Distribution of oxysterols in human serum: Characterization of 25hydroxycholesterol association with serum albumin

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Oxysterols, derived either from cholesterol autoxidation in vitro or through lipoprotein peroxidation in vivo are biologically active compounds implicated in the pathogenesis of atherosclerosis. Using 25-hydroxycholesterol (250HC) as a model and radiolabeled 250HC to trace mass, in this study we examined the transport of oxysterols in human serum. In contrast to cholesterol, which is associated exclusively with serum lipoproteins, 55 to 75% of 250HC added to serum in vitro was associated with the lipoprotein-deficient fraction of serum (LPDS, density >1.21 g/mL) over a wide concentration range. Upon sequential gel filtration, ion-exchange chromatography, and nondenaturing polyacrylamide gel electrophoresis, 250HC added to LPDS eluted in a single peak corresponding to a protein peak of molecular weight about 67 kD. Immunoprecipitation of serum albumin from LPDS also precipitated 250HC. 250HC added to albumin-depleted LPDS did not associate with any of the remaining serum proteins, suggesting that albumin is the sole protein with which 250HC associates to any significant extent in LPDS. Bovine serum albumin (BSA) was used to characterize the association of 250HC with albumin. Other oxysterols, including 19-hydroxycholesterol, 7P-hydroxycholesterol, and 7-ketocholesterol, effectively competed with 250HC for association with BSA, suggesting that they also associate with albumin in serum. The association of 2SOHC with albumin in serum is selective but of low affinity as calculated from Scatchard analysis. However, given the high concentration of albumin in serum a significant amount of serum oxysterols may be delivered to or removed from various tissues via albumin. (J. Nutr. Biochem. 6:618-625, 1995.)

Keywords: oxysterol; cholesterol: serum; lipoproteins; albumin

Introduction

Cholesterol undergoes autoxidation in air,^{1,2} forming oxidized derivatives known as oxysterols. Various oxysterols, including 7α - and 7β -hydroxycholesterol, 7-ketocholesterol, cholestane-3 β , 5 α , 6 β -triol, epoxycholesterol, 26hydroxycholesterol, and 25-hdyroxycholesterol (25OHC), have been detected in USP-grade cholesterol preparations³

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and cholesterol-containing foods, 5.7 especially in egg yolk and dairy products.^{5,7} The content of oxysterols in these products has been reported to be 3 to $6 \mu g$ of dry food and to increase after processing, particularly after deep frying.^{5,7} Oxysterols may enter the circulation as contaminants of cholesterol-containing food.' These oxysterols found in food have also been detected in human and animal plasma at concentrations of 20 to 1,200 nM (about 8 to 480 ng/ mL), $^{8-12}$ and their concentrations reportedly were correlated with elevated plasma cholesterol concentrations.¹² In addition, some oxysterols may be generated by peroxidation of lipoproteins.^{13,14} Administration of probucol, an antioxidant and hypocholesterolemic agent, to hypercholesterolemic rabbits decreased their plasma oxysterol concentrations, independent of its lipid-lowering effect.^{15,16}

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Many oxysterols exhibit atherogenic properties and an ability to modulate cholesterol metabolism. The atherogenicity of oxysterols has been suggested by animal feeding studies in which oxysterols, when fed at relatively high doses, were more potent than pure cholesterol in causing aortic endothelial damage and inducing atherosclero- $\sin^{3,17,18}$ One study showed that even at a low concentration the oxysterol cholestane-3 β , 5α , 6 β -triol enhanced cholesterol-induced atherosclerosis.¹⁹ Modulation of cholesterol metabolism by various oxysterols, including cholestane-3 β , 5 α , 6 β -triol, 26-hydroxycholesterol, and 25OHC, occurs via inhibition of the activity of 3-hydroxy-3-methylglutaryl coenzyme A (HMG-CoA) reductase, 20.21 enhancement of cholesterol esterification, $22-24$ and reduction of cellular cholesterol efflux.²⁵ In addition, numerou changes in cell structure and function have been reported following the incubation of cultured cells with an oxysterol such as 250HC, including changes in membrane composition and hospholipid packing, echinosis in human erythrocytes,²⁶ increased cell fragility,²⁷ and enhanced membrane permeability to extracellular Ca^{2+} . 28.29

