EVALUATION OF PROTHROMBIN GENE POLYMORPHISM FREQUENCY IN PATIENTS WITH ISCHEMIC STROKE IN IRAQI POPULATION

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Abstract

Stroke is one of the leading causes of death worldwide, and increasing evidence indicates that coagulation abnormalities cause stroke and the formation of a secondary ischemic stroke, either in the accompanying vessel or in the form of an embolus. Therefore, the balance between coagulation and fibrinolysis is a critical component of the ischemic stroke. The study population was included 75 ischemic stroke patients and 25 healthy subjects. The estimated incidence of ischemic stroke increased in the age group 65-74 years, with a significant difference (p<0.05) in comparison with the other groups. Results of coagulation tests indicated that no significant differences were found in the results of prothrombin time between the IS patients and control. A PCR-SSP technique was performed for detection prothrombin gene 20210 G/A (rs1799963) polymorphism, in ischemic stroke patients. The genotype distribution results of the 20210 G/A SNP of prothrombin gene were not showed significant differences (p>0.05) between controls (GG : n = 25, 100%; GA: n = 0; AA: n = 0) and ischemic stroke patients (GG : n = 66, 88%; GA: n = 8, 10.66%; AA : n = 1, 1.333%). The results of prothrombin time did not differ according to the prothrombin gene genotype and it statistically not significant (p>0.05) in ischemic stroke patients that carrier A allele or G allele than controls. Results showed that G allele of the 20210G/A and the mutant allele (A) was not statistically significantly linked with the ischemic stroke (OR = 0.251, 95% CI = 0.032-1.996, p = 0.16). However, the 20210 G/A mutant alleles are not entirely absent among Iraqi patients with ischemic stroke.

Key words: Stroke, Prothrombin gene, Iraq, 20210 G/A polymorphism

Introduction

Ischemic stroke (IS) is a neurological deficit attributed to an acute injury of the central nervous system by a vascular cause, it occurs as a result of obstruction by a blood clot (thrombus) or plugs within a blood vessel supplying blood to the brain- carotid or vertebral artery or less likely a cerebral vein (Al-Gazally et al., 2014; Hameed and Ahmed, 2018). This disease considered as the most common type of stroke in older adults with high mortality and morbidity rates (Al-Qaysi et al., 2015; Xu et al., 2018). Genetic factors that are associated with increased stroke risk are a family history of stroke and a number of single nucleotides polymorphisms SNPs (Baird, 2010). Hypercoagulability disorder is a hereditary or acquired condition that increases the risk of thrombosis (Linnemann & Hart, 2019). Hypercoagulability may have an important role in the pathogenesis of ischemic stroke (Carcaillon et al., 2011; Bona, 2016). Factor V Leiden (G1691A) and the prothrombin gene mutation (20210 G/ A; rs1799963) comprise the most common genetic associations with thrombosis and thus comprise the most commonly requested genetic thrombophilia investigations (Favaloro, 2019). Prothrombin (coagulation Factor II) is a complex glycoprotein that plays a central role in blood coagulation. It is the zymogen precursor to the protease thrombin that catalyzes the formation of the fibrin clot and regulates a multitude of other cellular responses related to coagulation and hemostasis (Wendeler et al., 2014). Prothrombin is activated to thrombin by the prothrombinase complex through sequential cleavage at two distinct sites (Adams & Huntington, 2016). Prothrombin is encoded by the FII gene, which is located on the short arm of chromosome 11 (Degen & Davie, 1987). Previous data have shown that the 20210G > A polymorphism of the Factor II gene is related to an increased prothrombin level, which may in turn lead to a procoagulant state (Sarecka-Hujar et al., 2017). This

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study aimed to assess the diagnostic prevalence and prognostic significance of prothrombin polymorphism: G20210A in ischemic stroke patients, and determine whether prothrombin time is associated with genotypes.

Materials and Methods

Study group consists of 75 patients with confirmed ischemic stroke. They were admitted to Alsader Specialized Center of the Middle Euphrates Neurosciences and 25 healthy subjects as controls. Tests were performed on 2ml of venous blood, which was collected from ischemic stroke patients and control group. 1ml was collected in sodium citrate tubes centrifuged at 5000 rpm for 5 minutes; plasma was used freshly for the Prothrombin time (PT). 1 ml was collected in tubes with anticoagulant EDTA and used for PCR test.

