



A Study of Glucocorticoid Receptor Gene Polymorphism (BclI) in Egyptian patients with Rheumatoid Arthritis

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ABSTRACT

Objective: The aim of this work was to investigate the relation of the glucocorticoid receptor gene polymorphism (BclI) with the rheumatoid arthritis (RA) activity. **Method:** This study included 20 volunteers controls and 70 patients with RA (35 in remission and 35 active). Serum α 1 antitrypsin was measured for all subjects by radial immunodiffusion (RID) plates. Other markers of inflammation and disease activity were also detected. All subjects were genotyped for glucocorticoid receptor gene polymorphism using PCR-RFLP. Serum α 1 antitrypsin was higher in active RA patients compared to inactive RA patients and controls. **Results:** Highly significant level of α 1 antitrypsin was detected with variant genotypes. Significant association was found between the investigated BclI polymorphism and the parameters of the RA activity. **Conclusions** This study suggesting the relationship between glucocorticoid receptor gene polymorphism and the activity of RA.

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INTRODUCTION

Rheumatoid arthritis (RA) happens when the immune system targets the joint linings. It is characterized by synovitis and autoantibody formation (1,2). Also, RA is a systemic inflammatory disorder, meaning it affects the entire body but primarily, characterized by progressive damage of synovial joints, but is common in the wrist and fingers. More women than men get rheumatoid arthritis and its incidence rises with increasing age. (3) Glucocorticoids(GC) perform their effects through binding to their cytosolic receptor

(GR) and the formed complex translocate into the nucleus where it interacts with the promoter GC-responsive elements (4,5). The control of target genes expression results from the previous interaction (6). The variability of GR gene may result from different forms of GR and in turn regulates several biological function that have an impact of the susceptibility to many diseases (7,8). GR gene consists of 9 exons and 8 introns and located on chromosome 5 (locus 5931) (7). BclI polymorphism consists of dialleles with 4.5 and 2.3 kb fragment lengths (9,10).

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Many of rheumatoid arthritis patients produce a group of antibodies reactive with the Fc fragment of IgG. Alpha1-antitrypsin (AAT) is a glycoprotein in blood known as a serine proteinase inhibitor. AAT inhibits a wide variety of proteases such as proteinase 3, neutrophil elastase, and cathepsin G. Alpha1-antitrypsin have anti-inflammatory, immunomodulatory, these proteases are also synthesized by monocytes (joint invading neutrophils) following inflammatory stimuli and involved in arthritis development. A lack of Alpha1-antitrypsin in rheumatoid arthritis patients could increase inflammation because of uninhibited lysosomal enzymes leads to a chronic uninhibited tissue breakdown (11).

In this study, we aimed to find the association of glucocorticoid receptor gene variation with the risk of RA and its activity, serum α 1 antitrypsin level and different rheumatoid inflammatory markers.

Patients and Methods

Study population

In this study three groups of patients was studied group I: 20 healthy volunteers serving as control group, a total of 70 RA patients were enrolled in this study, two groups were defined as follows: 35 of them were in remission (inactive group II) and the other 35 were in the active form of the disease (active group III). American Rheumatism Association (ARA) purposed a diagnostic criteria for RA (12), and the clinical activity was assessed according to the 28 count Disease Activity Score (DAS28). It was calculated with the following equation: $DAS28 = 0.56 \times \sqrt{28TJC} + 0.28 \times \sqrt{28SJC} + 0.7 \times \ln ESR + 0.014 \times GH$, where 28SJC and 28TJC are the swollen joint count and tender joint count from 28 joints and general health (GH) is the patient's global assessment on a 100-mm visual analog scale (VAS) (13). The included subjects were randomly chosen from Internal Medicine Department

of Tanta University Hospital, Egypt during the period from May2014 to April 2015.

Sampling

Peripheral blood was drawn from each volunteer, and was divided into 2 portions: 1ml of whole blood was collected into evacuated tubes containing EDTA for DNA extraction, and the remaining portion of the blood was used to separate serum immediately. Separated serum was stored at -20°C until further use.

