MPO G-463A Gene Polymorphism and Circulating Matrix Metalloproteinase-9, Myeloperoxidase Levels in Coronary Artery Disease Patients

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Abstract

Objective: Coronary artery disease (CAD) patients living in Elazığ region is to assess the levels of matrix metalloproteinase-9 (MMP-9) and myeloperoxidase (MPO), polymorphisms MPO gene.

Research Design and Methods: Eighty-eight patients with angiographically diagnosed coronary artery disease between the ages of 18-80 who applied to Department of Cardiology, Faculty of Medicine, Firat University were enrolled in this study. MPO, MMP-9 levels in plasma samples of individuals were determined by enzyme-linked immunosorbent assay (ELISA) according to the kit procedure. MPO polymorphism was studied using the PCR-RFLP technique.

Results: When plasma MPO and MMP-9 levels were compared in the control and CAD groups; MPO levels were statistically significant (p <0.05). MMP-9 levels were not statistically significant. MPO gene polymorphism was not found statistically significant in CAD.

Conclusion: MPO gene polymorphism is thought to be associated with greater population populations in CAD and may be used in the future as a biomarker for plasma MPO enzyme levels in CAD.

Key words: MPO, coronary artery disease, MMP-9

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Introduction

Oxidative stress plays an important role in the initiation and progression of cardiovascular disease (1). Oxidative stress biomarkers have been identified in coronary artery disease (CAD) (2-4). CAD has a multifactorial etiology, and it is thought to result from an interaction between genetic background and environmental factors (5-7). Myeloperoxidase (MPO) is a potent oxidative enzyme and it is expressed by neutrophils, monocytes, and some populations of macrophages. It catalyzes the modification of LDL in atherogenesis (8). MPO
has an important role in infectious diseases and in CAD (9). Several studies have shown that there is an association between MPO levels and CAD (8). Levels of MPO activity is associated with various genetic polymorphisms so that associated with cardiovascular risk (10). It has been shown that elevated MPO levels are associated with coronary artery disease (CAD) (11).

The gene coding for human myeloperoxidase, is located in chromosome 17q23.1. A few polymorphisms are described for the MPO gene that affects its transcription (12). In this polymorphism, the presence of a G rather than A at position 463 bp gives rise to three different genotypes: AA, AG and GG. To date, a number of studies have been suggested the potential association between the MPO G-463A polymorphism and CAD. However the results were inconsistent. Some of this studies have shown that GG genotype is a significant predictor of CAD and endothelial dysfunction when compared to AA/AG genotypes (13-16).

Matrix metalloproteinases (MMPs) are a family of zinc-containing endoproteinases which mainly secreted by monocyte-derived macrophage and vascular smooth muscle cells (19). MMP-9 is the most studied enzyme in the MMP-family. It is a key enzyme which determinant of extracellular matrix degradation, and collagen is the main substrate of MMP-9. MMP-9 concentrations and activity was higher in human atherosclerotic plaques (20-23). The serum level of MMP-9 might be a sensitive inflammation marker (24) and a predictor of cardiovascular mortality in patient with CAD (25). The purpose of this study was to test the hypothesis that MPO polymorphism ~463A/G is associated with CAD. In addition, as secondary objective we also investigated the association between CAD and MPO, MMP-9 plasma levels. Oxidative stress parameters have been evaluated in CAD frequently. However, there has not been much published work related to oxidative stress in CAD patients in Turkey. Similarly, not much literature is available on myeloperoxidase and MMP-9 level; MPO gene polymorphism and its association with lipids in CAD patients in Turkey. We, therefore evaluated these parameters in CAD patients.

Materials and Methods
The patient population and documentation of CAD severity
This was a case–control study. The cases and controls were the angiographically confirmed. The study sample comprised 176 persons who underwent coronary angiography for diagnostic purposes. The angiograms were assessed by two cardiologists who were unaware that the patients were to be included in the study. The three main arteries (left anterior descending artery, left circumflex artery, or right coronary artery) were classified as free of disease, or presenting ≥50 % stenosis. Patients having a normal angiogram with no atherosclerosis or lesions in coronary arteries were considered as CAD control subjects (n = 88). (Table 1). Written informed consent was obtained from all subjects, and a local ethical committee approved the study protocol.

Blood sample collection
Patients were fasted for at least 6 h before venous blood samples were drawn into 5 mL EDTA vacuum tubes and blood samples were separated in a refrigerated centrifuge within 15 min of collection for determination of plasma MPO and MMP-9. Plasma was divided into small aliquots, stored at −20°C until analysis. For determination of serum lipids, a blood sample was obtained from the cubital vein of each participant and the samples were sent for detection within 1 h. by using the Advia 1800 (Siemens Diagnostics, Tarrytown, NY, USA) autoanalyzer and reagents.

Determination of plasma levels of MPO and MMP-9
Plasma levels of MPO (Boster Biological Technology Co.,Ltd. USA; Cat no: EK0850) and MMP-9 (Boster Biological Technology Co.,Ltd. USA; Cat no: EK0465) were measured by enzyme-linked immunosorbent assay according to kit protocols.

