

Postprandial Lipemia and Associated Metabolic Disturbances in Healthy and Hyperlipemic Postmenopausal Women

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The increased risk for coronary artery disease observed in postmenopausal women is partly explained by a more atherogenic fasting lipoprotein profile. Moreover, natural menopause has been associated with an altered postprandial lipid profile. To better characterize the interaction between fasting and postprandial lipid profile after menopause, we examined postprandial changes in several lipid parameters in three age-matched groups of postmenopausal women (16 affected by mixed hyperlipemia, 17 by common hypercholesterolemia, and 17 normolipemic), who underwent a standardized oral fat-loading test. The magnitude of postprandial lipemia, expressed as 8-hour triglyceride incremental area under the curve, was greater in women with mixed hyperlipemia ($1,326 \pm 372 \text{ mg} \cdot \text{dL}^{-1} \cdot \text{h}^{-1}$) than in normal ($484 \pm 384 \text{ mg} \cdot \text{dL}^{-1} \cdot \text{h}^{-1}$) and hypercholesterolemic ($473 \pm 223 \text{ mg} \cdot \text{dL}^{-1} \cdot \text{h}^{-1}$; both $P < .0001$) women, and the differences held after adjustment for body mass index and fasting insulin. Women with mixed hyperlipemia showed a significant postprandial decrease in high-density lipoprotein 2 (HDL₂) cholesterol, lipoprotein (a), and low-density lipoprotein (LDL) particle size. Both hypercholesterolemic and normolipemic women showed a significant postprandial decrease in HDL cholesterol and lipoprotein (a) levels but not in LDL size. In a multiple linear regression analysis, fasting triglyceride levels, insulin level, and waist-hip ratio were all independent predictors of the magnitude of postprandial lipemia. In conclusion, postmenopausal women with mixed hyperlipemia show a greater postprandial triglyceride increase and a more pronounced reduction in HDL cholesterol level and LDL size than hypercholesterolemic and normolipemic subjects. The presence of the features of insulin resistance syndrome could contribute to the deterioration of postprandial lipemic response in these subjects.

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A RELATIONSHIP between postprandial lipemia and coronary artery disease (CAD) was initially observed in the 1950s and 1960s¹ and further investigated by Zilversmit, who proposed that atherogenesis was a postprandial phenomenon.² On the track of Zilversmit's hypothesis, several studies followed with the purpose to clarify the metabolic events occurring in the postprandial state and their association with cardiovascular disease.^{3,4} In this regard, a cross-sectional study by Patsch et al⁵ showed an association between plasma triglyceride levels measured late in the postprandial state and CAD in men with normal fasting triglyceridemia. Data on the postprandial lipid response as a CAD risk factor in normolipemic women are more conflicting.⁶⁻⁸

Postmenopausal estrogen deficiency has been associated with an adverse fasting⁹ and postprandial⁸ lipid profile, which may contribute to the increased CAD risk after menopause.^{10,11} A transient postprandial reduction in high-density lipoprotein (HDL) cholesterol levels has been described in normolipemic postmenopausal women.¹² However, postprandial changes in several lipid parameters, including HDL cholesterol subfractions, lipoprotein (a) [Lp(a)], and low-density lipoprotein (LDL) particle size, have not been investigated so far in women with dyslipidemia. Thus, we investigated changes in these parameters after a standardized oral fat-loading test in post-

menopausal women with hypercholesterolemia and mixed hyperlipemia and in a control group of normolipemic women.

