Biogenesis of apolipoprotein A-V and its impact on VLDL triglyceride secretion

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Running title: ApoA-V biogenesis and triglyceride secretion

Abbreviations:
apo, apolipoprotein; BSA, bovine serum albumin; Dox, doxycycline, ER, endoplasmic reticulum; FBS, fetal bovine serum; SDS-PAGE, Sodium dodecyl sulfate-polyacrylamide gel electrophoresis; TG, triglyceride; VLDL, very low density lipoprotein

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Abstract

Apolipoprotein A-V (apoA-V) is a potent regulator of intravascular triglyceride metabolism, yet its plasma concentration is very low compared to other apolipoproteins. To examine the basis for its low plasma concentration, the secretion efficiency of apoA-V was measured in stably transfected McA-RH7777 rat hepatoma cells. Pulse-chase experiments revealed that only ~20% of newly synthesized apoA-V is secreted into culture medium within 3 hours post-synthesis, and that ~65% undergoes presecretory turnover; similar results were obtained in transfected nonhepatic (CHO) cells. ApoA-V secreted by McA-RH7777 cells was not associated with cell surface heparin-competable binding sites. When stably transfected McA-RH7777 cells were treated with oleic acid, the resulting increase in triglyceride (TG) synthesis caused a reduction in apoA-V secretion, a reciprocal increase in cell-associated apoA-V, and movement of apoA-V onto cytosolic lipid droplets. In a stably transfected doxycycline-inducible McA-RH7777 cell line, apoA-V expression inhibited triglyceride secretion by ~50%, increased cellular TG, and reduced Z-average VLDL₁ particle diameter from 81 to 67 nm; however, no impact on apoB secretion was observed. These data demonstrate that apoA-V inefficiently traffics within the secretory pathway, that its intracellular itinerary can be regulated by changes in cellular TG accumulation, and that apoA-V synthesis can modulate VLDL TG mobilization and secretion.

Supplementary key words: lipoprotein assembly; triglyceride; lipid trafficking; very low density lipoprotein; apolipoprotein B
INTRODUCTION

Apolipoprotein A-V (apoA-V), a member of the exchangeable apolipoprotein family that is synthesized predominantly in the liver, is a potent regulator of intravascular triglyceride (TG) metabolism (1). When overexpressed in transgenic mice, apoA-V reduces plasma TG levels by 65%, whereas apoA-V gene inactivation increases plasma TG by 4-fold (2). The preponderance of current literature suggests that apoA-V affects plasma TG turnover by stimulating LPL-mediated lipolysis of TG-rich lipoproteins either directly or indirectly (3-7). ApoA-V has also been found to serve as a ligand for low density lipoprotein receptor family members and other potential lipoprotein receptors, and may thus contribute to the clearance of TG-rich lipoproteins and their remnants (8-11). However, recent studies have revealed that the impact of apoA-V on plasma TG concentration is complex and variable. In humans, several loss-of-function and null apoA-V alleles are associated with both reduced plasma apoA-V levels and elevated plasma TG (12, 13), yet other studies have found both positive and negative associations between plasma apoA-V and TG concentrations (7, 14, 15). Moreover, recent studies in mice found a positive correlation between plasma apoA-V and TG concentrations (16, 17).

Despite its apparent impact on intravascular TG-rich lipoprotein lipolysis and clearance, a peculiar characteristic of apoA-V is that its plasma concentration is in the range of 100-200 µg/L, which is ~10,000 fold lower than apoA-I, ~1000 fold lower than apoA-IV, and corresponds to ~1 molecule of apoA-V for every 1000 VLDL particles (18, 19). This presents a conundrum as to how an apolipoprotein circulating at such low levels could exert such a potent effect on plasma TG metabolism and concentration. Although it is certainly possible that apoA-V could function in plasma at extreme sub-stoichiometric concentrations relative to TG-rich lipoproteins, it has also been suggested that apoA-V might function within the hepatocyte to directly modulate hepatic TG...
metabolism and secretion (19, 20). Indeed, apoA-V was first identified as a gene that is up-regulated in rats following partial hepatectomy (21), suggesting that it could play a role in the conservation of intracellular lipids needed for liver regeneration. While an effect of apoA-V on TG production has not been observed in all studies (4, 5, 22), Schapp et al. (3) documented reduced hepatic TG production following adenovirus-mediated expression of human apoA-V in mouse liver. Most recently, the discovery that apoA-V may reside on cytosolic lipid droplets (23, 24) further supports the concept that apoA-V responds to and perhaps modulates aspects of intracellular hepatic TG metabolism.