PT determination was done by kit, which is BIOLABO REAGENT products (02160, Maizy, France/ REF: 13560). The clotting time was measured at 37°C in the presence of tissular thromboplastin and calcium, the PT (in sec) so measured and converted into PT(%) or INR.

Genomic DNA was isolated by using protocol of DNA Mini Kit, which designed for purifying DNA from Geneaid Biotech. Ltd., Taiwan Company (Cat. No. GS100, LOT. No. FE31708-B). Amplification the ofSNPs fragments in prothrombin gene was done by using a conventional PCR thermocycler (Labnet/USA). The amplification steps consisted of a first denaturation step where DNA was initially denatured for 10 minutes at 95°C and then 10 cycles were performed as follows: 94°C for 30 seconds, 60°C for 30 seconds and 72°C for 1 min. Then, 25 cycles were performed as follows: 94°C for 30 seconds, 55°C for 30 seconds and 72°C for 1 minute. The PCR amplification was completed by a final extension at 72°C for 7 minutes. Amplification of the prothrombin genes resulted in 340-bp products. DNA was amplified with primers specific for prothrombin 20210 G/A polymorphism for each sample. The primer-pairs which listed in table 1, were synthesized by Accu Oligo® Bioneer Corporation, USA. The primers sequences were published by Ranguelov et al., (2002) and Othman et al., (2010). PCR sequence-specific primers (PCR-SSP) technique was performed for detection and genotyping prothrombin single-nucleotide polymorphism in blood samples. The basis of this method is that a perfectly matched primer is more efficient in a PCR reaction than mismatched primers. So, specificity is determined by the use of sequence specific primers (SSP) in two reactions (in two tubes) with two sets of primers: one primer is specific for each allele (allele specific primer) and is paired with a second common primer. There is a reduction in the efficiency of Taq polymerase to amplify DNA when there is a 3’ single-base (3’ terminal nucleotide) mismatch inhibits the initiation of a non-specific reaction between the target DNA and the allele-specific primer so even primer pairs do not specifically anneal and are not efficiently amplified (Sadler et al., 1994; Ayyadevara et al., 2000). Therefore, only the required allele will be amplified. Then the amplified product in the two tubes can be detected by agarose gel electrophoresis for wild type (WT) and the mutant type (MT) separately. This method is a relatively simple and inexpensive procedure for prothrombin genotyping. The gel electrophoresis method, which included preparing the gel loading and running the gel, was done according to Sambrook and Russell (2001).

Statistical analysis

Statistical analyses of all results were carried out by the help of Statistical Package for the Social Sciences (SPSS) version 23 software statistical package using t-test and Chi-square test (with P value at level of significance less than 0.05) to compare values of results between groups. Result values were expressed as mean ± SE, number of patients, or percentages.

Results and Discussion

The clinical assessment revealed that the frequency of distribution of patients according to gender were 39 (52%) females and 36 (48%) males. The differences of gender were statistically not significant.

These results agree with Sohrabji et al., (2017) and Chauhan et al., (2017) studies that showed Women are affected by stroke, having a higher incidence and worse outcomes than men, also more women than men have strokes each year, in part because women live longer.

These findings are consistent with those of Dehlendorff et al., (2015) study that showed after the age of 60 years, women had more severe strokes than men.

These results are disagreement with Zafar et al.,

<table>
<thead>
<tr>
<th>Gene</th>
<th>Primers sequences</th>
<th>PCR product</th>
</tr>
</thead>
<tbody>
<tr>
<td>Prothrombin (PT) (20210 G/A)</td>
<td>PT common</td>
<td>TGTGGATGCTC-32</td>
</tr>
<tr>
<td></td>
<td>PT Wild</td>
<td>52-GACCTTGGAGC</td>
</tr>
<tr>
<td></td>
<td>(WT)</td>
<td>ATGGAGATATC-32</td>
</tr>
<tr>
<td></td>
<td>PT mutant</td>
<td>52-GACCTTGGAGC</td>
</tr>
<tr>
<td></td>
<td>(MT)</td>
<td>ATGGAGATAT-32</td>
</tr>
</tbody>
</table>
Evaluation of prothrombin gene polymorphism frequency in patients with ischemic stroke in Iraqi population