Genotyping
DNA was extracted from EDTA anti-coagulated whole blood from all subjects (controls and patients) and was prepared from DNA isolation kit (Sigma-Aldrich, GenElute Blood genomic DNA Kit, Catalog no: NA2010, Germany). The MPO 463 -A/G polymorphism was genotyped by restriction fragment length polymorphism (RFLP) assay. The polymorphic site at position -463 of the MPO gene was
amplified with use of forward primer (50-CGG TAT AGG CAC ACA ATG GTG AG-30) and reverse primer (50-GCA ATG GTT CAA GCG ATT CTT C-30) as described in the literature. Amplification was carried out in a volume of 20 μl containing 1 - 10μg of genomic DNA, 10 pmol of each primer, 0.5 mM of each dNTP, reaction buffer, and 1.0 U of Taq DNA polymerase. Amplified products were digested overnight with Aci I at 37 °C. The resulting fragments were 289 bp for A/A, 289, 169, 120 bp for A/G and allele and 169, 120 bp for GG genotype.

Statistical analysis
All of the statistical analyses were performed by using the SPSS 12.0 statistical package. We have presented normally distributed data as mean ± S.D. Categorical variables are presented by frequency of counts, and inter group comparisons analysed by a chi-squared analysis. We compared cases and controls on biochemical parameters and MPO levels using Student’s t-test. Anova test was used to investigate the plasma lipid, MPO and MMP-9 levels between the genotypes. When we analyzed the quantitative relationships between MPO and MMP-9, bivariate correlation coefficients were calculated, using Pearson’s for parametric data. For the comparisons of the values of different groups, P< 0.05 was considered statistically significant.

Results
Clinical characteristics of patients with and controls:
The clinical characteristics of the patients and controls are shown in Table 1. Total cholesterol and LDL cholesterol levels were significantly higher in patients (p<0.0001 and p<0.01 respectively). Levels of triglyceride and HDL cholesterol were not significantly different among groups. Sistol blood pressure and the frequency of smoking were higher in patients (p<0.05, p<0.01, respectively). MPO level was higher in CAD patients (p<0.05) although MMP-9 level was not significantly different among groups (figure 1).

Table 1. Clinical characteristics of individuals with CAD and controls

|                          | Control (n=88) | CAD (n=88) | p     |
|--------------------------|---------------|------------|-------|
| Gender (F/M)             |               |            | 0.224 |
| Age (years)              | 50.2±11.41    | 62.6±11.81 | 0.000**|
| Smoking, (%)             |               |            |       |
| %12,10                   |               | %38,59     | 0.008*|
| Systolic blood pressure (mmHg) | 119.8±15.43   | 126.8±23.84 | 0.032* |
| Diastolic blood pressure (mmHg) | 74.8±11.48    | 79.4±10.42 | 0.332 |
| Total cholesterol (mg/dL) | 175.4±51.17   | 206.5±51.01 | 0.000**|
| HDL-cholesterol (mg/dL)  | 45.2±9.10     | 42.4±14.84 | 0.130 |
| LDL-cholesterol (mg/dL)  | 114.8±41.16   | 132.4±40.64 | 0.007* |
| VLDL-cholesterol (mg/dL) | 32.4±27.64    | 31.7±22.76 | 0.529 |

Figure 1. Plasma MPO and MMP-9 levels in control and CAD patients.
**Genotype frequencies**

Genotypic and allelic frequencies for the study groups are shown in Table 2. No statistically significant difference found in the distribution of the MPO -463 G/A genotype and allele frequency in the patients and control groups.

| Table 2. Genotype and allele frequencies among groups |
|------------------------------------------------------|
| **CONTROL** | **CAD** |
| Genotype | n:88 | n:88 |
| GG Genotype | 63 (% 71,6) | 64 (% 72,7) |
| AG Genotype | 23 (% 26,1) | 21 (% 23,9) |
| AA Genotype | 2 (%2,3) | 3 (% 3,4) |
| Allele | | |
| G Allele | %84,66 | %84,66 |
| A Allele | %15,34 | %15,34 |

**Association of the MPO gene polymorphism with lipids, MPO and MMP-9 levels.**

Table 3 shows the association between the MPO gene polymorphism and lipid levels, Figure 2 shows the association between the MPO gene polymorphism and MPO, MMP-9 levels in controls and patients with CAD, respectively. The higher total cholesterol, LDL-cholesterol and HDL cholesterol level was shown in the AA genotype in controls and patients but this was not statistically significant. No statistically significant difference found in the levels of MPO and MMP-9 among genotypes of MPO in controls and patients with CAD.

| Table 3. Lipid levels of the study subjects according to MPO genotypes in controls and patients. |
|------------------------------------------------------|

![Figure 2](image-url)
in this study we found that in healthy groups with G allele had elevated MPO levels than A allele (p<0.05) and among subjects with G alleles, CAD patients with G allele had higher level of MPO compared with controls (p<0.005) (Figure 3).

