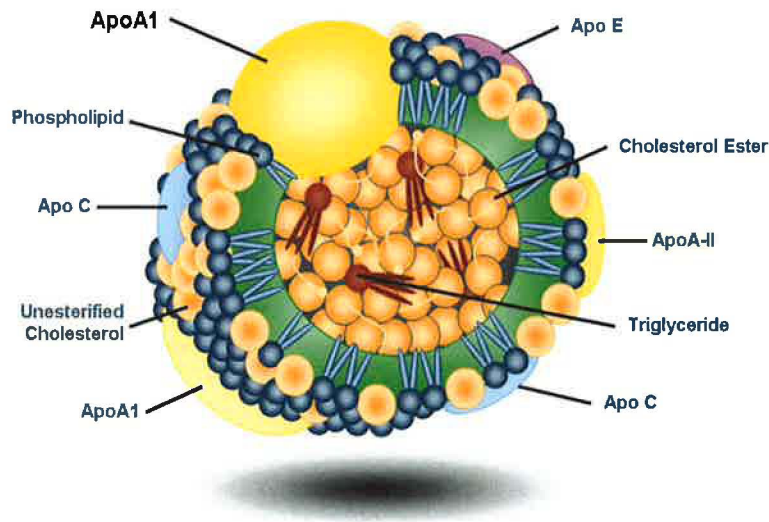


High-Density Lipoprotein - A Paucity of Therapy, A Paucity of Knowledge



**Anand Rohatgi, MD
Division of Cardiology
Internal Medicine Grand Rounds
UT Southwestern Medical School
July 9, 2010**

This is to acknowledge that Dr. Anand Rohatgi, M.D. has not disclosed any financial interests or other relationships with commercial concerns related directly or indirectly to this program. Dr. Anand Rohatgi will not be discussing off-label uses in his/her presentation.

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Introduction

Heart disease is the number one cause of death in the U.S., with coronary heart disease (CHD) and stroke accounting for the majority of these deaths. As of 2006, an estimated 17.6 million Americans had CHD and over 425,000 deaths in 2006 were directly attributable to CHD.¹ From 1980 to 2000, there has been a 50% reduction in the death rate from CHD, with about 24% of the reduction attributable to lowering of total cholesterol and consequent low-density lipoprotein cholesterol (LDL-C).² From all available epidemiologic evidence and randomized controlled trials of LDL-C lowering, the lower the baseline or achieved LDL-C, the lower the risk of CHD.^{3,4} Current guidelines reflect these associations by targeting LDL-C thresholds for initiating therapy and achieving therapeutic benefit.³

Despite these reductions, significant residual risk remains, and CHD remains the leading cause of death in men and women. It has long been known from several large epidemiologic studies *that high-density lipoprotein cholesterol, HDL-C, is inversely associated with CHD*.⁵ Four large American studies and one British study had been completed by the 1980s, with varying levels of risk factors but surprisingly similar levels of HDL-C among men and women separately. Taken together, *a 1 mg/dL increase in HDL-C was associated with 1.9-2.3% decrease in CHD risk in men and 3.2% decreased risk in women*. These risks were attenuated for CHD mortality in men except for one study but were magnified for CHD mortality in women. Adjustment for non-HDL-C (total – HDL-C) attenuated all associations between HDL and CHD risk but remained significant in 3 of the 5 studies (Figure 1).

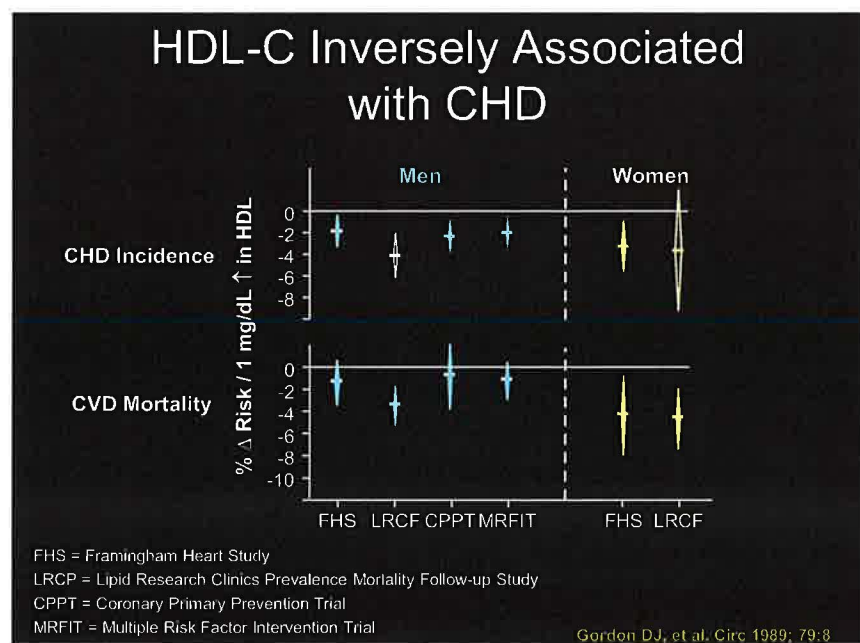


Figure 1

Similarly in patients with established CHD, low HDL-C is highly prevalent (>60%) despite statin use⁶ and is associated with increased risk of CHD events despite achievement of extremely low LDL-C levels $\leq 70\text{mg/dL}$ with statin therapy.⁷

These consistent observations have led to the inclusion of low HDL-C (<40mg/dL in men; <50mg/dL in women) as a major risk factor for CHD in most risk prediction algorithms, including the Adult Treatment Panel-III guidelines and the Framingham Risk Score.⁸ Unfortunately, whether HDL is causal in the development of or protection from CHD has not been proven in human studies. Importantly, despite classification of low HDL-C as a major CHD risk factor, evidence-based therapies targeting HDL-C that improve CHD risk are lacking, and improving HDL-C is considered a secondary aim in the ATP-III CHD risk reduction algorithm (Figure 2).⁸

Current ATP III-NCEP guidelines
2002

Treatment of low HDL cholesterol (<40 mg/dL)

- First reach LDL goal, then:
- Intensify weight management and increase physical activity.
- If triglycerides 200-499 mg/dL, achieve non-HDL goal.
- If triglycerides <200 mg/dL (isolated low HDL) in CHD or CHD equivalent, **consider** nicotinic acid or fibrate.

Figure 2

Recently, a novel class of drugs known as cholesteryl ester transfer protein inhibitors (CETP inhibitors) has been developed that markedly increase serum HDL-C by preventing transfer of cholesteryl ester out of HDL to other lipoprotein particles. In 2007-2008, the findings of several phase III trials of one such compound, torcetrapib, were published, showing increased risk of death and no improvement in progression of carotid and coronary atherosclerosis in patients with CHD or familial hypercholesterolemia (Figure 3).⁹⁻¹² Though there were several adverse effects of torcetrapib, including raised blood pressure and reduced potassium, the results of these studies have cast serious doubt on the strategy of raising HDL-C and have turned attention to measuring other aspects of HDL composition and function as better markers for CHD and targets for therapy.