In the current report, the secretory trafficking of apoA-V was examined in both hepatic and non-hepatic cells under basal conditions and during oleic acid-stimulated TG synthesis. These studies suggest that the low plasma concentrations of apoA-V may be due, in part, to its inherently inefficient exocytic trafficking, and that TG accumulation within hepatoma cells further antagonizes apoA-V secretion. Interestingly, these studies also revealed that in a stably transfected, inducible cell line, apoA-V gene expression reduces TG secretion, suggesting an extra-vascular mechanism by which apoA-V could modulate TG metabolism and plasma TG levels.

EXPERIMENTAL PROCEDURES

Cell culture

McA-RH7777 cell lines were grown in DMEM containing 4.5 g/L glucose and 10% FBS. CHO-K1 cells were maintained in DMEM-Ham's F-12 containing 10% FBS. All media were supplemented with 100 units/ml penicillin and 100 µg/ml streptomycin. Cells were grown in 100 mm dishes at 37°C in an atmosphere containing 5% CO₂.
Transfection and selection of stable clones

McA-RH7777 and CHO-K1 cells in 100 mm dishes were transfected at ~30% confluence with 16 µg of apoA-V expression plasmid (20) and 2 µg of pSV2-neo (25) using FuGENE 6 (Roche Molecular Biochemicals). Twenty-four hours post transfection, cells were subjected to selection with DMEM-10% FBS supplemented with 750 µg/ml G418 (Cellgro). Selection medium was replaced every 48 h for 10 days. Individual clones were isolated, expanded, and maintained in 250 µg/ml G418. To generate inducible cell clones, apoA-V was inserted into plasmid pTRE2hyg (Clontech), which was cotransfected into McA-RH7777 cells at a 1:1 ratio (20 µg total DNA) with plasmid pTet-On (Clontech). Twenty-four hours post-transfection, cells were subjected to selection with DMEM-10% FBS supplemented with 750 µg/ml G418 and 200 µg/ml hygromycin. Selection media was replaced every 48 h for ~14 days. Individual clones were selected and maintained in 250 µg/ml G418 and 100 µg/ml hygromycin. To induce apoA-V expression, cells were incubated for the times indicated with 1 µg/ml of doxycycline (Dox; BD Biosciences). Individual clones were analyzed for inducible expression by immunoblot analysis. Two clones (1 and 6) were characterized and clone 1 was used for the experiments described under Results.

Metabolic radiolabeling and analysis

Cells in 100 mm dishes were labeled for the indicated times with 100 µCi/ml of [35S]Met and Cys (EasyTag Express Protein Labeling Mix - Perkin Elmer) in Met and Cys deficient DMEM, followed in some cases by chase using complete DMEM containing 2.5 mM Met and 1 mM Cys. Unless otherwise indicated, all pulse and chase media contained 10% FBS. Following each labeling period, cells were placed on ice and cells and media were harvested, subjected to immunoprecipitation with anti-human apoA-V serum (Supplemental Fig. S1) or anti-human apoB (Academy Bio-medical, Houston,
TX), followed by SDS-PAGE (26). Band intensities were quantified using a Fujifilm BAS5000 phosphorimager. In some experiments cells were incubated with increasing concentrations of porcine heparin (Sigma No. H-7005). For labeling of lipids, 10 \( \mu \text{Ci/ml} \) \( [3\text{H}]\)oleate (Perkin Elmer) was added to cells for the times indicated. After washing monolayers, cells were lysed with 1% Triton X-100, 150 mM NaCl, 25 mM Tris, pH 7.4, 1 mM PMSF, 1 \( \mu \text{g/ml} \) leupeptin, and 1 \( \mu \text{g/ml} \) pepstatin. Both clarified cell lysate and media samples were supplemented with lipid markers and extracted with chloroform-methanol, as described (27, 28). Lipids were fractioned by thin layer chromatography in a neutral solvent tank (heptane:ether:acetic acid (90:30:1) (29). The TG-containing fraction was visualized by incubation in iodine vapor, cut from the plate, and quantified by liquid scintillation counting.

