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Reduction in Dietary Omega-6 Polyunsaturated Fatty Acids:Eicosapentaenoic Acid plus Docosahexaenoic Acid Ratio Minimizes Atherosclerotic Lesion Formation and Inflammatory Response in the LDL Receptor Null Mouse

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Abstract

Dietary very long chain omega (ω)-3 polyunsaturated fatty acids (PUFA) have been associated with reduced CVD risk, the mechanisms of which have yet to be fully elucidated. LDL receptor null mice (LDLr-/-) were used to assess the effect of different ratios of dietary ω -6 PUFA to eicosapentaenoic acid plus docosahexaenoic acid (@-6:EPA+DHA) on atherogenesis and inflammatory response. Mice were fed high saturated fat diets without EPA and DHA (HSF ω -6), or with ω -6:EPA+DHA at ratios of 20:1 (HSF R=20:1), 4:1 (HSF R=4:1), and 1:1 (HSF R=1:1) for 32 weeks. Mice fed the lowest ω-6:EPA+DHA ratio diet had lower circulating concentrations of non-HDL cholesterol (25%, P<0.05) and interleukin-6 (IL-6) (44%, P<0.05) compared to mice fed the HSF ω -6 diet. Aortic and elicited peritoneal macrophage (M ϕ) total cholesterol were 24% (P=0.07) and 25% (P<0.05) lower, respectively, in HSF R=1:1 compared to HSF @-6 fed mice. MCP-1 mRNA levels and secretion were 37% (P<0.05) and 38% (P<0.05) lower, respectively, in elicited peritoneal M ϕ isolated from HSF R=1:1 compared to HSF ω -6 fed mice. mRNA and protein levels of ATP-binding cassette A1, and mRNA levels of TNF α were significantly lower in elicited peritoneal M ϕ isolated from HSF R=1:1 fed mice, whereas there was no significant effect of diets with different ω-6:EPA+DHA ratios on CD36, Mø scavenger receptor 1, scavenger receptor B1 and IL-6 mRNA or protein levels. These data suggest that lower ω -6:EPA+DHA ratio diets lowered some measures of inflammation and M¢ cholesterol accumulation, which was associated with less aortic lesion formation in LDLr-/mice.

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Keywords

 ω -6:EPA+DHA ratio; ω -3 fatty acids; atherosclerosis; inflammation; macrophage cholesterol accumulation; LDLr-/- mouse; diet; elicited peritoneal M ϕ

Introduction

Cardiovascular disease (CVD) is the leading cause of death and disability in developed countries. Atherosclerosis is the major cause of CVD, accounting for about half of the attributable deaths [1]. One component of diet that appears to play a multifactorial role in CVD risk is very long chain omega (ω)-3 fatty acids, eicosapentaenoic acid (EPA, C20:5 ω -3) and docosahexaenoic acid (DHA, C22:6 ω -3). Observational data in humans suggest that consumption of 1 to 2 servings of fish per week is associated with decreased CVD mortality [2]. Concern has been raised that the current low absolute intakes of EPA+DHA and high ratios of ω -6 polyunsaturated fatty acids (ω -6 PUFA) to EPA+DHA (ω -6:EPA+DHA) may predispose some individuals to increased CVD risk [3].

Historically, plasma lipoprotein profiles and lipid accumulation in the blood vessel wall have been the major focus of research related to atherogenesis. More recently, a preponderance of evidence from clinical and experimental data suggests that inflammatory responses play a critical role [4]. M ϕ scavenger receptor 1 (MSR1) and CD36 are two scavenger receptors involved in cholesterol influx [5]. Two M ϕ membrane receptors involved in cholesterol efflux are ATP-transporter cassette A1 (ABCA1) and scavenger receptor B class 1 (SR-B1) [6]. When M ϕ cholesterol influx is greater than efflux, cholesterol homeostasis is disturbed and cholesteryl ester (CE) accumulates in cytoplasmic droplets. The resulting M ϕ -derived foam cells secrete pro-inflammatory factors, which amplify the local inflammatory reaction and produce reactive oxygen species, which in turn modify lipoproteins. Interleukin-6 (IL-6) and tumor necrosis factor alpha (TNF α) are major pro-inflammatory factors. IL-6 promotes the synthesis of acute-phase reactants and monocyte chemoattractant protein-1 (MCP-1) [7,8].

MCP-1 and its receptor, CC chemokine receptor 2 (CCR2), direct the migration of monocytes into the intima. Subsequent exposure of the monocytes to macrophage colony-stimulating factor promotes their differentiation to macrophages (M ϕ). M ϕ avidly take up modified apolipoprotein (apo) B-containing lipoproteins, resulting in the formation of M ϕ -derived foam cells. These cells secrete pro-inflammatory factors that amplify the local inflammatory response and produce reactive oxygen species, which modify lipoproteins resulting in increased uptake by scavenger receptors. Overexpression of MCP-1 is positively associated with accumulation of monocytes in fatty streaks [9]. It has been reported that dietary saturated fatty acids (SFA) increase and ω -3 PUFA decrease blood levels of these inflammatory biomarkers, whereas the effect of ω -6 PUFA has yet to be elucidated [10,11].

