

Vitamin D Receptor Gene Polymorphisms in Rheumatoid Arthritis

Romatoid Artritte D Vitamini Reseptörü Geni Polimorfizmleri

Ömer ATEŞ,¹ Bilgen DÖLEK,² Levent DALYAN,³ Ayşegül Topal-SARIKAYA³

¹Department of Medical Biology, Medical Faculty of Gaziosmanpaşa University, Tokat, Turkey;

²Department of Molecular Genetics, Düzen Laboratory, İstanbul, Turkey;

³Departments of Molecular Biology and Genetics, İstanbul University, İstanbul, Turkey

Objectives: In this study, we investigated the association of *BsmI*, *TaqI*, and *FokI* polymorphisms in the vitamin D receptor gene in rheumatoid arthritis patients with rheumatoid factor positivity and erosive disease of rheumatoid arthritis.

Patients and methods: In this study we analyzed *BsmI*, *TaqI* and *FokI* polymorphisms in the vitamin D receptor gene in 98 patients with RA (78 females, 20 males; mean age 50.8±12.3 years; range 38 to 63 years) and 122 healthy subjects (96 females, 26 males; mean age 57.1±6.1 years; range 51 to 63 years) by a polymerase chain reaction followed by enzymatic digestion between May 2006 and April 2008.

Results: The distributions of *BsmI*, *TaqI* and *FokI* alleles/genotypes frequencies were similar in patients and controls. There were no significant differences between the vitamin D receptor genotypes/alleles and the presence or absence of erosions and rheumatoid factor.

Conclusion: Our findings suggest that vitamin D receptor polymorphisms do not play a role in either rheumatoid arthritis susceptibility or in rheumatoid factor positivity and erosive disease of rheumatoid arthritis in the Turkish population.

Key words: Polymorphism; rheumatoid arthritis; vitamin D receptor.

Amaç: Bu çalışmada, romatoid artritli hastalarda D vitamini reseptörü genindeki *BsmI*, *TaqI* ve *FokI* polimorfizmleri ile romatoid faktör pozitifliği ve erozivite arasındaki ilişki araştırıldı.

Hastalar ve yöntemler: Bu çalışmada Mayıs 2006 - Nisan 2008 tarihleri arasında 98 romatoid artritli hasta (78 kadın, 20 erkek; ort. yaş 50.8±12.3 yıl; dağılım 38-63 yıl) ve 122 sağlıklı bireyde (96 kadın, 26 erkek; ort. yaş 57.1±6.1 yıl; dağılım 51-63 yıl), D vitamini reseptörü genindeki *BsmI*, *TaqI* ve *FokI* polimorfizmleri-polimeraz zincir reaksiyonu ve bunu takiben enzim kesim yöntemleri kullanılarak analiz edildi.

Bulgular: Hastalarda ve kontrollerde *BsmI*, *TaqI* ve *FokI* allel ve genotip frekansları benzer oranlarda idi. D vitamini reseptörü genotip ve alleli frekansları ile romatoid faktör pozitifliği ve erozivite arasında anlamlı fark bulunmadı.

Sonuç: Bulgularımız Türk toplumunda D vitamini reseptörü polimorfizmlerinin hem romatoid artrite yakınlıkta hem de romatoid faktör pozitifliğinde ve romatoid artrit eroziv hastalık oluşumunda rol oynamadığını göstermektedir.

Anahtar sözcükler: Polimorfizm; romatoid artrit; vitamin D reseptör.

Rheumatoid arthritis (RA) is a chronic inflammatory disease characterized by a distinctive pattern of joint involvement and joint destruction. Rheumatoid arthritis pathogenesis involves both genetic and environmental factors, but the complete etiological

picture remains unclear. The genetic basis of RA is quite complex. Genetic factors contribute 50 to 60% to the risk of developing RA.^[1]

Vitamin D (VD) is a potent regulator of calcium homeostasis and plays a role in immune regulation

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Correspondence: Ömer Ateş, PhD. Gaziosmanpaşa Üniversitesi Tıp Fakültesi Tıbbi Biyoloji Anabilim Dalı, 60100 Tokat, Turkey.

Tel: +90 356 - 212 95 00 e-mail: omerates27@yahoo.com

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and inflammation.^[2,3] A significant association between VD insufficiency and an increased incidence of autoimmune disorders has been determined.^[4] In addition, a significant clinical improvement was shown in the VD- treated patients with RA.^[5] Vitamin D receptor (VDR) has been demonstrated in the rheumatoid synovium and sites of cartilage erosion in patients with RA.^[4] Vitamin D receptor agonists have a critical physiological role in the regulation of the immune system.^[6,7] The role of VDR polymorphisms in RA has been studied by several investigators. There are controversial results in various reports due to ethnicities, extensive geographic variations and possibly study designs.^[8-12]

There is no data on frequencies of VDR polymorphisms in Turkish patients with RA. In this study, we aimed to investigate the association of *BsmI*, *TaqI*, and *FokI* polymorphisms in the VDR gene in RA patients with rheumatoid factor (RF) positivity and erosive disease of RA.

