Lipoprotein-associated phospholipase A₂, platelet-activating factor acetylhydrolase: a potential new risk factor for coronary artery disease

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Abstract

A specific and robust immunoassay for the lipoprotein-associated phospholipase A₂ (Lp-PLA₂), platelet-activating factor acetylhydrolase, is described for the first time. The immunoassay was used to evaluate possible links between plasma Lp-PLA₂ levels and atherosclerosis risk amongst susceptible individuals. Such an investigation was important because Lp-PLA₂ participates in the oxidative modification of low density lipoprotein by cleaving oxidised phosphatidylcholines, generating lysophosphatidylcholine and oxidised free fatty acids. The majority of Lp-PLA₂ was found associated with LDL (approximately 80%) and, as expected, enzyme levels were significantly positively correlated to LDL cholesterol. Plasma Lp-PLA₂ levels were significantly elevated in patients with angiographically proven coronary artery disease (CAD) when compared with age-matched controls, even though LDL cholesterol levels did not differ significantly. Indeed, when included in a general linear model with LDL cholesterol and other risk factors, Lp-PLA₂ appeared to be an independent predictor of disease status. We propose, therefore, that plasma Lp-PLA₂ mass should be viewed as a potential novel risk factor for CAD that provides information related to but additional to traditional lipoprotein measurements. © 2000 Elsevier Science Ireland Ltd. All rights reserved.

Keywords: Lipoprotein-associated phospholipase A₂; Potential new risk factor; Coronary artery disease

1. Introduction

Various dyslipidaemias exist that increase the predisposition of an individual to develop atherosclerotic coronary artery disease (CAD). Elevated plasma levels of low density lipoprotein (LDL) cholesterol have been shown in many studies to be associated with an increased incidence of CAD whereas plasma high density lipoprotein (HDL) levels are inversely related to risk [1,2]. Not surprisingly, plasma concentrations of apolipoprotein (apo) B and A-1, the major protein components of LDL and HDL, respectively, are also accepted risk factors for CAD. The precise pathophysiological mechanisms through which dyslipidaemias increase the likelihood of atherosclerosis remain poorly understood, but the oxidative modification of LDL within the arterial wall appears to play an important role [3,4].

Attention has recently focused on three lipoprotein-associated enzymes that can participate in the LDL oxidation process and as a consequence modulate the many pro-inflammatory activities of oxidised LDL (oxLDL). For instance, much of the protective activity of HDL against oxLDL has been attributed to the presence of paraoxonase and lecithin-cholesterol acyltransferase (LCAT), two enzymes found exclusively on HDL. Paraoxonase hydrolysates and inactivates pro-inflammatory lipid peroxides present in oxLDL [5,6],
and it was recently demonstrated that paraoxonase-null mice were more susceptible to atherosclerosis than wild-type littersmates [7]. LCAT, known for many years to be responsible for plasma cholesterol ester production, has also been shown to degrade both platelet activating factor (PAF) and oxidised polar phospholipids [8], two proposed atherogenic mediators [9].

The third enzyme thought to be intimately involved in LDL oxidation is the lipoprotein associated phospholipase A2 (Lp-PLA2), PAF acetylhydrolase. This enzyme, unlike the other two, is found predominately on LDL [10] and as such is better placed to participate directly in any oxidative modification process. Previous work has demonstrated that this phospholipase A2 is responsible for much if not all of the increased lysophosphatidylcholine content of oxidatively modified LDL [11–13]. It is also now recognised as a subtype of the growing phospholipase A2 superfamily [14] with a broad substrate preference because it can hydrolyse both PAF as well as more complex polar phosphatidylcholines [12,13,15]. The role of Lp-PLA2 in atherogenesis is currently under some debate because it can apparently play a dual function. Since much of the pro-atherogenic activity of oxidised LDL has been attributed to its increased lysophosphatidylcholine content, this PLA2 would appear to be pro-inflammatory in nature [16]; paradoxically, the ability of Lp-PLA2 to hydrolyse PAF and related phospholipids also suggests an anti-inflammatory role [9,17]. More recently, Lp-PLA2 has also been shown to release bioactive oxidised free fatty acids during the oxidation of LDL [13] thus providing further support for the pro-inflammatory argument.