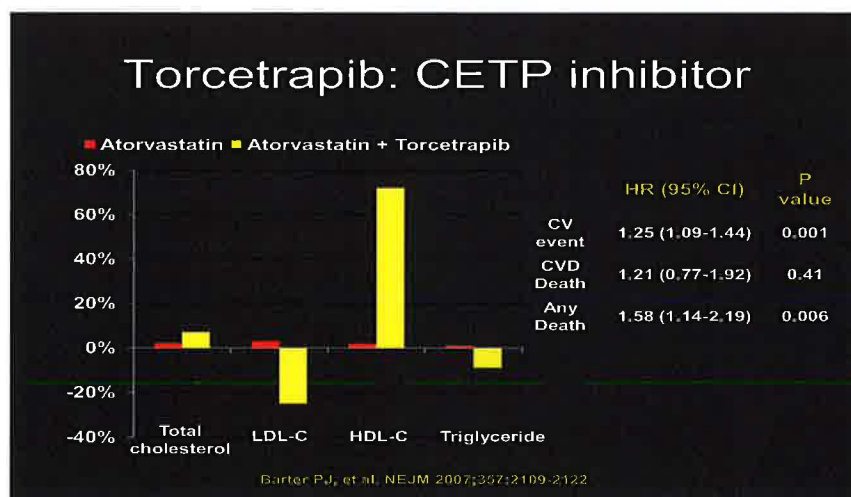


Figure 3

Role of HDL

Cells can produce the appropriate amount of cholesterol needed to maintain membrane and intracellular homeostasis. However, excess intra-cellular free cholesterol is toxic and is rapidly converted to oxysterols, stimulating export out of the cell. Macrophages can harbor oxidized and esterified cholesterol, resulting in the classic “foam cells” that characterize atherosclerotic lesions in arteries. The liver is the only organ that can degrade cholesterol and shuttle it out of the body through bile and feces. HDL participates as a universal plasma acceptor of cholesterol and phospholipid and directly transports cholesterol from peripheral tissues to the liver. This process, termed reverse cholesterol transport (RCT), is considered to be the main mechanism responsible for preventing the development of atherosclerotic lesions when excess lipid depositing occurs in the arterial walls.

I. HDL Biogenesis

A. Apolipoprotein A-I (apoA-I)

The HDL life-cycle is highly dynamic, initiating as protein and acquiring cholesterol from multiple acceptors with multiple delivery sites and exchange partners. **Apolipoprotein A-I (apoA-I)** is the main protein constituent of HDL (~70% of the apolipoprotein content) and is secreted from the liver and intestinal cells. Lipid-poor apoA-I accepts free cholesterol and phospholipids through transfer from lipoprotein lipase-mediated lipolysis of triglyceride-rich lipoproteins and cellular cholesterol efflux. Although it remains unclear whether HDL-C is causal in CHD, human apoA-I transgenic mice have increased HDL-C and are resistant to atherosclerosis,¹³ supporting the concept that apoA-I has a direct role in both HDL-C levels and atherosclerosis. Several studies have shown associations between apoA-I levels and CHD events, but the magnitude of these associations vary by population studied and adjustment for body size and other risk factors.^{14, 15} Multiple factors influence transcriptional control of the apoA-I gene, including dietary factors, endocrine effects, and various pharmacotherapies.^{16, 17} In the general population, apoA-I clearance is the most important determinant of plasma HDL-C and apoA-I levels,¹⁸ though in subjects with similar metabolic profiles apoA-I production rates have a larger effect on HDL concentration.¹⁹ Several other apolipoproteins are found in HDL particles, of which apoA-II is the most abundant (two-thirds of HDL particles). The exact physiologic role of apoA-II and other apolipoproteins, including apoA-IV, apoC (I-III), apoD, apoE, apoJ, apoL-I, and apoM have not been fully elucidated.

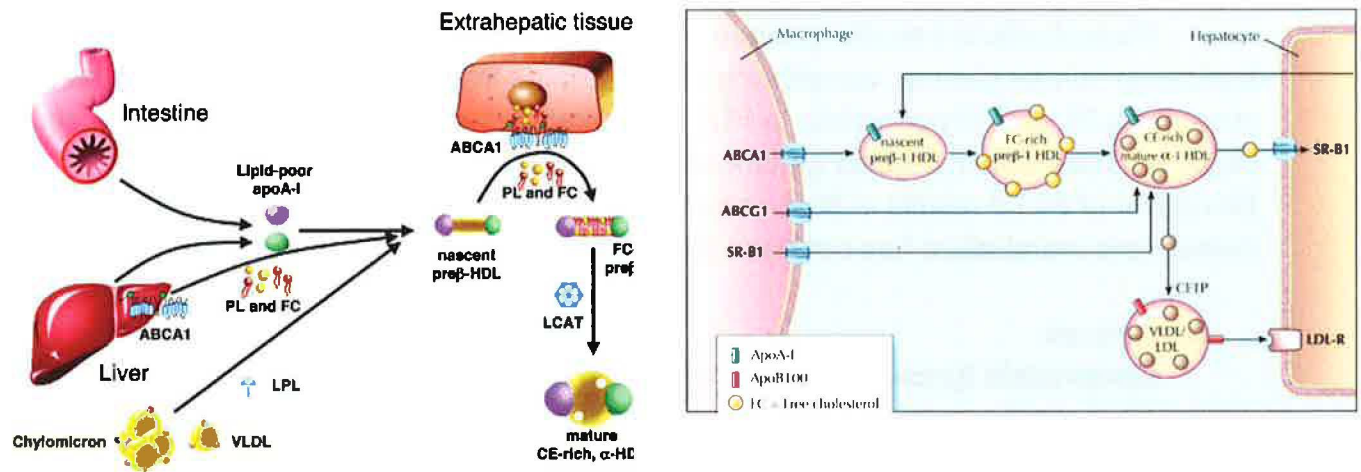
B. Formation of mature HDL and Reverse Cholesterol Transport (RCT)

Lipid-poor apoA-I secreted from the liver and intestinal cells quickly acquire a small amount of lipid in the plasma compartment to form pre-beta HDL particles (pre-beta migration on electrophoresis). Both lipid-poor apoA-I and pre-beta HDL accept cholesterol and phospholipid from peripheral tissues in a step-wise fashion to form mature HDL particles (Figure 4).

The first and key step of RCT and formation of mature HDL involves **ATP-binding cassette transporter-1 (ABCA1)**. In the extracellular space, lipid-poor ApoA-I interacts with ABCA1 on cellular membranes (mostly hepatocytes and macrophages) to mediate lipid transfer (cholesterol and phospholipid) from cell membranes to the ApoA-I particles, generating discoidal-shaped nascent pre-beta HDL particles. The critical importance of ABCA1 in HDL formation and cholesterol homeostasis was demonstrated by patients with Tangier disease who have a mutation in the ABCA1 gene, leading to intracellular cholesterol accumulation, profoundly low HDL-C, and increased risk of CHD, organ dysfunction, and characteristic yellow tonsils.²⁰ ABCA1 animal knockout models have extremely low HDL concentration as well,²¹ but the effects of ABCA1 are tissue-specific. Macrophage-specific ABCA1 deficiency has minimal effects on HDL concentration but profound effects on atherosclerosis.^{22, 23} Liver-specific ABCA1 deficiency, on the other hand, results in marked reductions in HDL concentration (~80%) and over-expression in the liver is associated with increased HDL concentration and resistance to atherosclerosis.^{24, 25}

The cholesterol acquired by HDL is esterified by **Lecithin:cholesterol acyltransferase (LCAT)**, forming a more spherical HDL particle with an inner hydrophobic core surrounded by ApoA-I molecules. These small alpha HDL particles (HDL3) can continue to accept lipid from macrophages via **ABCG1 and ABCG4**²⁶ as well as by aqueous diffusion,²⁷ generating larger, mature alpha HDL particles (HDL2). Like ABCA1, **Scavenger Receptor class B, Type I (SR-BI)** is found on both macrophages and hepatocytes but mediates cholesterol efflux solely to mature HDL instead of nascent HDL particles (Figure 4). Macrophage-specific RCT in vivo, the process theorized to be central to atherosclerosis, is dependent on the transporters ABCA1 and ABCG1 but not SR-BI.²⁸ Interestingly, SR-BI knockout models result in marked increases in HDL concentration but also increased atherosclerosis,²⁹ and macrophage-specific SR-BI deficient mice have normal HDL levels but increased atherosclerosis,³⁰ highlighting the discord between the effects of hepatic and macrophage-specific cholesterol efflux, HDL levels, and atherosclerosis.