(2016) study that showed IS to be significantly more common in males than females (p = 0.021). On the other hand, these results were in disagreement with Palm et al., (2012) study which reported that etiology of ischemic stroke differs between sexes, with large-artery atherosclerotic stroke and associated diseases (coronary artery disease and peripheral artery disease) being more common in men.

These results show that women diagnosed with ischemic stroke are more than men, largely due to increase in stroke incidence in older postmenopausal women, also lower stroke incidence observed in pre-menopausal women and robust preclinical evidence of neuroprotective and anti-inflammatory properties of estrogen (Koellhoffer & McCullough, 2013; Spychala et al., 2017).

The major risk factor of ischemic stroke in both genders was Hypertension, because its effect on alterations in cerebral artery structure and function that can impair blood flow (Pires et al., 2013), however, males were more likely to have a history of smoking, heart disease and dyslipidemia (Memis et al., 2016), (Wu et al., 2014), while in women more likely to have a past medical history of atrial fibrillation or hypertension and less likely to have a history of heart disease, dyslipidemia, or smoking (Reeves et al., 2009).

These results are disagreement with study done by Mohamed and Alshekhani (2016) that found Stroke incidence more common in male (59%) than female (41%).

These results are disagreement with Zafar et al., (2016) study showed IS to be significantly more common in males than females (p = 0.021). Women have more stroke events than men because of their longer life expectancy and much higher incidence at older ages (Reeves et al., 2008).

Assessment of age presentation of patients at diagnosis revealed that 7(9.3333%) in age group (35-44), 8 (10.666%) in age group (45-54), 9 (12%) in age group (55-64), 29 (38.666%) in age group (65-74) and 22(29.3333%) in age group (75-84) Fig. 1. Their ages ranged from 28 to 84 years, with a mean age of 66.933333 years.

The estimated incidence of ischemic stroke increased in the fourth age group (65-74), with a significant difference (p < 0.05) in comparison with the other groups.

These results are agreed with Ahangar et al., (2018) study that showed about 84.3% of stroke occurred in patients aged > 50 years.

These findings were comparable to the findings of Jung et al., (2012) study where the 46.098 patients included in their study with mean ± SD age was 66.1 ± 12.3 years and the increases in ischemic stroke disease prevalence were largest for persons aged 65 years or older.

These results showed that patients aged 65 years or older are more at risk than younger for ischemic stroke, maybe aging and growth of the population that led to an increase in the absolute number of strokes, especially in the elderly, that is expected to increase dramatically in the coming years (Béjot et al., 2019).

These results are agreed with Abd et al., (2017) study documented in Iraq, which showed the rate of stroke in Iraqi patients significantly associated with advance age. Soriano-Tárraga et al., A study (2016) showed that stroke is related to aging and corresponds to current findings.

These findings are consistent with the study of Abdul-Razak and Shati (2008) that showed that age is significantly related to the incidence of stroke, and advancing age (age more than 60 years) showed increased incidence of stroke. Advanced age is the most important predictor of poor outcome (Aksoy et al., 2013).

Several aging-related changes in the brain have been identified that are associated with an increase in vulnerability to ischemic stroke in the elderly (Chen et al., 2010).

Results of coagulation tests represented by prothrombin time and INR for IS patients were showed in table 2. No significant differences were found in the results of PT and INR between the IS patients and control.

The present study showed there is no significant difference in prothrombin time level between the IS patients and control group.

This finding was comparable to previous findings of Kravchenko and his colleagues (2018) who reported that

**Fig. 1:** Age distribution of patients presented with ischemic stroke (Group 1: 35-44 years, Group 2: 45-54 years, Group 3: 55-64 years, Group 4: 65-74 years, Group 5: 75-84 years).
Table 2: Coagulation tests (prothrombin time and INR) of ischemic stroke patients and control group.