Relatively little is known about the transport and metabolism of oxysterols in serum. In the early 198Os, three research groups examined the distribution of various oxysterols in primate serum. Among these, Peng et al.³⁰ reported that in squirrel monkeys ingested 250HC associated with serum lipoproteins, predominantly with very low density lipoprotein (VLDL) and low density lipoprotein (LDL). Javitt et al.³¹ reported that the small amount of 26hydroxycholesterol detected in human serum was associated mainly with LDL and high density lipoprotein (HDL). In contrast, Streuli et al.³² reported that about 11 to 50% of various oxysterols, including 7 α -hydroxycholesterol, 20 α hydroxycholesterol, and 250HC added to human serum in vitro, were associated with the nonlipoprotein fraction of serum (density > 1.21 g/mL). In our own preliminary studies, ingested 250HC in mildly hypercholesterolemic rabbit serum was distributed between the lipoproteins and the lipoprotein-deficient serum (Morel, unpublished data) (LPDS, $d > 1.21$ g/mL). Using 250HC as a model, in the current studies we investigated the distribution of oxysterols in human serum in more detail, focusing on the identification and characterization of the protein(s) in LPDS, with which oxysterols associate.

Methods and Materials

Materials

The following compounds were purchased from Sigma (St. Louis, MO USA): cholesterol, 25-hydroxycholesterol, 19-hydroxycholesterol, 7_B-hydroxycholesterol, 7-ketocholesterol, bovine semm albumin (BSA, essentially fatty acid-free), bilirubin, cholic acid, deoxycholic acid, oleic acid, palmitic acid, and Norit A activated charcoal. 25-[26, 27-³H]-hydroxycholesterol (77 Ci/ mmol) and $[4^{-14}C]$ -cholesterol (53.0 mCi/mmol) were obtained from NEN-DuPont (Wilmington, DE USA). Sephadex G-200, DEAE-Sepharose Fast Flow, Blue Sepharose 6 Fast Flow, and Dextran T-500 were from Pharmacia-LKB (Uppsala, Sweden). Goat anti-human albumin IgG and goat anti-human immunoglobulin IgG were purchased from Cappel Research Co. (Durham, NC USA).

Labeling serum in vitro

Fresh human sera were obtained from healthy volunteers between the ages of 24 and 36 years. [$3H$]250HC (20 μ Ci/mL) was first mixed with unlabeled 250HC (0.7 mg/mL) in ethanol to achieve a specific activity of 10 μ Ci/mg, and the solution was evaporated under nitrogen $(99.9\%$ pure). The dried $[³H]25OHC$ was then redissolved in 10 to 20 μ L of ethanol and dispersed into 3 mL of human sera in vitro to achieve final concentrations of 0 to 100 μ M (ethanol < 1%). In some experiments, serum was also labeled with $\lfloor^{14}C\rfloor$ cholesterol (20 μ Ci/mL) in a similar fashion but with no added unlabeled cholesterol. Following incubation at room temperature for 1 hr then at 4°C for 18 hr, solid potassium bromide (KBr) was added to the labeled serum to adjust the density of the serum to $d = 1.210$ g/mL. The labeled serum samples were subjected to ultracentrifugation at $50,000$ rpm $(4^{\circ}C)$ in a Beckman 50Ti rotor for 48 hr to isolate a single fraction of total lipoproteins and LPDS.³³ All LPDS preparations were isolated under the same conditions.