Determination of serum rheumatoid inflammatory parameters

Rheumatoid factor (RF) was measured by a quantitative immunonephelometry (Behring, Marburg, Germany). C-reactive protein (CRP) was measured by semiquantitative latex test (Omega, Avitex, UK). The erythrocyte sedimentation rate (ESR) was measured by the Westergren method; the reading of the 1st hour was included in the study (14). α 1- antitrypsin concentration in serum was assayed by radial immunodiffusion using commercial assay Kit. α 1- antitrypsin was obtained from KALLESTAD (15).

DNA extraction and genotyping

DNA was isolated and from whole blood according to the manufacturer instructions (Gene JET Genomic DNA) Purification Kit; Thermo Scientific, EU Lithuania). Isolated DNA was stored at -20°C until use. polymorphism was detected using the polymerase chain reaction -restriction fragment length polymorphism method (PCR-RFLP). The following primers were used: forward primer (5'dTAA GCC CAG ACC TGC TGT TG3'), and reverse primer 5'-(5'dGGG GTT TAC ACA ACC CGC TA3').

The PCR reaction was carried out in a total volume of 50 μl containing 50 ng of genomic DNA, 0.5 μM each primer (Biosearch Technologies, Petaluma, Ca USA), 100 μM dNTPs, 10 mM Tris-HCl (pH 8.3), 1.5 mM MgCl_2 , 50 mM KCl and

0.5 U of *Taq* polymerase. The amplification cycle was performed as follows: firstly, denaturation at 95 °C for 7 min, preannealing at 64 °C for 1 min, and then elongation at 72 °C for 2 min followed by 35 cycles of 30 s at 95 °C, 30 s at 64 °C and 1 min 15 s at 72 °C, and finally, elongation at 72 °C for 10 min. To investigate the Bcl1 polymorphism, 10 µl of the PCR products was digested with 4 U of restriction endonuclease Bcl1 (New England Biolabs Ltd, UK) for 6 h at 50 °C and the digested products were separated on 1% agarose gel in TBE buffer. The gel was stained with ethidium bromide and observed in UV-light. Digestion of the PCR product might give the following predicted fragment sizes: two bands of 90 and 116 bp in the case of homozygous individual for the shorter allele (G/G); a single band of 206 bp for homozygous individual for larger allele (C/C) and three bands of 90, 116, and 206 bp for heterozygous individuals (G/C) (Table 5) (16).

Statistical analysis

The results for quantitative variables were analyzed using one-way ANOVA tests and were expressed as the mean ± SD. Qualitative variables were expressed as percentages and were compared using the Chi-square test. A value of $P < 0.05$ was considered statistically significant. All data were evaluated using the statistical package for social sciences (SPSS).

Results and Discussion

Our study aim to identify the common genotype of GR gene Bcl1 polymorphism in some Egyptian RA patients using Polymerase chain reaction-restriction fragment length polymorphism (PCR-RFLP) and measuring clinical diagnostic parameters the erythrocyte sedimentation rate (ESR), α1 antitrypsin, C-reactive protein (CRP) and rheumatoid factor (RF), α1 antitrypsin.

In this study, the 3 included groups (healthy controls, active RA and inactive RA) were homogenous as regards age, sex,

levels of serum ESR and CRP were significantly higher in the active RA group compared to both inactive RA and control groups. RA patients had significantly higher levels of RF and DAS28 score rather than inactive patients. Alpha-1 antitrypsin significantly higher in the active and inactive RA groups compared to control groups. Also, There was no significant difference between both patient groups as regards Alpha-1 antitrypsin. (Table 1)

There were no significant differences between the genotypes as regards age, Levels of ESR, CRP, RF, DAS28 and alpha 1 antitrypsin were significantly increased in GG genotypes compared to CC genotypes. (Table 2). There were no significant differences between the genotypes as regards RF, ESR, CRP and DAS28 in active RA patients. (Table 4)