Figure 4



McPherson R. Mechanisms of Dyslipidemia; Mechanisms in Medicine Inc. 2004

deGoma, et al. JACC 2008;51:2199-2211

II. Remodeling of HDL (Figure 5)

A. Lipid transfer proteins

The composition of circulating HDL particles in the plasma compartment can be altered by bidirectional transport of cholesteryl ester and phospholipid mediated by lipid transfer factors as well as by lipid hydrolysis by lipolytic enzymes. Once nascent HDL particles acquire free cholesterol from peripheral tissue, the majority is esterified by **lecithin-cholesterol acyltransferase (LCAT)**,³¹ forming a hydrophobic core of cholesteryl ester surrounded by apolipoproteins. This step is critical in the formation of mature HDL particles; humans and mouse models with genetic LCAT deficiency syndromes have marked reduced HDL and apoA-I levels.³² Although critical to HDL formation, its effects on RCT are unclear as unesterified cholesterol in humans can be directly transferred to the liver and secreted in bile.³³

Cholesteryl ester transfer protein (CETP) circulates in the plasma bound to lipoproteins and mediates transfer of cholesteryl ester from HDL to apoB lipoproteins (very low, intermediate, and low-density lipoproteins: VLDL, IDL, LDL; chylomicrons; and remnants) in exchange for triglycerides.³⁴ The net effect of CETP is depletion of cholesteryl ester and reduction in size of HDL particles and HDL concentration. CETP-mediated transfer is accelerated in hypertriglyceridemia and the post-prandial state.^{35, 36} Observations of genetically CETP-deficient Japanese patients having extremely elevated HDL-C and that pharmacologic CETP inhibition markedly increases HDL-C in humans and reduces atherosclerosis in rabbit models have led to the development of a new class of lipid-modifying drugs. However, the effects of CETP-inhibition on atherosclerosis remain unclear given the recent failure of torcetrapib³⁷ and the observation that CETP-mediated transfer of cholesteryl ester from HDL to apoB-containing lipoproteins may serve an important role in RCT (a pathway blocked by CETP inhibition).

III. HDL-C uptake and HDL apolipoprotein catabolism

A. Scavenger receptor B, Type I (SR-BI)

SR-BI is highly expressed in liver, adrenal gland, and ovary and plays a major role in HDL selective lipid uptake in the liver and steroidogenic tissues.²⁹ HDL directly binds to SR-BI and lipid diffuses from the hydrophobic core to the plasma membrane, generating smaller, dense HDL particles. SR-BI knockout models result in markedly elevated HDL-C levels but not of plasma apoA-I and increased atherosclerosis. Conversely, SR-BI over-expression leads to reduced HDL-C and apoA-I levels.⁵² As with the lipases and transfer proteins mentioned above, HDL size and particle composition affect the interaction between HDL and SR-BI. Degradation and catabolism of HDL and apoA-I occur in the liver and kidney.

Disconnect between HDL cholesterol and CHD

The inverse relationship between HDL-C and CHD risk in the general population has not been consistently observed in individuals and families with rare monogenic disorders of HDL-C. Part of the reason for this discordance is the remarkable phenotypic variability in HDL-C, atherosclerosis, and CHD among these individuals. Rare mutations in ABCA1, apoA-I, and LCAT lead to markedly low HDL-C levels, whereas mutations in endothelial lipase (LIPG) and CETP lead to increased HDL-C.

I. Mutations associated with low HDL-C

A. ABCA1 mutations

Individuals homozygous (Tangier disease) or heterozygous (familial hypoalphalipoproteinemia) for ABCA1 mutations have extremely low levels of HDL-C, apoA-I, and impaired cellular cholesterol efflux. Though multiple reports suggest increased incidence of CHD,⁵³ the majority of patients do not develop clinical CHD over the same time periods of follow-up as unaffected family members.⁵⁴⁻⁵⁶ Patients with Tangier disease also have lower total cholesterol, LDL-C, and apoB, which may attenuate the atherogenic profile associated with low HDL-C.

Among those who do develop CHD, ABCA1 heterozygotes and homozygotes develop clinical CHD decades earlier with a dose-response allelic effect.^{54, 55} Surrogate measures of CHD, namely carotid intima media thickness (CIMT), are increased at an earlier age in subjects with heterozygous ABCA1 mutations.⁵⁷ However, consistently higher triglyceride levels associated with ABCA1 mutations may also mediate some of the increased risk of CHD in these individuals, diluting the role of isolated low HDL-C in CHD. In the largest cohort of ABCA1 heterozygotes to date, involving 109 ABCA1 heterozygotes in a total sample size of over 40,000 Danish subjects with over 6500

CHD cases, **ABCA1 heterozygotes had reduced HDL-C by 17 mg/dL but no increased risk of CHD (OR 0.93 95%CI [0.53-1.62]).**⁵⁸ In the overall population studied, a 17-mg/dL decrease in HDL-C was associated with a significantly increased odds ratio of 1.7, *emphasizing the discordant effects of genetically determined low HDL-C and population-based low HDL-C.*

B. ApoA-I mutations

ApoA-I is the major apolipoprotein on HDL and accounts for the majority of the effects of HDL on RCT. Plasma levels of apoA-I inversely correlate with CHD, and, when combined with apoB, outperform all other traditional measures of lipids in predicting CHD risk.^{14, 15} Multiple rare functional mutations in the apoA-I gene have been described that lead to decreased HDL-C, apoA-I, and cholesterol efflux. These mutations have been associated with premature CAD.^{59, 60}

In contrast to these apoA-I mutations, carriers of the apoA-I_{Milano} mutation have greater cholesterol efflux⁶¹ and protective effects against lipid oxidation⁶² than normal apoA-I despite significantly reduced HDL-C and apoA-I levels. The originally described kindred in Italy had a lack of clinically evident CHD despite an atherogenic profile of low HDL-C and high triglyceride levels.^{63, 64} Using CIMT, it was found that carriers of apoA-I_{Milano} have paradoxically similar atherosclerosis to controls,⁶⁵ but carriers of other apoA-I mutations had increased CIMT, as expected from life-long low HDL-C and apoA-I.^{65, 66}

C. LCAT mutations

Mutations in the LCAT gene lead to two distinct syndromes, familial LCAT deficiency with complete loss of LCAT activity and fish-eye disease characterized by loss of LCAT activity on HDL alone. Both syndromes lead to markedly reduced HDL-C, smaller, dense HDL particles, and corneal opacification. Studies of LCAT mutations and CHD in humans have been conflicting. Prior reports suggested no increased incidence of premature CHD in familial LCAT deficiency but increased risk in fish-eye disease.³² Several studies have shown increased CIMT in patients with genetically determined LCAT deficiency,^{67, 68} while other recent studies have not.^{69, 70} In the most recent and comprehensive study of 13 Italian families with LCAT deficiency, increasing copy number of LCAT mutations was associated with paradoxically decreased CIMT,^{69, 70} in stark contrast to prior reports but consistent with recent findings that LCAT deficiency does not affect cellular cholesterol efflux.⁷¹ Furthermore, in a nested case-control study, high plasma LCAT activity was not associated with decreased events but was associated with increased risk among subjects with high HDL-C.⁷²

II. Mutations associated with high HDL-C

A. Endothelial lipase mutations (LIPG)

The most frequently studied LIPG mutation in humans has been the common variant, Thr111Le. Studies have reported conflicting results on its association with HDL-C levels and no association with CHD.^{73, 74} On the other hand, in several large case-control cohorts, the rare mutation Asn396Ser was found to be associated with a 8 mg/dL increase in HDL-C, increased HDL particle size and number of large particles, and increased apoA-I levels.⁷³ Unlike other mutations that affect HDL-C, the Asn396Ser variant was not associated with any other lipids, including LDL, triglyceride, apoB, or other lipoprotein particle sizes. An analysis of the association between this mutation and CHD in a large pooled data set will be forthcoming to shed light on the causal role of HDL-C in CHD.

