MGP T-138C Polymorphism (TT Genotype) is Associated with Vascular Calcification Incidence in Indonesian Regular Hemodialysis Patients

Muhammad Hanif Wibowo¹, Riri Andri Muzasti¹,², Syafrizal Nasution¹,²

¹Department of Internal Medicine, Faculty of Medicine, Universitas Sumatera Utara/ Haji Adam Malik General Hospital, Jl. Dr. Mansyur No.5, Medan, Indonesia
²Division of Nephrology and Hypertension, Department of Internal Medicine, Faculty of Medicine, Universitas Sumatera Utara/ Haji Adam Malik General Hospital, Jl. Dr. Mansyur No.5, Medan, Indonesia

*Corresponding author. E-mail: riri.andri@usu.ac.id

Received date: Feb 5, 2020; Revised date: Aug 30, 2020; Accepted date: Sep 1, 2020

Abstract

BACKGROUND: Vascular calcification contributes greatly to the incidence of cardiovascular disease. Previously, vascular calcification was considered as a passive process caused by the mineral deposition from the circulation. Nowadays, researchers have found inhibitors and promoter factors from vascular calcification, one of which is the matrix gla protein (MGP). MGP levels depend on the gene that encodes them. The MGP T-138C polymorphism is one of the most common causes of vascular calcification.

METHODS: This was a case-control study involving 86 chronic kidney disease (CKD) patients who underwent regular hemodialysis in Rasyida Kidney Hospital, Medan, Indonesia. Vascular calcification was determined from lateral and posteroanterior abdominal X-ray. The MGP T-138 C polymorphism was analyzed using polymerase chain reaction restriction fragment length polymorphism (PCR-RFLP).

RESULTS: The TT/TC genotype associates with the incidence of vascular calcification with OR of 3.52, 95% CI: 1.23-10.106 (p-value<0.001).

CONCLUSION: There is an association between MGP T-138C polymorphism, particularly on T Allele with the incidence of vascular calcification in CKD patients undergoing regular hemodialysis.

KEYWORDS: hemodialysis, T-138 C polymorphism, vascular calcification, matrix GLA protein


Introduction

Vascular calcification contributes significantly to the incidence and the average mortality of the cardiovascular disease.(1) Based on the Indonesian Renal Registry in 2017, the highest mortality rate in hemodialysis (HD) patients is due to cardiovascular disease.(2) Vascular calcification was previously regarded as a passive process caused by circulatory deposition of minerals, mainly in patients with mineral imbalances. Currently, there are both inhibitory and promoting factors found from vascular calcification, one of which is the Matrix Gla Protein (MGP).(3-5) This protein is a vitamin K-dependent protein that gives a direct effect to blood vessels by inhibiting the formation of calcium crystal together with other calcification inhibitors such as fetuin-A. MGP also binds to bone morphogenetic protein-2 (BMP-2), a growth factor that converts vascular smooth muscle cells into osteoblast-like cells, and also binds with extracellular matrix components and act as an anti-apoptosis.(5-7) MGP is the first recognized protein to act as an in vivo calcification inhibitor. MGP is produced by bone and smooth muscle.(8) Several studies have identified polymorphisms in the MGP gene, and they are associated with various phenotypes.(9,10) Recent research has focused on the T-138 C polymorphism found in the promoter region.

The concentration of MGP levels depends on the genes that encode them.(8) The MGP T-138C polymorphism (rs
1800802) lies in the promoter region.(9) Based on previous studies it was found that the TT genotypes have 30% lower MGP serum levels in contrast to the CC genotypes, thus the TT genotypes accelerated the vascular calcification more prominently.(9) Conversely, another study concluded that the CC genotype of the MGP T-138C polymorphism was associated with slower progression of vascular calcification found in patients undergoing HD regularly.(10) This study set out to investigate the association of the MGP T-138C polymorphism with the incidence of vascular calcification in regular HD patients in Indonesia.

### Methods

This case-control study received ethical approval from the Health Research Ethical Committee, Faculty of Medicine, Universitas Sumatera Utara (No. 313/TGL/KEPK FK USU-RSUP HAM/2019). Patients who were undergoing regular HD at Haji Adam Malik Hospital, Medan, Indonesia between May to July 2019, were enrolled in this study as subjects. The inclusion criteria were chronic kidney disease (CKD) subjects who had undergone HD for more than 26 months and were 22-79 years old. Subjects with incomplete data were excluded from the study.

Eighty-six subjects were selected by the consecutive sampling method. The subjects were then divided into two groups; subjects with vascular calcification and without vascular calcification based on the results of the lateral and posteroanterior abdominal X-ray. Data of age, sex, duration of HD were obtained from medical records.

**MGP T-138C Polymorphism Test**

Blood samples from both groups were collected to obtain the genomic deoxyribonucleic acid (DNA) structure. The DNA extraction was examined by polymerase chain reaction (PCR) technique. The PCR restriction fragment length polymorphism (PCR-RFLP) was used for the analysis of the MGP T-138C Polymorphism in the Integrated Laboratory in the Faculty of Medicine, Universitas Sumatera Utara. The PCR technique was done using Veriti 96 Well Thermal Cylinder (Thermo Fisher Scientific, Waltham, MA, USA). The genotyping process was conducted based on previous study method with a slight modification.(9) Genomic DNA was amplified in the following primary sequence: 5′AAG CAT ACG ATG GCC AAA ACT TCT GCA 3′ (forward) and 5′GAA CTA GCA TTG GAA CTT TTC CCA ACC 3′ (reverse). The parameters of the amplification were as follows: firstly, denaturation was done for 5 minutes at 94°C. Thirty-six cycles for 1 minute at 94°C, 1 minute at 59°C, 1 minute at 72°C and 10 minutes for the final extension at 72°C. The PCR product (142 base pair-bp) was digested using the 5 U restriction enzyme of BsrI (New England Biolabs, Ipswich, MA, USA) at 65°C for 30 minutes and followed by thermal inactivation at 80°C for 20 minutes. The retention pattern for -138C alleles contains 142 bp fragments. Polymorphism of -138T was made into a frictional position and was digested into 118-24 bp fragment. The restriction product was then divided into 2.5% agarose gel and visualized by etching ethidium bromide.