To help resolve this apparent paradox we have developed a specific ELISA to measure Lp-PLA2 mass to evaluate the potential link between plasma Lp-PLA2 levels and individuals with atherosclerosis. Previous studies have shown that the plasma degradation of PAF is increased in patients with atherosclerosis [18–20] and stroke [21], but there was no information on the identity of the enzyme involved. By directly measuring Lp-PLA2 mass we demonstrate that plasma Lp-PLA2 is an independent predictor of CAD and provides information additional to LDL cholesterol.

2. Methods

2.1. Subjects

A total of 148 male subjects, aged 39–74 years, from Glasgow, UK were recruited into the study. They were drawn from to one of three groups. (1) Subjects (n = 48) with CAD were identified from those who attended for coronary angiogram at the Glasgow Royal Infirmary and met the entry criteria of a positive angiogram at age < 55 years. To be referred for angiography in local practice, individuals have symptoms of angina and a positive angiogram is recorded if stenotic disease is present. The severity of disease was not recorded in this study. The mean age at blood sampling was 55 ± 8 years. (2) Post myocardial infarction (PMI) patients (n = 46) were recruited from the coronary rehabilitation classes at the Glasgow Royal Infirmary. They were at least 1 year post event which had taken place at age < 55 years. The mean age at sampling was 54 ± 8 years. (3) Fifty-four normal aged matched subjects (53 ± 10 years) were recruited from the local community. Blood pressure measurements were taken and a record of history of smoking and medication was collected by questionnaire. No patients in the CAD and PMI group had diabetes but one in the control group was receiving insulin. Treatment with nitrates, aspirin and β-blockers were similar in diseased subjects, with nine in each group undergoing treatment. One subject in the PMI and two in the CAD groups were being treated with simvastatin and one in the CAD was taking ciprofibrate.

Blood samples (30 ml) were drawn after an overnight fast into K2 EDTA at a final concentration of 1 mg/ml and the plasma was analysed for lipids, lipoproteins and apolipoproteins immediately. Aliquots (1 ml) of plasma were fast frozen in liquid nitrogen and stored at −70°C. These frozen samples were transported monthly on dry ice to SmithKline Beecham Laboratories in Welwyn, Herts, for analysis of Lp-PLA2. The identity of the samples was blind to the workers analysing Lp-PLA2 levels.

2.2. Lipoprotein, lipid, and apolipoprotein measurements

Assays of plasma cholesterol, triglyceride, VLDL cholesterol, LDL cholesterol and HDL cholesterol were performed by a modification of the standard Lipid Research Clinic’s protocol [22]. Apo AI and B were measured by immunoturbidimetry using kits purchased from Orion Diagnostics, Espoo, Finland.

2.3. Monoclonal antibody (Mab) generation

Lp-PLA2 was purified to homogeneity from human LDL exactly as described previously [12]. Mice were immunised with 25 μg of purified, non-denatured, human Lp-PLA2 in Freund’s complete adjuvant followed by booster doses of 25 μg in Freund’s incomplete adjuvant after 4 weeks as well as two doses of 20 μg intraperitoneally in saline. Spleen cells were harvested one day after the last dose and fused with myeloma cells according to the method of Zola [23]. Positive hybridomas were cloned twice by the limiting dilution method and the clones expanded to 1-l spinner flasks.
for large scale production. Monoclonal antibodies (50–100 mg) were purified from the tissue culture medium by Protein A chromatography (Prosep-A, Bioprocessing, Consett, UK) and isotyped using a standard kit produced by Amersham (Amersham, UK). The monoclonal antibodies used in the assay were IgG and were named Mab 2C10 (first antibody) and Mab 2B4 (second antibody).

2.4. Reagents for immunoassay

Microtitre plates, the plate washer and fluorimeter were from Wallac (Turku, Finland). The plate-shaker incubator was purchased from Amersham. The coating buffer was 50 mM Na₂HPO₄, 150 mM NaCl, and 0.02% Kathon (Rohm and Haas, London, UK), pH 7.4. The blocking buffer was 50 mM Tris–HCl, 150 mM NaCl, 1% bovine serum albumin and 0.02% Kathon, pH 7.4. The wash solution contained 10 mM Tris–HCl, 150 mM NaCl, 0.05% Tween 20, and 0.02% Kathon, pH 7.4. The assay buffer used was 50 mM Tris–HCl, 150 mM NaCl, 0.5% bovine serum albumin, 0.05% bovine γ-globulin, 0.01% Tween 40, 20mM diethylenetriaminepentaacetic acid, and 0.02% Kathon, pH 7.4. Biotinylated anti-Lp-PLA₂ Mab 4B4 was prepared using a kit supplied by Amersham.