B. CETP mutations

In 1985 a 58 year-old Japanese man with an HDL-C of 301 mg/dL and his sister with HDL-C of 174 mg/dL were reported to have absence of CETP activity.⁷⁵ Several rare (Japanese) and common mutations in CETP are associated with increased HDL-C levels, but studies reporting association with CHD risk have been conflicting. In a recent meta-analysis involving 27,196 CHD cases and 55,338 controls, OR per allele for three common mutations were associated with modest increases in HDL-C and apoA-I (3-5%) and were modestly protective for CHD: OR 0.95 (95%CI 0.92-0.99) for TaqIb, 0.94 (95%CI 0.89-1.00) for I405V, and 0.95 (95%CI 0.91-1.00) for -629C>A.⁷⁶ Similarly in the Women's Genome Health Study, a genome wide scan revealed that the CETP mutation, 16q3, was the only mutation associated with both HDL-C and CHD: a per-allele increase in HDL of 3.1 mg/dL and a modest non-significant protective effect on incident CHD, HDL-adjusted HR of 0.84 (95%CI 0.68-1.03).⁷⁷ Interestingly half the 20 mutations in CETP in this study were significantly associated with higher HDL-C, as expected with loss of CETP function, and half were associated with significant paradoxical decreases in HDL-C.

In contrast to findings in genetically determined CETP deficiency states, pharmacologic CETP inhibition with torcetrapib had a much greater effect on increasing HDL-C (↑ 72%) and lowering LDL(↓ 24%) but was associated with a significant 25% increased risk of CV events and 58% increased risk of death.³⁷ Similarly contrasting findings were seen when plasma CETP activity was measured in the Framingham Heart Study; higher CETP activity was associated with reduced risk of CHD (OR per SD 0.86, 95%CI 0.76-0.97; HDL-adjusted),⁷⁸ calling into question the presumed cardiac benefits of CETP inhibition.

Summarizing findings from human subjects with genetically-determined low or high HDL-C, the effects on CHD are not uniform and sometimes paradoxical to the inverse associations seen between HDL-C and CHD in population studies. Except for the Asn396Ser endothelial lipase variant, most other variants affecting HDL-C levels

also affect other lipid parameters that presumably may either explain or attenuate associations with CHD.

III. Discordances between therapies that alter HDL-C and CHD

Most of the risk reduction in CHD due to lipid-modifying therapies has been attributed to LDL lowering, with statins being the most consistent and potent drug class showing this effect. As shown before, statins overall are associated with a 6-8% increase in HDL, and patients with significant LDL lowering as well as increases in HDL seem to benefit most when individual trials have been examined. Interestingly, the highest statin doses are associated with a slight reduction in HDL despite more CHD risk reduction than medium dose statins.⁷⁹ However, some therapies such as fibrates and the glitazones have shown variable effects on CHD despite consistent and more potent increases in HDL-C.^{80, 81} In addition, estrogens and torcetrapib are examples of drugs that significantly increase HDL but are associated with increased CHD risk.

A meta-regression study analyzed over 108 RCTs of lipid-modifying therapies to assess whether changes in HDL on therapy were associated with CHD events, independent of LDL and drug class.⁷⁹ The study found that change in LDL was associated with CHD risk reduction regardless of HDL level or drug class and explained a significant portion of the benefit. On the other hand, change in HDL did not significantly correlate with CHD risk adjusted for LDL and drug class.

When analyses were restricted to those therapies that are known to significantly raise HDL and excluded those associated with CHD harm, change in HDL was significantly associated with risk reduction but that association was completely abolished when adjusted for LDL and drug class, and, in fact, trended to increased harm. Restricting to studies of drugs specifically chosen to raise HDL (niacin combinations and fibrates) eliminated the trend to harm but still showed no association between change in HDL on therapy and CHD, unadjusted or adjusted for LDL and drug class.

The conclusion from this meta-regression of over 100 RCTs of lipid-modifying therapies was that change in HDL on treatment did not account for the CHD reduction in RCTs, when adjusted for LDL. Limitations include the omission of the Coronary Drug Project⁸² (niacin associated with CHD benefit but no HDL-C measured) and the Jupiter study⁸³ (rosuvastatin) as well as combining niacin and fibrate studies in the sensitivity analyses. The signal for increased harm with increased HDL adjusted for LDL will have to be further corroborated but does suggest again a limitation of using HDL-C as a measure of response to therapy.

Heterogenous HDL particle composition and function

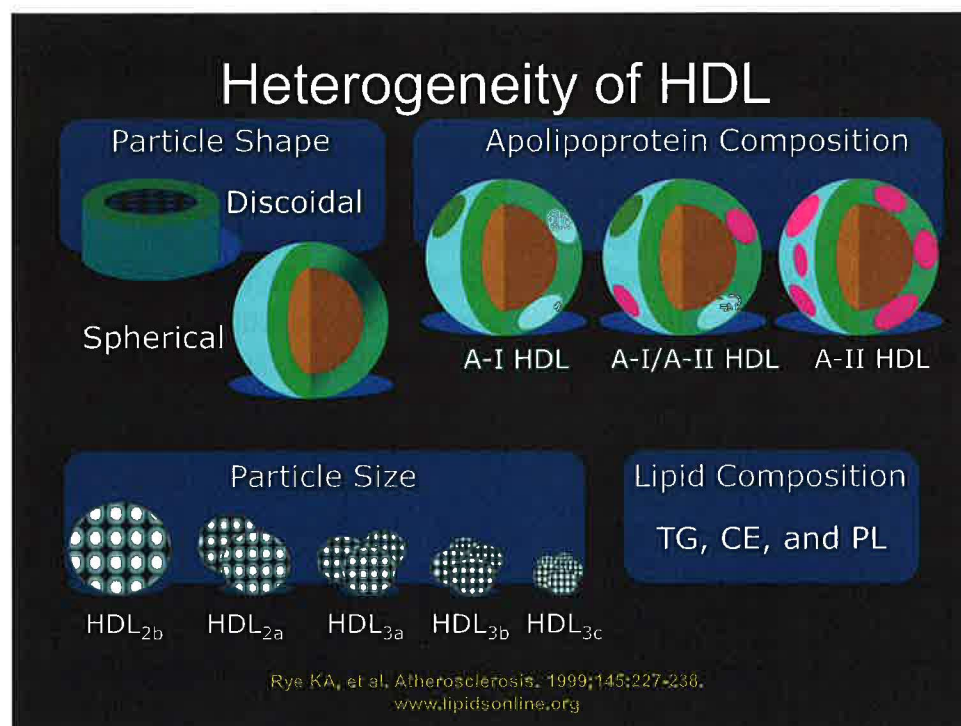


Figure 6

As mentioned above, HDL particles vary significantly according to shape, size, protein, and lipid composition (Figure 6).⁸⁴ Several methods exist for determining various HDL subspecies. Ultracentrifugation has long been the gold standard for determining lipoprotein subspecies by density (HDL: 1.063 – 1.21 g/mL). Gradient-gel electrophoresis discerns subspecies by size and NMR by proton signal, giving rise to 3-5 HDL species.⁸⁵

2-dimensional gel electrophoresis is a more recently developed method that allows more quantitative characterization of HDL subspecies. In this method, plasma or serum is electrophoresed in 2 phases, resulting in 12 distinct Apo-AI containing HDL subspecies that vary by size and charge (alpha and beta migrating).⁸⁶ In individuals with normal total and LDL levels, those with low HDL have significantly lower levels of apoA-I in the large mature alphaA1 particles and higher levels of apoA-I in the small, nascent pre-beta1 particles.⁸⁷ In the Framingham offspring cohort study, 2D-gel electrophoresis was performed in 169 CHD male cases and 1277 matched controls.⁸⁸ Despite matching by HDL-C, levels of HDL subspecies markedly differed, with less large mature HDL particles and more nascent pre-beta1 and small, dense mature HDL particles in subjects with CHD. The strongest HDL particle association was with large alpha1, showing an inverse correlation with prevalent CHD, adjusted for traditional risk factors and lipids levels, including triglyceride levels. However, there was some discordance in

subspecies measurements, where pre-alpha 1 was lower in pts w/ CHD but conferred increased risk with increasing levels when adjusted for risk factors and other lipids.

