.

Continuous-elution electrophoresis of the proteins (about 2.5 mg) was then performed with a Prep-Cell (Bio-Rad Laboratories, Milan, Italy) through a cylindrical gel (gel tube inside diameter 37 mm; gel length 50 mm) composed of 8.5% acrylamide, 2.7% N,N' **methylene-bis-acrylamide,** and 0.1% SDS. The buffers were **0.025** M Tris, 0.192 M glycine, and **0.1%** SDS, pH 8.4. Separation was achieved at 40 mA in about 8 h. Samples from the fractions collected every minute with a tube collector were dot-blotted with specific anti-apoH antibody (data not shown). The samples containing apoH, spanning the 34th to the 52nd fraction, were

Abbreviations: Tg, triglycerides; Chol, cholesterol; apoH, apolipoprotein H; HDL, high density lipoproteins; VLDL, very low density lipoproteins; SDSPAGE, sodium dodecyl sulfate polyacrylamide gel electrophoresis; DTT, dithiothreitol; HRP, horseradish peroxidase; OPD, o-phenylenediamine dihydrochloride; HPLC, high pressure liquid chromatography.

To whom correspondence should be addressed: Dipartimento di Medicina Interna, Corso A.M. Dagliotti **14,** 10126 Torino, Italy

BMB

OURNAL OF LIPID RESEARCH

Fig. 1. SDS-PAGE gel electrophoresis of apoH in the two purifica**tion steps. Lane 1: silver-stained 12% SDSPAGE gel of eluted apoH** fractions from affinity chromatography column. Lane 3: silver-stained **12% SDSPAGE gel of purified apoH by continuous elution electro phoresis. Lane 5: Western blot of duplicate gel of lane 3 stained with specific apoH antiserum, using a two-antibody method. In lane 2, molecular mass markers are (from top to bottom): 97.4, 66.2, 42.7, 31.0,21.5,14.4 kDa. In lane 4, molecular mass markers are (from top to bottom): 66.2, 42.7, 31.0, 14.4 kDa. Prestained molecular mass marked with black points (lane 6) are (from top to bottom): 106.0, 80.0.49.5,32.5,27.5, 18.5 kDa.**

pooled, dialyzed in 0.02 **M** phosphate buffer, 0.15 **M** NaCl, pH 7.4, and subjected in duplicate to 12% SDS-PAGE. One gel was silver-stained (Fig. 1, lane 3), the other was immunoblotted with the specific antibodies (Fig. 1, lane 5). These lanes show a single, approximately 50 kDa band corresponding to the pure protein. Purity **as** judged by SDS-PAGE was 98-100%. The isolated protein was also subjected to 12% SDS-PAGE after reduction with DTT (data not shown). There was an apparent increase in the molecular weight of the reduced form **as** reported by others (8, 9). The recovery of pure apoH from the protein mixture eluted from the column in the first purification step was 52%. The final yield from 20 ml starting plasma was about 34%, **as** in other studies (8).

The apoH thus separated was examined by reversephase chromatography on a C_{18} IP Beckman column connected to a Beckman HPLC system. Ten µg was dissolved in 0.1 **M** HsP04, 0.02 **M** triethylamine, 0.05 **M** NaC104, pH 3 (buffer B). Buffer C was acetonitrile.

The pure protein eluted **as** a single peak with a retention time of 33.7 min, using a 20-70% acetonitrile gradient for 35 min at 1 ml/min. Absorbance was measured at 2 14 nm.

Analysis of the N-terminal region with a Model 475A Gas-Phase Sequencer (Applied Biosystems, Foster City, CA) showed that its 10 amino acids were those published for apoH by Steinkasserer et al. (10), namely Gly-Arg-Thr-X-Pro-Lys-Pro-Asp-Asp-Leu.

