Dietary Fatty Acids Regulate Hepatic Low Density Lipoprotein (LDL) Transport by Altering LDL Receptor Protein and mRNA Levels

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Abstract

The concentration of LDL in plasma is strongly influenced by the amount and the type of lipid in the diet. Recent studies in the hamster have shown that dietary fatty acids differentially affect circulating LDL levels primarily by altering receptor-dependent LDL uptake in the liver. To investigate the mechanistic basis of this effect, rates of receptor-dependent LDL transport in the liver were correlated with LDL receptor protein and mRNA levels in hamsters fed safflower oil or coconut oil and varying amounts of cholesterol. Hepatic LDL receptor activity was significantly lower in animals fed coconut oil than in animals fed safflower oil at all levels of cholesterol intake (26, 53, and 61% lower at cholesterol intakes of 0, 0.06, and 0.12%, respectively). These fatty acid–induced changes in hepatic LDL receptor activity were accompanied by parallel changes in hepatic LDL receptor protein and mRNA levels, suggesting that dietary fatty acids regulate the LDL receptor pathway largely at the mRNA level. (*J. Clin. Invest.* 1993. 92:743–749.) Key words: dietary fatty acids • LDL transport • liver • LDL receptor mRNA

Introduction

A number of risk factors are known to contribute to the genesis of atherosclerosis and coronary heart disease. Of these, an elevated concentration of LDL in the plasma appears to be one of the most important (1). The concentration of LDL in plasma is determined by the rate at which LDL enters the plasma relative to the rate at which it is cleared from plasma by the various tissues of the body. LDL are formed in plasma during the metabolism of VLDL, which in turn are secreted by the liver (2). Tissues take up LDL from plasma by at least two mechanisms. One of these, termed receptor-dependent transport, involves the interaction of LDL particles with cell surface receptors, followed by endocytosis and catabolism of the LDL particle in the lysosomal compartment (3, 4). Tissues also take up LDL by a nonsaturable, receptor-independent process that is thought to represent bulk fluid phase endocytosis (5). In normal animals and humans, receptor-dependent mechanisms account for 70–80% of total LDL turnover (6–9) and the vast majority of receptor-dependent LDL uptake occurs in the liver (10–12). Thus, changes in the concentration of LDL in plasma

are generally due to changes in the rate of LDL production, changes in receptor-dependent LDL uptake by the liver, or changes in both of these processes.

The concentration of total and LDL cholesterol in plasma is strongly influenced by the amount and type of lipid in the diet (13–16). Thus, dietary triglycerides containing predominantly saturated fatty acids raise plasma LDL concentrations when compared with dietary triglycerides containing predominantly unsaturated fatty acids. Dietary cholesterol also raises plasma LDL concentrations, although individual responses vary widely (17). We have previously shown in the hamster that dietary lipids alter plasma LDL concentrations primarily by altering the rate of receptor-dependent LDL uptake by the liver (18, 19). When added to a low fat diet, cholesterol modestly suppresses receptor-dependent LDL uptake by the liver and raises circulating LDL levels. Dietary triglycerides containing predominantly saturated fatty acids greatly augment the suppressive effect of dietary cholesterol on receptor-dependent LDL uptake by the liver, whereas unsaturated fatty acids partially restore receptor activity. Thus, at any level of dietary cholesterol, receptor-dependent LDL uptake by the liver is always higher in animals fed unsaturated fatty acids than in animals fed saturated fatty acids.

The major form of regulation of the LDL receptor pathway demonstrated to date is sterol-mediated feedback repression of LDL receptor gene transcription (20–22). Dietary cholesterol presumably suppresses hepatic LDL receptor activity via this mechanism (23) and, indeed, dietary cholesterol has been shown to reduce hepatic LDL receptor mRNA levels in nonhuman primates (24). How dietary fatty acids regulate receptor-dependent LDL uptake by the liver has not been established. In these studies we examined this question by quantifying hepatic LDL receptor protein and mRNA levels and rates of receptor-dependent LDL transport in hamsters fed triglycerides containing predominantly saturated or unsaturated fatty acids. These studies demonstrate that regulation of hepatic LDL transport by dietary fatty acids is due largely, if not entirely, to changes in LDL receptor protein and mRNA levels.

Methods

Animals and diets. Male Golden Syrian hamsters (Sasco, Inc., Omaha, NE) were housed in colony cages and subjected to light cycling for at least 3 wk before introduction of the experimental diets. The control semisynthetic diet used in these studies contained 20% soy protein, 0.3% DL-methionine, 10% cellulose, 8.5% salt mix, 1% vitamin mix, 0.2% choline bitartrate, 2% corn oil, and 58% corn starch. The experimental diets were prepared by replacing corn starch with the desired amount of triglyceride on a cal/cal basis, assuming 4 cal/g of corn starch and 9 cal/g of triglyceride. The triglycerides used in these studies were safflower oil and hydrogenated coconut oil. The fatty acid composition of the safflower oil, as determined by capillary gas-liquid chromatography, was 7% as 16:0, 2% as 18:0, 12% as 18:1, and 78% as 18:2. The hydrogenated coconut oil contained 9% as 8:0, 6% as 10:0, 51% as

12:0, 17% as 14:0, 8% as 16:0, and 8% as 18:0. The diets were fed ad lib and all studies were carried out during the mid-dark phase of the light cycle.