The protein make-up of HDL particles is also quite heterogeneous. Most of the apolipoprotein on the surface of HDL is apolipoprotein A-I (apoA-I). Various other apolipoproteins make up a smaller fraction of the remaining apolipoprotein component and have diverse biological activities as well. The recent advent of protein analysis by mass spectroscopy, or proteomics, has allowed remarkable discrimination in detecting proteins on the surface of HDL. The HDL proteome appears to have as many as 30 different proteins, many of which are not standard apolipoproteins known to be associated with HDL.^{89, 90} These types of analyses suggest that the heterogeneous protein makeup of HDL reflects heterogeneous HDL functionality, including lipid metabolism, protease action, and immunity. Investigations into these diverse HDL functions are yielding new insights into how HDL may exert its anti-atherogenic effects and how therapeutic modulation can alter these functions in relation to HDL-C levels.

HDL Function

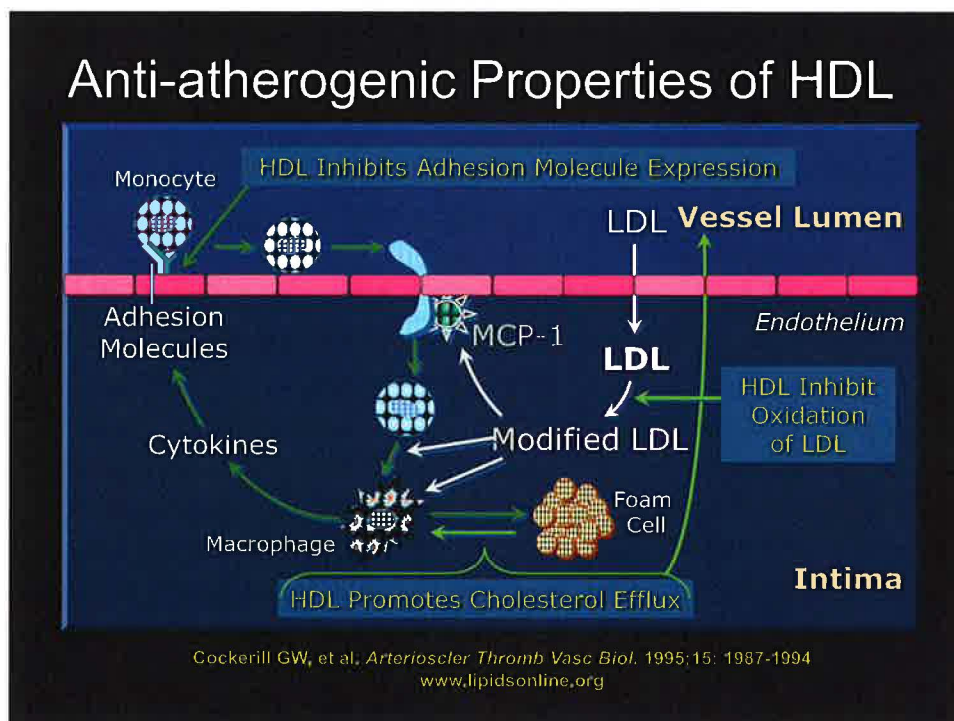


Figure 7

The process of atherosclerotic plaque development in the arterial wall involves multiple steps, including uptake of oxidized LDL by macrophages in the intima, increased cytokine production leading to endothelial cell adhesion molecule expression and recruitment of monocytes to the endothelial surface, and transmigration into the intimal space and foam cell formation. The effects of HDL on these steps are

considered to be anti-atherogenic but can vary significantly in various disease states and in response to therapy (Figure 7).

A. Cholesterol Efflux (Reverse Cholesterol Transport – RCT)

The concept of reverse cholesterol transport involves movement of cholesterol from peripheral tissue into the plasma compartment, delivered to the liver and excreted through bile and feces. Because this is a highly dynamic process, quantifying cholesterol efflux may give better insight into the functional activity of HDL. There are several ways of quantifying cholesterol efflux, and only recently have systems been developed and optimized to study human HDL. By loading or incubating cells with radiolabeled cholesterol, movement of this labeled pool into various compartments can be measured, including an extracellular medium, the plasma compartment, and feces. In terms of testing human blood and HDL, the ex vivo cellular efflux models have developed scientific momentum in assessing the RCT transport properties of HDL.

Ex vivo cellular cholesterol efflux models have used 3 main cell lines, macrophages, hepatoma, and fibroblast cells, to quantify the movement of radiolabeled cholesterol from within the cells to an extracellular acceptor (Figure 8).²⁷ Acceptors that have been studied include purified or human isolated HDL or apoA-I, whole serum or plasma. Since the predominant mechanism related to atherosclerosis is dependent on macrophage-specific cholesterol efflux, that cellular model has potential promise for human investigation.

