Methylene tetrahydrofolate reductase genotype and the risk and extent of coronary artery disease in a population with low plasma folate

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Abstract

Objective—To determine the effects of the thermolabile methylene tetrahydrofolate reductase (MTHFR) mutation on the presence and extent of coronary atherosclerosis in a population with low plasma folate.

Methods—242 consecutive patients undergoing coronary angiography were prospectively evaluated for conventional risk factors, plasma homocysteine, vitamin B-12, and folate, and MTHFR genotype. The severity of coronary atherosclerosis was determined by the Leaman score.

Results-Mean (SD) plasma homocysteine was 15.6 (10) µmol/l in controls and 18.5 (11) µmol/l in patients with coronary arterv disease (p > 0.05).homocysteine concentrations Plasma above 15 µmol/l were a risk factor for coronary artery disease (p = 0.03, risk ratio 2.1, 95% confidence interval (CI) 1.07 to 4.4). Homocysteine remained an independent risk factor on multivariate analysis when conventional risk factors were taken into account (p = 0.04). Homocysteine concentrations above 15 µmol/l were correlated with the extent of atherosclerosis (p = 0.04, risk ratio 3.2, 95% CI 1.3 to 8.2). Homocysteine had no effect on other lipid variables (p > 0.05). Plasma folate was 15.8 (7.2) nmol/l in controls and 11.5 (2.9) nmol/l in patients with coronary artery disease. Plasma folate concentrations below 12.9 nmol/l (5.7 ng/ml) conferred a risk for coronary artery disease (p = 0.03, risk ratio 2.42, 95% CI 1.05 to 5.59). When the MTHFR genotype was determined, the TT genotype was present in 7.4% of patients and 5.2% of controls (p > 0.05). The prevalence of alleles was within the Hardy-Weinberg equilibrium $(TT 7, CT 40, CC 53, \chi^2 = 2.3, p = 0.3)$. The highest homocysteine concentrations were found in patients with the TT genotype and folate below the median of the population (p = 0.01). The extent of coronary atherosclerosis judged by the Leaman score was significantly higher in patients with the TT genotype (p = 0.03). Conclusions-Plasma homocysteine over 15 µmol/l was a significant risk factor for the presence and extent of coronary artery disease. The mean plasma folate of the population was low and correlated negatively with homocysteine. Although TT

genotype was not an independent predictor of coronary artery disease, it was an important predictor of the extent of coronary atherosclerosis and plasma homocysteine, especially in the presence of plasma folate values below the median of the population. These findings may have important implications for folate replacement in patients with the TT genotype. (*Heart* 1999;81:518–522)

Keywords: coronary artery disease; vitamin B-12; folate; MTHFR mutation

Raised plasma homocysteine has emerged as a new independent risk factor for coronary artery disease.¹ Evidence from several studies involving over 3000 patients has shown that a mild to moderate rise in plasma homocysteine is strongly associated with the development of atherosclerosis.2 The major determinants of plasma homocysteine concentrations are nutritional and genetic factors.^{3 4} A common single base pair exchange of cytosine to thymidine at nucleotide 677 of methylene tetrahydrofolate reductase (MTHFR) gene leads to thermolabile MTHFR with reduced enzyme activity, resulting in raised plasma homocysteine concentrations.5 It has been reported that this mutation is associated with the development of coronary artery disease.6 However, studies in different populations have not been in agreement about the importance of this mutation.7 It is claimed that the mutation may be important in subjects with lower folate concentrations.8

The Turkish population has a high prevalence of coronary artery disease, even though Turkey is a Mediterranean country and Turks have relatively low plasma cholesterol concentrations.⁹⁻¹¹ We were therefore interested to examine the relations between plasma homocysteine, the thermolabile MTHFR mutation, and the development of coronary artery disease in this population.

Methods

SUBJECTS

The study population consisted of 242 consecutive patients undergoing coronary angiography for diagnostic purposes. The indication for angiography was either chest pain or the evaluation of the coronary arteries in patients scheduled for valve surgery. A detailed questionnaire was completed in all subjects before cardiac catheterisation, together with a physical examination, ECG, and

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Table 1 Distribution of risk factors among patients with coronary artery disease (CAD) and controls

