

Involvement of the 5-HTTLPR polymorphism of the serotonin transporter gene SLC6A4 in the onset of migraine

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INTRODUCTION

Migraine is a complex neurological disorder characterized by long-lasting unilateral headaches. The genetic component seems to play a crucial role in the susceptibility and manifestation of the disease, particularly genetic alterations that affect the functioning of the serotonergic system and its components.

OBJECTIVE

The study aims to study a possible correlation between the 5-HTTLPR polymorphism of the serotonin transporter gene SLC6A4 in the onset of migraine

METHODS

The study is carried out within the molecular biology unit - Hematology Laboratory of the main military training hospital of Tunis

5-HTTLPR polymorphism

Insertion/deletion of 43 bp in the polymorphic 5-HTTLPR region located in the promoter sequence of the SLC6A4 gene, results in two allelic forms with different sizes and activities:

- **The long (L)** variant allele with the 43 bp insertion (512 bp) which increases the mRNA expression level and density of presynaptic 5-HTT (transporter) and therefore eliminates more serotonin from the synaptic cleft (maintain cerebral 5-HT homeostasis).

- **The short (S)** variant allele without 43 bp (469 bp) which produces lower levels of transporter mRNA, decreases its presynaptic density, and thereby reduces its activity.

Case-control study carried out on 60 migraine patients and 60 controls

1 • Genomic DNA extraction

2 • DNA amplification (PCR)

3 • Migration on agarose gel

4 • Calculate and compare the distribution of genotype and allele frequencies

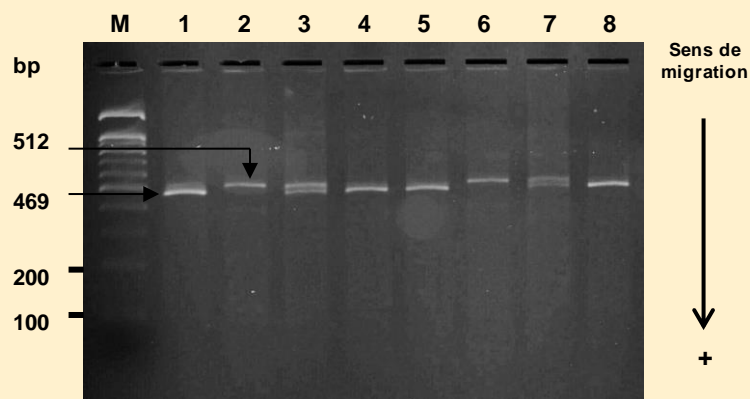


Fig: Electrophoretic profile of 5-HTTLPR polymorphism genotyping

M: Size marker (100 bp); 1, 4, 5, 8: Homozygous SS genotype (469 bp); 2, 6: Homozygous LL genotype (512 bp); 3, 7: Heterozygous SL genotype (469 bp + 512 bp)

Table: Comparison of genotypic and allelic frequencies between patients and controls

	Patients %	Controls %	χ^2	OR	[CI 95%]	P
SS	53,33	58,33	0,30	0,81	[0,39-1,67]	0,58
LL	46,66	38,33	0,85	1,40	[0,68-2,91]	0,35
SL	0	3,33	2,03	0	indéfini	0,15
S	53,33	60	1,08	0,76	[0,45-1,27]	0,29
L	46,66	40	1,08	1,31	[0,78-2,18]	0,29

Chi-deux (χ^2), OR: Odds Ratio, CI: Confidence interval, p: P Value (significant if $p < 0.05$).

Analysis of genotype frequencies revealed that the homozygous SS genotype is common in both groups, with a frequency of 53.33% among patients and 58.33% among controls.

The LL genotype is more prevalent among migraine patients (46.66%) compared to controls (38.33%). Conversely, the heterozygous SL genotype is the least common in both groups (absent in migraine patients and very rarely present 3.33% in the control group)

Examination of allele frequencies indicated that the S allele is significantly more frequent in the control group (60%), while the L allele is more prevalent in migraine patients (48%).

The study of genotypic and allelic frequencies as well as the application of different statistical tests showed a statistically non-significant association between the 5-HTTLPR polymorphism and migraine with a p -value > 0.05 and a χ^2 value < 3.84 .

RESULTS

The electrophoretic profile (fig) showed the presence of:

- A band of **469 bp** for the **homozygous SS** genotype
- A band of **512 bp** for the **homozygous LL** genotype
- Two bands **469 bp and 512 bp**, for individuals with the **heterozygous SL** genotype

CONCLUSION

The molecular study of the SLC6A4 gene of the serotonin transporter, showed a statistically insignificant correlation for the polymorphisms, 5-HTTLPR, so it does not represent a risk factor for migraine.