**Figure 3.** MPO levels in CAD patients and control subjects according to MPO alleles.

**Correlations between MPO and MMP-9 in total population**

Figure 4 shows the correlations between MPO and MMP-9 level in total population. There was a significant positive correlation between MPO and MMP-9 level ($r = 0.243$, $P = 0.01$).

**Figure 4.** Correlations between MPO and MMP-9 in total population

**Discussion**

In this study we assessed plasma levels of MPO, MMP-9 and MPO gene polymorphism in CAD patients and control subjects in whom the presence of CAD was defined by coronary angiographic examination. The main findings of this study are as follows: (i) In patients with angiographically proven CAD, plasma MPO levels are elevated as compared with subjects without CAD; (ii) In patients with angiographically proven CAD, plasma MMP-9 levels are not difference as compared with subjects without CAD; (iii) We did not observed any significant differences between MPO genotypes and study groups and (iv) A positive correlation between MPO and MMP-9 level was found.
CAD remains a heterogeneous disease with a wide range of clinical presentations and outcomes. Various systemic markers of inflammation have been investigated associated with CAD and cardiovascular events (26). The importance of PMN degranulation of MPO in the coronary circulation have emerged as powerful predictors of adverse outcome in patients with CAD (27–29). MPO is a redox-active hemoprotein released by activated neutrophils and is associated with inflammation and oxidative stress. Its role in catalyzing the formation of oxidizing agents. Hypochlorous acid, tyrosyl radical, and nitrogen dioxide are the products of MPO and they can contribute to oxidative damage of host lipids and proteins, leading to atherosclerosis. Brennan and colleagues (28) assessed the predictive value of serum MPO levels in patients with chest pain at a hospital emergency department. Serum MPO levels were significantly higher in patients who subsequently had a myocardial infarction (MI) within 16 hours of presentation (28). In a case-control study of 158 patients with CAD and 175 controls, Zhang et al. (11) have shown that elevated levels of blood MPO were associated with the presence of CAD. In the present study, we showed that corroborates the findings by Zhang et al. (11) in a series of patients in Turkey.

A few polymorphisms are described for the MPO gene, including the −638C/A, the V53F and the −463A/G (30). These polymorphisms could be genetically associated with the pathophysiology of cardiovascular diseases. We chose the −463A/G due to continuous literatures associated with coronary artery disease (13–16). Some studies have shown that GG genotype is associated with increased incidence of CAD (13–16), however, others failed to demonstrate a relation between MPO polymorphisms and atherosclerosis. We did not observed any significant differences between MPO genotypes and study groups. Furthermore, according to MPO −463 G/A allele in healthy groups with G allele had elevated MPO levels than A allele (p<0.05) and subjects with G alleles, CAD patients with G genotype had higher level of MPO compared with controls (p<0.005).

Our study and the results have present certain limitations, since the study-group was obtained from a single hospital in East of Turkey, and the samples might not include all the characteristics of patients from other centers. It is also accepted that CAD is a disease effected by multiple genes and environmental factors. Furthermore, since the MPO AA genotype is relatively infrequent, the actual value of genotype detection in the primary prevention of CAD may be disputable. However, to date, a number of studies have been conducted to assess the potential association between the MPO G−463A polymorphism and CAD. However the results were inconsistent.

MMPs had been identified in vascular remodeling process. They play an important role in the pathogenesis and progression of atherosclerosis (31). Several studies have shown the importance of these enzymes in development and progression of heart failure (32–34).

Some cross-sectional studies have reported that the plasma MMP-9 level was significantly higher in CAD patients (35,36). Blankenberg et al. (2003) showed that plasma MMP-9 concentration was higher in patients with CAD and might be a novel predictor of cardiovascular disease (25). In the present study, we investigated the association of the MMP-9 levels with coronary artery disease. We did not observed any significant differences in MMP-9 levels between controls and CAD patients. However, some other cohort studies suggested that serum MMP-9 is not a strong predictor of CAD (37,38). These inconsistent results can be explained as; plasma MMP-9 level is affected by many factors, such as genetic polymorphisms of MMP-9, drugs, and other factors. Lacchini et al. (2010) found that there are interethenic differences in MMP-9 genetic polymorphisms (39). Furthermore, in our study, MPO levels were correlated with the increase in MMP-9 concentration (P = 0.01), supporting the previous mentioned hypothesis, that neutrophils could be the major contributors for increased MMP-9 levels.

Conclusions
Observations from this study suggest that MPO level was significantly higher in CAD patients but there was no significant association between MMP-9 level and CAD. We found that there was no significant differences between MPO genotypes and study groups. It is also generally accepted that CAD is a disease produced by both multiple genes and environmental factors, and many other genes could be potential candidates for CAD. Our results have present certain limitations, since the study-group was obtained from a single hospital in east of Turkey, and the samples might not include all the characteristics of patients from other centers. However, further studies may be carried out in a larger population.

Conflict of Interest
The authors declare no conflicts of interests.
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