To better understand the impact of dietary ω -6:EPA+DHA ratios in atherosclerotic lesion formation, LDL receptor null (LDLr-/-) mice were fed diets high in saturated fat and cholesterol, differing only in ω -6:EPA+DHA ratios. The effect of these diets on aortic lipid accumulation, serum lipoprotein profile, plasma inflammatory biomarkers, and elicited peritoneal M ϕ fatty acid composition, cholesterol content, and secretion of inflammatory factors was measured. mRNA and protein levels of genes involved in cholesterol accumulation and inflammation in elicited peritoneal M ϕ were simultaneously determined.

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Materials and Methods

Animals and diets

Forty-one eight-week old, male LDLr-/- mice (Jackson Laboratory, Bar Harbor, Maine) initially weighing 20.1 ± 2.5 g were placed in individual cages with stainless-steel wire bottoms in a windowless room maintained at 22 to 24° C, 45% relative humidity and a daily 10/14 hour light/dark cycle with the light period from 06:00 to 16:00. After 1-week of acclimatization, mice were weighed and randomly assigned to one of four groups. Mice were fed high saturated fat (HSF; 20% fat, w/w) and cholesterol (0.2%, w/w) diets ad libitum, differing only in the ω-6:EPA+DHA ratios: no EPA+DHA (HSF ω-6), and ω-6:EPA+DHA ratio of 20:1 (HSF R=20:1), 4:1 (HSF R=4:1), and 1:1 (HSF R=1:1) (Tables 1). The HSF ω -6 diet has previously been shown to induce atherosclerotic lesion formation in the LDLr-/- mouse [12]. The ratio of ω-6:EPA+DHA in the diets was manipulated by adding different amounts of fish oil (Omega Protein Inc., Houston, TX) and safflower oil. The endogenous cholesterol in the fish oil was compensated for by adding purified cholesterol so the final amount was equivalent among diets. The experimental diets were designed so that the saturated fatty acid (SFA), monounsaturated fatty acid (MUFA) and PUFA content, with the exception of ω -6 PUFA, EPA and DHA, was equivalent. This was achieved by altering the amounts of butter fat, safflower oil and fish oil used to prepare the diets (Table 2). Body weight and food intake were monitored weekly over the 32-week feeding period. In order to ensure that adequate quantities of M¢ would be available for all measurements, five days prior to killing mice were given an intraperitoneal injection of 1.0 ml Brewer thioglycollate broth (4.05 g/100 ml) to elicit M ϕ accumulation. Elicited peritoneal Mo were collected as previously described [13]. After a 16-18 hour fast the mice were anesthetized with CO_2 and killed by exsanguination. Blood was collected by retro-orbital bleeding to harvest serum at week 0, 12 and 32, and by cardiac puncture to harvest plasma at week 32. Serum and plasma were obtained by blood centrifugation at $1,500 \times g$ at 4°C for 25 minutes.

Atherosclerotic lesion quantitation

Mouse hearts were perfused in situ for 1 minute with diethylpyrocarbonate treated-phosphate buffered saline (PBS) containing 1.5 µmol/l aprotinin and 0.1 mmol/l phenylmethylsulfonyl fluoride through a cannula inserted into the left ventricle. Aortas were dissected from the aortic root to the iliac bifurcation using a stereoscopic zoom microscope. As a measure of atherosclerotic lesion formation, total cholesterol (TC) and free cholesterol (FC) were quantified (8 aortas/group) as previously described [14,15]. CE was calculated as the difference between the two measures. The residual delipidated aortic tissue was digested in 1N NaOH and total protein was determined using a bicinchoninic acid (BCA) kit (Pierce Ins., Rockford, IL).

Immunohistochemistry

In two randomly chosen mice from each diet group, aortic arches were isolated from the left aortic valve to the right subclavian artery branch and embedded in OCT compound (Tissue-Tek 4583; Sakura Finetek), snap-frozen in liquid nitrogen and stored at -80° C until sectioning. Aortic arches were sectioned at a 5µm thickness and stained in the pathology facility at the New England Medical Center Hospital. Rat anti-mouse monocyte/M ϕ antibody (MCA519G; Serotec, Raleigh, NC) was used to identify monocyte/M ϕ .

Serum lipid and lipoprotein concentrations

Serum TC, HDL-cholesterol (HDL-C) and triglyceride (TG) concentrations were measured using an Olympus AU400 analyzer with enzymatic reagents (Olympus America, Melville, NY). Non HDL-C was calculated as the difference between TC and HDL-C [16].

Plasma concentrations of inflammatory factors

Plasma TNFα, IL-6 and MCP-1 concentrations were measured using Quantikine® ELISA kits (R&D Systems, Minneapolis, MN).

Elicited peritoneal Mp culture and stimulation

Elicited peritoneal M ϕ were cultured in RPMI1640 medium (ATCC, Manassas, VA) containing 2% heat-inactivated fetal bovine serum (Invitrogen, Carlsbad, CA) for 2 hours (1×10⁶ cells per ml). One million cells were stimulated with 1 µg/ml lipopolysaccharide (LPS) for 15 hours and inflammatory factors released into the culture medium were measured. The remaining cells were washed with PBS three times and used for the measurement of cholesterol content, fatty acid profile, mRNA and protein levels of genes involved in inflammation and cholesterol accumulation.

Cholesterol content and fatty acid profile—Elicited peritoneal M ϕ TC, FC and protein concentrations were determined as described for the aorta. M ϕ fatty acid profiles were determined as previously described [17].