Purification of 250HC carrier-protein in humans LPDS

LPDS, isolated from unlabeled human serum and then dialyzed in phosphate buffer saline (PBS) containing 2 mM ethylene-bis (oxyethylenenitriol) tetraacetic acid (EDTA), was labeled with $[{}^{3}H]25OHC$ (20 µCi/mL) as described above except that, in order to increase the $[^3H]25OHC$ specific radioactivity in the labeled LPDS, no unlabeled 250HC mass was added. After incubation at room temperature for 1 hr, then at 4°C for 18 hr, the labeled LPDS was gel filtered (Sephadex G-200, 800×30 mm, eluted with PBS containing 2 mM EDTA and 0.05% sodium azide, 1 mL/hr). Blue Dextran-2000 (2,000 kD), BSA (67 kD), cytochrome c (12.5 kD), and NaCl (57.5 D) were used to calibrate the column. Elution of Blue Dextran-2000, the protein standards, and LPDS protein were monitored by the absorbance at 280 nm, NaCl was monitored by conductivity, and ${}^{3}H$ radiolabeled was quantitated by liquid scintillation counting. Fractions (2.7 mL) containing radioactivity were pooled and dialyzed in 0.025 M Tris-HCl buffer, pH 8.0, then applied (about 40 mL) to a DEAE-Sepharose ion exchange column (500 \times 25 mm) equilibrated with a 0.025 M Tris-HCl buffer. Elution was carried out with a linear-gradient of NaCl, from 0.05 to 0.14 M in 0.025 M Tris-HCl buffer. Eluate fractions (3.0 mL) in which radioactivity was detected were then pooled and further purified by nondenaturing polyacrylamide gel electrophoresis (PAGE). All procedures were performed at 4°C.

Nondenaturing polyacrylamide gel electrophoresis was conducted; 34 one set of the gels was stained with 0.025% Coomassie Blue R-250 while another set in duplicate was sliced with a stainless steel gel slicer into small equal pieces (about 1 mm thick). Each gel slice was then immersed in 0.5 mL of 10% sodium dodecyl sulfate (SDS) solution at room temperature for 72 hr to allow diffusion of the trapped proteins into the SDS solution. The solution was then assessed for ³H radioactivity by liquid scintillation counting, protein content using the method described by Lowry et al., 35 and protein purity by SDS-polyacrylamide gel electrophoresis (SDS-PAGE).

Preparation of albumin-deficient human LPDS

Human LPDS, after dialysis in 0.025 M Tris-HCl buffer, was depleted of albumin by elution through a Blue Sepharose 6 Fast Flow affinity column (100 \times 30 mm, in 0.025 M Tris-HCl buffer with 2 mM EDTA and 0.05% sodium azide, 15 mL/hr). As described by Travis et al., 37 this method has been shown to quantitatively and selectively remove albumin from serum without altering other serum proteins. Aliquots of the LPDS sample before

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and after chromatography were assessed by SDS-PAGE to demonstrate depletion of serum albumin. No changes in the quantities of other serum proteins were detectable. The eluate from the affinity column was collected, lyophilized, and stored at -20° C. Albumin-deficient LPDS was reconstituted with PBS and labeled in vitro with $[{}^3H]25OHC$ prior to use.

Immunoprecipitation of serum albumin

Immunoprecipitation was performed according to Marsh et al.³⁸ In brief, $10 \mu L$ of the humans LPDS prelabeled with $[3H]25OHC$ was mixed with increasing volumes $(0 \text{ to } 1,200 \text{ }\mu\text{L})$ of goat antihuman albumin IgG $(21.3 \text{ mg of protein/mL})$. The mixture (total volume 1.2 mL) was incubated at room temperature for 1 hr with shaking at 10 min intervals and at 4°C for 18 hr. The precipitate was pelleted by centrifugation $(1,000g)$ for 20 min); the pellets were washed twice with cold PBS containing 0.1% SDS and 0.15% dithiothreitol (DTT) and once with cold PBS only. The lipids were extracted twice from the pellets with 2 mL of ethanol: acetone (1:1 vol/vol) and aliquots were taken for liquid scintillation counting. The remaining protein precipitate was dried under N_2 , dissolved in 0.5 mL of 0.1 N NaOH, and quantitated according to Lowry et al.³⁵ As a negative control, goat anti-human immunoglobulin IgG (20.7 mg of protein/ml) was used instead of anti-human albumin IgG. In a parallel study, increasing amounts of $[3H]25OHC$ -labeled LPDS were added to a constant volume of anti-albumin IgG, and the experiment was carried out in a similar fashion.