Determination of hepatic LDL uptake rates in vivo. Plasma was obtained from normocholesterolemic hamster and human donors. The LDL was isolated from plasma by preparative ultracentrifugation in the density range of 1.020–1.055 g/ml and labeled with ^{125}I- or ^{131}I-tyramine cellobiose as previously described (25). The human LDL was also reductively methylated to completely eliminate its recognition by the LDL receptor (26). Rates of hepatic LDL uptake were measured using a primed infusion of ^{125}I-tyramine cellobiose–labeled LDL. The infusions of ^{125}I-tyramine cellobiose–labeled LDL were continued for 4 h, at which time each animal was administered a bolus of ^{131}I-tyramine cellobiose–labeled LDL as a volume marker and killed 10 min later by exsanguination through the abdominal aorta. Samples of the liver along with aliquots of plasma were assayed for radioactivity in a gamma counter (Packard Instruments, Meriden, CT). The amount of labeled LDL in the liver at 10 min (^{131}I disintegrations per minute per gram of liver divided by the specific activity of ^{131}I in plasma) and at 4 h (^{125}I disintegrations per minute per gram of liver divided by the specific activity of ^{125}I in plasma) was then calculated. The increase in the tissue content of LDL cholesterol or LDL protein with time represents the rate of LDL uptake in micrograms of LDL cholesterol or LDL protein taken up per hour per gram of tissue.

Since receptor-dependent LDL uptake by the liver is saturable and since plasma LDL concentrations varied widely among the different experimental groups, changes in receptor-dependent LDL uptake could not be directly equated with changes in LDL receptor activity (27). To relate changes in receptor-dependent LDL uptake to changes in LDL receptor activity, the experimentally determined uptake rates were superimposed on kinetic curves describing the relationship between hepatic LDL uptake and circulating LDL concentrations in normal animals. By relating the rates of receptor-dependent and receptor-independent LDL uptake in the experimental animals to these normal kinetic curves, it was possible to determine how the various dietary manipulations affected LDL receptor activity (defined as the rate of receptor-dependent LDL uptake in an experimental animal relative to the rate of receptor-dependent LDL uptake seen in control animals at the same plasma LDL concentration).

Determination of LDL receptor protein levels. Liver membrane proteins were solubilized essentially as described by Schneider et al. (28) and solubilized LDL receptor protein determined by immunoblotting. Samples of hamster liver were homogenized in 5 vol of homogenization buffer (50 mM Tris/maleate, pH 6.5, 150 mM NaCl, 2 mM CaCl$_2$, 1 mM PMSF, and 2 μM leupeptin) using a Dounce homogenizer (10 strokes with the loose and 5 with the tight pestle). The homogenate was centrifuged at 1,000 g for 10 min and the supernatant was centrifuged at 8,000 g for 20 min. The 8,000-g supernatent was centrifuged at 104,000 g for 60 min. The pellet was suspended in a buffer containing 250 mM Tris/maleate, pH 6.5, 2 mM CaCl$_2$, 1 mM PMSF, and 2.5 μM leupeptin. An equal volume of a buffer containing 2% Triton X-100, 320 mM NaCl, and 2 mM CaCl$_2$ was then added. The suspension was stirred at 4°C for 15 min and then centrifuged at 104,000 g for 1 h. Aliquots of the supernatent containing 500 μg protein were adjusted to 2% SDS and 0.2 M sucrose and loaded onto a 7.5% polyacrylamide gel. After electrophoresis (29), proteins were transferred to polyvinyldifluoride (PVDF) membrane (Immobilon-P; Millipore Corp., Bedford, MA) with 50 V for 14 h in buffer containing 25 mM Tris, pH 8.3, 192 mM glycine, 0.01% SDS, and 20% (vol/vol) methanol. After electrophoretic transfer, membranes were blocked with 5% nonfat dried milk and incubated with a 1:200 dilution of LDL receptor antiserum. Polyclonal, monospecific antiserum was prepared in New Zealand White rabbits against a synthetic peptide corresponding to the COOH-terminal 13 amino acids of the hamster LDL receptor (30). After incubation with the primary antibody, the PVDF membrane was incubated with 0.5 μCi/ml ^{125}I-labeled donkey anti–rabbit antibody (Amersham Corp., Arlington Heights, IL). The radiolabeled bands were identified by autoradiography, excised, and assayed for radioactivity in a gamma counter. Identically sized pieces of membrane from above and below the radiolabeled bands were also counted as a measure of nonspecific background radioactivity.