In order to evaluate the ability of apoH to bind negatively charged phospholipid, we incubated apoH at different dilutions with a constant concentration of cardiolipin. For binding experiments, apoH was labeled with horseradish peroxidase (HRP) by the two-step glutaraldehyde method. Briefly, glutaraldehyde was diluted in 0.1 **M** phosphate buffer, pH 6.8, to a final concentration of 1.25%. Five mg HRP was dissolved in 0.1 ml glutaraldehyde solution and allowed to incubate overnight at room temperature. The HRP-glutaraldehyde mix was added to an apoH solution (400 μ g in 200 μ l 0.1 **M** carbonate/bicarbonate buffer, pH 9.5) and incubated overnight at room temperature. The remaining sites were blocked with 0.2 **M** ethanolamine, pH 7, for 2 h at 4°C. Labeled apoH was dialyzed in 0.05 **M** Tris, 0.15 **M** NaCI, pH 7.4, overnight at 4°C. The unconjugated HRP molecules were removed by gel filtration.

Microtiter wells (Sigma Chemical, St. Louis, MO) were coated with 100 **pl** of cardyolipin (Sigma Chemical) at 30 μ g/ml in ethanol, dried under vacuum, and blocked with 350 μ l of 1% milk powder/0.3% gelatine in phosphate buffer-saline, pH 7.6. HRP-apoH was diluted 100- to 204,800-fold in washing buffer (0.1 **M** bicarbonate, 0.5 **M** NaC1, 0.1% milk powder, 0.1% Tween-20, pH 8.1). One hundred **pl** from each dilution was added to each well in duplicate, incubated for 2 h at 37"C, and then washed four times with washing buffer.

The plate was washed six times with assay buffer solution. One hundred **pl** of a freshly prepared enzyme substrate solution made by dissolving **an** OPD tablet in 0.05 **M** phosphatecitrated buffer, pH 5.0, to a final concentration of 1 mg/ml was added to the wells. The plate was briefly agitated, covered, and left in the dark at room temperature for 20 min. The reaction was then stopped by adding 50 p13 **M** HC1 to each well, and the plate was agitated to ensure thorough mixing. Well absorbance was measured with a Bio-Rad 3550 reader at 490 nm. HRP-apoH was able to bind to wells coated with cardiolipin in a dose-dependent manner. Binding of apoH was found for 1:100,1:200, and 1:400 dilutions, corresponding to 840, 420, and 210 ng/well, respectively.

DISCUSSION

Apolipoproteins play a key role in lipid metabolism **as** the structural or coenzymatic units of the lipoproteins. The levels of the main plasma lipids, too, are influenced by changes in the quantity or phenotype of some apolipoproteins (7).

Recent work **(2)** has shown that apoH's genic polymorphism influences plasma lipid levels, though their relation to its own plasma level has not been established. Isolation of apoH in a pure, structurally uncleaved form is thus a matter of importance, as impairment has been detected in commercial preparations (11).

The technique currently used for this purpose requires three or more chromatographic steps. Our procedure is much simpler and avoids all precipitating steps that could lead to protein loss. We found that pure apoH could not be obtained with a single affinity chromatography step. Higher molecular weight proteins, in fact, coeluted with apoH (Fig. 1, lane 1). They were also recognized, albeit with a weaker signal, by the anti-apoH antibodies in a Western blot analysis (data not shown), probably because of the polyclonality of the antibody binding apoH in the column, or because apoH formed indirect bonds with other proteins due to its strong electric charge, as reported by other authors (8). Subsequent purification by means of continuous-elution electrophoresis, on the other hand, gave 98-100% pure apoH (Fig. 1, lanes 3 and 5). The purified material displayed only one band reacting with the antisera, suggesting that our procedure included conditions for dissociation of the apoH complexes in plasma or, better still, was selective for free apoH. The final yield after the two purification steps was in agreement with that reported by McNally et al. **(S),** who observed binding of apoH to many biological substances by hydrophilic and other interactions.