**Immunofluorescence microscopy**

Stably transfected McA-RH7777 cells were plated on poly-L-Lysine coated cover slips and 24 hours later fixed in 3.7% formaldehyde in PBS for 20 min, followed by incubation for 1 h in PBS containing 10 mM glycine, 0.1% saponin, and 3% BSA. Cells were then incubated for 1 h in primary antibody diluted in PBS containing 0.1% saponin and 1% BSA. The following dilutions were used: rabbit anti-human apoA-V serum, 1:300; mouse anti-human ADRP (Fitzgerald Industries International, Inc.), 1:50. Cells were then incubated with rhodamine-conjugated goat anti-rabbit IgG, and FITC-conjugated goat anti-mouse IgG, (1:20; Jackson ImmunoResearch) for 1 hour in PBS containing 0.1% saponin and 1% BSA. Cells were post-fixed with 3.7% formaldehyde in PBS, mounted using ProLong Gold antifade reagent, (Invitrogen), and viewed using a Zeiss Axioplan 2 microscope with a 63x oil objective.
Isolation of lipid droplets

McA-RH7777 cells were cultured with and without 0.8 mM oleate complexed to 1.5% BSA in DMEM-10% FBS. After 24 hours the cells were homogenized, and lipid droplets isolated by sucrose gradient centrifugation, as described (26). Twelve, 1 ml fractions, collected from the top of the gradient using an Autodensiflow gradient fractionator (Labconco), were subjected to TCA precipitation, SDS-PAGE, and immunoblot analysis using anti-human apoA-V serum. TG content of the lipid droplet fraction was determined by enzymatic assay (Wako).

VLDL particle diameter distribution

The hydrodynamic diameter of VLDL produced by transfected McA-RH7777 cells was measured using a Zetasizer Nano-S® Model ZEN1600 dynamic laser light scattering instrument (Malvern Instruments) at a wavelength of 633 nm. Following density gradient ultracentrifugation, as described (30, 31), gradients were fractionated into 12, 1 ml fractions from the top using the Auto Densi-Flow gradient fractionator. The VLDL₁ (Sₗ > 100) and VLDL₂ (Sₗ 20–100) fractions were transferred to a quartz cuvette and light scattering readings were performed at 20 °C. Volume-adjusted size distributions are displayed. Gradient samples were subsequently subjected to immunoprecipitation with anti-apoB antibodies and analyzed by SDS-PAGE.
RESULTS

Analysis of apoA-V secretion kinetics in hepatic and non-hepatic cells

Previous studies documented inefficient secretion of apoA-V from transiently transfected COS cells undergoing continuous metabolic radiolabeling with $[^{35}S]$Met and Cys (20). However, as endogenous apoA-V expression is predominantly limited to hepatocytes, we compared the secretory behavior of apoA-V in stably transfected rat hepatoma cells (McA-RH7777) and CHO cells. To examine secretion quantitatively, pulse-chase analyses were performed. After a 10-minute pulse with $[^{35}S]$Met and Cys and a 120 min chase, only 20% of newly synthesized apoA-V was secreted from McA-RH7777 cells (Fig 1A) and 38% from CHO (Fig. 1B). In addition to limited recovery of apoA-V in media, little cell-associated apoA-V remained after the 120 min chase (~10%). Hence, in addition to its limited secretion the majority of newly synthesized apoA-V appeared to undergo rapid presecretory turnover.

As apoA-V can associate with heparin sulfate proteoglycans (6, 32), the limited recovery of apoA-V in media fractions could result from binding of newly secreted apoA-V to the cell surface. To determine if newly secreted apoA-V associated with heparin competable binding sites on McA-RH7777 cells, transfected cells were radiolabeled with $[^{35}S]$Met and Cys in the presence and absence of heparin, and the distribution of cell-associated and media apoA-V was examined. As shown in Fig. 2A, the relative distribution of apoA-V present in cell and media fractions was unaffected by the presence of heparin at concentrations up to 10 U/ml. To determine if higher concentrations of heparin might affect cell association, the experiment was repeated using concentrations of up to 100 U/ml; however, as shown in Fig. 2B, no significant change in cell-associated apoA-V was detected. These results suggest that the low apoA-V secretion efficiency observed in Fig. 1 is not due to cell surface proteoglycan-mediated sequestration. To explore whether apoA-V bound to the surface of McA-
RH7777 cells via some other mechanism, live cell staining with anti-apoA-V antibody was performed. No cell staining was observed unless cells were first permeabilized with saponin (Supplemental Fig. S2).