Determination of LDL receptor mRNA levels. Hepatic LDL receptor and glyceraldehyde-3-phosphate dehydrogenase (GAPDH,[1] used as an invariant control) mRNA levels were determined by nuclease protection as previously described (31, 32). cDNA probes were not available for Syrian hamster LDL receptor or GAPDH. Therefore, after reverse transcriptase synthesis of the complementary DNA, the PCR was used to amplify sequences encoding fragments of the LDL receptor and GAPDH cDNAs from Syrian hamster liver RNA. Oligonucleotide primers used to amplify the LDL receptor (5'-AAAGGA-TCCGTAGACTGGATCCATGGCAACATCTACTGGAC-3' and 5'-AAAGAATTCATAGATGGCCAAGGAGAAGGGGTG-3') and GAPDH (5'-AAAGGATCCACTGGCGTCTTCACCACCATG-GAG-3' and 5'-AAAGAATTCGTCATGGATGACCTTGGC-CAGGGG-3') sequences were selected from areas of 100% homology in the published rat (33) and Chinese hamster (30) sequences (LDL receptor) or rat (34), human (34), and European hamster (35) sequences (GAPDH). The PCR was carried out sequentially for 5 min at 55°C, 2 min at 72°C, and 45 s at 95°C for 30 cycles in a programmable thermal controller (MJ Research, Inc., Watertown, MA). The oligonucleotide primers were synthesized with restriction sites (BamHI and EcoRI) to allow direct subcloning of the amplified DNA into the plasmid pGEM (Promega Corp., Madison, WI) for sequencing and into the bacteriophage M13 (Pharmacia LKB Biotechnology Inc., Piscataway, NJ) for the preparation of ^{32}P-labeled single-stranded probes. Probes were synthesized as previously described (32) using 0.5 μM [^{32}P]dCTP and 1 μM (LDL receptor) or 300 μM (GAPDH) unlabeled dCTP. The sizes of the full-length, undigested, single-stranded cDNA probes, including M13 sequences, were 500 nucleotides for the LDL receptor and 240 nucleotides for GAPDH.

Samples of hamster liver were homogenized in guanidinium isothiocyanate and the RNA was isolated by centrifugation on cesium chloride. Total RNA (40 μg) was hybridized with the ^{32}P-labeled cDNA probes simultaneously at 48°C overnight. Unhybridized probe, present in excess relative to the amount of specific mRNA, was then digested with 40 U of mung bean nuclease (GIBCO BRL, Gaithersburg, MD). The mRNA-protected ^{32}P-labeled probes were separated on 7 M urea, 6% polyacrylamide gels together with ^{32}P-labeled MspI-digested pBR322 size standards and identified by autoradiography. The radiolabeled bands were excised and assayed for radioactivity by liquid scintillation spectroscopy. Identically sized bands from samples containing no RNA were also counted as a measure of nonspecific background radioactivity. The levels of GAPDH mRNA did not vary with dietary changes and were used to correct for any procedural losses.

Determination of liver and plasma cholesterol distribution. Hepatic esterified and unesterified cholesterol were separated using silicic acid/celite columns and quantified by capillary gas-liquid chromatography (36). The cholesterol distribution in plasma was determined by gel filtration chromatography using a Superose 6 column (Sigma Chemical Co., St. Louis, MO). 2-ml aliquots were collected and assayed for cholesterol using an enzymatic kit (Boehringer Mannheim Corp., Indianapolis, IN).

Results

Dietary fatty acids affect plasma LDL concentrations primarily by altering receptor-dependent LDL uptake in the liver. These studies were undertaken to determine the extent to which changes in LDL receptor number and mRNA levels contribute to these fatty acid–induced changes in hepatic LDL transport. Groups of animals were fed diets containing predominantly

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saturated (coconut oil) or unsaturated (safflower oil) fatty acids and varying amounts of cholesterol (0, 0.06, or 0.12%) for 6 wk. At the end of the feeding period, rates of receptor-dependent LDL uptake by the liver were correlated with LDL receptor protein and mRNA levels. Fig. 1 shows the effect of the experimental diets on plasma LDL cholesterol concentrations. On the cholesterol-free diet, plasma LDL cholesterol concentrations were significantly higher in animals fed the saturated triglyceride than in animals fed the unsaturated triglyceride (55 vs 31 mg/dl, $P < 0.01$). The differential effect of saturated and unsaturated lipids was greater when modest amounts of cholesterol were added to the diet. In animals fed 0.06% cholesterol ($\sim$ 120 mg/kcal), mean plasma LDL-cholesterol concentrations equaled 148 and 65 mg/dl ($P < 0.01$) on the saturated and unsaturated triglyceride diets, respectively. In animals fed 0.12% cholesterol ($\sim$ 240 mg/kcal), the corresponding values were 262 and 104 mg/dl ($P < 0.01$).