	$\begin{array}{l} Controls \\ (n = 91) \end{array}$	CAD (n = 151)	p Value	Odds ratio	95% CI
Total cholesterol (mmol/l)	5.36 (1.22)	5.57 (1.17)	0.09	1.78	0.93 to 3.41
Triglycerides (mmol/l)	1.70 (0.79)	1.99 (0.90)	0.009	3.10	1.33 to 7.23
HDL (mmol/l)	1.06 (0.28)	1.03 (0.23)	1.0	1.04	0.45 to 2.39
LDL (mmol/l)	3.31 (0.85)	3.75 (0.75)	0.01	2.60	1.23 to 5.46
Total cholesterol/HDL	5.04	5.37	0.015	2.70	1.25 to 5.84
Smoking (%)	28	52	0.002	2.84	1.47 to 5.47
Hypertension (%)	25	43	0.015	2.34	1.19 to 4.60
Diabetes mellitus (%)	0.6	18	0.025	3.42	1.12 to 10.46
Positive family history (%)	43	31	0.1	1.71	0.82 to 3.54

CI, confidence interval; HDL, high density lipoprotein; LDL, low density lipoprotein.

chest *x* ray. The questionnaire provided information about coronary risk factors such as smoking, diabetes mellitus, hypertension, and family history, as well as the presence of angina. Previous myocardial infarction was diagnosed according to conventional criteria described by the World Health Organisation (WHO). Patients with an acute illness such as myocardial infarction within the past three months, or a chronic disease such as chronic renal failure, psoriasis, and collagen vascular disease, were not included.

Cardiac catheterisation was performed by the Judkins technique. A left ventriculogram was performed in the 30° right anterior oblique and 60° left anterior oblique positions, followed by selective coronary angiography. Blinded assessment of coronary angiograms identified coronary artery disease as the presence of > 50% stenosis in any coronary vessel. According to this definition, 151 subjects were found to have coronary artery disease (patient group), while 91 had a completely normal coronary anatomy (control group). The mean (SD) age of the patient group was 57 (11) years and of the control group, 52 (11) years. The coronary score was determined subjectively by two different observers using the Leaman scoring system.¹² In this system, the coronary arterial tree is divided into segments. Each lesion is given a score according to the location and severity of stenosis. Using this system, it is possible to score the extent of coronary atherosclerosis more accurately than it is when only the number of diseased vessels is stated.

MEASUREMENT OF LIPIDS

Blood was drawn in the fasting state on the day of coronary angiography. Plasma samples were analysed for fasting glucose, lipid profile, blood urea nitrogen, creatinine, and liver enzymes. The total cholesterol, high density lipoprotein (HDL) cholesterol, and triglycerides were determined by conventional methods.

Table 2 Distribution of genotypes among patients with coronary artery disease and controls

	Genotypes			Alleles		
	TT	CT	CC	T	С	
Coronary artery disease Controls 95% Confidence interval p Value	7.4% 5.2% 0.32 to 5.31 0.701	46.8% 43.1% 0.56 to 2.19 0.765	45.8% 51.7% 0.42 to 1.66 0.622	30.9% 26.7% 0.37 to 1 0.635	69.1% 73.3% 1.62	

MEASUREMENT OF PLASMA HOMOCYSTEINE,

VITAMIN B-12, AND FOLATE CONCENTRATIONS Total homocysteine in blood samples was determined by an immunoassay (Axis Biochemicals, Oslo, Norway). The assay method was based on enzymatic conversion of homocysteine to S-adenosyl-L-homocysteine, followed by quantification of S-adenosyl-Lhomocysteine by an enzyme linked immunoassay.¹³ Plasma vitamin B-12 and folate concentrations were both determined by microparticle enzyme immunoassay (Abbott Laboratories, Abbott Park, Illinois, USA).

GENETIC ANALYSIS

The polymerase chain reaction primers for the amplification of MTHFR have been described previously.¹⁴ The primers generate a 198 base pair (bp) fragment. The MTHFR polymorphism, a C to T substitution at bp 677, produces a Hinf I digestion site, which creates fragments of 175 bp and 23 bp. The fragments are analysed by polyacrylamide gel electrophoresis.

STATISTICS

The frequencies of the allele genotypes among the cases and controls were counted and compared by the χ^2 test with the values predicted by the Hardy–Weinberg equilibrium in the sample. Odds ratios were calculated as a measure of the association of the MTHFR genotype with the phenotype. For each odds ratio, two tailed p values and 95% confidence intervals (CI) were calculated. To determine the associations of the alleles with phenotype, multiple logistic regression analysis was performed. Leuvene's two tailed *t* test was used to compare the means of variables.