Assay for $\int^3 H/25$ -hydroxycholesterol binding

Binding of $[3H]25OHC$ to albumin was assessed using a modification of the density-gradient method of Haddad et al.³⁹ $[^3H]25OHC$ in 20 μL of ethanol was added to 2.0 mL of BSA (essentially fatty acid free, 30 nM) to reach final concentrations of 0 to 300 nM (0 to 10 molar excess) in the presence or absence of excess (40 fold) unlabeled 250HC. The mixture was incubated at 37°C for 18 hr before the density of the sample was adjusted to 1.21 g/mL by adding solid KBr. After ultracentrifugation $(50,000)$ rpm for 48 hr at 5° C), the samples were collected along the gradient and analyzed for radioactivity and protein content. Virtually all unbound 250HC was found in a thin layer (3 to 4 mm) floating on the top of the solution, whereas BSA-bound 250HC precipitated after centrifugation. The recovery of total $[^3H]25OHC$ radiolabeled was 70 to 88% in the absence of a 40 fold excess of unlabeled 250HC and 89 to 100% in the presence of excess. To compensate for low counts in the bottom (bound) fraction, bound $[{}^{3}H125OHC$ was calculated by subtracting free $[{}^{3}H]25OHC$ in the top fraction from total $[{}^{3}H]25OHC$ added to the samples. Specific binding was calculated by subtracting the values for $\binom{3}{12}$ 30HC binding in the presence of unlabeled 250HC from those in the absence of unlabeled 250HC.

Competition for association with serum albumin

BSA (essentially fatty acid free) was prelabeled with 250HC by adding $[3H]25OHC$ (0.1 mM, 20 μ Ci/mmol) to 0.01 mM BSA then incubating at $4^{\circ}C$ for 18 hr. A high (nonphysiological) concentration of $[{}^{3}H]25OHC$ was used in these experiments to maximize the ratio of 250HC to BSA (1O:l) and to minimize the nonspecific association of a small amount of $[3H]25OHC$ with the W_{u} of t_{u} and T_{u} of the tubes. Increasing concentrations of unlabeled putative wall be the three concentrations of unidered phalix $\frac{1}{2}$ albuming bilirubin and palmitic and palmitic actual particle actual particle actual particle solutions were dried under N, then dissolved in 20 PL of 0.1 N N OH and addedded in 20 PL of 0.1 N N OH and a 20 PL of 0.1 N N t_1 solution. Unlabeled absolute the labeled oxysterols were dissolved or d_1 solved oxysterols were dissolved to the fabeled BSA solution. Of addeded oxysteriors were dissolved

 1% cholate. Deoxycholate and cholate were dissolved in 20 μ L of PBS and added directly to prelabeled albumin. All these compounds were soluble at the concentrations used. The total volume of the mixture was restricted to 1.2 mL, and all mixtures were prepared in duplicate. After incubation at 20°C for 16 hr, unbound [\textdegree H]25OHC was adsorbed by dextran-coated charcoal (100 μ L) according to Taylor et al., 40° and pelleted by centrifugation at $1,000g$ for 20 min. The $[^3H]25OHC$ remaining in the supernatan (BSA-bound) was quantitated by liquid scintillation counting.

Results

Distribution of 25OHC in serum

Figure 1 shows that $[^{3}H125OHC$ radiolabel added in vitro to human serum was found associated not only with serum lipoproteins but also with LPDS $(d > 1.21 \text{ g/mL})$ over a wide range of 25OHC concentrations. Within the concentration range (10 to 1,000 nM) that oxysterols including 250HC have been detected in vivo, $8-10$ about 55 to 75% of [³H]25OHC radiolabel was found in the LPDS fraction. A similar distribution was seen even at a high nonphysiological concentration of 100 μ M (data not shown). In contrast, when $[14C]$ cholesterol radiolabel was added to serum to trace the distribution of endogenous serum cholesterol, virtually no $[{}^{14}$ C]cholesterol radiolabel was found in the LPDS fraction. Thus, unlike cholesterol that associated exclusively with the serum lipoproteins, 25OHC associated with both lipoproteins and LPDS.