Absolute rates of total and receptor-independent LDL uptake by the liver were measured in vivo using homologous and methylated human LDL, respectively. Receptor-dependent LDL uptake was taken as the difference between total and receptor-independent uptake. Since receptor-dependent LDL uptake is saturable, and since mean plasma LDL concentrations varied by nearly 10-fold among the different experimental groups, changes in absolute rates of LDL uptake by the liver could not be equated directly with changes in receptor activity. To calculate changes in hepatic LDL receptor activity, it was necessary to relate the changes in absolute rates of LDL uptake in the experimental animals to kinetic curves describing the relationship between LDL uptake and circulating LDL concentrations in control animals.

Fig. 2 shows the kinetic curves for normal hepatic LDL transport in control hamsters. The shaded areas represent the relationship between total (stippled) and receptor-independent (hatched) LDL uptake and plasma LDL concentrations over the range of LDL concentrations observed in these studies. These kinetic curves were previously established by quantifying rates of total and receptor-independent LDL uptake in control animals under conditions where plasma LDL concentrations were acutely raised and maintained at various levels by infusions of unlabeled LDL (27). Superimposed on these standard kinetic curves are the mean rates of total and receptor-independent LDL uptake determined in these studies. Values for animals fed safflower oil are shown in the top panel of Fig. 2 and values for animals fed coconut oil are shown in the bottom panel.
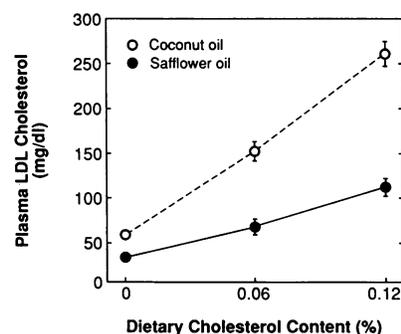
Figure 1. Plasma LDL cholesterol concentrations in animals fed saturated and unsaturated fatty acids. Animals were fed a semisynthetic diet supplemented with 20% (by weight) safflower oil or coconut oil and varying amounts of cholesterol for 6 wk. Each value represents the mean±SE for data obtained in 10–12 animals. Differences between the safflower and coconut oil groups were significant ($P < 0.01$) at each level of dietary cholesterol.
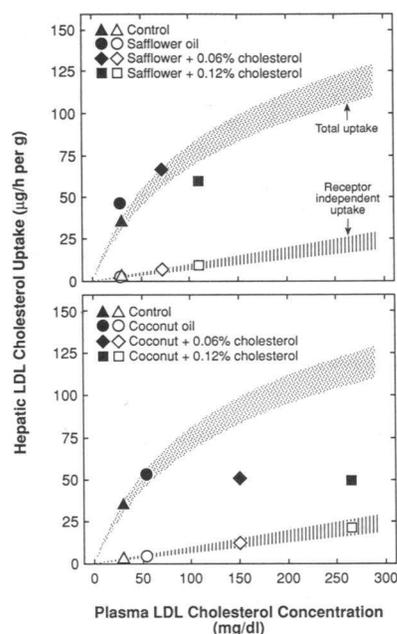
Figure 2. Determination of hepatic receptor-dependent LDL transport. The shaded areas represent the kinetic curves for total (stippled) and receptor-independent (hatched) LDL cholesterol uptake determined in control animals as described in Methods. Superimposed on these standard kinetic curves are the absolute rates of total and receptor-independent LDL cholesterol uptake in animals fed the safflower oil (top) or coconut oil (bottom) diets plotted as a function of the plasma LDL cholesterol concentration in the same animals. Each point represents the mean for data obtained in 10–12 animals.

In control animals, rates of total (solid triangles) and receptor-independent (open triangles) LDL cholesterol uptake equaled 36 and 3 μg/h per g, respectively, at a plasma LDL cholesterol concentration of 31 mg/dl. These values fell on the standard kinetic curves for LDL transport, and hepatic LDL receptor activity in these animals was assigned a value of 100%. Hepatic LDL receptor activity in each of the experimental groups was then calculated by dividing the rate of receptor-dependent LDL uptake in the experimental animals by the rate of receptor-dependent LDL uptake that would be seen in normal animals at the same plasma LDL cholesterol concentration. For example, as illustrated in the bottom panel of Fig. 2, rates of total (solid circle) and receptor-independent (open circle) LDL cholesterol uptake in animals fed coconut oil equaled 53 and 4 μg/h per g, respectively, at a plasma LDL cholesterol concentration of 55 mg/dl. These values were not displaced from the standard kinetic curves, indicating no change in the receptor-dependent or -independent pathways. In contrast, the rate of total LDL cholesterol uptake in animals fed coconut oil plus 0.12% cholesterol (solid square) equaled 50 μg/h per g at a plasma LDL cholesterol concentration of 278 mg/dl, whereas normal animals would transport $\sim$ 118 μg/h per g at this LDL concentration. Since receptor-independent LDL transport was normal in these animals (21 μg/h per g), the reduction in total LDL cholesterol uptake could be attributed entirely to a reduction in receptor-dependent transport. Thus, the rate of receptor-dependent LDL uptake in animals fed coconut oil and 0.12% cholesterol ($50 - 21 = 29$ μg/h per g) was suppressed by 70% relative to the rate that would be seen in control animals at the same plasma LDL concentration ($118 - 21 = 97$ μg/h per g).