Results

CHARACTERISTICS OF THE STUDY POPULATION

Our study population consisted of 242 patients undergoing coronary angiography. Of these 151 were found to have coronary artery disease and 91 had normal coronary arteries. Of the patients with coronary artery disease, 58% had a previous myocardial infarction. None of the controls had a myocardial infarction. The distribution of several risk factors is shown in table 1 and reflects the expected differences in the prevalence of known risk factors for coronary artery disease. We found the following conventional risk factors to be significant risk factors for coronary artery disease in our study population (table 1): LDL cholesterol (p = 0.01, risk ratio 2.6, 95% CI 1.23 to 5.46),triglycerides (p = 0.009, risk ratio 3.1, 95% CI 1.33 to 7.23), total cholesterol to HDL ratio (p = 0.015, risk ratio 2.7, 95% CI 1.25 to 5.84), smoking (p = 0.002, risk ratio 2.84, 95%) CI 1.47 to 5.47), hypertension (p = 0.015, risk ratio 2.34, 95% CI 1.19 to 4.6), and diabetes mellitus (p = 0.025, risk ratio 3.42, 95% CI 1.12 to 10.4). The patients with coronary artery disease had a mean (SD) Leaman score of 13.5 (12).

 Table 3
 Distribution of plasma homocysteine levels according to genotype and plasma folate when population is divided according to median folate concentration

	CC	CT	TT	p Value
Plasma homocysteine (µmol/l)	14.5 (7.0)	16.9 (9.0)	23.3 (15.0)	0.001
Plasma homocysteine in patients with folate <12.9 nmol/l (µmol/l)	15.2 (8.0)	19.5 (10.0)	28.4 (15.0)	0.01
Plasma homocysteine in patients with folate >12.9 nmol/l (µmol/l)	14.0 (6.0)	14.5 (8.0)	12.0 (4.0)	NS

Values are mean (SD).

PLASMA HOMOCYSTEINE, VITAMIN B-12, AND FOLATE CONCENTRATIONS

The mean (SD) homocysteine concentrations were 15.6 (10) µmol/l in controls and 18.5 (11) µmol/l in patients with coronary artery disease (p > 0.05). Plasma homocysteine concentration above 15 µmol/l was found to be a significant risk factor for coronary artery disease (p = 0.03, risk ratio 2.1, 95% CI 1.07 to 4.4). On multivariate analysis when all other conventional risk factors were accounted for, plasma homocysteine concentrations remained an independent significant predictor of coronary artery disease risk (p = 0.04). The homocysteine concentrations were not different between smokers and non-smokers, at 17 (9) v16 (9) μ mol/l, p > 0.05. In addition, homocysteine concentrations above 15 µmol/l were an important risk factor for the severity of atherosclerosis as assessed by the Leaman score (p = 0.014, risk ratio 3.2, 95% CI 1.3 to 8.2).

The plasma vitamin B-12 concentrations were 238 (100) pg/ml in patients with coronary artery disease and 244 (111) pg/ml in control subjects (p > 0.05). There was no association between plasma vitamin B-12 and plasma homocysteine concentrations (p > 0.05). In addition, we found no association between the presence and extent of coronary artery disease and plasma vitamin B-12 concentrations (p > 0.05).

The folate concentrations were 11.5 (2.9) nmol/l in patients with coronary artery disease and 15.8 (7.2) nmol/l in control subjects



Figure 1 The distribution of plasma homocysteine concentrations according to genotype. *p = 0.001, CC v TT.



Figure 2 The distribution of Learnan scores according to genotype. *p = 0.03, CC v TT.

 Table 4
 Predictors of coronary artery disease in multiple logistic regression

Variable	p Value	
Smoking	0.02	
Diabetes	0.005	
Hypertension	0.002	
LDL cholesterol	0.003	
Total cholesterol/HDL ratio	0.015	
Homocysteine	0.04	
Folate	0.03	
TT genotype	0.25	

HDL, high density lipoprotein; LDL, low density lipoprotein.

(p = 0.04). There was a significant negative correlation between plasma homocysteine and folate concentrations (p = 0.03, r = -0.2). Plasma folate concentrations below the median conferred a risk of coronary artery disease (risk ratio 2.42, p = 0.03, 95% CI 1.0 to 5.5). In addition, on subgroup analysis, smokers were found to have significantly lower folate concentrations than non-smokers (p = 0.04).

When the effects of homocysteine on lipid variables were investigated, no effect was found on total cholesterol or triglyceride concentrations (p > 0.05). However, a weak negative correlation between HDL and plasma homocysteine was found (r = -0.037, p > 0.05).

FREQUENCY OF ALLELES AND GENOTYPE

The distribution of genotypes is shown in table 2. TT genotype was present in 7.4% of patients with coronary artery disease and 5.2% of controls. The prevalence of alleles did not differ significantly (TT 7%, CT 40%, CC 53%) from those predicted by the Hardy–Weinberg equilibrium ($\chi^2 = 2.3$, p = 0.3).