**Effect of oleate on apoA-V secretion efficiency and subcellular localization**

ApoA-V synthesis is up-regulated in hepatic regeneration (21), a condition associated with hepatic TG accumulation (33-36). Hence, we explored whether the intracellular trafficking of apoA-V was affected by alterations in cellular TG synthesis and accumulation. Stably transfected McA-RH7777 cells were incubated with and without oleate before and during pulse-chase analyses. Inclusion of oleate caused a marked (~46%) reduction in the secretion of apoA-V from stably transfected McA-RH7777 cells (Fig. 3A and B) and a corresponding (~36%) increase in cell-associated apoA-V (Panel C). The oleate-induced alteration in secretion occurred exclusively at the level of trafficking and subcellular localization, as oleate had no effect on the high percentage of apoA-V that underwent pre-secretory turnover during the 120 min chase (Panel D).

Shu et al. demonstrated that an apoA-V-GFP fusion protein and native apoA-V could associate with cytosolic lipid droplets in oleate treated McA-RH7777 cells (23, 24). Presumably, this localization arises from the retrotranslocation (or other mode of trafficking) of apoA-V from the ER lumen into the cytosol whereupon it associates with lipid droplets (37). We therefore hypothesized that the reduced secretion of apoA-V observed with oleate treatment was caused by a corresponding increase in delivery of apoA-V to lipid droplets. To explore this possibility, stably transfected McA-RH7777 cells were incubated with and without oleate for 24 hours. The addition of oleate promoted increased synthesis and accumulation of neutral lipids, as evidenced by an increased intensity of Nile red staining (Fig.4A). Indirect immunofluorescence microscopy demonstrated that oleate treatment was accompanied by increased apoA-V binding to
lipid droplet structures (Fig. 4B), which were identified based on the colocalization with the lipid droplet binding protein, ADRP (38). The oleate-induced relocalization of apoA-V onto lipid droplets was further confirmed biochemically by isolating lipid droplets via cell homogenization and sucrose density centrifugation (39). Relative to controls, oleate treatment resulted in a ~16-fold increase in TG contained in the lipid droplet fraction (Fig. 4C, fraction 1; 5.42 versus 88.18 µg TG in control and oleate treated cells, respectively). When gradient fractions were subjected to SDS-PAGE and immunoblotting, the apoA-V content in fraction 1 also increased by ~10-fold in oleate versus control cells. These data indicate that the movement of apoA-V onto lipid droplets may directly compete with the exocytic trafficking of apoA-V.

**Effect of apoA-V expression on apoB and triglyceride secretion**

Based on its low concentration in plasma, the possibility that apoA-V modulates intracellular hepatic TG metabolism has been proposed (3, 19, 20). Hence, we explored whether the expression of apoA-V could affect TG secretion in McA-RH7777 hepatoma cells. The apoA-V stably transfected McA-RH7777 cells used in Figs. 1-4 responded to oleate by increasing TG secretion and apoB mass by ~2-fold (data not shown). However, to rule out possible phenotypic variability associated with clonally selected cell lines, we generated McA-RH7777 cell lines that expressed apoA-V under the control of a doxycycline (Dox)-inducible promoter. As shown in Fig. 5A, these cells displayed undetectable basal apoA-V expression and a robust induction when cells were incubated with Dox. To examine the consequences of apoA-V expression on TG secretion, cells were labeled with [³H]oleate in the absence and presence of Dox for 24 h. Induction of apoA-V expression resulted in a ~50% increase in cellular TG content (Fig. 5B) and a roughly corresponding decrease in TG secretion (Fig. 5C). To assess the impact of apoA-V on apoB, control and induced cells were subjected to radiolabel pulse-chase
analysis with [³⁵S]Met and Cys (Fig. 5D). As shown in Figs. 5D and E, apoA-V expression appeared to have no impact on the secretion or intracellular stability of apoB.