From the type of analysis illustrated in Fig. 2, hepatic LDL receptor activity (defined as the rate of receptor-dependent LDL uptake in experimental animals relative to the rate of receptor-dependent LDL uptake in control animals at the same plasma LDL cholesterol concentration) was calculated for
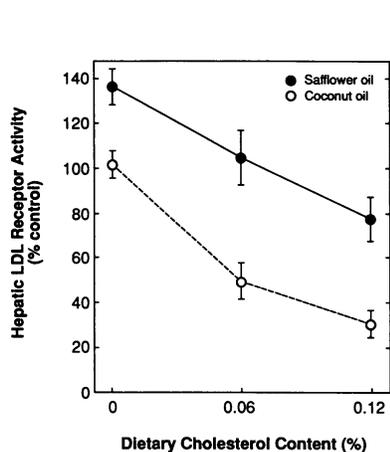
Figure 3. Regulation of hepatic LDL receptor activity by dietary saturated and unsaturated fatty acids. Hepatic LDL receptor activity was determined as illustrated in Fig. 2. Each point represents the mean±1 SE for data obtained in 10–12 animals. Differences between the safflower and coconut oil groups were significant ($P < 0.01$) at each level of dietary cholesterol.

Figure 4. Immunoblotting of LDL receptor from solubilized liver membranes. Hepatic membranes were prepared from animals fed a semisynthetic diet supplemented with 20% (by weight) safflower oil or coconut oil and 0.06% cholesterol for 6 wk. Solubilized liver membrane proteins (500 μg/lane) were separated on 7.5% polyacrylamide gels. The proteins were transferred electrophoretically to PVDF membrane and incubated with a 1:200 dilution of antiserum raised against a synthetic peptide corresponding to the COOH-terminal 13 amino acids of the hamster LDL receptor. After incubation with the primary antibody, the PVDF paper was incubated with 0.5 μCi/ml ^{125}I-labeled donkey anti–rabbit antibody and the dried membranes were subjected to autoradiography.

each of the experimental groups and these values are shown in Fig. 3. When added to a cholesterol-free diet, safflower oil increased hepatic LDL receptor activity by ∼ 35%, whereas coconut oil had no effect. Dietary cholesterol suppressed hepatic LDL receptor activity in both groups; however, the effect of cholesterol was much greater in animals fed coconut oil than in animals fed safflower oil. On the 0.06% cholesterol diet, hepatic LDL receptor activity equaled 49 and 103% of the control value in animals fed coconut oil and safflower oil, respectively. On the 0.12% cholesterol diet, hepatic LDL receptor activity equaled 30 and 77% of the control value in animals fed coconut oil and safflower oil, respectively. Thus, hepatic LDL receptor activity was significantly lower ($P < 0.01$) in animals fed coconut oil than in animals fed safflower oil at all levels of cholesterol intake (26, 53, and 61% lower at cholesterol intakes of 0, 0.06, and 0.12%, respectively).

In parallel studies, identically treated groups of animals were used to prepare hepatic membranes for the determination of LDL receptor protein and to obtain liver samples for the isolation of total RNA. A representative autoradiogram depicting changes in LDL receptor protein in animals fed saturated and unsaturated triglycerides is shown in Fig. 4, and the mean changes in LDL receptor protein are summarized in Fig. 5. On a cholesterol-free diet, LDL receptor protein in solubilized liver membranes was 16% higher in animals fed safflower oil than in animals fed coconut oil ($P > 0.05$). As with rates of receptor-dependent LDL transport, dietary cholesterol suppressed hepatic LDL receptor protein in both groups; however, the effect of cholesterol was much greater in animals fed the coconut oil than in animals fed the safflower oil. When the diet contained 0.06% cholesterol, hepatic LDL receptor protein equaled 52 and 93% of the control value in animals fed coconut oil and safflower oil, respectively. On the 0.12% cholesterol diet, hepatic LDL receptor protein equaled 36 and 80% of the control value in animals fed coconut oil and safflower oil, respectively. Thus, relative to the unsaturated triglyceride, the saturated triglyceride reduced hepatic LDL receptor protein by 44 ($P < 0.01$) and 55% ($P < 0.01$) in animals fed 0.06 and 0.12% cholesterol, respectively.