Secretion of inflammatory factors—TNF α , IL-6 and MCP-1 secretion by LPSstimulated elicited peritoneal M ϕ were measured using DuoSet® ELISA kits (R&D Systems, Minneapolis, MN). Elicited peritoneal M ϕ attached to the culture plates were digested by 0.5N NaOH and total protein was measured using BCA kits (Pierce Ins., Rockford, IL).

Western blot—Protein was extracted from elicited peritoneal M ϕ using a RIPA kit (Santa Cruz, Santa Cruz, CA). Western blotting was performed as previously described [16] with the following primary antibodies, MSR1 (Serotec, South San Francisco, CA), SR-B1 (Novus Biologicals, Littleton, CO), ABCA1 (Novus Biologicals, Littleton, CO), CD36 (Cascade Bioscience, Winchester, MA) and β -actin (Sigma, St. Louis, MO). Signals were visualized by chemiluminescence (Amersham Biosciences, Piscataway, NJ), quantified using a GS-800 calibrated densitometer (Bio-Rad, Hercules, CA), and normalized by β -actin.

Real-time PCR—RNA was extracted from elicited peritoneal M ϕ using the RNeasy mini kit (Qiagen, Valencia, CA). Single strand cDNA was synthesized from RNA using SuperScriptTM [] reverse transcriptase according to the manufacturer's instructions (Invitrogen, Carlsbad, CA). Primers for peroxisomal proliferator activated receptor (PPAR) β , PPAR γ , MSR1, CD36, SR-B1, ABCA1, MCP-1, TNF α and β -actin (Table 3) were designed using Primer Express version 2.0 (Applied Biosystems, Foster City, CA). β -actin was used as an endogenous control. Primer amplification efficiency and specificity were verified for each set of primers. cDNA levels of the genes of interest were measured using power SYBR green master mix on real-time PCR 7300 (Applied Biosystems, Foster City, CA). cDNA levels of IL-6 and PPAR α were measured using Taqman® primers and Taqman® PreAmp master mix kit (Applied Biosystems, Foster City, CA). cDNA levels of IL-6 and PPAR α were measured using Taqman® primers and Taqman® PreAmp master mix kit (Applied Biosystems, Foster City, CA). cDNA levels of 10 minutes, 40 cycles of 95°C for 15 seconds and 60°C for 1 minute, 1 cycle of dissociation stage. mRNA fold change was calculated using the 2(-Delta Delta C(T)) method [18].

Statistical methods

Data were checked for normality and appropriate transformations were performed when necessary prior to statistical analysis (PROC UNIVARIATE; SAS version 9.1, SAS Institute Inc, Cary, NC). An analysis of variance (PROC GLM) followed by Tukey's post hoc test was performed to compare multiple group means. Differences were considered significant at P<0.05. Untransformed data are presented in text, figures and tables as mean \pm standard deviation (SD).

Results

Animal body weight and food intake

There were no significant differences in mouse body weight or food intake among the groups at baseline and throughout the 32-week feeding period (data not shown).

Aortic lesion composition

Aortic TC and FC content was lower in the HSF R=1:1 compared to HSF ω -6 diet fed mice, whereas content was intermediate in the other diet groups (Figures 1A and 1B). The dietary ω -6:EPA+DHA ratios had no significant effect on aortic CE content. There was less staining for M ϕ in the cross sections of aortic arches isolated from lower ratio diet fed mice relative to HSF ω -6 diet fed mice (Figure 1C).

Serum lipid and lipoprotein concentrations

Serum TC (data no shown) and non HDL-C concentrations were both approximately 25% lower in the HSF R=4:1 and HSF R=1:1 compared with the HSF ω -6 diet group (P<0.05) (Figure 2A). There was no significant difference in serum ratios of TC to HDL-C among the different diet groups. Serum TG concentrations were 30% and 40% lower in the mice fed the HSF R=1:1 and HSF R=4:1 diets, respectively, compared to mice fed the HSF ω -6 diet at 12 weeks. These differences did not reach statistical significance at 32 weeks due to high degree of variability in response among mice.

Plasma concentrations of inflammatory factors

IL-6 concentrations were 44% lower in the HSF R=1:1 compared to HSF ω -6 diet fed mice (P<0.05), whereas concentrations were intermediate in the other diet groups (Figure 3). Due to high degree of variability in response among mice, there was no significant difference in MCP-1 and TNF α concentrations among the diet groups.

Elicited peritoneal Mo fatty acid profiles and cholesterol content

The fatty acid profile of the elicited peritoneal M ϕ reflected that of the diets (Table 4). As the ratio of ω -6:EPA+DHA in the diet was lower, the proportion of EPA, docosapentaenoic acid (DPA, C22:5) and DHA in the elicited peritoneal M ϕ was higher. These changes were concurrent with a decrease in the proportion of ω -6 fatty acids, resulting in a progressive decrease in the ω -6:EPA+DHA ratio to 4.9, 3.2, 1.7 and 0.9 in elicited peritoneal M ϕ isolated from HSF ω -6, HSF R=20:1, HSF R=4:1 and HSF R=1:1 diet fed mice, respectively.