PATIENTS AND METHODS

Patients

We studied VDR genetic polymorphisms in 98 RA patients (78 females and 20 males; mean age 50.8±12.3 years; range 38 to 63 years) and 122 healthy hospital workers (96 females and 26 males; mean age 57.1±6.1 years; range 51 to 63 years) who had taken part in our previous studies.^[13,14] All patients were registered at the outpatient clinic of İstanbul University, Cerrahpaşa Medical Faculty, Rheumatology Department, and they fulfilled the American College of Rheumatology (ACR) classification criteria for the diagnosis of RA.^[15] In general, patients with elevated RF and anti-cyclic citrullinated peptide (anti-CCP) have more severe disease. Although it is known that anti-CCP is more sensitive and has a more specific parameter than RF, we have only the RF results as a part of our previous study. Rheumatoid factor was known in 92 RA patients

and, among these, 58 (63%) had positive RF. Hand and feet X-rays were available in 74 of the patients and, among these, 51 (69%) had erosions. The healthy group had no history of any rheumatologic disorder. This study protocol was approved by the Ethics Committee of İstanbul University, Cerrahpaşa Medical Faculty and written informed consent was obtained from each subject.

Genetic analysis

Genomic deoxyribonucleic acid (DNA) was isolated from frozen whole blood-EDTA according to a standard procedure.^[16] The primer sequences were used previously.^[17] Polymerase chain reaction (PCR) conditions were 96 °C for 1 min in all reactions, followed by 30 cycles at 94 °C for 45 s, 60 °C for 45 s, 72 °C for 45 s (*FokI*); 30 cycles at 94 °C for 1 min, 55 °C for 1 min, 72 °C for 1 min (*TaqI*); and 30 cycles at 94 °C for 1 min, 60 °C for 1 min, 70 °C for 1 min (*BsmI*). All PCR products were analyzed by restriction fragment length polymorphism (RFLP) and gel electrophoresis.

Statistical analysis

Statistical analysis was performed by Epi Info Software Version 3.2.2 (CDC, Atlanta GA, USA). The distributions of *BsmI*, *TaqI* and *FokI* polymorphisms in the VDR gene between RA patients and healthy controls were compared using the chi-square or Fischer's exact test. *P* value less than 0.05 was considered significant. The genotype distribution and Hardy-Weinberg equilibrium were tested with the chi-square test for quality of fit, Arlequin software version 2000, (University of Geneva, Switzerland).

RESULTS

The demographic characteristics of RA patients and healthy controls are given in Table 1. The distributions of alleles and genotypes frequencies of *BsmI*, *TaqI* and *FokI* polymorphisms of VDR gene in RA and controls

Demographic features	RA patients (n=98)			Healthy controls (n=122)		
	n	%	Mean±SD	n	%	Mean±SD
Age (years)			50.84±12.3			57.1±6.1
Sex						
Female	78			96		
Male	20			26		
Patients with erosions	51	69			N/A	
Patients with RF positivity	58	63			N/A	

RA: Rheumatoid arthritis; SD: Standard deviation; N/A: Not applicable; RF: Rheumatoid factor.

are given in Table 2. The frequencies of F, T and B allele were higher while that of the f, t, b allele were lower among RA patients in comparison to controls, but no associations were found between these alleles and RA ($p=0.10$, 0.06 , and 0.11 , respectively). No significant differences in the distribution of the VDR genotypes between RA and controls were observed. The frequencies of *BsmI*, *TaqI* and *FokI* alleles/genotypes were similar among patients regardless of RF positivity and presence or absence of erosions (Table 3). There were no significant differences between the VDR genotypes/alleles and presence or absence of erosions and RF.

DISCUSSION

There have been several studies on the possible association of RA, clinical manifestations of RA, and VDR polymorphisms. The results have been controversial due to ethnicities and geographic variations.^[8-12] In addition, the age groups of the subjects and the sample sizes of these studies may account for the contradictory results. The present study evaluated for the first time the possible association of VDR polymorphisms among a group of Turkish RA patients.

The study results showed that only 63% of the patients with RA were positive for RF. Similarly, in the Baran et al.^[18] study, the results were 59 percent. These results could be explained as RF may present less sensitive results in the activity of disease. Anti-CCP can be a better prognostic factor as shown in various studies.^[19-20] However, we have only the RF results as a part of our previous study.