Polymorphism of Glucocorticoid Receptor (GR) gene, resulting from variability of GR gene, can affect many biological functions regulation, such as GC responsiveness and regulation of hypothalamic-pituitary-adrenal axis, thereby underlying susceptibility to many diseases. The Bcl1 site was found to be in either the first or second intron of the GR gene (17). There are three Bcl1 sites, one in intron 1 and two in intron 2. Only the site within intron 2 has Bcl1 polymorphism where at 647 bp from exon/intron junction (18). In case of presence of wild type restriction site of Bcl1 (TGATCA), fragments of 2.2 kb will be generated after enzymatic digestion. On the other hand, fragments of 3.9 kb will be produced in case of polymorphic Bcl1 site (TCATCA), where the bold base indicated the G/C transition. Our primers were designed to flank the Bcl1 site at position +647, giving a fragment of 206 bp. To study the Bcl1 polymorphism among our patients, the DNA product was digested with Bcl1 restriction endonuclease and the genotype

was determined according to the pattern of digested fragments (see Table 5).

Alpha1-antitrypsin is a very effective inhibitor of proteolytic enzyme, responsible for almost 90% of the total serum trypsin inhibitory capacity. Increased inflammation and activity of lysosomal enzyme in rheumatoid arthritis may force the concentration of alpha1-antitrypsin to rise (11).

This study showed there was no significant difference between both active and inactive patient groups as regards Alpha-1 antitrypsin which were significantly increased in GG genotypes compared to CC genotypes.

Brackertz *et al.*, compared the levels of proteinase inhibitors between normal individuals and patients of probable and classical rheumatoid arthritis and found out that there was a significant rise in the levels of alpha1-antitrypsin inhibitor in serum as well as in synovial fluid of rheumatoid arthritis patients (19). The immunochemically determined levels of alpha-1-antitrypsin were found to be raised in the patients but their proteinase inhibitory capacity was depressed. This implied to the fact that, though the alpha1-antitrypsin levels were raised in the rheumatoid arthritis patients, they were partially inactivated with a decreased specific elastase inhibitory activity (20). In another study, the quantitative radial immuno diffusion method was used to study the concentration of alpha1-antitrypsin in the serum and synovial fluid. The levels were much higher in the rheumatoid arthritis patients than the healthy controls individuals (21).

On the contrary, other studies reported that the minor alleles of the BclII and N363 S polymorphisms are associated with decreased susceptibility to RA development. These opposite associations suggest that determined GC resistance may predispose to development of autoimmune disorders, at least in RA, and vice versa (22).

Conclusions

Glucocorticoid receptor gene is important for RA and could use as clinical manegment of RA. Further studies on large individuals groups may cause more association between gene polymorphism and RA patient degree.

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Table.1 Clinical and biochemical characteristics of RA patients and healthy control

Parameters	Healthy Controls (group I) n = 20	Inactive RA cases (group II) n = 35	Active RA cases (group III) n = 35	P1	P2	P3
Age	42.03 ± 8.77	43.03 ± 10.48	44.46 ± 9.94	0.666	0.282	0.560
Sex M/F	5 / 15	5 / 30	7 / 28	0.322	0.666	0.526
DAS		1.78 ± 0.51	4.71 ± 1.02			0.001*
CRP	3.38 ± 1.2	4.86 ± 2.83	48.29 ± 26.08	0.031*	0.001*	0.001*
RF		26.89 ± 15.77	105.91 ± 37.68			0.001*
ESR	12.13 ± 2.49	16.46 ± 9.68	78.51 ± 18.11	0.056	0.001*	0.001*
Alpha-1 antitrypsin	149.7 ± 52.6	209.4 ± 35.13	247.1 ± 54.45	0.010*	0.001*	0.092

Table.2 Relation between BclI polymorphism and the clinical and biochemical characteristics of all RA patients (n = 70)

Parameters	Healthy Controls (group I) n = 20	CC V (n=22)	GG IV (n=48)	P1	P2	P3
Age	42.03 ± 8.77	41.05 ± 10.91	43.88 ± 9.82	0.752	0.468	0.284
Sex M/F	5 / 15	2 / 20	8 / 40	0.167	0.426	0.401
DAS		1.82 ± 0.53	3.89 ± 1.63			0.001*
CRP	3.38 ± 1.2	5.18 ± 2.99	36.40 ± 29.71	0.016*	0.001*	0.001*
RF		26.77 ± 15.53	87.75 ± 46.73			0.001*
ESR	12.13 ± 2.49	35.05 ± 32.21	63.31 ± 31.0	0.003*	0.001*	0.001*
Alpha-1 antitrypsin	149.7 ± 52.6	275.6 ± 65.6	221.31 ± 55.4	0.001*	0.010*	0.046*