PATIENTS AND METHODS

Patients and Controls

Three age-matched groups of postmenopausal women were selected for the study. Seventeen women were affected by polygenic hypercholesterolemia, 16 by mixed hyperlipemia, and 17 normolipemic women served as control subjects. Secondary hyperlipemia was excluded in all subjects by appropriate thyroid, renal, and liver function tests. Menopausal status was defined by questionnaire as the absence of menses for at least 1 year. Women with uncertain menopausal status, as well as subjects in menopause for less than 1 year, were excluded from the study. All subjects with diabetes mellitus and history of CAD were excluded from the study. No subject was taking hormone-replacement therapy, lipid-lowering drugs, or other drugs affecting lipid or glucidic metabolism. Polygenic hypercholesterolemia was defined as LDL cholesterol $> 160 \text{ mg/dL}$ in the study subject and in one or more family members in the absence of xanthomas in the family. Mixed hyperlipemia was defined as LDL cholesterol level $\geq 160 \text{ mg/dL}$ and triglycerides $\geq 200 \text{ mg/dL}$; 6 of the 16 subjects fulfilled all the criteria for diagnosis of familial combined hyperlipemia.¹³ Normolipemic subjects had LDL cholesterol levels $< 160 \text{ mg/dL}$ and triglyceride levels $< 180 \text{ mg/dL}$. The diet regimen before the test was homogeneous in the three groups. Hyperlipemic subjects followed a lipid-lowering American Heart Association step II diet (total calories as fats $< 30\%$, saturated fats $< 7\%$, cholesterol 200 mg/d). Limitations for monosaccharides and alcohol were prescribed in subjects with mixed hyperlipemia. Normolipemic subjects had a normocaloric hypolipidic diet with the same characteristics for at least 2 weeks before the oral fat-tolerance test. All subjects were asked to abstain from alcohol for at least 24 hours before the test.

Procedures

Participants started the test at 8 AM at the Lipid Clinic of our institution. The standardized high-fat meal consisted of whipping cream, liquid chocolate, and nonfat dry milk and contained 65 g of fat, 25 g of carbohydrates, 6 g of proteins per m^2 body surface area.^{5,14,15}

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The corresponding caloric intake was 700 kcal/m², 83% derived from fat, 14% from carbohydrates, and 3% from proteins. The cholesterol content was 240 mg, and the ratio of polyunsaturated to saturated fats was 0.06. In each subject, a baseline venous blood sample was collected at 8 AM after a 14-hour fast. Study subjects consumed the test meal at 8:30 AM. Thereafter, participants were offered water throughout the postprandial period, but no food or caloric drink was allowed. At baseline and 4, 6, and 8 hours after the meal, the following variables were determined: total cholesterol level, triglyceride levels, levels of HDL cholesterol and its HDL₂ and HDL₃ subfractions, LDL cholesterol level, LDL particle size, and Lp(a) level. Serum insulin and lipoprotein lipase *HindIII* genotype were tested only at baseline.

Total cholesterol, triglyceride, and HDL cholesterol levels were determined by a colorimetric method (Dimension Autoanalyzer; DADE Inc., Newark, NJ), and LDL cholesterol level was calculated from the Friedewald equation.¹⁶ Plasma Lp(a) concentration was measured by enzyme-linked immunosorbent assay. HDL₂ and HDL₃ cholesterol concentrations were determined after precipitation with two different reagents containing polyethylene glycol (Immuno AG, Vienna, Austria). An aliquot of plasma (blood anticoagulated with EDTA) was used to determine LDL size by gradient gel electrophoresis according to the method of Rainwater et al.¹⁷ A 2% to 16% polyacrylamide gel was prepared; samples for seeding were preincubated and stained with Sudan Black, then seeded with thyroglobulin, ferritin, catalase, lacticodehydrogenase, albumin, and latex of a known size to be used as standard migration distances. The LDL and standard migration distances were read on a densitometry (590 nm). A quadratic equation (polynomial regression of Stokes) was used to convert migration distance into particle diameter; the estimated diameter of the greatest peak in each scan was identified as the LDL particle diameter. Insulin level was determined in duplicate by radioimmunoassay. Lipoprotein lipase *HindIII* polymorphism genotyping was performed by DNA extraction, polymerase chain reaction, and *HindIII* restriction enzyme digestion of the amplified products.¹⁸ Informed consent was obtained by all participants, and the study was approved by the local ethics committee.