We could not observe any association between the following *BsmI*, *TaqI*, *FokI* polymorphisms in VDR gene and RA with RF positivity and erosive disease of RA. These findings are parallel with previous studies conducted on the association of RA and VDR with negative results. Goertz et al.^[11] failed to show a significant association between RA susceptibility or bone turnover, family history, the presence of RF and *FokI*, *TaqI*, *BsmI* genotypes in VDR gene. Lee et al.^[12] observed no association with VDR *TaqI*, *BsmI* polymorphisms and RA susceptibility and bone erosion. Garcia-Lozano et al.^[8] showed no significant difference between *TaqI*, *BsmI* ApaI polymorphisms of VDR gene and RA. The study results published by Rass et al.^[21] which indicated that there was no relationship between *BsmI* polymorphism and RA confirms our findings in this study. However, the fact that there is a significant correlation between RF positivity and severity of RA observed in the Rass et al. study contraindicates the results of our study.^[21] In two reports, an association was found between *BsmI* polymorphism of the VDR gene and RA severity.^[8-10] Our results agree with the results of Gomez-Vaquero et al.^[10] that reported no association between the VDR genotypes/alleles and the presence or absence of erosions and RF. Masi et al.^[22] found no relationship between RA and *FokI* genotype, but their results suggest that *FokI* polymorphism may lead to higher risk for loose bone mass. Gough et al.^[23] reported a significant association between the presence of the VDR gene t allele (*TaqI*) in female patients with RA and accelerated bone loss. An earlier study suggested discordance between *TaqI*, *BsmI* polymorphisms and RA but a positive correlation between *FokI* (F allele and F/F genotype) and RA.^[9]

In summary, in several studies, similar to our findings, no significant associations have been found between RA and VDR polymorphisms except for *FokI* polymorphism which was significantly associated with RA in a French population.^[9] These results reflect that *TaqI*, *FokI*, *BsmI* and ApaI polymorphisms in the VDR gene may not be responsible for the primary

Table 2. Distribution of vitamin D receptor polymorphisms in patients and controls

Vitamin D receptor locus, polymorphism	RA patients (n=98)		Healthy controls (n=122)	
	n	%	n	%
Fok				
F/F	51	52	53	43
F/f	40	41	57	47
f/f	7	7	12	10
Allele				
F	142	72	163	67
f	54	28	81	33
Taq				
T/T	45	46	44	36
T/t	43	44	60	49
t/t	10	10	18	15
Allele				
T	133	68	148	61
t	63	32	96	39
Bsm				
B/B	12	12	7	6
B/b	46	47	60	49
b/b	40	41	55	45
Allele				
B	70	36	74	30
b	126	64	170	70

RA: Rheumatoid arthritis.

Table 3. Relationship between clinical characteristics of patients with rheumatoid arthritis and vitamin D receptor polymorphisms

Vitamin D receptor locus, polymorphism	Rheumatoid factor				Erosive disease				<i>p</i>	
	(+)		(-)		(+)		(-)			
	(n=58)		(n=34)		(n=51)		(n=23)			
	n	%	n	%	n	%	n	%		
Fok										
F/F	29	50	18	53	25	49	11	48		
F/f	25	43	13	38	23	45	10	43	0.84 ^a , 0.85 ^b	
f/f	4	7	3	9	3	6	2	9		
Allele										
F	83	72	49	72	73	72	32	70		
f	32	28	19	28	29	28	14	30	0.47 ^a , 0.40 ^b	
Taq										
T/T	26	45	16	47	28	55	12	52		
T/t	27	46	14	41	20	39	8	35	0.82 ^a , 0.59 ^b	
t/t	5	9	4	12	3	6	3	13		
Allele										
T	79	68	46	68	76	75	32	70		
t	37	32	22	32	26	25	14	30	0.47 ^a , 0.26 ^b	
Bsm										
B/B	8	14	4	12	5	10	3	13		
B/b	24	41	17	50	26	51	10	43	0.72 ^a , 0.81 ^b	
b/b	26	45	13	38	20	39	10	43		
Allele										
B	40	34	25	37	36	35	16	35		
b	76	66	43	63	66	65	30	65	0.37 ^a , 0.47 ^b	

a: *p* value, RF (+) vs RF (-); b: *p* value, erosive (+) vs erosive (-).

disease association. Because there are controversial results in various reports, additional studies are needed to elucidate the possible association between the VDR gene and clinical features of RA including other polymorphisms in the VDR gene such as Cdx2 polymorphism. These controversial results could be due to heterogeneity between populations and the small number of samples used in the studies. Not forgetting the small sample size of our study, the present data suggests that VDR polymorphisms do not play a role either in RA susceptibility or in RF positivity and erosive disease of RA in a group of Turkish population.

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Declaration of conflicting interests

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