### Data Analysis

Univariate data analysis was conducted to determine the frequency distribution. Categorical variables were presented as frequencies (n) and percentages (%), while numerical variables were presented as mean and standard deviation for normally distributed data and as categorical variables for not normally distributed data. A \( p \)-value<0.05 was considered statistically significant. Data analysis was done by using SPSS Ver. 24 (IBM Corporation, Armonk, NY, USA).

### Results

Eighty-six subjects who met the inclusion criteria were included, 42 subjects were male (48.8%), and 44 subjects of them were female (51.2%). The median age was 59 years (32-79 years). The majority of all 86 subjects were Bataknese (65.1%), followed by 17 Malays (19.8%), 3 Chinese (3.5%), and 10 Javanese (11.6%). The median duration of HD was 45 months (27-229 months). The majority of subjects had hypertension as the comorbid disease in 60 people (69.8%), followed by diabetes mellitus in 18 people (20.9%).

From the results of RFLP examination on the MGP T-138C polymorphism, the majority of subjects had TT genotype found in 40 subjects (46.5%), followed by TC genotype in 28 subjects (32.6%) and CC subjects in 18 people (20.9%) (Table 1). As gender matching was arranged, the same number of female and male was obtained, which was 21 male patients (50.0%) in the case group and 22 female patients (50.0%) in the control group (Table 2). Mean age was 54.40±1.5 years and 58.67±1.661 years in the case and the control group, respectively. In the case group, the average HD duration was 45 months and 46 months in the control group.

The first set of analysis was to determined whether there was any significant differences between the characteristics
Table 1. Subject’s characteristics.

<table>
<thead>
<tr>
<th>Characteristics</th>
<th>n = 86 (100%)</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Sex, n (%)</strong></td>
<td></td>
</tr>
<tr>
<td>Male</td>
<td>42 (48.8%)</td>
</tr>
<tr>
<td>Female</td>
<td>44 (51.2%)</td>
</tr>
<tr>
<td><strong>Age (years), Median (Min-Max)</strong></td>
<td>59 (32-79)</td>
</tr>
<tr>
<td><strong>Ethnic Group, n (%)</strong></td>
<td></td>
</tr>
<tr>
<td>Batakense</td>
<td>56 (65.1%)</td>
</tr>
<tr>
<td>Malay</td>
<td>17 (19.8%)</td>
</tr>
<tr>
<td>Tionghoa</td>
<td>3 (3.5%)</td>
</tr>
<tr>
<td>Javanese</td>
<td>10 (11.6%)</td>
</tr>
<tr>
<td><strong>HD Duration (months), Median (Min-Max)</strong></td>
<td>45 (27-229)</td>
</tr>
<tr>
<td><strong>Hypertension (%)</strong></td>
<td></td>
</tr>
<tr>
<td>Yes</td>
<td>60 (69.8%)</td>
</tr>
<tr>
<td>No</td>
<td>26 (30.2%)</td>
</tr>
<tr>
<td><strong>Diabetes Mellitus, n (%)</strong></td>
<td></td>
</tr>
<tr>
<td>Yes</td>
<td>18 (20.9%)</td>
</tr>
<tr>
<td>No</td>
<td>68 (79.1%)</td>
</tr>
<tr>
<td><strong>T-138 C Polymorphism, n (%)</strong></td>
<td></td>
</tr>
<tr>
<td>TT</td>
<td>40 (46.5%)</td>
</tr>
<tr>
<td>TC</td>
<td>28 (32.6%)</td>
</tr>
<tr>
<td>CC</td>
<td>18 (20.9%)</td>
</tr>
</tbody>
</table>

of the study population of those with vascular calcification in the case group and those without in the control group. Based on the characteristics of the study population shown in Table 2, there was no evidence that each of the characteristic variables had any significant differences. The characteristics included age, HD duration, ethnic group and comorbid disease such as hypertension and diabetes mellitus. None of these variables was statistically proven with \( p \)-value\( \geq 0.05 \).

The next section of the survey was concerned with the association between the MGP T-138 C polymorphism of TT, CC and CC genotypes with the incidence of vascular calcification. Stepwise multiple regression analysis was performed to investigate the significance of possible predictors of vascular calcification as assessed by correlation analysis. The results, as shown in Table 3, indicated that there was an insignificant association between the TT, TC and CC genotypes with the incidence of vascular calcification with \( p \)-value\( =0.06 \). Another set of groups tested for association was by grouping the subjects into two different set of genotypes which was TT genotype versus TC/CC genotype and TT/TC genotype versus CC genotype. No significant association found in first grouping with \( p \)-value\( =0.195 \) while the next group showed a significant association between the TT/TC genotypes versus CC genotype with the incidence of vascular calcification with \( p \)-value\( =0.001 \) (PR: 7.14 with 95% CI: 1.88-27.01) (Table 3).

### Discussion

The majority of the subjects consisted of female (51.2%) with a median age of 59 years. The age characteristics of...
Conclusion

It can be concluded that there was a significant association between MGP T-138 C gene polymorphism, particularly on T Allele with the incidence of vascular calcification in CKD patients undergoing regular HD.
MGP T-138C Polymorphism is Associated with Vascular Calcification Incidence (Wibowo MH, et al.)
DOI: 10.18585/inabj.v12i4.1162

References