The ability to bind a negatively charged phospholipid was evaluated by incubating apoH at different dilutions with a constant concentration of cardiolipin. Only native apoH and hence the active form is suitable for experiments designed to test whether purified apoH has retained lipid-binding activity. When clipped between Lys-317 and Thr-318, apoH lost its ability to bind negatively charged phospholipid. It is possible that such a clip could result in a conformational change in lipid-binding sites and alter the ability of apoH to bind lipid (11). Our apoH preparation satisfied all these criteria and bound successfully to cardiolipin in microplate wells. Our purification technique allowed the isolation of an uncleaved protein without other contaminants, such as a protease, which may be copurified with apoH as frequently reported.

In conclusion, this combination of two relatively simple methods was found to be an effective way of separating useful amounts of high-purity apoH for study of its role in lipid metabolism. **WI**

This work was supported in part by **grants** from Consiglio Nazionale delle Ricerche, Roma (Grant No. 93.00432.PF40, Progetto Finalizzato "INVECCHIAMENTO") and "Ministero dell'Universitá e della Ricerca Scientifica e Tecnologica" **(Quota** 60%), Roma, Italy. We are indebted to Prof. A. Conti (CNR) for help in amino acid analysis.

Manuscript received 16 August 1995 and in reuisedfonn 4 December 1995.

REFERENCES

- 1. Kamboh, M. I., R. E. Ferrel, and B. Sepehrnia. 1988. Genetic studies of human apolipoproteins. **IV.** Structural heterogeneity of apolipoprotein H (β 2-glycoprotein I). *Am. J. Hum. Genet.* **42:** 452-457.
- *2.* Sepehrnia, B., M. I. Kamboh, L. L. Adams-Campbell, C. H. Bunker, M. Nwankwo, P. P. Majumder, and R. E. Ferrel. 1989. Genetic studies of human apolipoproteins. VIII. Role of the apolipoprotein H polymorphism in relation to serum lipoprotein concentrations. *Hum. Genet.* **82:** 118-122.
- 3. Schousboe, I. 1985. P2-glycoprotein I: a plasma inhibitor of the contact activation of the intrinsic blood coagulation pathway. *Blood.* 66: 1086-1091.
- 4. McNeil, P. H., R. J. Simpson, C. N. Chesterman, and S. A. Krilis. 1990. Anti-phospholipid antibodies are directed against a complex antigen that includes a lipid-binding inhibitor of coagulation: β 2-glycoprotein I (apolipoprotein H). *Proc. Natl. Acad. Sci. USA.* **87:** 4120-4124.
- 5. Polz, E., and G. M. Kostner. 1979. The binding of β 2-glycoprotein-I to human serum lipoproteins. Distribution among density fractions. *FEBS Lett.* **102:** 183-186.
- **6.** Nakaya, Y. E., E. J. Schaefer, and H. B. Brewer, Jr. 1980. Activation of human post heparin lipoprotein lipase by apolipoprotein H (P2-glycoprotein-I). *Biochem. Biophys. Res. Commun.* **95:** 1168-1 172.
- **7.** Breslow, J. L. 1988. Apolipoprotein genetic variation and human disease. *Physiol. Rev. 58* 85-132.
- 8. McNally, T., I. J. Mackie, D. A. Isenberg, and S. J. Machin. 1993. Immunoelectrophoresis and ELISA techniques for assay of plasma β 2-glycoprotein-I and the influence of plasma lipids. *Thromb.* Res. **72:** 275-286.
- **9.** Day, J. R., M. Berkaw, and P. Arnaud. 1989. Isolation and characterization of the protein and cDNA for β 2-glycoprotein4 (apolipoprotein H). *Protides. Biol. Fluids. 36* 423-429.
- 10. Steinkasserer, A., C. Estaller, E. H. Weiss, and R. B. Sim. 1991. Complete nucleotide and deduced amino acid sequence of human P2-glycoprotein I. *Biochem.* J. **277:** 387-391.
- 11. Hunt, J. E., R. J. Simpson, and S. A. Krilis. 1993. Identification of a region of β 2-glycoprotein I critical for lipid binding and anti-cardiolipin antibody cofactor activity. *Proc. Natl. Acad.* Sci. *USA.* **90:** 2141-2145.

OURNAL OF LIPID RESEARCH