<table>
<thead>
<tr>
<th>Parameter</th>
<th>Ischemic stroke patients (n=75)</th>
<th>Control (n=25)</th>
<th>P-Values</th>
</tr>
</thead>
<tbody>
<tr>
<td>Prothrombin time (sec)</td>
<td>15.61 ±2.913</td>
<td>15.6153846 ±1.514</td>
<td>0.9</td>
</tr>
<tr>
<td>INR</td>
<td>1.15 ±0.312</td>
<td>1.156153846 ±0.118</td>
<td>0.9</td>
</tr>
</tbody>
</table>

Results values were expressed as mean± SE, *: p< 0.05 or significant differences between mean values. Abbreviations: PTT= partial thromboplastin time, INR= international normalized ratio.

Table 3: The results of genotypic frequencies of 20210 G/A at prothrombin gene in patients and controls.

<table>
<thead>
<tr>
<th>Genotypes</th>
<th>Healthy controls (N = 25)</th>
<th>Ischemic stroke patients (N = 75)</th>
</tr>
</thead>
<tbody>
<tr>
<td>GG</td>
<td>25 (100%)</td>
<td>66 (88%)</td>
</tr>
<tr>
<td>GA</td>
<td>0</td>
<td>8 (10.666%)</td>
</tr>
<tr>
<td>AA</td>
<td>0</td>
<td>1 (1.333%)</td>
</tr>
<tr>
<td>P-value</td>
<td>0.192*</td>
<td></td>
</tr>
<tr>
<td>Alleles frequency</td>
<td>N(%)</td>
<td>N(%)</td>
</tr>
<tr>
<td>G allele</td>
<td>50 (100%)</td>
<td>140 (92.666%)</td>
</tr>
<tr>
<td>A allele</td>
<td>0</td>
<td>10 (7.333%)</td>
</tr>
<tr>
<td>X²</td>
<td>1.976</td>
<td></td>
</tr>
<tr>
<td>P-value</td>
<td>0.16*</td>
<td></td>
</tr>
<tr>
<td>OR (95%CI)</td>
<td>0.251(0.032-1.996)</td>
<td></td>
</tr>
</tbody>
</table>

Data were expressed as number and a percentage (N%). *P >0.05 not significant. Abbreviations: X²= chi-square, OR = odds ratio, CI = confidence interval.

2. That means the frequencies of 20210 G/A of prothrombin gene in the 75 Iraqi ischemic stroke patients in Al-Najaf province were not with significant differences with that of the 25 healthy controls group (p>0.05).

This result showed that G allele of the 20210G/A and the mutant allele (A) was not statistically significantly linked with the ischemic stroke (OR = 0.251, 95% CI = 0.032-1.996, p = 0.16).

Fig. 2: The electrophoresis image of PCR-SSP analysis of prothrombin geneSNP(20210G/ A) (Lane 1: 100 bp DNA Ladder; Lane 2&3 (sample 1): Heterozygous genotype (GA); 340-bp in each tube); Lane 4&5(sample 2): homozygous mutant (AA) genotype (tube 1 for wild type there was no band, tube 2 for mutant type 340-bp ); Lane 6&7(sample 3): homozygous wild (GG) genotype (tube 1 for wild type 340-bp, tube 2 for mutant type there was no band); Lane 8&9(sample 4): Heterozygous genotype (GA); and Lane 10&11(sample 5): Heterozygous genotype (GA).
This conclusion indicates the 20210 G/A SNP did not show associations with ischemic stroke risk.

Ridker et al., (1999) study found that the G20210A prothrombin gene variant was not associated with increased risk of stroke. This study is consistent with the current study.

The study done by Chang and Zhang (2000) found that Factor II gene G20210A mutant allele is absent in Chinese patients with ischemic stroke and normal subjects and this mutation may not be a major risk factor for thrombogenesis in Chinese people. This study is compatible with the present study.

Lopaciuk et al., (2001) study indicated that prothrombin G20210A genotype, was not associated with an increased risk for ischemic stroke in young adults. This study is consistent with the current study.

These results are consistent with Weischer et al., (2010) study which showed that there was no association between G20210A prothrombin heterogeneity and IS risk.