2.5. Lp-PLA₂ two site immunoassay procedure

Microtitre plate wells were coated with the first Mab 2C10 (4 μg/ml, 100 μl/well) overnight at 4°C after which the contents of the well were aspirated and 250 μl per well of blocking buffer added for 60 min at 22°C. The wells were washed four times and 50 μl biotin labelled second monoclonal antibody 4B4 (2 μg/ml in assay buffer) and 50 μl purified recombinant Lp-PLA₂ standards (0.3–30 ng/ml), plasma (diluted 1/100 in assay buffer) or SMART fraction (diluted 1/10 in assay buffer), was added and the wells incubated for 60 min at 22°C. The plate wells were washed four times and 100 μl per well of Eu-streptavidin (200 ng/ml, Wallac) was added and incubated for 30 min at 22°C. After the immunoreaction was complete the wells were washed four times and 200 μl per well of enhancement solution (Wallac) added and after a 5 min incubation the fluorescence was measured. The fluorimeter software was used for curve fitting and result calculations.

2.6. Lipoprotein fractionation

Plasma samples were thawed, placed on ice, and filtered using a 0.3 μM sterile filter. Plasma was fractionated into its lipoprotein components by gel filtration on a Pharmacia FPLC SMART system. Routinely 30 μl of filtered plasma was analysed using a Superose 6PC 3.2/30 column equilibrated at 15°C with 150 mM NaCl, 1 mM EDTA, and 0.05% Na azide, pH 8. Lipoproteins were eluted at a flow rate of 30 μl/min and the fractions corresponding to VLDL, LDL, and HDL pooled based upon protein (absorbance at 280 nm), cholesterol and Lp-PLA₂ activity profiles.

2.7. Lp-PLA₂ assays

Either 50 μl aliquots of whole plasma or 30 μl of the fractionated lipoprotein pools were assayed using 50 μM PAF as described previously [12].

2.8. Statistics

Group differences were explored by ANOVA using Tukey’s and Fishers method for pairwise comparison. Regression analysis and correlation coefficients were obtained for continuous variables. All statistical tests were performed using Minitab version 10.0. The ability of a parameter to predict disease was tested by entering variables into a general linear model.

3. Results

3.1. Lp-PLA₂ two-site immunoassay

Fig. 1 shows that the Lp-PLA₂ calibration curve was linear over the range 0.3–30 ng/ml. The lowest calibrator had counts approximately five times above background. Excellent linearity of dilution was observed and the within- and between-assay coefficient of variation was 4 and 9%, respectively (n = 27). The recovery of Lp-PLA₂ added to 10 different patient samples was 95 ± 8%. Zero cross-reactivity in the assay was noted at a 1000-fold excess with three different human recombi-
nattant PLA₂’s which were generated at SmithKline Beecham Laboratories (data not shown): 14 kDa PLA₂ (CHO-derived), 85 kDa PLA₂ (baculovirus-derived), and a recently described related serine-dependent PLA₂ (baculovirus-derived). Both the 14 kDa PLA₂ and 85 kDa PLA₂ are calcium-dependent arachidonic acid selective enzymes [14], whereas the serine-dependent enzyme is calcium-independent and has 40% amino acid identity with Lp-PLA₂ [24].

Fig. 2 demonstrates, as expected, a clear co-elution of Lp-PLA₂ activity with immunoassay derived Lp-PLA₂ mass following separation of plasma lipoprotein fractions by FPLC. Most of Lp-PLA₂ is in the LDL size range.

3.2. Analysis of patient data

3.2.1. Lipids, lipoproteins, and apolipoproteins

There were no significant differences between normals and the control groups in age or blood pressure (Table 1). However, the number of smokers was higher in the diseased groups compared to control with 42% of the PMI group being smokers (ANOVA P = 0.014). There were no significant differences between normals and diseased groups in plasma cholesterol, triglyceride, VLDL cholesterol, LDL cholesterol and apo AI (Table 2). HDL cholesterol was lower in the diseased groups when compared to normal (P = 0.031) and both CAD and PMI groups were significantly different from control (P < 0.01). Analysis by ANOVA highlighted a significantly higher (P = 0.042) apo B in the diseased groups and this was due to the CAD group which gave a significance of P < 0.01 when compared to the normal group. The lipoprotein values were intermediate in the PMI group with higher levels in the CAD group except for HDLc which was lowest in CAD.