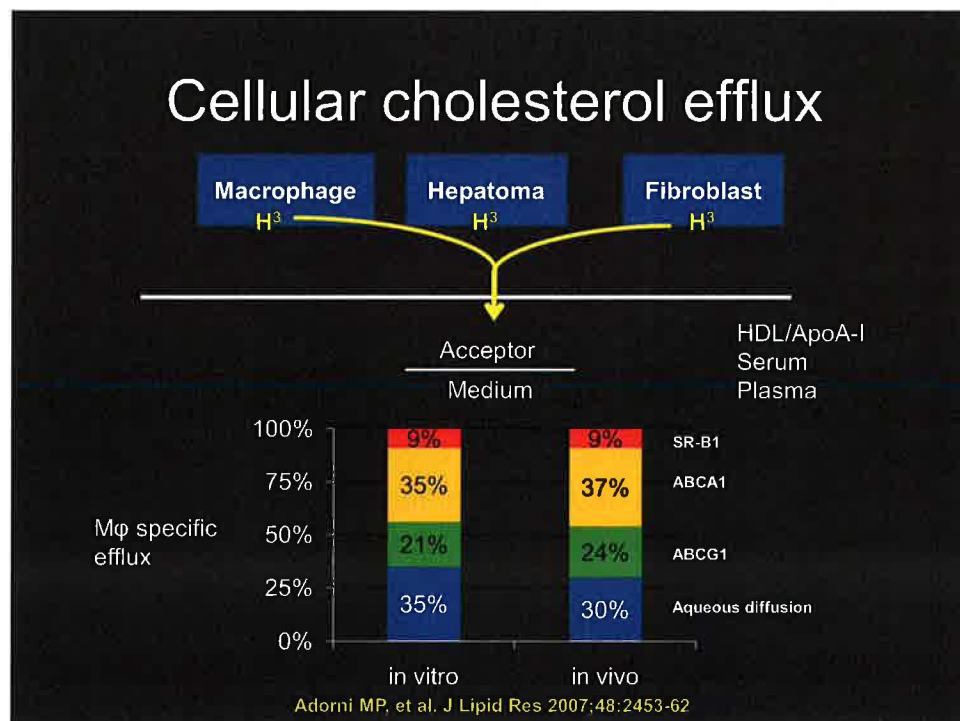


Figure 8

As discussed before, macrophages contain several different transporters that can mediate cholesterol efflux, including ABCA1, the predominant transporter of intracellular free cholesterol to nascent HDL, and ABCG1 and SRB1, which transport cholesterol to mature HDL particles. Reassuringly, the ex vivo system correlates well with in vivo murine models of macrophage cholesterol efflux, indicating that ABCA1 and ABCG1 contribute significantly to macrophage-specific efflux, but not SR-B1. In addition, unmeasured efflux, presumably reflecting passive aqueous diffusion, also contributes to efflux when cells are loaded with cholesterol, and is responsible for the majority of macrophage efflux in cholesterol normal cells. The advantage of measuring macrophage RCT is that it better correlates with atherosclerosis than HDL-C levels in animal models. In mouse studies, genetic manipulations and pharmacologic interventions leading to increased macrophage RCT led to decreased atherosclerosis and vice versa, except for ABCG1 KO models.⁹¹ In these same mice, the relationship between changes in HDL-C levels and RCT was not as robust, and only the Apo-A1 models showed concordant relationships between HDL-C levels and RCT. Studies using HDL from human samples show that there is a wide distribution of macrophage-specific cholesterol efflux in subjects with similar HDL-C levels,⁹² corroborating the discordance seen in mouse models. Whether macrophage-specific cholesterol efflux is a better correlate with atherosclerosis in humans has yet to be determined.

B. Vascular Inflammation - Cell-Adhesion Molecules

The increased expression of intercellular adhesion molecule-1 (ICAM-1), vascular cell adhesion molecule-1 (VCAM-1), and E-selectin on endothelial cells is a necessary component of the chronic inflammatory process of atherosclerosis.⁹³ Adhesion molecules tether monocytes in the circulation, causing them to roll along the luminal surface of endothelial cells and transmigrate into the sub-endothelial space where they can acquire free cholesterol and become resident foam cells. The foam cells and oxidized lipids stimulate cytokines that contribute to further increased expression of adhesion molecules, setting up a vicious cycle of inflammation in the arterial wall.

The effects of HDL on the endothelium and vascular inflammation have been recently characterized. In a pivotal study by Philip Barter's group in Australia, human umbilical vein endothelial cells were incubated with HDL isolated from healthy individuals, and adhesion molecule expression was assessed in response to TNF-alpha activation.⁹⁴ They found that the cytokine-induced expression of endothelial cell adhesion molecules was reduced in a dose-dependent manner with increasing concentrations of HDL, and, at physiologic levels, was almost completely inhibited. The same group went on to show that the in vitro protective effects of HDL on endothelial adhesion molecule expression were also seen in an in vivo model of vascular inflammation by inserting compressive collar rings around the carotid arteries of anesthetized rabbits and then measuring the amount of adhesion molecule expression

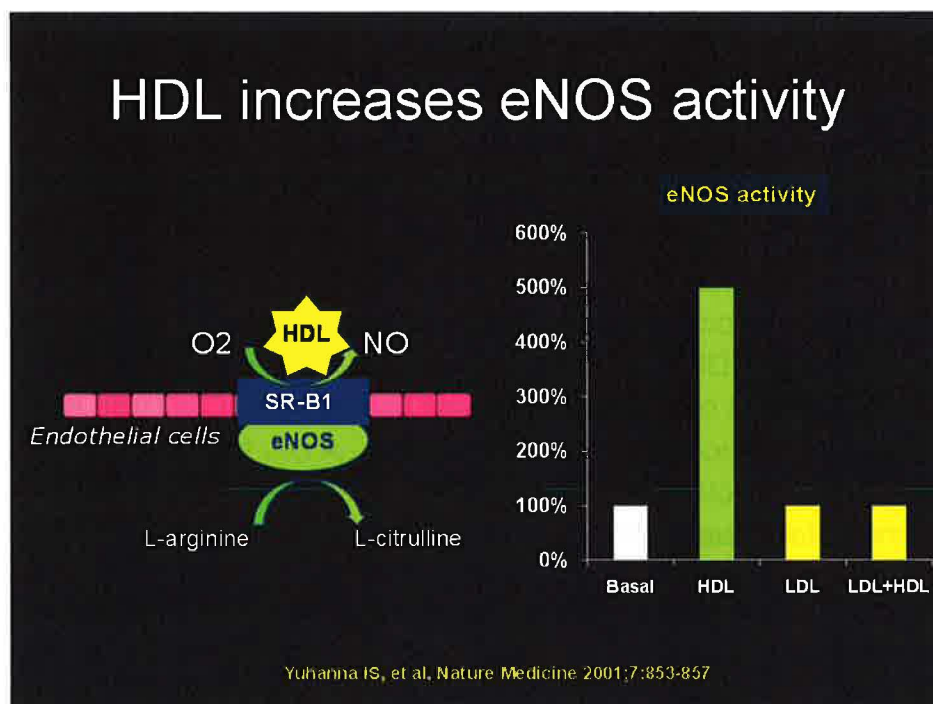
after sacrifice.⁹⁵ Insertion of the collar induced a robust neutrophil response in the carotid vessel wall that was abolished by infusion of recombinant HDL (rHDL). Similarly, the markedly increased expression of VCAM-1 and ICAM-1 was also abolished with infusion of rHDL.

Further confirming the role of HDL in inhibiting vascular inflammation, Fogelman's group in UCLA developed a monocyte chemotaxis assay assessing the migration of human-derived monocytes across human aortic endothelial cells.⁹⁶ They found that the increased monocyte migration in response to LDL was inhibited in a dose-dependent fashion by HDL from normal subjects but was unaffected by HDL from patients with CHD. They went on to show that even patients with CHD with high HDL had impaired inhibition of monocyte chemotactic activity. A cell-free assay has been developed assessing the effects of HDL on lipid oxidation which shows similar results to the monocyte chemotaxis assay and may be amenable for large-scale clinical testing if replicated and validated in larger cohorts.⁹⁶

C. Endothelial Nitric Oxide (NO)

Nitric oxide (NO) produced by the endothelium is a potent vasodilator and has multiple effects on endothelium and vascular smooth muscle. In atherosclerotic vascular disease, levels of endothelium-derived NO are significantly reduced, leading to increased neutrophil adherence, enhanced smooth muscle cell proliferation, and platelet aggregation and adhesion.⁹⁷ NO is produced by endothelial nitric oxide synthase (eNOS), and mouse models of NOS antagonism or genetic NOS deficiency lead to accelerated atherosclerosis, supporting the concept that NO is protective and that NO deficiency serves a critical role in atherosclerosis.⁹⁷ Much of the biology of the effects of HDL on endothelium and eNOS has been worked out here at UTSW in the labs of Philip Shaul, Chieko Mineo and others.

Figure 9



In a cell-based system, endothelial production of nitrite and nitrate is generated by incubating endothelial cells with L-arginine, a substrate of nitric oxide synthase (NOS). NO synthase activity can be characterized by measuring the conversion of labeled L-arginine to L-citrulline. Increased eNOS activity in this cell-based system would suggest protective effects on the endothelium and atherosclerosis and may serve as a way of testing the effects of agents on the capacity of human HDL to promote eNOS activity. When HDL from healthy subjects was tested, there was a 5-fold increase in eNOS activity compared to the basal state, and this effect was completely blunted by either LDL alone or adding LDL to HDL (Figure 9).⁹⁸

Shaul and colleagues also assessed the endothelial effects of HDL in an in vivo system and found that HDL caused direct relaxation of phenylephrine precontracted thoracic aorta rings from mice, an effect that was blunted by a NOS antagonist and by denuding the endothelium.⁹⁸ They also determined that the effects of HDL on NOS activation and endothelium-dependent vasorelaxation were mediated by SR-B1 in endothelial cells, the same receptor in liver cells that accepts free cholesterol from mature HDL particles and delivers cholesterol to mature HDL particles from macrophages as well. In mice, this cell-based system has been used to test the effects of novel HDL therapeutics such as apoA-1 mimetic peptides and has been correlated with vasodilatory response in intact animal models.