Figure 1 Distribution of [³H]25OHC between lipoproteins and lipoprotein-deficient serum (LPDS) in human serum. As described in $t_{\rm H}$ Methods, increasing and \sim $t_{\rm H}$ and \sim $t_{\rm H}$ \sim $t_{\rm H}$ \sim $t_{\rm H}$ $\frac{1}{2}$ materials and meanous, increasing amounts of $\frac{1}{2}$ ripportion wormig, note added to o life of mathem soremic fraction (distance α is a non-magnetized into a hpoproton coordinating natural ($\alpha \prec$ g/mL) and a LPDS fraction ($d > 1.21$ g/mL) by density ultracentrif-
ugation. Data shown are mean \pm SD for triplicate samples.

Purification of protein with which 25OHC associates in human LFDS

To determine whether oxysterols in LPDS were free or protein-associated in solution, $[3H]25OHC$ -labeled human LPDS was subjected to gel filtration on a Sephadex G-200 column (Figure 2A). The $[^{3}H]25OHC$ radiolabeled eluted in a single peak corresponding to a protein peak with a molecular weight of about 60 to '70 kD, which is close to that of a BSA standard. The recovery of $[3H]25OHC$ radiolabel in this protein peak was 81%, suggesting that most of [³H]25OHC in LPDS was protein-associated. To further characterize the protein(s) with which 250HC associated, the fractions containing $[3H]25OHC$ radioactivity were pooled, dialyzed, and then eluted through an ion exchange column (DEAE-Sepharose) under a linear gradient of NaCl. Again, a single peak of $\binom{3}{125}$ H $\binom{3}{125}$ C radiolabel was observed (Figure $2\tilde{B}$). The purity of the samples after each of these steps was examined by SDS-PAGE as shown in Figure 2C. The predominant constituent of the fractions with which 250HC associated was a protein with molecular weight of approximately 67 kD.

The fractions with which 250HC associated were further analyzed by nondenaturing PAGE. As illustrated in Figure 3, $[{}^{3}H]25OHC$ radiolabel was detected in three slices along the gel (slices 9 to 11). In a parallel set of gels, BSA migrated to a similar point in the gel (data not shown). Upon analysis by SDS-PAGE, each slice contained a single protein band with molecular weight of approximately 67 kD.

The comigration of an approximately 67 kD protein with 250HC through gel filtration and ion exchange chromatography as well as in an electric field suggested that this protein might be a carrier protein for 250HC in serum. The similarity in its apparent molecular weight with that of serum albumin (67 kD) suggested that this protein might be albumin.

Immunoprecipitation of protein-associated 25OHC

To ask directly if albumin is a carrier protein for 250HC in human LPDS, a goat-derived anti-human albumin IgG was

 \blacksquare (A) Gel filtration of human LPDS labeled in vitro with $\frac{1}{2}$ $\frac{1}{2}$ $\frac{1}{2}$ $\frac{1}{2}$ $\frac{1}{2}$ continuation is a separated elu-vivo with t_{in} and t_{out} (believes of Ebe), because α are α and α and α are α show by arrows. Elias Dexition 2000 (2,000 help and 2001 (07 help shown by arrows. Elution of protein was monitored by absorbance at 280 nm and [³H]25OHC radioactivity by liquid scintillation counting. The recoveries of 3H radioactivity and protein were 81% and $82.$ The focusities of $\frac{1}{2}$, reduced vity and protein volume of $\frac{1}{2}$ 260 to 290 mL) from gel filtration chromatography were collected, pooled, and they have get infration conditionally were concerned pooled, and dialyzed in 0.025 M Tris-HCl $(\text{pH} = 8.0)$ buffer. (B) Dialyzed sample (20 mL) was then applied to a DEAE-Sepharose ion exchange column (500 \times 25 mm) pre-equilibrated with the same buffer. Elution was carried out with a linear-gradient of NaCl, from 0.05 to 0.14 M, in the above buffer. The recoveries of ${}^{3}H$ radioactivity and protein were 82% and 79%, respectively. (C) SDS-PAGE of human serum samples during steps of 25OHC-carrier protein purification. Lanes 1 and 10: molecular weight standards; lanes 2 and 3: duplicate samples of human LPDS; lanes 4 and 5: [³H]25OHC peak fractions obtained after gel filtration; lanes 6 and 7: dialyzed gel filtration samples; and lanes 8 and 9 : $[³H]25OHC$ peak fractions after ion exchange chromatography.