The pattern of differences in the TC content of the elicited peritoneal M ϕ harvested from the mice fed the different diets were similar to that seen in the aortas of the respective mice (Figures 1 and 4). M ϕ TC content was12%, 22% and 26% lower in the HSF R=20:1, HSF R=4:1, HSF R=1:1, respectively, compared to the HSF ω -6 fed mice. In contrast to that observed in the aorta, the difference in TC content was in the CE rather than FC fraction.

mRNA and protein levels of genes involved in cholesterol accumulation in elicited peritoneal $\ensuremath{\mathsf{M}}\xspace\varphi$

To investigate the mechanism(s) underlying the lower CE content in the elicited peritoneal M ϕ harvested from the mice fed the lower ω -6:EPA+DHA ratio diets, mRNA and protein levels of genes involved in M ϕ cholesterol accumulation were measured. M ϕ ABCA1 mRNA levels were 8%, 35% and 30% lower, and protein levels were 2%, 26% and 30% lower, in the HSF R=20:1, HSF R=4:1, HSF R=1:1 fed mice, respectively, compared to the HSF ω -6 fed mice (Figure 5A and 5B). MSR1, SR-B1 and CD36 mRNA and protein levels were similar among

diet groups. There was no significant effect of dietary treatment on mRNA levels of MSR1, PPAR α , PPAR β/δ , PPAR γ (data not shown).

mRNA levels and secretion of inflammatory factors in elicited peritoneal Mo

To explore whether plasma concentrations of inflammatory factors depend on M ϕ secretion, we measured MCP-1, TNF α and IL-6 mRNA levels and secretion in elicited peritoneal M ϕ . M ϕ MCP-1 mRNA levels were 1%, 20% and 27% lower, and protein secretion were 17%, 34% and 38% lower, in the HSF R=20:1, HSF R=4:1, HSF R=1:1 fed mice, respectively, compared to the HSF ω -6 fed mice (Figure 5C and 5D). A similar pattern was seen in TNF α mRNA levels but not secretion. There was no significant effect of dietary ω -6:EPA+DHA ratios on IL-6 secretion and mRNA levels.

Discussion

Observational data are somewhat inconsistent with respect to the relationship between dietary very long chain ω -3 fatty acids (EPA and DHA) and risk of developing CVD [19-21]. This inconsistency has spawned a debate as to whether the critical variable is the ratio of ω -6:EPA +DHA or the absolute level of EPA and DHA in the diet [22]. The former issue was addressed in the current study using LDLr-/- mice fed an atherogenic diet containing different ratios of ω -6:EPA+DHA as an experimental model. The results demonstrated that the diet with the lowest ratio of ω -6:EPA+DHA, HSF R=1:1, resulted in a less atherogenic plasma lipid profile, less M ϕ deposition in the aortic wall, and lower plasma IL-6 concentrations, with intermediate outcomes in the HSF R=20:1 and HSF R=4:1 diet groups. In addition, elicited peritoneal M ϕ isolated from HSF R=1:1 diet fed mice had fatty acid profiles that mirrored the dietary fatty acids, lower cholesterol content, and lower MCP-1 mRNA levels and secretion in response to stimulation. These observations were associated with less atherosclerotic lesion development.

The experiment diets were designed to have equivalent compositions with respect to SFA, MUFA, and total PUFA. The only difference among the diets was the ratio of ω -6 PUFA to EPA+DHA. Currently recommendations for individuals with CVD and hypertriglyceridemia are to consume 1 gram to 3 grams of EPA+DHA, respectively [2]. For a person consuming 2200 kcal that is equivalent to approximately 0.4% to 1% energy from. Five grams of EPA +DHA per kilogram diet is equivalent to 1% energy in the HSF R=1:1 diet. The amount of EPA+DHA in the HSF R=4:1 diet and HSF R=20:1 diets is 0.5% and 0.1% energy, respectively.

Mice fed the lowest ratio of ω -6:EPA+DHA diet had lower serum TC and non HDL-C concentrations and lower aortic TC and FC accumulation compared to mice fed the HSF ω -6 diet. Changes in LDLr-/- mice aortic cholesterol content in response to an atherogenic diet have previously been reported in both FC and CE fractions [23]. The smaller difference and higher variability in the CE fraction among mice might have accounted for larger differences in FC than CE accumulation in this study. Two previous studies have reported that LDLr-/- mice fed fish oil had less atherosclerotic lesion area and lower plasma TC concentrations compared to mice fed olive oil or corn oil [24,25]. Apolipoprotein E and LDL receptor double null mice fed a low ω -6: α -linolenic acid ratio diet compared to a high ω -6: α -linolenic acid ratio diet also resulted in less atherosclerotic lesion area and lower plasma TC concentrations [26]. Mice have been reported to have a higher capacity to convert α -linolenic acid to EPA than humans, albeit under more extreme dietary conditions prior to and during the period of assessment [27,28]. Under these circumstances the authors speculated that it was the EPA and DHA, rather than the α -linolenic acid that resulted in lowered plasma TC concentrations.