The finding that apoA-V impacts TG but not apoB secretion suggests an effect on particle size but not particle number. To assess apoB particle characteristics in the presence and absence of apoA-V expression, stably transfected McA-RH7777 cells treated with and without Dox were metabolically radiolabeled with [³⁵S]Met and Cys for 4 h, and media were subjected to cumulative rate density gradient ultracentrifugation (30, 31). No apparent change in apoB density distribution was observed in response to apoA-V expression (Fig. 6A); however, when the VLDL₁ fraction (Sₙ > 100) was analyzed by dynamic laser light scattering, the Dox-treated cells displayed a 26 nm reduction in peak VLDL₁ particle size diameter from 66 to 42 nm and a 14 nm reduction in Z-average diameter from 81 to 67 nm (Fig. 6B); apoA-V had no impact on the VLDL₂ (Sₙ 20–100) peak diameter of ~31 nm, as expected. These data indicate that the reduced TG secretion observed upon induction of apoA-V expression is due primarily to attenuation of second step particle maturation, essential for the formation of TG-rich VLDL.

DISCUSSION

One of the signature characteristics of apoA-V is that its plasma concentration is extremely low compared to its homologous relatives in the exchangeable apolipoprotein family, apoA-I and apoA-IV, which circulate at levels that are 10,000- and 1,000-fold higher (1, 14, 15, 18, 19). Although little is known about the catabolic fate of plasma apoA-V, our observation that only a small fraction of newly synthesized apoA-V is secreted from McA-RH7777 hepatoma cells into media suggests that inefficient secretion of apoA-V from the liver into the plasma compartment may, in part, contribute to its low plasma concentration. Although Shu et al. (32) recently estimated that the
mass of plasma apoA-V in human apoA-V transgenic mice was 4-fold greater than the amount present in the entire liver, this relationship does not necessarily reflect secretion efficiency, as many factors, including presecretory turnover and plasma residence time, can affect this ratio.

The basis for the inefficient secretion of apoA-V is unknown. It is possible that apoA-V lacks effective anterograde transport properties, critical for exit from the ER, or that its hydrophobicity and insolubility in the absence of lipid (20, 40) affects its folding and transport competence. In either case, prolonged residence time in the ER may promote apoA-V’s retrograde translocation into the cytosol (41). Although many such dislocated proteins are targeted for turnover, apoA-V’s affinity for lipid droplets may protect a population from degradation, particularly as cytosolic lipid droplet formation may provide an escape route from the ER to the cytosol for some ER-localized proteins (37). It is therefore likely that within the cell, apoA-V must continually associate with lipids to maintain its solubility, and thus its trafficking may be particularly sensitive to lipid fluxes from the ER membrane, whether into the ER lumen, or into the cytosol on the surface of lipid droplets (42).

The present data also provide evidence of a linkage between intracellular apoA-V trafficking and TG metabolism. Treatment of stably transfected McA-RH7777 hepatoma cells with oleate, which increased intracellular TG accumulation, caused a dramatic inhibition of apoA-V secretion and a reciprocal increase in intracellular apoA-V. Fluorescence confocal microscopy established that this was due to the association of apoA-V with the surface of cytosolic lipid droplets, as has been observed previously (23, 24, 32). Hence, it appears that hepatic TG synthesis or accumulation may drive a dynamic competition between apoA-V secretion and lipid droplet association.

The consequence of apoA-V expression and localization was explored using a regulatable apoA-V expression system. When transfected apoA-V expression was
induced with Dox, neither the secretion nor density distribution of apoB in McA-RH7777 cells was altered dramatically. This finding agrees with observations of Shu et al, who also noted no change in apoB secretion or density gradient distribution in Hep3B hepatoma cells stably transfected with human apoA-V (23). However, in the current study, apoA-V expression was associated with a ~50% reduction in TG secretion and a corresponding increase in cellular TG content. To explore the basis for this observation, particle size analysis was performed. While the VLDL₂ fraction displayed no significant size change in Dox treated cells, the VLDL₁ peak particle diameter was reduced by ~26 nm upon Dox-mediated induction of apoA-V expression (Fig. 6). Assuming a VLDL₁:VLDL₂ ratio of 1:3 (Fig. 6A), this change in particle diameter corresponds to an ~40% reduction in VLDL volume, a value in reasonable agreement with the ~50% reduction in radiolabeled TG secretion observed in Fig. 5C. Hence, the reduced TG content of media in apoA-V expressing McA-RH7777 cells appears to arise from a reduction in VLDL particle size distribution but not particle number. These data are consistent with findings of Schaap at al., who found that adenovirus-mediated expression of human apoA-V inhibited VLDL-TG secretion, also without affecting particle number (3). While other studies have failed to establish a link between apoA-V expression and VLDL secretion in transgenic mice (5) it is worth noting that upon induction of apoA-V in vivo during liver regeneration (21), hepatic TG synthesis and accumulation are up-regulated without an accompanying increase in VLDL secretion (43).