Figure 3 Nondenaturing PAGE of protein(s) with which 25OHC associates. Fractions 120 to 145 collected after ion exchange (DEAE-Sepharose) were pooled and subjected to electrophoresis in a 10% polyacrylamide tube gel. The gels were sliced and immersed in 10% SDS solution to let diffusion of trapped proteins and [3H]250HC to the solution. Aliquots were assayed by liquid scintillation counting and absorbance at 280 nm. Purity of protein was checked by SDS-PAGE. Values for protein and radioactivity are means from duplicate samples. The recoveries for 3H radioactivity and protein were 84% and 79%, respectively. Insert: SDS-PAGE. Lane 1: molecular weight standards; lanes 2 to 6: samples from gel fractions 8 to 12.

used to immunoprecipitate serum albumin from the LPDS labeled with $\int_0^3 H$]250HC. As illustrated in *Figure 4*, this antibody, but not an anti-immunoglobulin IgG, precipitated $[3H]25OHC$ from LPDS in an antibody dose-dependent manner. In this experiment, at a maximum, about 73% of the total 3H radiolabel in the mixture was precipitated by anti-albumin IgG, and virtually no albumin remained in the supematant (SDS-PAGE; gel not shown). In another experiment, increasing amounts of the antigen $(I³H)25OHC$ labeled LPDS) added to a constant mass of the antibody (anti-albumin IgG) precipitated at maximum more than 95% of [³H]25OHC radiolabel (data not shown). These data confirm that the protein with which 250HC associates in LPDS is serum albumin.

Association of 25OHC with albumin-depleted LPDS

To determine whether the high concentration of albumin in serum might mask a potential association between 250HC and any other serum proteins, human LPDS was first depleted of albumin by elution through a Blue Sepharose affinity column then labeled with $[3H]25OHC$ in vitro and gel filtered (Sephadex G-200). As shown in Figure 5A (insert), the Blue Sepharose-treated LPDS was markedly depleted of albumin. Upon gel filtration of the labeled albumindepleted LPDS, only 6% of the applied $\binom{3}{12}$ 50HC radioactivity was recovered in the eluate, suggesting that even in

Figure 4 Immunoprecipitation of serum albumin from human LPDS labeled with [3H]250HC. Ten microliters of [3H]250HClabeled human LPDS (45 mg of protein/ml) was mixed with increasing volumes of goat-derived IgG against either human-albumin or human-immunoglobulins (21.3 and 20.7 mg of protein/ml) as indicated. Protein precipitate (immune complex) was pelleted by centrifugation, and both the pellet and the supernatant were assayed for [3H]250HC radioactivities. Data represent the percent of total $[3H]25OHC$ in each fraction and are means \pm SD for triplicate samples.

the absence of albumin, 250HC did not associate to any significant extent with other serum proteins.

To determine whether serum albumin could selectively remove $[3H]25OHC$ retained in the column, the column was washed with a mixture of proteins of different molecular weights containing BSA (67 kD) and ovalbumin (43 kD). As shown in Figure 5B, the $[3H]25OHC$ retained in the column was completely eluted in two peaks; about 73% of the total 3H radiolabel eluted in a peak corresponding to a protein peak of molecular weight approximately 120 to 130 kD, while the remaining 27% of 3H radiolabel eluted with a protein peak of approximately 60 to 70 kD. No 3 H radiolabel was associated with the 40 to 50 kD peak of ovalbumin. Upon SDS-PAGE (Figure 5C), both protein peaks with which $[3H]25OHC$ associated contained a predominant band with a molecular weight identical to that of BSA. Thus, the higher molecular weight (120 to 130 kD) protein with which $[3H]25OHC$ associated was a dimer of BSA.

