Association Between Singe Nucleotide Polymorphism of Toll-like Receptors (TLR)-7 and Behçet Uveitis in a Turkish Population

Hilal Eser Ozturk¹, Sengul Tural², Nurten Kara², Yuksel Sullu¹

¹Ondokuz Mayıs University, Ophthalmology Department

²Ondokuz Mayis University, Department of Medical Biology Section of Medical Genetics

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Background

- Behçet disease has multifactorial pathogenesis
- Inflammatory response is triggered by various environmental factors in genetically susceptible individuals.
- BD has common features with autoinflammatory diseases which is believed that the innate immune system has an important role in initiation.
- TLRs are the key components of innate immune system.
- Single nucleotide polymorphisms (SNPs) in *TLR-7* gene were analyzed in patients with Behçet uveitis.

Patients & Methods

- Prospective cross-sectional clinical and genetic study
- Genomic DNA obtained from 198 individuals:
 - 89 patients with BU
 - 109 healthy controls
- 2 SNP in *TLR-7* gene were determined by using PCR with spesific primers.
 - rs179008
 - rs 179009
- DNA sequencing and bioinformatic analysis were done.

SNP	Genotype/allel	Patients (n=89) (%)	Controls (n=109) (%)	χ²	P value	OR (%95CI)	
TLR-7 rs179009	TT	75(84,0)	92(84,0)	0.621*	0.73		
	TC	2(2,00)	1(1,00)				
	CC	12(14,0)	16(15,0)				
	CC+TC:TT	14 : 75	17 : 92	0.006	0.97	1.01 (0.459-2.198	
	TC+TT:CC	77 : 12	93 : 16	0.057	0.81	0.90 (0.390-2.043)	
	С	26	33	0.021	0.88	0.959 (0.545-1,66)	
	Т	152	185				
TLR-7 rs179008	AA	75 (84,30)	89 (81,6)	2,983	0.39		
	AT	2 (2,30)	7 (6,4)				
	TT	12(13,40)	13(12,0)				
	AT+TT:AA	14:75	20 :89	0.236	0.62	0,831 (0,385-1,762)	
	AA+AT:AA	12: 77	13: 96	0.107	0.74	1,150 (0,486-1,702)	
	Т	26	33	0.020	0.88	0,959 (0,545-1,676)	
	Α	152	185				

SNP-SNP	SNP Genotypes	Patients n (%)	Controls n (%)	χ²	P value
	TLR-7				
rs 179009 (T→C)	AATT	72 (45,9)	85 (54,1)		
rs 179008 (A → T)	AACC	3 (42,9)	4 (57,1)		
	ATTC	2 (66,7)	1 (33,3)	9,355	0,09
	TTTT	3 (100)	0 (0.0)		
	TTGG	9 (40,9)	13 (59,1)		
	ATTT	0 (0,0)	6 (100)		
Total		89	109		

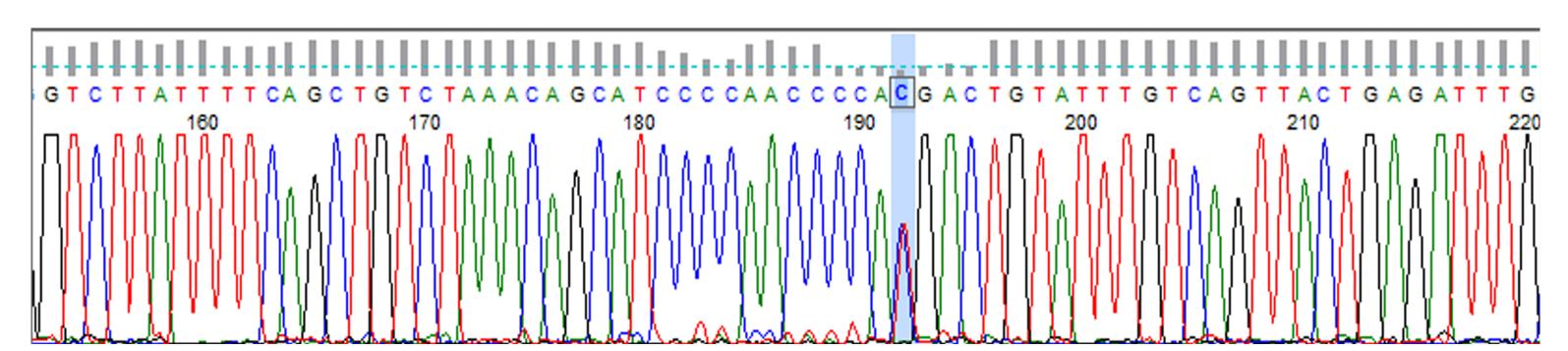


Figure: TLR-7 rs 179009 alelle T/C

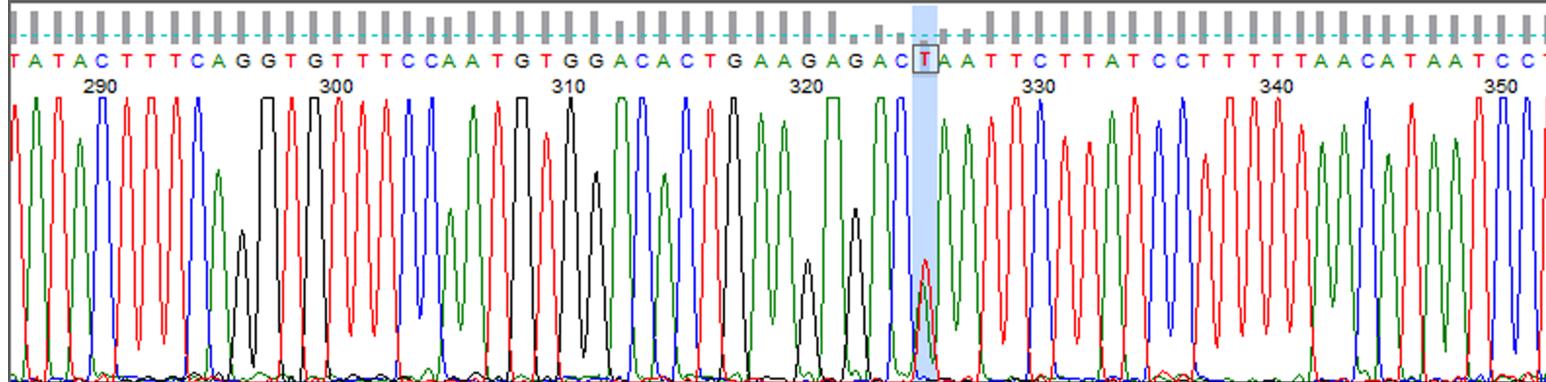


Figure: *TLR-7* rs 179008 allele A/T

Results

- Patient group:
 - 65 (%73) male, 24 (%27) female
 - Mean age 35.1 ± 12.35 (11-63)
- Control group
 - 67 (%61) male, 42 (%39) female
 - Mean age 33.75 ± 14.48 (21-55)
- There was no statistically significant difference in genotype distribution for *TLR-7* rs179009 T/C and *TLR-7* rs 179008 A/T genotypes in patient and control group (p=0.73, p=039 respectively)
- In the combined genotype analysis, there was no statistically significant difference (p=0.09)

Conclusion

- This genetic study did not find any relationship between two SNPs at the TLR7 gene loci and Behçet uveitis in a Turkish population
- No effect of these genotypes on the other clinical features of BD including oral ulcer, genital ulcer, skin involvement, pathergy results, joint involvement, neurological involvement and thromboembolism were detected as well.

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