LDL receptor mRNA levels were quantified by nuclease protection using single-stranded cDNA probes specific for the Golden Syrian hamster. A representative autoradiogram depicting the changes in hepatic LDL receptor mRNA levels in animals fed saturated and unsaturated triglycerides is shown in Fig. 6, and the mean changes in LDL receptor mRNA levels are

summarized in Fig. 7. On a cholesterol-free diet, hepatic LDL receptor activity was 27% higher in animals fed safflower oil than in animals fed coconut oil ($P < 0.025$). Dietary cholesterol suppressed hepatic LDL receptor mRNA levels in both groups, but again the effect of cholesterol was much greater in animals fed the coconut oil than in animals fed the safflower oil. When the diet contained 0.06% cholesterol, hepatic LDL receptor mRNA equaled 58 and 102% of the control value in animals fed coconut oil and safflower oil, respectively. On the 0.12% cholesterol diet, hepatic LDL receptor mRNA equaled 38 and 85% of the control value in animals fed coconut oil and safflower oil, respectively. Thus, relative to the unsaturated triglyceride, the saturated triglyceride reduced hepatic LDL receptor mRNA by 43 ($P < 0.01$) and 55% ($P < 0.01$) in animals fed 0.06 and 0.12% cholesterol, respectively. Overall, the changes in hepatic LDL receptor activity shown in Fig. 3 could be accounted for largely by changes in LDL receptor protein and mRNA, suggesting that dietary fatty acids may regulate the LDL receptor pathway at the transcriptional level. As shown in Fig. 8, there was a significant correlation between the changes
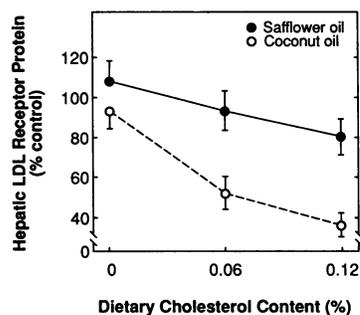
Figure 5. Regulation of hepatic LDL receptor protein levels by dietary saturated and unsaturated fatty acids. Each point represents the mean±1 SE for data obtained from 9–12 animals. Differences between the safflower and coconut oil groups were significant ($P < 0.01$) in animals fed 0.06 and 0.12% cholesterol.
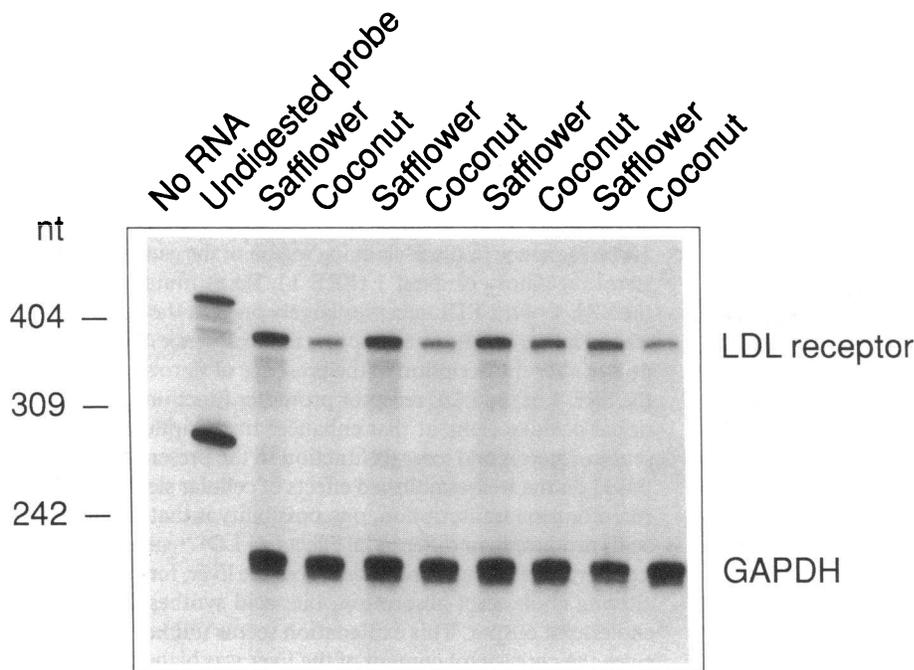
Figure 6. Measurement of hepatic LDL receptor mRNA levels. Hepatic RNA was isolated from animals fed a semi-synthetic diet supplemented with 20% (by weight) safflower oil or coconut oil and 0.06% cholesterol for 6 wk. Total RNA (40 μg) was hybridized with ^{32}P-labeled probes and the LDL receptor and GAPDH bands resistant to mung bean nuclease digestion were analyzed by polyacrylamide gel electrophoresis followed by autoradiography as described in Methods.

in LDL receptor protein and the changes in LDL receptor mRNA levels when both parameters were measured in the same liver ($r = 0.67$).