Statistical Analysis

The Kolmogorov-Smirnov algorithm was used to determine whether each variable had a normal distribution, and only the distribution of

Lp(a) was significantly different from the normal distribution. Parameters are expressed as means \pm SD except for Lp(a), which is expressed as median (25th to 75th percentile). ANOVA, corrected with Tukey's post hoc test, tested the between-group differences in demographic, lipid, and metabolic variables. ANOVA for repeated measurements was used to assess differences among various times of observation.

The magnitude of postprandial triglyceride changes was estimated as 8-hour incremental area under the curve (AUC).⁵ Triglycerides were plotted against time, and the AUC from the beginning of the test to the 8th hour was calculated by the trapezoidal method after subtraction of the baseline values. The differences among the three groups in triglyceride incremental AUC were assessed by analysis of covariance (ANCOVA) after adjustment for body mass index and fasting insulin level. For Lp(a), statistical differences between groups and intragroup variations from baseline values were calculated by use of Kruskal-Wallis and Friedman tests and Tukey's post hoc test for multiple comparisons. Univariate correlations were obtained with Pearson's correlation test, and stepwise multiple linear regression tested the relationship of several variables to triglyceride incremental AUC.

RESULTS

The three groups did not differ by age and smoking habits (Table 1). Insulin level, waist-hip ratio, body mass index, and plasma triglyceride level were higher, and HDL cholesterol and HDL₂ cholesterol levels were significantly lower, in women with mixed hyperlipemia than in the other two groups. By selection, total and LDL cholesterol were greater in both hyperlipemic groups than in normal women; LDL particle size and HDL₃ cholesterol level were lower in subjects with mixed hyperlipemia than in normal subjects.

Postprandial lipid parameters after the oral fat load are reported in Table 2. Subjects with mixed hyperlipemia showed a significant increase in serum triglycerides after 4 hours, and the maximal value was reached 6 hours after the meal. A significant reduction was observed in HDL and HDL₂ cholesterol levels. Plasma Lp(a) levels decreased after 4 and 6 hours, in concomitance with the increase in triglyceride levels. A significant decrease in LDL size was observed at the 4th and

Table 1. Demographic Characteristics and Fasting Laboratory Parameters of the Study Population

	Normal (n = 17)	Hypercholesterolemia (n = 17)	Mixed Hyperlipemia (n = 16)
Age (yr)	57 \pm 8	54 \pm 6	55 \pm 5
Body mass index (kg/m ²)	26.7 \pm 1.3	25.9 \pm 2.3	27.9 \pm 1.6*†
Waist-hip ratio	0.84 \pm 0.08	0.89 \pm 0.05	0.97 \pm 0.05*†
Insulin (mU/L)	7.6 \pm 4.9	9.2 \pm 6.6	22.5 \pm 9.5*†
Triglycerides (mg/dL)	80 \pm 35	113 \pm 48	211 \pm 43*†
Total cholesterol (mg/dL)	205 \pm 22	269 \pm 34*	260 \pm 35*
LDL cholesterol (mg/dL)	129 \pm 16	187 \pm 34*	173 \pm 31*
HDL cholesterol (mg/dL)	59 \pm 18	58 \pm 15	43 \pm 11*†
HDL ₂ cholesterol (mg/dL)	22 \pm 11	20 \pm 9	11 \pm 3*†
HDL ₃ cholesterol (mg/dL)	39 \pm 8	38 \pm 7	32 \pm 9*
LDL size (nm)	27.0 \pm 1.1	26.6 \pm 1.6	25.4 \pm 1.1*
Lp(a) (mg/dL)	7.3 (4.4-18.7)	20.7 (17.1-37.9)	11.4 (8.3-24.4)
Lipoprotein lipase <i>HindIII</i> genotype			
H ⁺ /+	8/17	5/17	10/16
H ⁺ /-	5/17	9/17	6/16
H ⁻ /-	4/17	3/17	0/16

NOTE. Values are expressed as mean \pm SD; Lp(a) is expressed as median (25% to 75%).

* $P < .05$ v normal subjects.

† $P < .05$ v hypercholesterolemia.

