To the Editor

Venous thrombosis can be the result of various acquired and genetic factors. Several alterations in the genetics of hemostatic factors that contribute to thrombosis have been described [1]. Inherited anticoagulant protein deficiency-related hypercoagulability is responsible for 5%-10% of all cases of venous thromboembolism (VTE) [2]. An important recognized inherited cause of VTE is resistance to activated protein C (APC), which is mostly due to a single point mutation in the blood coagulation factor V gene (G1691A) [3].

Furthermore, a genetic factor that increases the risk of VTE G20210A variation in the prothrombin gene associated with elevated plasma prothrombin level has been described [4].

These 2 genetic prothrombotic factors have heterogeneous geographic distribution patterns. Factor V Leiden and prothrombin G20210A mutation allele frequencies in Europeans are been reported to be 1.4%-7.0% and 3.0%, respectively [5,6]. There may be important differences in the allelic frequency of these mutations in different European populations, and they may exhibit regional variations in Europe [7-10]. The aim of the present study was to determine the frequency of factor V Leiden and prothrombin G20210A mutations in an Albanian population. Such data may help elucidate the ethnic and/or geographic differences in the incidence of thrombotic diseases.

We assessed the genetic frequency of the 2 thrombotic risk polymorphisms in a group of 225 randomly selected healthy Albanians, which included a mixture of all cultural groups in Albania, according to a previously reported method [11]. Blood samples were collected at a referral center that serviced people from different regions of Albania. The control group was in Hardy-Weinberg equilibrium. The participants were not chosen from any specific population. The additional members of the same family and those with a family history of any kind of thrombosis were excluded. Written informed consent was obtained from every participant.

Factor V Leiden and prothrombin G20210A prothrombotic alleles are inherited prothrombotic mutations with population-dependent frequencies. Differences in geographic distribution primarily account for regional variation in the incidence of thromboembolism [11]. The high incidence rate in
Europeans, as a risk factor for thromboembolic diseases, suggests that screening populations for this mutation is of great importance. The distribution of prothrombin G20210A, the second most frequent prothrombotic polymorphism in humans, has a north-south gradient in Europe. A carrier frequency of 3.0% has been reported in southern Europe, nearly 2-fold greater than that in northern Europe [6]. It has a very high prevalence in Mediterranean countries. In addition, some microheterogeneity is observed within populations [12].

Factor V Leiden and prothrombin G20210A mutations have not been previously reported in Albanians. To the best of our knowledge this is the first study to examine the frequency of 2 thrombosis-related polymorphisms in healthy Albanians. Factor V Leiden and prothrombin G20210A mutation was observed in 1.8% and 3.1% of the participants, respectively (Table 1).

The present study’s results show that prothrombin gene 20210GA variation in the Albanian study population did not differ from that reported in other European studies. The 1691GA mutation is not frequent in the Albanian population and its prevalence is similar to that reported from Italy and Spain, and lower than in Greece [13-14].

The present study determined the frequency of 2 thrombophilic polymorphisms in an Albanian population. Determination of the heterogeneity of the prevalence of these 2 common risk factors can help elucidate the geographic differences in the incidence of thromboembolic diseases. Additional research on thrombotic Albanian patients is warranted, as the results may aid in the improvement of evidence-based management.

**Conflict of interest statement**

The authors of this paper have no conflicts of interest, including specific financial interests, relationships, and/or affiliations relevant to the subject matter or materials included.

**References**

1. Lane DA, Grant GJ. Role of hemostatic gene polymorphisms in venous and arterial thrombotic disease, Blood 2000;95:1517-32.

**Table 1. The prevalence of Factor V Leiden and prothrombin G20210A mutations in an Albanian population**

<table>
<thead>
<tr>
<th></th>
<th>Total</th>
<th>n</th>
<th>%</th>
<th>Frequency</th>
</tr>
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<tbody>
<tr>
<td>FV 1691A</td>
<td>225</td>
<td>4</td>
<td>1.8</td>
<td>4/450 (0.009)</td>
</tr>
<tr>
<td>PT 20210A</td>
<td>225</td>
<td>7</td>
<td>3.1</td>
<td>7/450 (0.016)</td>
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