The major form of regulation of the LDL receptor pathway demonstrated to date is feedback repression of receptor gene transcription by cellular sterols. One possibility is that dietary fatty acids regulate LDL receptor mRNA levels by altering the content or distribution of sterols in the liver. Fig. 9 shows the changes in hepatic esterified and unesterified cholesterol levels in animals fed the various experimental diets. On a cholesterol-free diet, the cholesteryl ester content of the liver was significantly higher in animals fed safflower oil than in animals fed coconut oil (0.8 vs 0.4 mg/g, $P < 0.01$). Dietary cholesterol increased hepatic cholesteryl esters in both groups; however, the increase was much greater in animals fed safflower oil than in animals fed coconut oil. These data indicate that fatty acids do not regulate the LDL receptor pathway simply by altering the cholesterol content of the liver, since unsaturated fatty acids increased LDL receptor mRNA levels under circumstances where the total and esterified cholesterol content of the liver was also increased. As shown in the bottom panel, the two

dietary triglycerides did not differentially affect the total unesterified cholesterol content of the liver.

Discussion

The concentration of LDL in plasma is strongly influenced by the amount and type of triglyceride in the diet. Dietary triglycerides containing predominantly saturated fatty acids increase total and LDL cholesterol concentrations relative to the same amount of an unsaturated triglyceride in both animals and humans. The effects of dietary triglycerides on the major transport processes that control circulating LDL levels have been examined in detail in the hamster (18, 19). These studies have shown that dietary fatty acids produce their differential effects on circulating LDL levels primarily by altering receptor-dependent LDL uptake in the liver. Thus, relative to triglycerides containing predominantly saturated fatty acids, triglycerides containing predominantly unsaturated fatty acids accelerate the rate of receptor-dependent LDL uptake in the liver, thereby lowering plasma LDL concentrations. The present studies show that these fatty acid–induced changes in receptor-depen-
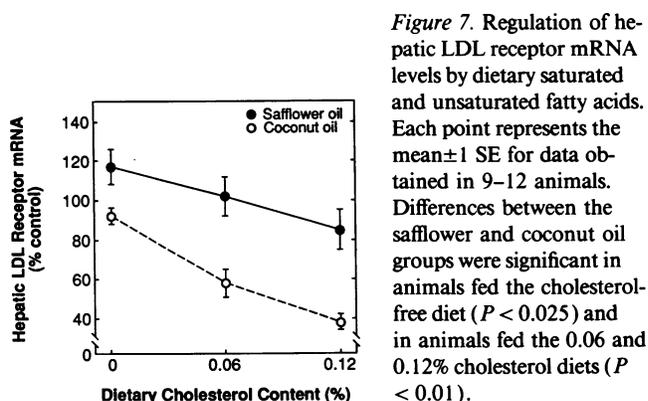
Figure 7. Regulation of hepatic LDL receptor mRNA levels by dietary saturated and unsaturated fatty acids. Each point represents the mean±1 SE for data obtained in 9–12 animals. Differences between the safflower and coconut oil groups were significant in animals fed the cholesterol-free diet ($P < 0.025$) and in animals fed the 0.06 and 0.12% cholesterol diets ($P < 0.01$).
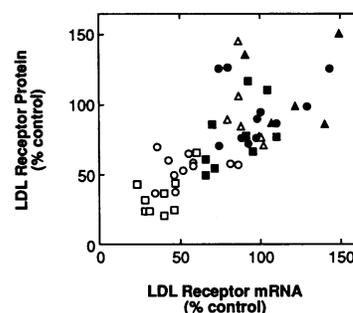
Figure 8. Correlation between hepatic LDL receptor protein and mRNA levels. The change in LDL receptor protein is plotted as a function of the change in LDL receptor mRNA for animals where both parameters were determined in the same liver ($r = 0.67$). The open and closed symbols represent animals fed coconut and safflower oil, respectively, with either 0 (*triangles*), 0.06 (*circles*), or 0.12% (*squares*) cholesterol.
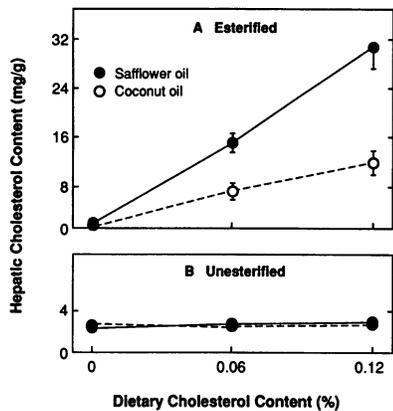
Figure 9. Regulation of hepatic cholesterol levels by dietary saturated and unsaturated fatty acids. Each point represents the mean±1 SE for data obtained in 10–12 animals. Differences between the safflower and coconut oil groups were significant ($P < 0.01$) at each level of dietary cholesterol.

dent LDL transport in the liver are accompanied by parallel changes in LDL receptor protein and mRNA levels. These data suggest that dietary fatty acids regulate the LDL receptor pathway largely at the mRNA level.