Table.3 Frequencies of BclI genotypes in RA patients and healthy controls

		Inactive RA cases (group II) n = 35	Active RA cases (group III) n = 35	Total
CC	N	32	7	39
	%	91.0%	20.0%	21.4%
GG	N	3	28	31
	%	9.0%	80.0%	78.6%
Chi-square	X ²	36.192		
	P-value	0.001*		

Table.4 Relation between Bcl1 polymorphism and the parameters of the disease activity in active RA patients (n =35)

Parameters	VI-GG (n=25)	VII-CC (n=10)	p. value
Age	44.0 ± 10.61	45.60 ± 8.44	0.674
Sex M/F	6 / 19	1 / 9	0.350
DAS	4.81 ± 1.04	4.44 ± 0.96	0.336
CRP	49.76 ± 26.55	44.60 ± 25.85	0.604
RF	99.56 ± 20.61	121.80 ± 61.96	0.116
ESR	80.40 ± 18.72	73.80 ± 16.44	0.338

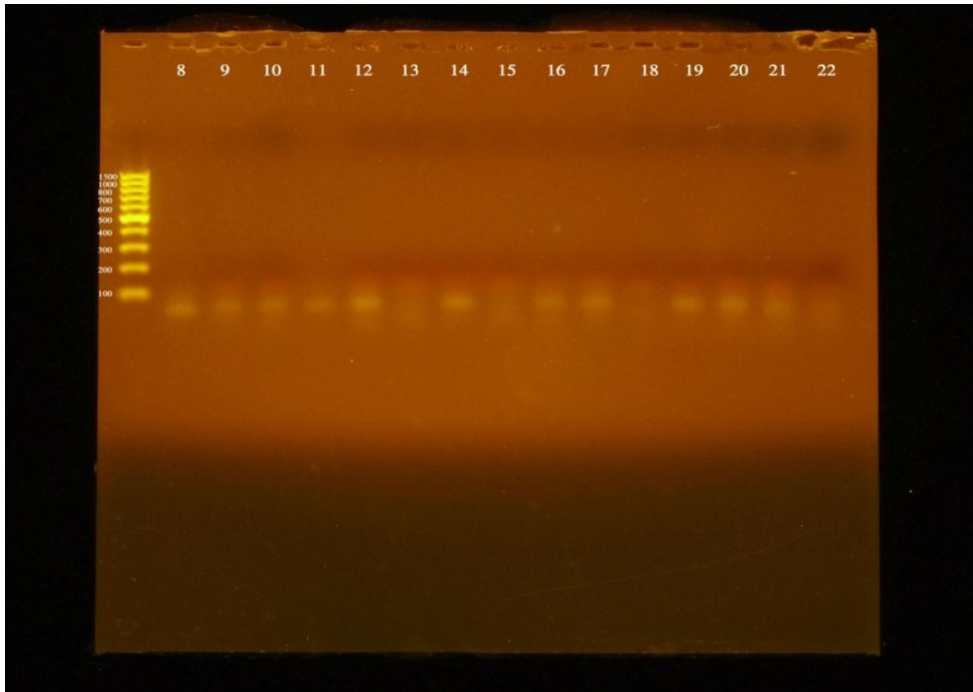


Fig.(1):Agarose gel Electrophoresis of Bcl1 gene polymorphism.Lane 1: DNA ladder;lane 8 healthy control, lanes 9-22: PCR product digestion of 15 patients. The digestion pattern shows a homozygous individual for the smaller allele. The digestion of PCR product by the Bcl1 enzyme produced two fragments for homozygous G/G.