M\u00f6 play a major role in the uptake of oxidized LDL, and accumulation and deposition of cholesterol in the arterial wall [29,30]. Based on the results from previous studies [13,31,32],

the expression and production of inflammatory factors is comparable between resident and elicited peritoneal $M\phi$. In addition, there was no significant difference in the response of nutritional interventions between the two types of peritoneal $M\phi$. In our study, elicited peritoneal $M\phi$ were used to ensure an adequate number of cells were harvested. The fatty acid profile of the diet was reflected in the fatty acid profile of the elicited peritoneal $M\phi$ isolated from the respective groups of mice. The TC content of these $M\phi$ followed a similar pattern to that of the corresponding aortas. Lower $M\phi$ TC and CE content in the mice fed the HSF R=1:1 diet suggests lower uptake and accumulation of modified LDL and/or more cholesterol efflux.

In order to understand the mechanism causing the difference in M ϕ cholesterol content, the expression of two proteins involved in cholesterol influx, MSR1 and CD36, and two proteins involved in cholesterol efflux, SR-B1 and ABCA1, was assessed. Mice fed diets with ω -6 and the highest ratio of ω -6:EPA+DHA had the highest ABCA1 mRNA and protein levels compared to mice fed diets with the lowest ratio. There was no significant difference in mRNA and protein levels of MSR1, SR-B1 and CD36. It has been reported that ABCA1 expression is upregulated in response to cholesterol enrichment in M ϕ and fibroblasts *in vitro* [33,34] and in mouse M\u00f6 in vivo [35]. Consistent with these observations, in the current study, ABCA1 mRNA and protein levels were higher in elicited peritoneal M ϕ isolated from the HSF ω -6 and highest ω -6:EPA+DHA ratio diet fed LDLr-/- mice, which had the highest M ϕ cholesterol content. Some studies have shown that ABCA1 expression is regulated by liver-X-receptors (LXR) at a transcriptional level [36-38]. We speculate that the higher ω -6:EPA+DHA ratio diets resulted in increased production of LXR ligands in M ϕ , such as 24(S),25-epoxycholesterol and 27-hydroxycholesterol, which would have activated LXR. Ligand activated-LXR can upregulate ABCA1 gene expression and enhance ABCA1-mediated cholesterol efflux from accumulated in peritoneal cavity, we could not test the above hypothesis in this study. These observations suggest that the greater cholesterol accumulation in the elicited peritoneal $M\phi$ isolated from mice fed the HSF ω -6 diet is a result of higher plasma TC and non HDL-C

PUFA regulate M ϕ ABCA1 protein amounts at both transcription and post-translation levels. In our study, EPA and DHA content was higher in elicited peritoneal M ϕ isolated from mice fed lower ω -6:EPA+DHA ratio diets. Previous data have shown that PUFA, including EPA and DHA, lower ABCA1 protein levels in murine M ϕ through increasing ABCA1 protein degradation [41]. EPA, DHA and their metabolites are ligands for PPAR. The ligand-activated PPAR can prevent LXR from binding to the ABCA1 promoter, and lower ABCA1 expression [39,40,42-44]. More work is required to compare the effect of arachidonic acids (AA) and linoleic acids (LA), two major ω -6 PUFA in elicited peritoneal M ϕ , to EPA and DHA on ABCA1 expression through PPAR-LXR pathway.

The inflammatory process plays a major role in the development of atherosclerotic plaque [4, 45]. IL-6, MCP-1 and TNF α are important inflammatory biomarkers for atherosclerosis [7, 46,47]. Epidemiological observations have suggested a positive association between Western type diets, and plasma MCP-1, IL-6 and TNF α concentrations [48-50], and an inverse association between dietary EPA+DHA and plasma concentrations of C-reactive protein and IL-6 [51-53]. Manipulating the ω -6:EPA+DHA ratios had variable effects on markers of inflammation both in plasma concentrations and as secreted by elicited peritoneal M ϕ . Mice fed the HSF R=1:1 diet tended to have lower plasma IL-6, MCP-1 and TNF α concentrations than mice fed the HSF ω -6 diet, although the later two factors did not reach statistical significance. In contrast, stimulated elicited peritoneal M ϕ had secretion rates and mRNA levels of IL-6 and TNF α indistinguishable among diet groups. Taken together, these data suggest that the major source of plasma IL-6 and TNF α is other than that produced by M ϕ , perhaps released from endothelial cells, adipocytes or T cells.

MCP-1, a major inflammatory factor released from $M\phi$, functions to recruit monocytes into the arterial intima [54]. Overexpression of MCP-1 has been positively related to monocyte accumulation in fatty streaks [9,55]. MCP-1 and LDL receptor double null mice had lower monocyte and vessel wall lipid accumulation compared to LDLr-/- mice [56,57]. In the present from stimulated elicited peritoneal M ϕ were significantly lower in mice fed the HSF R=1:1 diet than the HSF ω -6 diet, with intermediate levels in the other two diet groups. This is consistent with in vitro studies showing that EPA and DHA decrease MCP-1 mRNA levels and secretion in/from endothelial cells and human kidney cells [58,59]. In addition to $M\phi$, MCP-1 is also produced by endothelial cells [60], smooth muscle cells [61] and adjocytes [62]. It has been reported that EPA upregulates MCP-1 mRNA levels compared to arachidonic acid (C20:4, ω -6) in endothelial cells, and might further increase MCP-1 secretion [60]. This may cause a smaller difference in plasma MCP-1 concentrations and explain the disparity between plasma MCP-1 concentrations and elicited peritoneal Mø secretion. The data suggest that diets with lower ratios of ω -6:EPA+DHA result in lower plasma MCP-1 concentrations and its secretion from elicited peritoneal Mø. EPA and DHA may downregulate MCP-1 expression through inhibiting nuclear factor-kappa B activation via an activated PPARdependent pathway [59].