These findings are in accordance with results reported by Bondarenko et al., (2011) who showed that there were no association between prothrombin gene polymorphism and the risk of ischemic stroke development.

The study done by Içli et al., (2013) that showed there was no statistically significant difference in prothrombin G20210A variant among stroke patients and control groups. This study corresponds to the current study.

In the study conducted by Jiang et al., (2014) observed that the association of the prothrombin G20210A mutation with ischemic stroke did not achieve statistical significance (OR = 2.5, 95% CI = 0.9-6.5, p = 0.07). This study is consistent with the current study.

These findings consistent with Pirhoushiaran et al., (2014) study conducted in Iran which observed that FII G20210A SNP was not significant differences between IS patient and control groups.

The study of Erten et al., (2015) observed that mutation of prothrombin gene G20210A is not associated with stroke and is not important in the case of ischemic stroke. This study agrees with the current study.

Recently the study by Ayas et al., (2019) showed that no significant difference was found in Prothrombin 20210 G/A polymorphism with ischemic stroke risk. This result close to the present study.

The present study inconsistent with the study of De Stefano et al., (1998) which found that prothrombin mutation G20210A was significantly higher in stroke patients than in the control group.

Lalouschek et al., (2005) that found the frequency of the FII G20210A mutation was significantly higher in patients compared with controls (6% versus 1%; adjusted OR, 6.1; 95% CI, 1.3 to 28.3; P = 0.021). These results disagreement with the current study.

A study by Eterovic and his colleagues (2007) showed that G20210A mutations in prothrombin gene were significantly higher in IS patients than controls. This study is incompatible with the current study.

The results of prothrombin time did not differ according to the prothrombin gene genotype and it statistically not significant (p > 0.05) in ischemic stroke patients that carrier A allele or G allele than controls table 4.

Bauer, (1993) reported that the elevated levels of plasma prothrombin could lead to a higher occurrence of thrombin formation and subsequently more incidence of fibrin clotting.

The G20210A point mutation in factor II or prothrombin is associated with increased plasma levels of prothrombin and increased risk of venous thrombosis in its heterozygous form (Poort et al., 1996; Van Cott and Laposata, 1998).

Soria et al., study (2000) demonstrated that the polymorphism in prothrombin gene G20210A associated with increased prothrombin levels as well as increased susceptibility to thrombosis.

In a study conducted by Jadaon (2011) mentioned that the mutation of Prothrombin G20210A causes higher levels of the clotting factor prothrombin in the blood and creating a higher tendency towards blood clotting (hypercoagulability).

The mutation in the prothrombin gene may enhance the thrombogenic potential (Dobrynina et al., 2012).

**Table 4:** The results of prothrombin time (sec.) according to genotype of gene polymorphism (20210 G/ A) in ischemic stroke patients and controls.

<table>
<thead>
<tr>
<th>Genotype</th>
<th>Prothrombin time (sec)</th>
<th>P values</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Healthy controls(N=25)</td>
<td>Ischemic stroke patients(N=75)</td>
</tr>
<tr>
<td>GG</td>
<td>15.615 ± 1.514</td>
<td>15.522 ± 2.993</td>
</tr>
<tr>
<td>GA</td>
<td>0</td>
<td>16.38 ± 2.4</td>
</tr>
<tr>
<td>AA</td>
<td>0</td>
<td>15.3</td>
</tr>
</tbody>
</table>

Results values were expressed as mean± SE, *: p> 0.05 or no significant differences between mean values.
Ghatak et al., (2013) study demonstrated that this mutation in FII 20210 G/A is associated with an increased plasma concentration of prothrombin, which leads to an increased potential for thrombin generation.

Sarecka-Hujar et al., study (2017) showed that the 20210G > A polymorphism of the prothrombin gene is related to an increased prothrombin level, which may in turn lead to a procoagulant state in ischemic stroke patients.

Conclusions

The genotype distribution results of the 20210 G/A SNP of prothrombin gene were not showed significant differences between controls and ischemic stroke patients. However, the 20210 G/A mutant alleles are not entirely absent among Iraqi patients with ischemic stroke.

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References


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