Considering these observations, we speculate that depending on the relative rates of TG, and VLDL synthesis, apoA-V either enters the secretory pathway, possibly on the surface of nascent VLDL, becomes subject to presecretory degradation, or buds from the cytoplasmic side of the ER membrane on the surface of lipid droplets (27). Thus, under specific metabolic conditions, apoA-V could modulate VLDL assembly by
facilitating the diversion and sequestration of a pool of metabolically active TG within the ER membrane towards lipid droplet formation and away from apoB lipidation. As ablation of apoA-V expression does not appear to increase hepatic TG secretion (44), this suggests that apoA-V may exert mainly an inhibitory effect on VLDL-TG secretion, which becomes robust only as hepatic TG synthesis increases.

Observations from animal and human studies provide further evidence of the linkage between apoA-V gene expression and hepatic TG synthesis, storage, and secretion. Shu et al. (32) observed that both hepatic TG content and apoA-V lipid droplet association were increased in human apoA-V transgenic mice, whereas inactivation of the mouse apoA-V gene had little effect. Although the impact on plasma TG was not examined in that study, Pamir et al., found that when human apoA-V transgenic mice were fed a high fat and high sucrose diet, fasting plasma TG levels fell instead of increasing (45), suggesting that apoA-V gene expression had inhibited diet-induced hepatic VLDL-TG secretion. Werner et al. observed that hepatic steatosis induced by essential fatty acid deficiency is accompanied by increased apoA-V gene expression (46); yet, Huang et al. found that plasma apoA-V levels were 45% lower in obese, insulin resistant, dyslipidemic subjects (47), suggesting that hepatic steatosis, which is a concomitant of the metabolic syndrome, reduced hepatic apoA-V secretion. At present the mechanism of the linkage between apoA-V and TG metabolism is not well understood, but it is relevant that the apoA-V gene contains two E-box elements that can bind SREBP1c (48), a nuclear factor that plays a central role regulating hepatic TG synthesis (49, 50), and that the human apoA-V promoter contains a PPARα response element (51, 52), and thus is up-regulated by PPARα agonists, which are also potent modulators of hepatic TG metabolism (50).

In summary, our data from a hepatic cell line model establishes that apoA-V is inefficiently secreted, that stimulation of TG synthesis significantly inhibits apoA-V
secretion and redirects the trafficking of apoA-V to the surface of cytosolic lipid droplets, and that up-regulation of apoA-V gene expression reduces VLDL-TG with little apparent effect on apoB secretion itself. These data suggest that, in addition to its well established function in regulating plasma TG levels by catalyzing the peripheral lipolysis and clearance of TG-rich lipoproteins, apoA-V may also play a critical role in modulating hepatic lipoprotein secretion and TG storage. ApoA-V may thus stand at the crossroads between hepatic lipid export and storage, and may be an important factor in determining the susceptibility to hepatic steatosis, lipotoxicity, and insulin sensitivity.

Acknowledgments
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Protein 1 Plays a Critical Role in the Lipolytic Processing of Chylomicrons. Cell Metab. 5: 279-291.


FIGURE LEGENDS

Fig. 1. Kinetics of apoA-V secretion. A: McA-RH7777 and B: CHO-K1 cells, stably transfected with human apoA-V, were pulse radiolabeled with [35S]Met and Cys for 10 min and chased for the times indicated. After each chase time, apoA-V in cells and media was recovered by immunoprecipitation and analyzed by SDS-PAGE and phosphorimager analysis. Percent secretion is calculated as the amount of radiolabeled apoA-V present at each chase time point as a percent of apoA-V synthesized during the 10 min pulse (0 min chase).