Table 2. Postprandial Lipid Parameters After the Oral Fat Load

	Triglycerides (mg/dL)	LDL Size (nm)	HDL-C (mg/dL)	HDL ₂ -C (mg/dL)	HDL ₃ -C (mg/dL)	Lp(a) (mg/dL)
Controls						
0 h	80 ± 35	27.0 ± 1.1	59.2 ± 18	22.5 ± 11	38.7 ± 8	7.3 (3-36)
4 h	162 ± 112*	27.5 ± 0.6	58.0 ± 20	23.5 ± 10	37.7 ± 11	6.5* (2-32)
6 h	177 ± 95*	27.2 ± 1.3	56.9 ± 20*	23.6 ± 10	36.9 ± 12	6.5* (2-29)
8 h	116 ± 52*	27.2 ± 1.1	59.3 ± 19	24.6 ± 10	37.2 ± 11	6.9 (2-28)
Hypercholesterolemia						
0 h	113 ± 48	26.6 ± 1.6	57.8 ± 15	20.2 ± 9	38.8 ± 7	20.7 (11-52)
4 h	209 ± 94*	26.7 ± 1.9	57.1 ± 10*	19.1 ± 8*	38.9 ± 8	20.0* (10-5)
6 h	205 ± 11*	26.6 ± 1.9	55.5 ± 14*	19.4 ± 9	37.7 ± 8	21.9 (10-48)
8 h	148 ± 80*	26.8 ± 1.9	57.8 ± 14	20.6 ± 8	38.8 ± 8	21.9 (10-49)
Mixed hyperlipemia						
0 h	211 ± 43	25.4 ± 1.1	43.6 ± 11	11.5 ± 3	32.2 ± 9	11.4 (8-25)
4 h	402 ± 99*	24.7 ± 1.3*	41.3 ± 11*	9.2 ± 2	32.2 ± 8	10.7* (6-21)
6 h	460 ± 107*	24.7 ± 1.4*	39.7 ± 10*	8.5 ± 2*	31.1 ± 8	10.0* (6-21)
8 h	371 ± 123*	24.8 ± 3.5	41.3 ± 11*	10.0 ± 3	31.3 ± 10	10.5 (6-22)

Abbreviations: HDL-C, HDL cholesterol; HDL₂-C, HDL₂ cholesterol; HDL₃-C, HDL₃ cholesterol.

* $P < .05$ v baseline values.

6th hours (Fig 1). In women with hypercholesterolemia, triglyceride levels peaked at the 4th and 6th hours, with a concomitant decrease in HDL and HDL₂ cholesterol levels. Lp(a) had a significant decrease after 4 hours, and no variation in LDL size was observed. In normolipemic subjects, triglyceride levels increased at the 4th and 6th hours. A concomitant decrease in HDL cholesterol level was observed at the 6th hour. Lp(a) levels showed a concomitant decrease at the 4th and 6th hours. No other significant variations in HDL₂ and HDL₃ cholesterol levels or LDL size after the fat load were observed.

The incremental triglyceride AUC was greater in women with mixed hyperlipemia ($1,326 \pm 372 \text{ mg} \cdot \text{dL}^{-1} \cdot \text{h}^{-1}$) than in normal women ($484 \pm 384 \text{ mg} \cdot \text{dL}^{-1} \cdot \text{h}^{-1}$; $P < .0001$) and in women with hypercholesterolemia ($473 \pm 223 \text{ mg} \cdot \text{dL}^{-1} \cdot \text{h}^{-1}$; $P < .0001$). The differences remained significant after

adjustment for baseline differences among groups in body mass index and insulin levels ($P < .003$ v controls; $P < .001$ v hypercholesterolemia). No differences in triglyceride incremental AUC were observed between hypercholesterolemic and normolipemic subjects. The allelic distribution of lipoprotein lipase *HindIII* polymorphism did not differ in the three groups ($P = .20$; Table 1). When the study population was divided on the basis of the presence or absence of at least one H⁺ allele, triglyceride incremental AUC was significantly higher in H⁺ allele carriers ($n = 43$, $801 \pm 79 \text{ mg} \cdot \text{dL}^{-1} \cdot \text{h}^{-1}$) than in subjects with H⁻/H⁻ genotype ($n = 7$, $393 \pm 74 \text{ mg} \cdot \text{dL}^{-1} \cdot \text{h}^{-1}$; $P < .01$).