3.2.2. Lp-PLA₂ mass and activity

There was a strong positive correlation (r = 0.86, P < 0.001) between mass and activity of plasma Lp-PLA₂ (Fig. 3A). The same was true in each of the lipoprotein fractions for PLA₂ mass versus activity, VLDL (r = 0.45, P < 0.001), LDL (r = 0.60, P < 0.001) and HDL (r = 0.71, P < 0.001). Of the total Lp-PLA₂ mass associated with the lipoproteins, 5.8% was found in VLDL, 83.3% in LDL and 10.9% in HDL. Similar findings were observed for PLA₂ activity with 7% in VLDL, 80.4% in LDL and 12.6% in HDL. Recoveries of plasma Lp-PLA₂ mass, plasma Lp-PLA₂ activity and total plasma cholesterol from the lipoproteins fractionated by FPLC were all 95%.

Plasma Lp-PLA₂ mass was significantly correlated with cholesterol (r = 0.59, P < 0.001), triglyceride (r = 0.23, P < 0.01), VLDLc (r = 0.30, P < 0.001), LDLc (r = 0.57, P < 0.001), HDLc (r = −0.23, P < 0.01) and apo B (r = 0.45, P < 0.001) as shown in Table 3. Likewise PLA₂ mass in LDL was significantly correlated with cholesterol (r = 0.56, P < 0.001), LDLc (r = 0.58, P < 0.001) and apo B (r = 0.39, P < 0.001). The linear relationship between Lp-PLA₂ mass and LDLc is illustrated in Fig. 3B.

Plasma Lp-PLA₂ mass differed significantly between the groups (ANOVA P = 0.009) with highest levels in CAD which differed significantly from control (P <

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**Table 1**

Differences between groups

<table>
<thead>
<tr>
<th>Parameters</th>
<th>Control</th>
<th>CAD</th>
<th>PMI</th>
<th>P</th>
</tr>
</thead>
<tbody>
<tr>
<td>N</td>
<td>52</td>
<td>43</td>
<td>43</td>
<td></td>
</tr>
<tr>
<td>Age (years)</td>
<td>53.6 (1.4)</td>
<td>55.1 (1.2)</td>
<td>53.0 (1.3)</td>
<td>0.491</td>
</tr>
<tr>
<td>BP systolic</td>
<td>131.9 (2.7)</td>
<td>132.4 (3.2)</td>
<td>131.9 (2.7)</td>
<td></td>
</tr>
<tr>
<td>BP diastolic</td>
<td>80.5 (2.1)</td>
<td>80.8 (1.5)</td>
<td></td>
<td>0.375</td>
</tr>
<tr>
<td>Smokers (%)</td>
<td>20</td>
<td>25</td>
<td>42</td>
<td>0.014</td>
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</table>

* Values are mean (S.E.M.).

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**Table 2**

Lipids and lipoproteins

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<th>CAD</th>
<th>PMI</th>
<th>P</th>
</tr>
</thead>
<tbody>
<tr>
<td>Cholesterol (mmol/l)</td>
<td>5.61 (0.13)</td>
<td>5.92 (0.16)</td>
<td>5.46 (0.15)</td>
<td>0.084</td>
</tr>
<tr>
<td>Triglyceride (mmol/l)</td>
<td>1.58 (0.10)</td>
<td>2.18 (0.26)</td>
<td>1.88 (0.15)</td>
<td>0.053</td>
</tr>
<tr>
<td>VLDLc (mmol/l)</td>
<td>0.725 (0.05)</td>
<td>1.00 (0.12)</td>
<td>0.85 (0.06)</td>
<td>0.052</td>
</tr>
<tr>
<td>LDLc (mmol/l)</td>
<td>3.80 (0.13)</td>
<td>3.96 (0.14)</td>
<td>3.65 (0.14)</td>
<td>0.292</td>
</tr>
<tr>
<td>HDLc (mmol/l)</td>
<td>1.13 (0.04)</td>
<td>0.99* (0.04)</td>
<td>1.00* (0.05)</td>
<td>0.031</td>
</tr>
<tr>
<td>Apo AI (g/l)</td>
<td>1.23 (0.03)</td>
<td>1.20 (0.05)</td>
<td>1.17 (0.07)</td>
<td>0.710</td>
</tr>
<tr>
<td>Apo B (g/l)</td>
<td>1.08 (0.04)</td>
<td>1.25* (0.06)</td>
<td>1.20 (0.06)</td>
<td>0.042</td>
</tr>
</tbody>
</table>