The diets used in these studies were rich in triglycerides containing almost exclusively saturated or unsaturated fatty acids. These diets were specifically chosen to maximize the differential effects of dietary fatty acids on hepatic LDL transport and thereby provide the best opportunity for examining the mechanism of this effect. In addition, varying amounts of cholesterol were added to the diet since an interaction between dietary cholesterol and fatty acids exists whereby small amounts of cholesterol magnify the differential effects of saturated and unsaturated fatty acids (19). The two triglycerides used in these studies produced significant differences in receptor-dependent LDL transport at all levels of cholesterol intake and these differences in receptor activity could be accounted for largely by changes in LDL receptor protein and mRNA levels. Whether a similar mechanism accounts for the smaller changes in hepatic LDL receptor activity seen with less drastic alterations in dietary fatty acids is not known. About one-third to one-half of the fatty acids present in typical Western diets are saturated. If half of these saturated fatty acids were replaced by unsaturated fatty acids or carbohydrate, hepatic LDL receptor activity would increase by 25–30% and plasma LDL concentrations would fall by 20–25% (37), changes that if sustained could reduce the risk of mortality from coronary heart disease by as much as 50% in humans (38). Unfortunately, the changes in hepatic LDL receptor activity that occur under these clinically relevant conditions are quite small and it may not be possible to reliably investigate the underlying mechanisms using currently available techniques.

Previous studies investigating the effect of dietary triglycerides on hepatic LDL receptor mRNA levels have yielded mixed results. Two groups have shown that saturated and unsaturated triglycerides produce no differential effects on hepatic LDL receptor mRNA levels in monkeys whether added to low or high cholesterol diets (24, 39). A third group reported a significant differential effect of saturated and unsaturated triglycerides on hepatic LDL receptor mRNA levels in one subgroup of baboons; however, in this study the dietary triglycerides produced no differential effects on plasma β-lipoprotein or apo B concentrations (40). Of note, in none of these studies were receptor mRNA levels correlated with actual rates of receptor-dependent LDL uptake in the liver. Thus, it is possible that the fatty acid–induced changes in receptor-dependent transport

were smaller than anticipated, making changes in LDL receptor mRNA levels difficult to detect.

The mechanism whereby dietary fatty acids differentially regulate LDL receptor mRNA levels is not known. The principal mechanism of regulation of the LDL receptor pathway demonstrated to date is feedback repression of LDL receptor gene transcription by cellular sterols (20, 21). Sterol-mediated regulation of the LDL receptor promoter has been localized to a 10-bp sequence in the 5′-flanking region of the gene termed the sterol regulatory element 1 (SRE 1). Point mutations within the SRE 1 of the LDL receptor largely prevent the induction of transcription that normally occurs in the absence of sterols, but do not alter transcription in the presence of sterols (22). Thus, the SRE 1 of the LDL receptor promoter functions as a conditional positive element that enhances transcription in the absence of sterols and loses its function in the presence of sterols. Based on the well-established effects of cellular sterols on LDL receptor gene transcription, one possibility is that dietary fatty acids produce their differential effects on LDL receptor mRNA levels by altering sterol balance across the liver, for example, by altering cholesterol absorption, bile acid synthesis, or biliary cholesterol output. This explanation seems unlikely, however, since the cholesterol content of the liver was higher in animals fed unsaturated fatty acids than in animals fed saturated fatty acids.

Alternatively, dietary fatty acids may alter the distribution of cholesterol within the hepatocyte. The fraction of total cellular cholesterol that is metabolically active and involved in sterol-mediated regulation of LDL receptor gene expression is likely to be very small. This small regulatory pool of unesterified cholesterol is in equilibrium with a much larger pool of unesterified structural cholesterol located primarily in plasma membranes and with a storage pool of esterified cholesterol located in the cytoplasm. The activity of the cholesterol esterifying enzyme acyl-CoA:cholesterol acyltransferase (ACAT) is known to be higher in animals fed unsaturated fatty acids than in animals fed saturated fatty acids (41). Thus, saturated fatty acids may inhibit ACAT activity, resulting in an expansion of the putative regulatory pool of free cholesterol (or related derivative) that mediates feedback repression of the LDL receptor pathway. Conversely, unsaturated fatty acids may accelerate cholesterol esterification leading to depletion of this regulatory pool of cholesterol. That changes in ACAT activity may lead to regulation of the LDL receptor pathway has been demonstrated in studies with ACAT inhibitors. Thus, preincubation of cultured HepG2 cells (42), cultured rat hepatocytes (43), or cultured J774 macrophages (44) with specific inhibitors of ACAT markedly enhances the sensitivity of the LDL receptor pathway to downregulation by lipoprotein cholesterol.

Finally, free fatty acids (45) and fatty acyl CoAs (46) have been shown to differentially regulate the expression of several genes, including genes encoding transcription factors. Thus, it is possible that dietary fatty acids may regulate LDL receptor mRNA levels through a mechanism that does not involve classic sterol-mediated regulation of gene transcription (47).

Acknowledgments

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