Fig. 2. Effect of heparin on apoA-V cell association. A: Stably transfected McA-RH7777 cells were radiolabeled with [35S]Met and Cys for 2 h, followed by addition of the indicated concentration of porcine heparin and further incubation for 6 h. Cell lysates (C) and media (M) samples were immunoprecipitated with anti-human apoA-V serum and subjected to SDS-PAGE and fluorography. B: The experiment in panel A was repeated in triplicate using the indicated concentration of porcine heparin. Radioactive band intensities were quantified by phosphorimager analysis and expressed as the percentage of total radiolabeled apoA-V associated with the cell pellet (mean ±SEM, N=3).
**Fig. 3.** Oleate-induced triglyceride synthesis reduces apoA-V secretion efficiency. **A:** Stably transfected McA-RH7777 cells were pretreated without (-OA) and with oleate (+OA) (0.8 mM oleate complexed to 1.5% BSA) for 2 h and subjected to pulse-chase analysis, as described in Fig. 1, in the continued absence and presence of OA. **B:** Duplicate dishes of cells incubated without and with OA were subjected to pulse radiolabeling for 10 min followed by 120 min chase. The percentages of initial (0 min chase) cell-associated radiolabeled apoA-V recovered in media (B) and cell fractions (C) at the end of the 120 min are shown. **D:** Percentage of initial radiolabeled apoA-V lost during the 120 min chase. For B-D, error bars depict data range.

**Fig. 4.** Effect of oleate-induced triglyceride accumulation on apoA-V subcellular localization. McA-RH7777 cells, stably transfected with human apoA-V, were incubated with or without OA, as described for Fig. 3, in the presence of 10% FBS for 16 h. **A:** Cells were stained with Nile red, as described (53) and examined by fluorescence microscopy. **B:** Cells were fixed, and stained with anti apoA-V and anti-ADRP antibodies as described under Materials and Methods. Areas of colocalization of apoA-V (Rhodamine; red) and ADRP (FITC-green) are detected in the overlay (yellow). **C:** Cells were homogenized and subjected to sucrose gradient centrifugation (39). Twelve, one ml fractions were collected and subjected to precipitation with trichloroacetic acid, SDS-PAGE, and immunoblot analysis using anti-human apoA-V serum. Fraction 1 (top) contains the cellular lipid droplet fraction (39).
**Fig. 5.** Effect of apoA-V on apoB and triglyceride secretion. A McA-RH7777 cell line that expresses human apoA-V under the control of a Dox-inducible promoter was incubated without (-) or with (+) 1 µg/ml Dox for 48 h. **A:** cells were labeled with [35S]Met and Cys for 4 h, also - and + Dox, and cell lysates were subjected to immunoprecipitation with anti apoA-V serum, SDS-PAGE and phosphorimager analysis. **B** and **C:** Cells were incubated in media containing 20% FBS and 0.4 mM oleate complexed to 0.75% BSA, - or + 1 µg/ml Dox for 24 h, followed by radiolabeling with [3H]oleate (10 µCi/ml) in the same medium for 24 h. Cells (B) and media (C) were extracted with chloroform-methanol and the lipid extract was fractionated by TLC. TG bands were quantified by liquid scintillation counting. **D:** Cells cultured for 48 h, as described for B and C, in the absence or presence of Dox were pulse radiolabeled with [35S]Met and Cys for 30 min and chased with unlabeled media for 0 or 2 hours, as indicated. ApoB in cell and media samples was immunoprecipitated and subjected to SDS-PAGE and phosphorimager analysis. **E:** Percentage of newly synthesized apoB (0 h chase) recovered in media after 2 h chase. Control (-Dox) efficiency was set to 100% (B, C, and E are Mean ±SEM; N=3). All data were analyzed using a paired student t test (*P<0.05).

**Fig. 6.** ApoA-V expression impacts lipoprotein particle size distribution. Inducible McA-RH7777 cells were incubated in the absence (-) or presence (+) of Dox for 48 h, and radiolabeled as described for Fig. 5A. **A:** Media were harvested and subjected to cumulative rate flotation ultracentrifugation, as described (30, 31). Size distribution of particles contained in the VLDL$_1$ (B) and VLDL$_2$ (C) fractions was determined using dynamic laser light scattering analysis.
Fig. 1

A

McA

B

CHO-K1

% Initial Counts

0 20 40 60 80 100 120

Time (min)

Cell

Media

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Fig. 2

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B

% Cell Associated

![Bar graph showing % Cell Associated vs Heparin (U/ml)](image)
Fig. 5

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B

Cellular $[^{3}\text{H}]\text{TG}$ (DPM/mg protein $\times 10^4$)

C

Media $[^{3}\text{H}]\text{TG}$ (DPM/mg protein $\times 10^3$)

D

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E

apoB secretion (% control)

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### A

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### B

- **VLDL₁**
  - - Dox
  - + Dox

### C

- **VLDL₂**
  - - Dox
  - + Dox