The highest plasma homocysteine concentrations were found in the patients with the TT genotype compared with other genotypes (fig 1). On linear multivariate analysis, when corrected for vitamin B-12 and folate concentrations, TT genotype remained a significant predictor of plasma homocysteine (p = 0.004, 95% CI 2.55 to 12.7). For further analysis, the group was divided according to the median plasma folate concentration of the population, which was 12.9 nmol/l. Those with folate concentrations below the median and the TT genotype had the highest homocysteine concentrations (p = 0.01) (table 3). Furthermore, patients with the TT genotype had the highest Leaman scores compared with other genotypes (fig 2). The TT genotype was not associated with other conventional risk factors such as smoking, lipid concentrations, or hypertension (p > 0.05). When the predictors of coronary artery disease were evaluated on multiple logistic regression analysis, in addition to conventional risk factors, plasma homocysteine and folate concentrations were independent predictors, whereas the TT genotype was not found to be an independent predictor of coronary artery disease (table 4).

Discussion

A growing body of published evidence suggests that raised concentrations of homocysteine are an independent risk factor for coronary artery disease.15 In a meta-analysis, a rise of 5 µmol/l of homocysteine was shown to be equivalent to a 0.5 µmol/l increase in total cholesterol in its effect on coronary risk.2 Our study was of interest because, despite a high prevalence of coronary morbidity and mortality in Turkey, the total cholesterol concentrations are relatively low and are not the major cause for the development of atherosclerosis. A recent epidemiological survey has shown that the prevalence of coronary artery disease is 4% in our adult population, with total cholesterol concentrations of 4.7 and 5.1 mmol/l for men and women, respectively. Furthermore, the prevalence of congenital neural tube defects associated with folate deficiency is significantly higher than in other countries.^{16 17} Several recent studies including the large study by the European COMAC group have shown that lower folate concentrations are associated with atherosclerotic vascular disease and increased coronary risk.18 19 The WHO has proposed a lower limit of 13.6 nmol/l (6 ng/ml) for folate concentrations.²⁰ It has been shown that with values below 12.5 nmol/l (5.5 ng/ml) homocysteine concentrations are actually increased.²¹ Most of our subjects in both the patient and the control groups had low folate concentrations according to these criteria. In the light of these findings, we thought that raised homocysteine concentrations might be an important risk factor for coronary artery disease in Turkey.

We found that plasma homocysteine concentrations above 15 µmol/l were a significant and independent risk factor for coronary artery disease, with a risk ratio of 2.1. This is in accordance with previous studies in other populations.²² Moreover, increased concentrations of homocysteine also predicted the extent of atherosclerosis as judged by the Leaman score. Several in vitro studies have shown that raised plasma homocysteine promotes both atherosclerosis and thrombosis by causing endothelial injury, smooth muscle cell prolifdensity lipoprotein eration, and low oxidation.23-25 The effects of high plasma homocysteine concentrations on the risk of coronary artery disease have not been shown to be mediated by the known risk factors.²⁶ Our findings support this theory. There was no association between increased homocysteine concentrations and the other risk factors examined in our study. One weak inverse association with HDL was statistically nonsignificant. The two predictors of plasma homocysteine concentrations were found to be the TT genotype and plasma folate.

The importance of the TT genotype has been questioned in previous studies conducted in well nourished populations.²⁷ Our subjects had significantly lower folate concentrations than similar groups studied in the United States and Australia. A recent study in a US population showed that the TT genotype increased homocysteine concentrations only in patients with folate concentrations below the median.²⁸ Our study is one of the very few to look at genotype and plasma homocysteine, vitamin B-12, and folate concentrations simultaneously in the same population. We chose to evaluate subjects assessed by coronary angiography to be sure that the control group did not have substantial but clinically silent coronary atherosclerosis. In our population, the TT genotype significantly predicted the extent of atherosclerosis in patients with coronary artery disease. However, this genotype was not an independent predictor of the presence of coronary artery disease, but an important predictor of plasma homocysteine concentrations. In this group of Turkish patients, the highest plasma homocysteine concentrations were present in patients where the TT genotype and folate deficiency coexisted.

This may have important clinical implications. Dietary and environmental factors may modify hyperhomocysteinaemia arising from genetic causes, offering the potential for therapeutic intervention. Recent studies have shown that an increased intake of vitamin B-6, vitamin B-12, and folate can reduce plasma homocysteine concentrations, even in patients with the TT genotype. With 1000 µg folate supplementation, a 21% reduction in homocysteine level has been observed.²⁹ However, it remains to be determined whether treatment of hyperhomocysteinaemia will lead to a reduction in vascular disease. Although no appropriate studies have been undertaken so far, a decrease in homocysteine concentrations with vitamin supplementation, especially in people with the TT genotype, may prevent atherosclerosis in these individuals. Randomised trials are needed to determine the effect of vitamins in decreasing vascular disease risk.

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