Using multiple linear regression analysis, we tested the independent relationship of several variables to triglyceride incremental AUC in the whole study population. Fasting insulin ($P < .0001$) and triglyceride ($P < .03$) levels and an elevated waist-hip ratio ($P < .04$) were all independent predictors of a greater incremental AUC. The resulting equation was triglyceride incremental AUC ($\text{mg} \cdot \text{dL}^{-1} \cdot \text{h}^{-1}$) = $-708 + (23.5 \times \text{insulin [mU/L]}) + (1.91 \times \text{triglycerides [mg/dL]}) + (1,518 \times \text{waist-hip ratio})$. Overall, 74% of the observed variation in triglyceride incremental AUC was explained by the model (multiple $r = 0.86$). Age, body mass index, HDL and HDL₂ cholesterol levels, LDL size, and Lp(a) level failed to enter the multivariate equation.

DISCUSSION

The main finding of the present study is that postprandial lipid changes after a standardized oral fat-loading test showed marked differences in hyperlipemic and normolipemic postmenopausal women. Indeed, subjects with mixed hyperlipemia had a greater postprandial triglyceride response than hypercholesterolemic women and normolipemic control subjects. This result confirms and extends to the female sex previous studies reporting exaggerated postprandial triglyceridemia in men with endogenous hypertriglyceridemia,¹⁹ which is largely attributable to increased retention of large VLDL particles as a

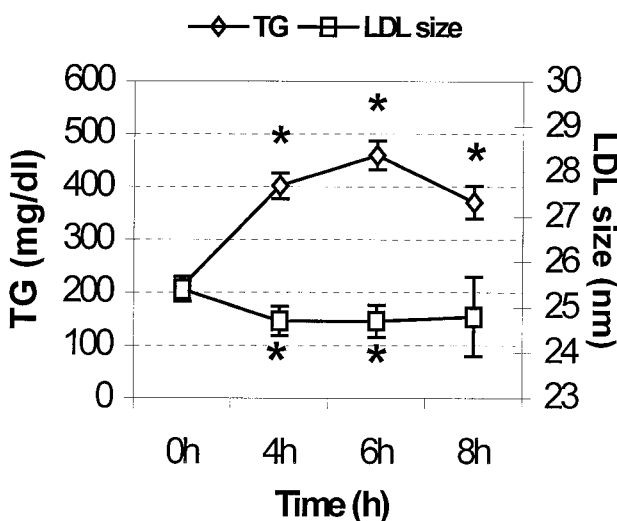


Fig 1. Postprandial serum triglyceride (TG) level and LDL particle size in 16 postmenopausal women affected by mixed hyperlipemia. Error bars represent SEM. * $P < .05$ compared with fasting values.

consequence of a removal defect.²⁰ However, a decreased chylomicron clearance capacity has also been observed in postmenopausal women because of the lack of endogenous estrogen.²¹

The postprandial triglyceride increase in women with mixed hyperlipemia was mirrored by a concomitant decrease in HDL and HDL₂ cholesterol and plasma Lp(a) concentrations. Normal and hypercholesterolemic women showed a significant decrease in HDL cholesterol level and a transient reduction in plasma Lp(a) levels. In this regard, Westerveld et al reported reduced plasma HDL cholesterol levels 3 to 8 hours after ingestion of an oral fat load in normolipemic postmenopausal women.¹² Moreover, an inverse relationship has been reported between fasting Lp(a) and triglyceride levels, suggesting that Lp(a) could form a metastable complex with triglyceride-rich lipoproteins, which could be removed at a higher rate,^{22,23} thus providing a possible explanation for the postprandial decrease in plasma Lp(a) levels.