A limitation of this work is that we could not address questions related to the absolute level of EPA and DHA in the diet and subsequent effects on the outcome variables assessed. The study was designed to assess the effect of an extreme range of ω -6:EPA+DHA ratios on atherogenesis and inflammatory response, not chronic exposure as might be experienced by humans during a lifetime. This study was not designed to assess the effects of linoleic acid or α -linolenic acid on atherogenesis. These fatty acids have been shown, under a variety of conditions, to be associated with decreased CVD in humans. The LDLr-/- mouse is a good model for human atherosclerosis, allowing for an assessment of aortic lesion formation. However, plasma lipid and lipoprotein profiles and the composition of the lesion itself were not similar to that observed in humans. This is likely due to the lack of cholesteryl ester transfer protein in mice. No animal model perfectly mimics humans, including inflammatory response and plasma lipid and lipoprotein profiles, in response to the Western diet, fish oil, or one modified in fatty acid profile.

In summary, LDLr-/- mice fed diets with the lowest ratio of ω -6:DHA+EPA (HSF R=1:1) had the lowest concentrations of serum atherogenic lipids and plasma IL-6, and the least elicited peritoneal M ϕ cholesterol accumulation and MCP-1 expression, which in turn were associated with the lowest aortic lesion formation. These data suggest that the effect of different ratios of ω -6:DHA+EPA on atherogenesis is multifactorial.

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Figure 1.

Representative aorta from each diet group, white areas denote atherosclerotic lesion (A), aortic cholesterol content in whole aortas (B), and M ϕ immunostaining by an antibody to mouse M ϕ surface marker in the cross-sections of aortic arches (C) in LDLr-/- mice. Magnified (200×) images of aortic arch cross-sections. The intensity of the brown color corresponds to the presence levels of M ϕ . n =11 in HSF R=4:1 group, n=10 in other groups. Values are mean \pm SD of untransformed data. Prior to statistical analysis, the following transformation was made, aortic FC (Log). An analysis of variance (PROC GLM) followed by Tukey's post hoc test was performed to compare group means. Bars without a common superscript differ, P<0.05.



Figure 2.

Serum lipid and lipoprotein concentrations in LDLr-/- mice. Serum was obtained at 0-, 12- and 32-week post-diet interventions. n =11 in HSF R=4:1 group, n=10 in other groups. Values are mean \pm SD of untransformed data. Prior to statistical analysis, HDL-C at 32-week point was transformed (inverse). An analysis of variance (PROC GLM) followed by Tukey's post hoc test was performed for multiple comparisons. Bars without a common superscript differ, P<0.05.



Figure 3.

Plasma concentrations of inflammatory factors in LDLr-/- mice. n =11 in HSF R=4:1 group, n=10 in all other groups. Values are mean \pm SD of untransformed data. Prior to statistical analysis, the following transformations were made, plasma MCP-1 (log), plasma TNF α (log), plasma IL-6 (inverse). An analysis of variance (PROC GLM) followed by Tukey's post hoc test was performed for multiple comparisons. Bars without a common superscript differ, P<0.05.



Figure 4.

Elicited peritoneal M ϕ cholesterol content in LDLr-/- mice. n =11 in HSF R=4:1 group, n=10 in all other groups. Values are mean ± SD of untransformed data. Prior to statistical analysis, the following transformations were made, M ϕ FC (square root), M ϕ CE (square). An analysis of variance (PROC GLM) followed by Tukey's post hoc test was performed for multiple comparisons. Bars without a common superscript differ, P<0.05.



Figure 5.

mRNA (A) and protein (B) levels of ABCA1, SR-B1, CD36; mRNA levels (C) and secretion (D) of MCP-1, TNF α and IL-6 in/from elicited peritoneal M ϕ harvested from LDLr-/- mice at the end of the 32-week feeding period. n =11 in HSF R=4:1 group, n=10 in all other groups. Values are mean ± SD of untransformed data. Prior to statistical analysis, the following transformations were made, mRNA levels of ABCA1 (log), MCP-1 (log), TNF α (square root), IL-6 (log) and MCP-1 secretion (square root). An analysis of variance (PROC GLM) followed by Tukey's post hoc test was performed for multiple comparisons. Bars without a common superscript differ, P<0.05.

Table 1

Composition of the experimental diets

	Diet										
Ingredient ^a	HSF ^b ω-6	HSF R ^c =20:1	HSF R=4:1	HSF R=1:1							
	g/kg diet										
Casein	240	240	240	240							
L-methionine	3.6	3.6	3.6	3.6							
Cornstarch	150	150	150	150							
Sucrose	299	299	299	299							
Cellulose	50	50	50	50							
Mineral mix, AIN-76	42	42	42	42							
Vitamin mix	12	12	12	12							
Calcium carbonate	3.6	3.6	3.6	3.6							
Lipid	201.5	201.5	201.5	201.5							
Cholesterol	1.5	1.5	1.5	1.5							
Anhydrous milk fat	191.0	190.0	185.5	179.0							
Safflower oil	9.0	8.1	6.0	0.1							
Fish oil ^d	0.0	1.9	8.5	20.9							
SFA	125.3	125.0	123.6	122.0							
MUFA	59.0	59.0	58.7	58.5							
PUFA	12.5	12.5	13.2	13.3							
ω-6 PUFA	12.5	11.8	10.6	6.9							

 a Harlan Teklad (Madison, WI), with the exception of the fish oil. Vitamin mix was Teklad 40060. Anhydrous milk fat provided an additional 0.05% of cholesterol, the total cholesterol was 0.2%.