As an example of how measuring NO production may be used in humans, a recent small study of about 40 mostly male subjects tested the effects of HDL on endothelial NO production (by a different method) in subjects with diabetes compared to healthy controls.⁹⁹ They found that the HDL from patients with diabetes produced

much less endothelial NO and resulted in reduced relaxation of mouse aortic rings (Figure 10).

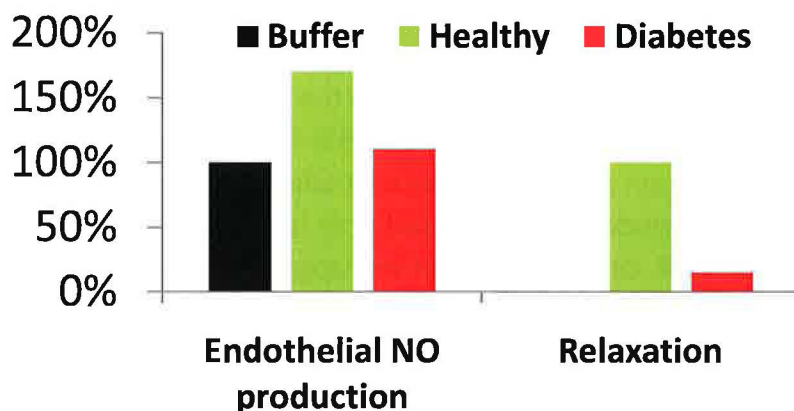


Figure 10

In summary, HDL has significant effects on cholesterol transport and vascular health, including inhibiting inflammation and oxidation and directly promoting endothelial nitric oxide activity, all of which are purported to exert anti-atherogenic effects in healthy individuals and are diminished in diseased states. Quantifying these HDL functions may serve as better biomarkers for predicting CHD and predicting response to therapies that modify HDL. Other effects of HDL include anti-apoptotic and antithrombotic effects, and an increasing understanding of the role of HDL in innate immunity and protection from infection.¹⁰⁰

Novel HDL-modifying therapies

Given the lack of drugs that specifically target HDL metabolism (except for niacin), and the recent failure of the first CETP-inhibitor, torcetrapib, multiple novel compounds are being studied to test the hypothesis that improving HDL-C levels and HDL function will reduce the risk of CHD. Two classes of drugs will be briefly reviewed here, HDL mimetics and CETP-inhibitors, with a concluding summary of the current level of evidence for niacin.

A. HDL Mimetics

Recombinant HDL (rHDL)

Recombinant HDL has been formulated using human apoA-1 complexed with phospholipid at fixed molar ratios, usually infused via peripheral IV at a dose of 80mg/kg over four hours. Several studies have shown beneficial effects of rHDL in humans. In one study of patients with hypercholesterolemia, infusion of rHDL led to restoration of endothelial function as measured by forearm blood flow.¹⁰¹ In another study of 20 patients presenting for peripheral arterial revascularization, patients were randomized to a single 4-hour infusion of 80mg/kg rHDL and then underwent intervention 1 week later.¹⁰² HDL from patients randomized to rHDL exhibited significantly greater

cholesterol efflux, which correlated with markedly reduced cholesterol content and macrophage accumulation in excised arterial plaques. The Melbourne group studied the effects of rHDL in 13 male patients with diabetes.¹⁰³ A single 4-hour infusion resulted in reduced endothelial cell expression of adhesion molecules ex-vivo and reduced platelet aggregation. The infusion resulted in increased capacity of HDL from patients to promote cholesterol efflux from platelets. In contrast to the receptor-dependent effects of HDL on cholesterol efflux via ABCA1, ACBG1, and SR-B1 and on endothelial NOS activation via SR-B1, the effects on platelet cholesterol efflux and platelet activation were found to be receptor-independent and affected only by the phospholipid and cholesterol concentration of rHDL or HDL, and not the apoA-I content. These studies highlight the diverse effects of HDL and rHDL through multiple different pathways.

ApoA-I_{Milano}

Based on the finding of a rare ApoA-I mutant in Italy that led to reduced HDL but no CHD based on a single cysteine to arginine substitution,⁶³ a recombinant form was developed and showed marked reductions in atherosclerosis in animal models, even after a single infusion. In the first human study of ApoA-I_{Milano}, patients who recently had acute coronary syndromes were randomized to receive recombinant ApoA-I_{Milano} or placebo for 5 weekly infusions. The primary endpoint, change in percent atheroma volume by intravascular ultrasound during coronary angiography, was modestly but significantly reduced from baseline in the intervention arm, but the difference was not significant compared to placebo, perhaps because of the small numbers of patients.¹⁰⁴

In a similar study design of patients following acute coronary syndromes, a recombinant form of HDL using apo-A1 derived from wild-type human plasma complexed to soybean-derived phospholipid was tested using intravascular ultrasound derived characteristics as the primary endpoints. In this study 4 weekly infusions were administered, but the high dose arm (80mg/kg) was discontinued due to asymptomatic increases in liver enzymes. Though there were improvements from baseline in coronary plaque characteristics in those randomized to rHDL, these changes were not consistently different from placebo.¹⁰⁵

Autologous HDL delipidation

Another interesting IV HDL therapy is autologous HDL delipidation, whereby the patient's blood is plasmapheresed and returned to the patient after the cholesterol from HDL is stripped using various solvents. In a pilot study of ACS patients assessing tolerability, 7 weekly delipidation sessions led to non-significant reductions in coronary plaque characteristics, but similar in magnitude to the Apo-A1_{milano} trial. The authors suggested that because these effects are larger than those seen with high-dose statins, larger clinical trials investigating clinical endpoints may show potential benefit of such therapies in the acute setting.

In summary, IV infusions of recombinant HDL or autologous HDL delipidation have led to non-significant trends toward improved coronary plaque volume and other characteristics in high-risk patients. Larger trials investigating clinical endpoints will determine whether these resource- and cost-intensive therapies will translate into clinical benefit.

Oral ApoA-I mimetic: D4F

The large size of ApoA-I at 243 amino acids has made it expensive and difficult to synthesize, necessitating IV administration and severely restricting its use to acute and inpatient settings. The development of an 18aa peptide that mimics the helical structure of apoA-I has led to the development of several oral apo-AI mimetics, of which the D4F version has been tested in humans.¹⁰⁶ In animal models, D4F leads to reduced atherosclerosis synergistically with statins and reduced inflammation. In the first human study, a single dose given to patients with CHD or otherwise high-risk patients was tolerated well without significant side effects and led to improvements in inflammatory indices, such as monocyte chemotaxis, compared to baseline.¹⁰⁷ Whether these effects of oral or IV mimetic peptides will translate into true clinical benefit will have to await larger clinical trials.

B. CETP Inhibitors

Torcetrapib was the first CETP-inhibitor to be tested in Phase III clinical trials. As described above, randomization to torcetrapib was associated with increased all-cause and coronary death and no significant change in coronary or carotid atherosclerosis.^{9-11, 37} Despite these disappointing findings, post-hoc analyses suggested that the subjects who achieved the highest HDL-C levels on torcetrapib actually had coronary regression and that other CETP inhibitors without such deleterious adverse events would have potential cardiac benefits.¹⁰⁸ Dalcetrapib is another CETP inhibitor without the deleterious effects on blood pressure and the rennin-angiotensin-aldosterone system associated with torcetrapib. Enrollment has completed in a phase III, double-blind, placebo-controlled multi-center trial of 15,600 patients with acute coronary syndromes. Patients were randomized to dalcetrapib 600mg once daily vs. placebo on top of statin therapy (clinicaltrials.gov). Hard clinical endpoints will be assessed as the primary endpoint, with results projected in 2013. Other CETP inhibitors such as anacetrapib and JTT-705 (vaccine) are in smaller phase II-III clinical studies and also do not have the adverse side effect profile of torcetrapib. All of the CETP inhibitors markedly increase HDL-C levels by 60-100%, increase large HDL particles, and improve cholesterol efflux. Whether these changes in HDL metabolism will lead to clinical benefit will be determined with the results of large clinical trials.