We also found a transient postprandial decrease in LDL particle size in women with mixed hyperlipemia (Fig 1), but not in hypercholesterolemic and normolipemic subjects. Fasting small LDL particles have been considered predictive of postprandial hyperlipemia,²⁴ but to our knowledge this is the first study to document a transient reduction of LDL particle size during a fat-loading test. Our findings suggest that a moderate postprandial increase in plasma triglyceride levels, as observed in normal and hypercholesterolemic women, may not be sufficient to induce a significant decrease of LDL particle size. In contrast, a more substantial degree of postprandial hypertriglyceridemia, as observed in subjects with mixed hyperlipemia, is able to shift the distribution of circulating LDL toward smaller and denser species. In addition, a significant univariate correlation was found between serum triglyceride and fasting insulin levels and LDL size, both in the fasting and the postprandial states (results not shown). Thus, our data provide support for the notion that serum triglycerides²⁵ and insulin²⁶ are important determinants of the distribution of LDL subfractions.

In the present study, postmenopausal women with mixed hyperlipemia showed higher fasting insulinemia and several markers of the insulin-resistance syndrome, including greater body mass index and waist-hip ratio, lower HDL cholesterol level, and smaller LDL size than in normal and hypercholesterolemic subjects. The combination of multiple markers of insulin resistance in patients with mixed hyperlipemia is well documented²⁷⁻²⁹ and might have influenced the postprandial triglyceride response in our study. However, triglyceride incremental AUC remained significantly greater in women with mixed hyperlipemia than in the other two groups after adjustment for body mass index and fasting insulin, thus supporting

the view of an independent association between fasting hyperlipemia and postprandial hypertriglyceridemia.

In our study, fasting insulinemia and waist-hip ratio were both independent determinants of triglyceride incremental AUC, along with fasting triglyceride levels. This finding confirms previous studies, carried out in men or in combined groups of men and women with an age distribution including both premenopausal and postmenopausal women, in which insulinemia^{29,30} and abdominal fat accumulation³¹ were both determinants of postprandial triglyceride response. Taken together, these data support the view that fasting hyperinsulinemia and visceral obesity, which may both be considered as markers of insulin resistance,³² may contribute to postprandial hyperlipemia, probably through an increase in postprandial delivery of fatty acids to the circulation.^{33,34}

We also assessed the role of *HindIII* polymorphism of lipoprotein lipase. Lipoprotein lipase is a key enzyme involved in the first catabolic step of triglyceride-rich lipoproteins in plasma, and different mutations and polymorphisms have been described that impair the function of the enzyme.³⁵ *HindIII* polymorphism of lipoprotein lipase plays an important role in the determination of basal triglyceridemia, with a dose-dependent effect. Data on the relation between *HindIII* polymorphism and postprandial lipemia are scarce. In the only report on this topic,³⁶ no significant correlation was found between the presence of the H⁺ allele and postprandial triglyceride levels. In our study, the distribution of *HindIII* genotypes did not differ in the three groups; however, the subjects with at least one H⁺ allele had a greater triglyceride incremental AUC than H⁻ subjects, thus indicating a possible influence of H⁺ allele on postprandial lipemia. Additional data are needed to support the hypothesis that *HindIII* lipoprotein lipase polymorphism can influence postprandial lipemia.

In summary, the present results indicate that after a fat-loading test, both normolipemic and hyperlipemic postmenopausal women are characterized by multiple disturbances of the lipoprotein metabolism, including abnormal elevation of triglyceride levels and reduction in HDL cholesterol and HDL₂ cholesterol concentrations. Postmenopausal women with mixed hyperlipemia exhibited an amplification of these abnormalities and a transient reduction in LDL particle size, which might be additional markers of cardiovascular risk. Fasting plasma triglyceride levels, insulin levels, and abdominal fat accumulation were independent determinants of these metabolic disturbances. The presence of other features of the insulin resistance syndrome, including reduced HDL cholesterol level, smaller LDL particle size, and lipoprotein lipase *HindIII* polymorphism, could contribute to the deterioration of the postprandial lipemic response in these subjects.

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