 $^b\mathrm{HSF:}$ high saturated fatty acid and high cholesterol diet

 C R: ratio of ω -6:EPA+DHA

^dOmega Protein Inc. (Houston, TX). To compensate for the small amounts of cholesterol in the fish oil (0.24g/100g fish oil), 1.48 g/kg and 1.45 g/kg of cholesterol was added to HSF R=4:1 and HSF R=1:1 diets, respectively.

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Table 2

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Fatty acid profile of the experimental diets

	HSF R=1:1		66.85 ± 0.00	3.18 ± 0.01	11.10 ± 0.06	28.65 ± 0.09	20.93 ± 0.03	0.07 ± 0.00	26.47 ± 0.04	2.33 ± 0.02	18.36 ± 0.04	4.87 ± 0.05	$0.04{\pm}0.00$	$6.67{\pm}0.14$	2.85 ± 0.02	1.80 ± 0.00	0.46 ± 0.02	$0.07{\pm}0.00$	0.15 ± 0.01	0.25 ± 0.00	0.12 ± 0.00	3.82 ± 0.06	0.86 ± 0.01	1.25 ± 0.02	0.39 ± 0.01	1.32±0.02
	HSF R=4:1		67.88 ± 0.07	3.17 ± 0.02	10.93 ± 0.03	28.98 ± 0.03	21.58 ± 0.11	0.07 ± 0.00	25.62 ± 0.07	1.74 ± 0.00	18.60 ± 0.04	4.33±0.05	0.03 ± 0.00	6.50 ± 0.01	4.42 ± 0.00	3.42 ± 0.01	0.47 ± 0.03	0.05 ± 0.00	0.09 ± 0.00	0.26 ± 0.02	0.13 ± 0.00	2.07 ± 0.00	0.78 ± 0.00	0.54 ± 0.00	0.23 ± 0.00	0.52±0.00
Dict	HSF R=20:1	0 ∕0	68.42 ± 0.26	3.33 ± 0.07	10.92 ± 0.02	28.75 ± 0.13	21.46 ± 0.28	0.07±0.00	25.32 ± 0.25	1.44 ± 0.00	$18.67{\pm}0.20$	4.24 ± 0.05	0.01 ± 0.00	6.26 ± 0.01	5.07 ± 0.00	4.16 ± 0.04	0.48 ± 0.00	0.05 ± 0.00	0.06 ± 0.00	0.21±0.03	0.11 ± 0.01	1.19 ± 0.01	0.74 ± 0.00	0.18 ± 0.00	0.14 ± 0.00	0.12±0.00
	HSF @-6		68.30 ± 0.29	3.25 ± 0.07	10.83 ± 0.06	28.87±0.06	21.84 ± 0.14	0.06 ± 0.00	25.37 ± 0.18	1.33 ± 0.03	18.83 ± 0.10	4.26 ± 0.05	$0.01{\pm}0.00$	6.32 ± 0.11	$5.36{\pm}0.10$	$4.47{\pm}0.05$	0.45 ± 0.01	$0.04{\pm}0.00$	$0.09{\pm}0.04$	$0.20{\pm}0.01$	0.11 ± 0.02	0.96 ± 0.09	$0.74{\pm}0.00$	0.10 ± 0.00	0.13 ± 0.00	0.01 ± 0.00
	Fatty acid		SFA	C12:0	C14:0	C16:0	C18:0	C24:0	MUFA	C16:1 @-7	C18:1 @-9	C18:1 00-7	C24:1	PUFA	9-0)	C18:2	C18:3	C20:3	C20:4	C22:2	C22:5	00-3	C18:3 00-3	C20:5 @-3	C22:5 00-3	C22:6 @-3

Table 3

Mouse oligonucleotide sequences of primers

Accession No.	Forward primer	Reverse primer
<u>NM_013454</u>	CCTGCTAAAATACCGGCAAGG	GTAACCCGTTCCCAACTGGTTT
<u>NM_016741</u>	TGGAACGGACTCAGCAAGATC	AATTCCAGCGAGGA TTCGG
<u>NM_007643</u>	ATTAATGGCACAGACGCAGC	CCGAACACAGCGTAGATAGACC
<u>AF203781</u>	TCTACAGCAAAGCAACAGGAGG	TCCACGTGCGCTTGTTCTT
<u>NM_013693</u>	TGTAGCCCACGTCGTAGCAAA	GCTGGCACCACTAGTTGGTTGT
<u>NM_011333</u>	TCTCTCTTCCTCCACCACCATG	GCGTTAACTGCATCTGGCTGA
<u>NM_011145</u>	AGTGCGATCGGATCTGCAAGA	TCCAAAGCGGATAGCGTTGTG
<u>NM_011146</u>	TCTTAACTGCCGGATCCACAAA	CCAAACCTGATGGCATTGTGA
<u>NM_007393</u>	CTTTTCCAGCCTTCCTTCTTGG	CAGCACTGTGTTGGCATAGAGG
	Accession No. NM_013454 NM_016741 NM_007643 AF203781 NM_013693 NM_011333 NM_011145 NM_011146 NM_007393	Accession No. Forward primer NM_013454 CCTGCTAAAATACCGGCAAGG NM_016741 TGGAACGGACTCAGCAAGATC NM_007643 ATTAATGGCACAGACGCAGC AF203781 TCTACAGCAAAGCAACAGGAGGG NM_013693 TGTAGCCCACGTCGTAGCAAAA NM_011333 TCTCTCTTCCTCCACCACGAGA NM_011145 AGTGCGATCGGATCTGCAAAA NM_011146 TCTTAACTGCCGGATCCACAAAA