Characterization of the association between 2SOHC and albumin

A number of oxysterols, including 19-hydroxycholesterol, 7B-hydroxycholesterol, and 7-ketocholesterol, were tested for their abilities to compete with $[{}^{3}H]25OHC$ for association with BSA. There results are shown in Figure 6A. Each

Figure 5 Gel filtration of albumin-deficient human LPDS after labeling with $[{}^{3}H]25OHC$ (50 μ Ci/mL). (A) Elution with PBS containing 2 mM EDTA and 0.01% sodium azide. Insert: SDS-PAGE of human LPDS before (lane 1) and after (lane 2) blue Sepharose affinity chromatography. Lane 3: molecular weight standards. (B) Elution with 10 mL of 2% BSA and 2% ovalbumin in PBS. Protein content was monitored by absorbance at 280 nm and [³H]25OHC radioactivity by liquid scintillation counting. (C) SDS-PAGE of protein-peak samples from the BSA/ovalbumin elution: molecular weight standards in lanes 1 and 10; pooled fractions from the BSA/ovalbumin elution in lanes 3 and 4 (325 to 375 mL), lanes 5 and 6 (275 to 324 mL), and lanes 7 to 9 (220 to 274 mL).

oxysterol effectively decreased the association of [3H]250HC with BSA in a dose-dependent manner; maximal competition (10% if origin) was seen at a 10 fold molar excess of unlabeled oxysterol. It is thus likely that other oxysterols exhibit a similar distribution in serum, i.e., they also associate with albumin. We also tested whether the putative iigands of albumin, including free fatty acids, bilirubin, and bile acids (cholate and deoxycholate, also biological detergents) could limit the association of $[^3H]25OHC$ with BSA. As shown in Figure 6B, bilirubin reduced the association of $[{}^{3}H]25OHC$ with albumin in a dose-dependent manner to 20% of its original value. In contrast, free fatty acids (palmitic or oleic) and bile salts (cholate or deoxycholate), up to a 10 fold molar excess, had no effect on the association of radiolabeled 250HC with

Figure 6 Competition for association with albumin by other oxysterols, albumin ligands and biological detergents. [3H]250HC (0.1 mM) in ethanol was added to BSA (essentially fatty acid free, 0.01 mM) in vitro to yield a 250HC to BSA ratio of IO: 1. Other oxysterols and known albumin ligands were then added to (as described in the Methods and materials section) 1 mL of [³H]25OHC-labeled BSA to achieve the indicated competitor-to-[³H]25OHC molar ratios. Charcoal-dextran was used to adsorb free [3H]250HC and competing ligands and removed by centrifugation (1,OOOg). The 3H radiolabel remaining in the supernatant was BSA-associated. Data shown are means \pm range for duplicate samples.

BSA. Progesterone, estradiol, cortisone, corticosterone, and hydrocortisone were also ineffective competitors (data not shown).

To further examine the nature of the interaction between 250HC and albumin, a binding assay was performed as described in the Methods and Materials section. The association of increasing concentrations of labeled 25OHC with fatty acid-free BSA in the presence and absence of a 40 fold excess of unlabeled 250HC was examined. Scatchard analysis⁴¹ of these data suggested a K_d of 5×10^{-4} M and a B_{max} of 0.11 mol of 25OHC/mol of albumin, values indicative of a low affinity association with a maximum binding of one molecule of 250HC per ten molecules of BSA. Other methods of assessing 250HC binding, such as dextran-charcoal precipitation of unbound oxysterol as described by Dawson et al 42 and a solid-phase binding assay,⁴³ yielded similar binding curves and Scatchard analysis results (data not shown).