* Values are mean (S.E.M.), P from ANOVA.
* Significant by Fishers pair wise comparison P < 0.05.
0.001) (Table 4). The level of the enzyme in LDL followed the same trend (CAD versus control, \( P < 0.01 \)) but that in VLDL and HDL did not differ between the groups.

General linear models (GLMs) were used to test for predictors of disease. Controls were compared with all disease subjects (CAD and PMI) in model A, with those with CAD in model B and with those with PMI in model C Table 5. Plasma Lp-PLA2 mass was a significant predictor of risk in models A and B and showed a trend in model C. Models A and B indicate that Lp-PLA2 was an independent predictor of risk when the common risk factors, i.e. LDLc, HDLc, smoking and systolic blood pressure, were included in the multivariate models. In a GLM with apo B and Lp-PLA2, the mass of the enzyme was still a significant predictor of CAD \((r^2 = 0.91, P = 0.047)\) but not of PMI status. When HDLc, smoking and systolic BP were added to the model the trend towards higher Lp-PLA2 was still evident but was no longer significant. Apo B, like LDLc, was not a significant predictor in any GLM model when HDLc, smoking and systolic BP were included in the model.

4. Discussion

In view of the known positive correlation between plasma Lp-PLA2 activity and LDLc levels \([25–27]\) we attempted to ensure that the group of male subjects studied were well matched for fasting cholesterol levels \((5.5–5.9 \text{ mmol/l})\) and LDLc levels \((3.7–4.0 \text{ mmol/l})\). Thus, we intentionally chose PMI patients who were at least 1 year post event because acute MI is associated with depression of plasma lipoprotein concentrations and plasma Lp-PLA2 activity \([28,29]\). Cholesterol and LDLc did not differ significantly between the three groups although plasma triglycerides \((1.6–2.2 \text{ mmol/l})\) did show a tendency to be higher in the two diseased groups, CAD and PMI. HDLc, however, was significantly decreased in both patient groups \((1.0 \text{ versus } 1.13 \text{ mmol/l}, P < 0.01)\) and apo B significantly elevated in the CAD but not the PMI group. Contrary to earlier findings, PMI subjects were closer to normal than were the CAD group. We attribute to greater intervention and lifestyle changes in the year following the MI. The CAD subjects studied were attending a risk factor clinic for the first time and so were likely not to have modified their risk factors.

Several workers have reported \([10,30–32]\) that Lp-PLA2 (or PAF-AH) is transported in plasma predominantly associated with LDL and, in agreement, we

<table>
<thead>
<tr>
<th>Table 3</th>
<th>Correlations between Lp-PLA2 mass and lipoproteins</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Plasma PLA2</td>
</tr>
<tr>
<td>Cholesterol</td>
<td>0.592***</td>
</tr>
<tr>
<td>Triglyceride</td>
<td>0.233***</td>
</tr>
<tr>
<td>VLDLc</td>
<td>0.299***</td>
</tr>
<tr>
<td>LDLc</td>
<td>0.574***</td>
</tr>
<tr>
<td>HDLc</td>
<td>−0.226**</td>
</tr>
<tr>
<td>Apo AI</td>
<td>−0.003</td>
</tr>
<tr>
<td>Apo B</td>
<td>0.450***</td>
</tr>
</tbody>
</table>

* \( P < 0.05 \).
** \( P < 0.01 \).
*** \( P < 0.001 \).