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Table 4

Membrane fatty acid profile in elicited peritoneal $M\phi^a$

	Diet									
Fatty acid	HSF ω-6	HSF R=20:1	HSF R=4:1	HSF R=1:1						
		0,	6							
SFA	34.97±5.97	33.30±2.30	34.00±3.16	33.49±2.53						
C12:0	0.38±0.34	0.29±0.20	0.31±0.13	0.27±0.09						
C14:0	1.88±0.19	1.67±0.47	1.98±0.28	1.74±0.33						
C16:0	21.67±3.23	21.91±4.41	21.29±3.21	21.62±2.04						
C18:0	10.66±3.03	9.10±3.09	10.01±0.92	9.47±0.86						
C20:0	$0.14{\pm}0.07$	0.12±0.05	0.10±0.03	0.10±0.03						
C22:0	0.11±0.07	0.09±0.04	$0.10{\pm}0.02$	0.11±0.02						
C24:0	$0.14{\pm}0.04$	0.13±0.05	0.20±0.23	0.18 ± 0.05						
MUFA	42.13±5.55	46.00±5.72	42.80±2.57	43.08±2.37						
C16:1 ω-9	1.31±0.07	1.19±0.27	1.55±0.26	1.55±0.32						
C16:1 ω-7	4.07±1.27	4.17±1.37	4.67±0.93	4.77±0.79						
C18:1 ω-9	28.28±4.76	32.03±8.43	28.45±3.59	29.05±2.93						
C18:1 ω-7	$5.50{\pm}1.05$	5.71±1.52	5.20±0.76	4.88±0.41						
C20:1	1.14±0.19	1.10±0.35	1.03±0.15	0.81±0.13						
C22:1	0.31±0.12	0.32±0.14	0.32±0.19	0.31±0.13						
C24:1	0.67 ± 0.28	0.63±0.28	0.63±0.13	0.66±0.12						
PUFA	22.90±2.51	20.70±6.16	23.20±4.86	23.44±1.17						
ω-6	16.75±1.91 ^a	13.31±3.90 ^b	11.21 ± 1.74^{b}	8.46±1.14 ^c						
C18:2	4.72±0.90 ^a	3.88 ± 0.82^{a}	$3.80{\pm}0.68^{a}$	1.67 ± 0.46^{b}						
C18:3	0.02±0.01	0.02±0.03	0.03±0.03	0.02±0.03						
C20:2	1.04 ± 0.14	0.84±0.24	0.54 ± 0.05	0.35±0.04						
C20:3	1.32±0.30	1.32±0.43	1.12±0.19	0.90±0.14						
C20:4	6.48±1.18 ^a	$5.34{\pm}2.02^{ab}$	4.47 ± 0.85^{b}	4.14 ± 0.77^{b}						
C22:2	$1.34{\pm}1.42$	0.64 ± 0.42	0.50±0.34	0.69±0.57						
C22:5	0.15±0.19	0.07 ± 0.05	0.13±0.09	0.19 ± 0.08						
C22:4	1.68 ± 0.44	1.21±0.50	0.64±0.16	0.51±0.10						
ω-3	6.16±0.68 ^c	7.39±2.27°	11.98±3.21 ^b	14.98±0.69 ^a						
C18:3 ω-3	0.19 ± 0.04	0.20 ± 0.05	0.21±0.05	0.19 ± 0.02						
C20:5 ω-3	1.03 ± 0.73^{b}	0.93 ± 0.35^{b}	1.98±0.56 ^a	3.29±0.72 ^a						
C22:5 ω-3	2.47 ± 0.37^{b}	2.99±1.02 ^b	4.89±1.21 ^a	5.33±2.10 ^a						
C22:6 ω-3	2.46 ± 0.71^{b}	3.27 ± 0.99^{ab}	4.89±1.48 ^a	6.17 ± 1.42^{a}						
ω-6:EPA+DHA	4.88±0.78 ^a	3.18±0.20 ^a	1.74±0.46 ^b	0.91±0.11 ^c						

 a Values are mean ± SD of untransformed data, n = 11 in HSF R=4:1 group, n=10 in all other groups. Prior to the statistical analysis the following transformations were made, C20:5 (square root), C22:5 (square), C22:6 (log), ω -6:EPA+DHA (inverse), MUFA (log), PUFA (log) and ω -6 PUFA (log). An analysis of variance (PROC GLM) followed by Tukey's post hoc test was performed for multiple comparison. Within a row means without a common superscript differ, P<0.05.