C. Niacin

Clinical Trials

Niacin is a B vitamin and available over the counter. It also comes as an extended release prescription (ER-Niacin, Abbott). The most common side effect is flushing which has been reported in over 60% of subjects with the immediate-release formulation, leading to high rates of discontinuation.

Niacin is the most potent HDL-raising drug currently available for use. On background statin therapy, niacin raises HDL by about 20% and lowers total and LDL by 10-15% and triglycerides by 20-30%.¹⁰⁹ Very few randomized controlled trials have studied the effect of niacin on cardiovascular endpoints. The only study reported in the literature to date that was designed to study actual clinical events was the Coronary Drug Project, completed in 1975.⁸² Several lipid agents were studied in men with existing CHD, including niacin at a dose of 3 grams/d. After 5 years, there was a significant 27% reduction in non-fatal CHD events compared to placebo but no difference in mortality. After 15 years follow-up, after the study had terminated almost a decade before, there was a significant 11% reduction in mortality in those who had been originally randomized to niacin compared to placebo.¹¹⁰

Several other studies have randomized patients to niacin (HATS, ARBITER-2, ARBITER-6),^{109, 111, 112} assessing changes in atherosclerosis. These studies have also been in patients with existing CHD but enrolled relatively small numbers compared to the CDP. Reported coronary events in the niacin group were lower than the comparator groups with borderline significance. It is difficult to extrapolate too much from these data given the low numbers of absolute events. However, of note, both ARBITER studies showed a reduction in events in those randomized to niacin in addition to background low to medium-dose background statin therapy.

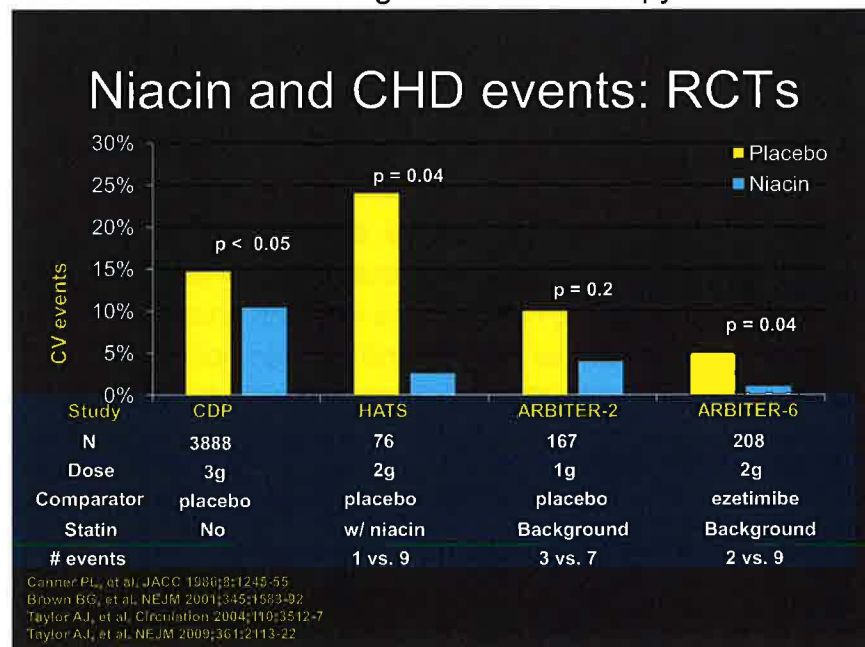


Figure 11

The ARBITER trials assessed changes in CIMT by ultrasound in patients randomized to niacin vs. a comparator in patients with CHD. A true placebo arm without statin showed progression,¹⁰⁹ ezetimibe with statin showed no significant change,¹¹² and niacin with statin showed regression of CIMT.^{109, 112} Another small study of patients with CHD using MRI instead of ultrasound showed similar regression of carotid wall area with niacin compared to placebo on top of statin.¹¹³ The HATS study showed significant regression in coronary stenosis in patients randomized to simvastatin plus niacin compared to placebo.¹¹¹

Mechanisms of Niacin's benefits

The effects of niacin on apolipoprotein B containing lipoproteins are believed to be primarily mediated through inhibition of triglyceride synthesis in the liver, leading to decreased production of all apoB-containing compounds, including LDL, VLDL, and Lp(a).¹¹⁴ The mechanism of action leading to increased HDL is less well worked out, but is believed to be due to decreased catabolic rate of HDL and ApoA-1, possibly by blocking whole particle uptake by the liver and extending the half-life of circulating HDL and apoA-1.¹¹⁴

Not only does niacin reduce atherogenic lipoprotein levels and increase HDL-C, it also improves multiple indices of HDL function. Niacin has been shown to increase ABCA1 expression in vitro in the liver and human-derived monocytes and reduce cellular cholesterol content.¹¹⁵ Niacin has also been shown to increase cholesterol efflux by as much as 28% compared to placebo, likely due primarily to increased HDL-C.¹¹⁶ A recent study extensively evaluated the effects of niacin on endothelial function in patients with diabetes. In this study, 30 subjects with diabetes were randomized to placebo or ER-niacin titrated up to 1500mg/d for 3 months. Compared to placebo, niacin was associated with a 22% increase in HDL-C and a significant 50% increase in the effect of patient HDL to promote endothelial NO production. These findings were correlated with a significant increase in flow-mediated dilation of the radial artery in these same subjects as measured by a non-invasive ultrasound device pre- and post-arterial occlusion.⁹⁹

Niacin also directly inhibits acute vascular inflammation. In another recently published study, the carotid collar model in rabbits was used to test the effect of orally administered niacin on acute vascular inflammation.¹¹⁷ Niacin markedly decreased expression of endothelial cell adhesion molecules and inhibited intima-media neutrophil recruitment and myeloperoxidase accumulation. These changes were independent of plasma lipids and HDL levels and were similar when cells were directly incubated with niacin, suggesting a lipid-independent effect. These studies in aggregate support the hypothesis that niacin's beneficial role in atherosclerosis is likely due to both direct HDL

raising and improved HDL function. Whether these changes will translate into consistent clinical benefits on top of statin therapy will be determined by on-going clinical trials.

D. Active Clinical Trials involving Niacin and CETP inhibitors

	AIM-HIGH	HPS2-THRIVE	DAL-Outcomes
# pts	3300	25000	15600
Age	>45	50-80	>45
Pts	Vasc Dx; metsyn	Vasc Dz	ACS
Dose	ER-niacin 2g	ER-niacin 2g	Dalcetrapib 600mg
Statin	Simva	Simva 40 ± zetia	Yes - Standard care
Outcomes	MACE	MACE	MACE
Results	2011??	2013	2013

Summary and Future directions

Despite inverse associations between HDL-C levels and CHD in population-based studies, genetic syndromes in humans leading to either low or high HDL-C levels do not support a causal role of HDL in CHD. Furthermore, several therapies that raise HDL do not translate into clinical benefit, including torcetrapib (CETP inhibitor), fibrates, and estrogen. These discordances may be explained by the lack of HDL-C levels to capture the heterogeneous composition and function of HDL. Efforts to elucidate the diverse functions of HDL and correlate them to CHD and responses to therapy will facilitate understanding of HDL metabolism and perhaps lead to better therapeutic targets than HDL-C. Ultimately, active randomized clinical trials of HDL-modifying therapies will determine the clinical benefit of modulating HDL.

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