<table>
<thead>
<tr>
<th>Table 4</th>
<th>Lp-PLA2 mass and patient group*</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Control</td>
</tr>
<tr>
<td>Plasma PLA2 (ng/ml)</td>
<td>1028.1 (38.5)</td>
</tr>
<tr>
<td>VLDL-PLA2 (ng/ml)</td>
<td>64.6 (4.6)</td>
</tr>
<tr>
<td>LDL-PLA2 (ng/ml)</td>
<td>817.7 (36.8)</td>
</tr>
<tr>
<td>HDL-PLA2 (ng/ml)</td>
<td>104.4 (12.8)</td>
</tr>
</tbody>
</table>

* Values are mean (S.E.M.), \( P \) from ANOVA.
* Significant by Fishers pair wise comparison \( P < 0.05 \).
Table 5
Predictors of coronary disease using a general linear model

<table>
<thead>
<tr>
<th></th>
<th>Model A (CAD and PMI)</th>
<th>Model B (CAD)</th>
<th>Model C (PMI)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>$r^2$</td>
<td>$P$</td>
<td>$r^2$</td>
</tr>
<tr>
<td>LDLc</td>
<td>0.0015</td>
<td>0.207</td>
<td>0.18</td>
</tr>
<tr>
<td>HDLc</td>
<td>1.35</td>
<td>0.041</td>
<td>1.31</td>
</tr>
<tr>
<td>Lp-PLA2 mass</td>
<td>1.64</td>
<td>0.01</td>
<td>1.38</td>
</tr>
<tr>
<td>Smoking</td>
<td>2.13</td>
<td>0.002</td>
<td>1.61</td>
</tr>
<tr>
<td>Systolic BP</td>
<td>0.12</td>
<td>0.444</td>
<td>0.02</td>
</tr>
<tr>
<td>Overall</td>
<td>26.1</td>
<td></td>
<td>18.3</td>
</tr>
</tbody>
</table>

* $r^2$ is the estimated independent contribution of the variable.

observed that 80% of the PAF-hydrolysing activity within plasma was associated with LDL. Additionally, through the development of a robust and selective immunoassay for Lp-PLA2, we showed for the first time that the lipoprotein distribution of Lp-PLA2 mass matches exactly that of PAF-AH activity. The strong correlation ($r = 0.86, P < 0.001$) between Lp-PLA2 mass and activity in plasma clearly shows that Lp-PLA2 contributes the majority of the PAF-hydrolysing capability of plasma.

We found a strong correlation between Lp-PLA2 mass and plasma LDLc in agreement with earlier studies [25–27] which had based their observations on enzyme activity measurements. Similarly, we noted a strong correlation between plasma Lp-PLA2 mass and plasma cholesterol, triglyceride, VLDLc, and apo B (Table 3) in agreement with Guerra et al. [27] who also demonstrated an association between PAF-AH activity and apo B, cholesterol, and triglyceride. The significant negative correlation observed by others [25–27] between the PAF-degrading capacity of plasma and HDLc was also corroborated in our Lp-PLA2 mass measurement studies. We therefore propose that the immunoassay developed offers a simple, sensitive, and reliable method for the detection of Lp-PLA2 in plasma compared to activity measurements.

Our observation that plasma Lp-PLA2 mass levels were significantly elevated in the CAD group supports and confirms previous reports [18–20] suggesting a link between this lipoprotein-bound enzyme and individuals with atherosclerosis. This positive association was made all the more striking since LDL cholesterol levels were not significantly different between normals and CAD. A comparison of risk factors in CAD and normal groups indicated that plasma Lp-PLA2 mass was a clearer marker of risk on a case control basis than either plasma cholesterol or LDL cholesterol. Even though the enzyme exhibited a strong relationship with LDL this was not the sole cause of the risk association since it was an independent predictor in the linear model. Indeed, plasma Lp-PLA2 was a stronger predictor of disease in the CAD group than apo B. Part of the reason why elevated Lp-PLA2 was such a good predictor of disease may, as suggested by Guerra et al. [27], be due to it reflecting the presence of LDL particles that are cleared slowly from the circulation. They speculated that the rate of removal of LDL from the circulation would determine the clearance rate of Lp-PLA2 and therefore the plasma mass of the enzyme for any individual. Consistent with this idea has been the observation that Lp-PLA2 is found enriched in the highly atherogenic small dense LDL [33], a LDL subtype known to have a prolonged plasma half-life [34]. Moreover, this association would tend to align Lp-PLA2 with a predominantly pro-inflammatory role since these particles are considered pro-atherogenic because they preferentially seed in the arterial intima and, being older, they are prone to oxidative modification [35].

In summary, the studies strongly support plasma Lp-PLA2 as a potential novel risk factor for CAD which provides information related to and additional to that obtained from traditional lipid analyses.

Acknowledgements

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