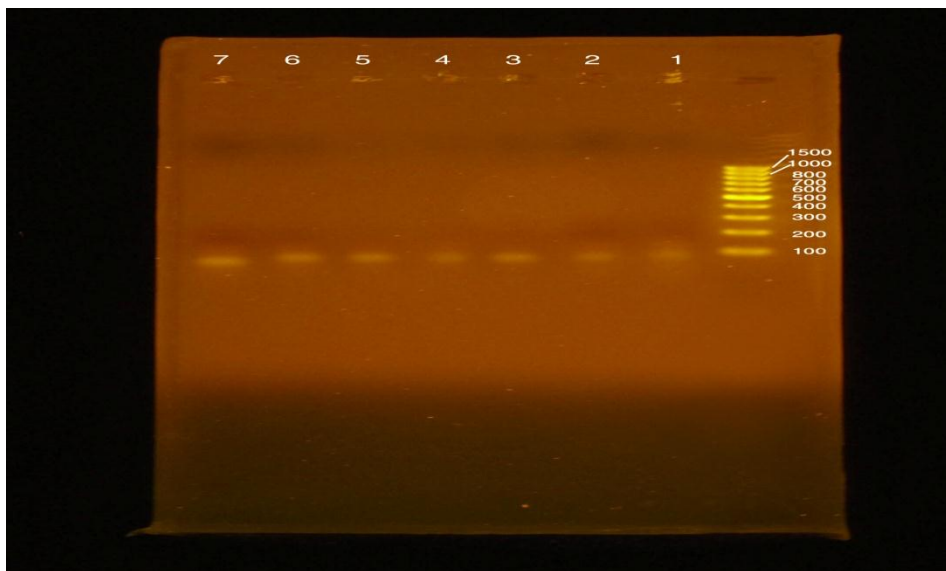


Fig.(2):Agarose gel Electrophoresis of Bcl1 gene polymorphism.Lane 1: DNA ladder;lane 1healthy control, lanes 2-7: PCR product digestion of 15 patients. The digestion pattern shows a homozygous individual for the larger allele. The digestion of PCR product by the Bcl1 enzyme produced two fragments for homozygous c/c.

A-



B-

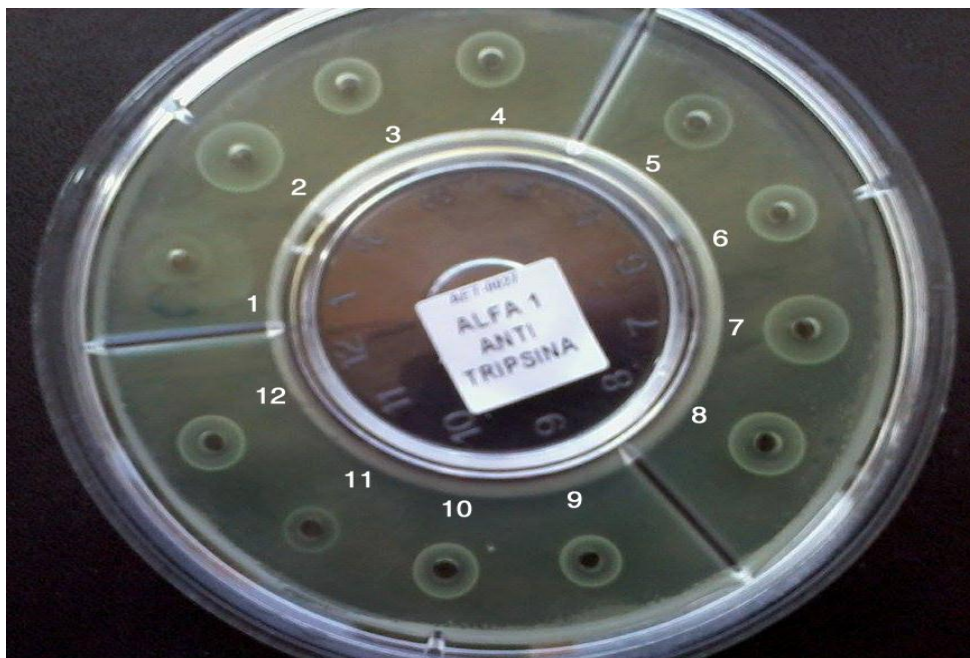


Fig. (3): Photogram showing the immunoprecipitation ring of serum α 1 antitrypsin in RA patients, well 1 healthy control. Photo A, B.