Discussion

These studies show that 25-hydroxycholesterol, a biologically active oxysterol, distributes between lipoproteins and

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albumin in serum. These findings agree with and extend the observations of Streuli et a1.32 that oxysterols associate in vitro with the "nonlipoprotein serum" fraction. The identification of albumin as the sole protein in LPDS with which 250HC associates suggests that albumin may serve as a transport protein for oxysterols. Moreover, given the ability of other oxysterols to compete with labeled 250HC for association with BSA, it is likely that they too associate with albumin in serum.

The distribution of 25OHC differs markedly from that of cholesterol, which is associated exclusively with lipoprotein in serum. Part of this difference may be explained by the greater polarity of 250HC and other oxysterols. Their greater solubilities in an aqueous environment may allow them to interact with albumin more readily. Despite the apparent low affinity, the association so formed is stable to numerous manipulations including ultracentrifugation and electrophoresis.

Despite the structural similarities between oxysterols and other steroids, such as corticosteroids and vitamin D (1,25 dihydroxycholecalciferol) which bind to specific transportproteins in serum, $44,45$ our findings suggest that albumin is the sole nonlipoprotein plasma protein with which 250HC associates to any significant extent. From this point of view, the association of 250HC with albumin is selective and specific. It is known that serum albumin associates with many hydrophobic compounds in circulation.⁴⁶ Among these, long-chain free fatty acids and bilirubin bind to albumin with high affinity and low capacity,⁴⁶ while others such as progesterone and testosterone bind to albumin with low affinity and high capacity. 44 Given the structural similarity between steroids and oxysterols, it is not entirely surprising that the interaction between albumin and 250HC is of low affinity. This interaction likely depends on hydrophobic attraction between the oxysterol molecule and the hydrophobic pocket (or domain) of the albumin molecule. This speculation is supported by the apparent lack of stereospecificity for oxysterol association with albumin as well as by the observation that bilirubin, a molecule distinct in structure from that of oxysterols which binds to a specific binding site on albumin, 46 also inhibited the association of oxysterol with albumin. It is likely that the interaction between oxysterol and albumin occurs near the bilirubinbinding region, or that the binding of bilirubin to albumin alters its conformation such that oxysterol can no longer associate. However, exactly how approximately one molecule of 250HC actually interacts with ten molecules of albumin on a molecular basis is not clear. Apparently, though, this interaction between oxysterols and hydrophobic domain(s) of albumin remains stable even in the presence of biological detergents such as cholate and deoxycholate.

From our chromatography studies, 250HC appeared to bind more readily to the dimeric form of albumin than to the monomer. In many animal species, including rat, rabbit, guinea pig, and human, preferential binding of serum sterols to multimeric forms of their binding proteins has been reported. $47,48$ While the reason for this is not clear, it is possible that the number of ligand-binding sites is greater in the multimer than in the monomer, as has been reported for the binding of progesterone to BSA.⁴⁹

Very recently, Bjorkhem et al.⁵⁰ have provided evidence showing that cholesterol loading in human alveolar macrophages effectively enhanced the conversion of cholesterol to oxysterols and the secretion of oxysterols into the medium. This secretion of cellular oxysterols required the presence of serum proteins in the culture medium. Based on these observations, the investigators predicted the presence of an oxysterol-carrier protein in serum that could facilitate the removal of cellular cholesterol in the form of oxysterols. Our findings suggest that albumin is this oxysterol-carrier protein and, in addition, other data from our laboratory²⁵ have demonstrated that BSA can stimulate efflux of 250HC from a variety of cells in culture. Given the high concentration of albumin in serum, the capacity of albumin for oxysterol association is significant. It is possible that a significant amount of oxysterol converted from cholesterol is secreted and removed by albumin. Thus, it may be, as hypothesized by Bjorkhem et al.,⁵⁰ that conversion of cellular cholesterol to oxysterols represents a general defense mechanism to counteract elevated intracellular cholesterol levels. However, it is not clear whether such reverse transport of oxysterols to the liver for further catabolism occurs in vivo and functions to limit atherosclerosis, or whether the other proatherogenic effects of oxysterols, including cytotoxicity, 5^{1-53} enhancement of cholesterol esterification,²²⁻²⁴ and reduction of cholesterol efflux²⁵ prevail.

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