

Correlation between HLA haplotypes and the development of antidrug antibodies in a cohort of patients with rheumatic diseases

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Introduction: The aim of this study was to investigate the correlation between human leukocyte antigen (HLA) haplotypes and the development of antidrug antibodies (ADAs) in a cohort of patients with rheumatic diseases.

Patients and methods: We evaluated the presence of ADAs in 248 patients with inflammatory rheumatic diseases after 6 months of treatment with anti-TNF drugs: 26 patients were treated with infliximab (IFX; three with rheumatoid arthritis [RA], 13 with ankylosing spondylitis [AS], 10 with psoriatic arthritis [PsA]); 83 treated with adalimumab (ADA; 24 with RA, 36 with AS, 23 with PsA); 88 treated with etanercept (ETA; 35 with RA, 27 with AS, 26 with PsA); 32 treated with certolizumab (CERT; 25 with RA, two with AS, five with PsA); and 19 treated with golimumab (GOL; three with RA, seven with AS, nine with PsA). Serum drug and ADA levels were determined using Lisa-Tracker Duo, the ADA-positive samples underwent an inhibition test, and the true-positive samples underwent genetic HLA typing. To have a homogeneous control population, we also performed genetic HLA typing of 11 ADA-negative patients.

Results: After inhibition test, the frequency of ADAs was 2/26 patients treated with IFX (7.69%), 4/83 treated with ADA (4.81%), 0/88 treated with ETA (0%), 4/32 treated with CERT (12.5%), and 1/19 treated with GOL (5.26%). The frequency of HLA alleles in the examined patients was HLA-DR β -11 0.636, HLA-DQ-03 0.636, and HLA-DQ-05 0.727. The estimated relative risks between the ADA-positive patients and the ADA-negative patients were HLA-DR β -11 2.528 (95% CI 0.336–19.036), HLA-DQ-03 1.750 (95% CI 0.289–10.581), and HLA-DQ-05 2.424 (95% CI 0.308–15.449).

Conclusion: This is the first study that shows an association between HLA and genetic factors associated with the occurrence of ADAs in patients with rheumatic diseases, but the number of samples is too small to draw any definite conclusion.

Keywords: antidrug antibodies, HLA haplotypes, rheumatic diseases

Introduction

The introduction of anti-TNF drugs has benefited many patients with rheumatoid arthritis (RA), but, in some cases, the drugs trigger an immune response that leads to the development of antidrug antibodies (ADAs), one of the causes of a clinical non-response to treatment with infliximab (IFX) or adalimumab (ADA) that has also been described in patients with Crohn's disease.^{1,2} Many factors that can affect immunogenicity include disease activity, drug dose and dosing schedule, the route of administration, concomitant medications (including immunosuppressants), and genetic background.³

It is difficult to identify the genetic factors associated with the development of ADAs because of the large number of patients necessary for this type of study, and

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only one study of the influence of genetic variation on immunogenicity of anti-TNF agents has been published so far. This study found a correlation between the *IL10* genotype and the production of ADA ADAs, but the causality of this relationship requires further investigation.⁴ Drug-related factors can also influence the immunogenicity of a biological agent as differences in amino acid sequences between therapeutic and endogenous proteins promote the formation of ADAs, drugs with epitopes that bind to B or T cells have greater immunogenic potential, and the structural properties of a therapeutic protein can also increase the risk of an immune reaction.⁵⁻⁷

The aim of this study was to analyze the association between human leukocyte antigen (HLA) haplotypes and the development of ADAs in a cohort of patients with rheumatic diseases.

Patients and methods

We evaluated 248 Caucasian rheumatic patients after 6 months of treatment with anti-TNF drugs, of whom 26 were treated with IFX (three with RA, 13 with ankylosing spondylitis [AS], 10 with psoriatic arthritis [PsA]); 83 treated with ADA (24 with RA, 36 with AS, 23 with PsA); 88 treated with etanercept (ETA; 35 with RA, 27 with AS, 26 with PsA); 32 treated with certolizumab (CERT; 25 with RA, two with AS, five with PsA); and 19 treated with golimumab (GOL; three with RA, seven with AS, nine with PsA). Characteristics of patients at baseline with disease activity score for the respective three diseases and the concomitant use of Disease Modifying Anti-Rheumatic Drug (DMARD) are specified in Table 1. Serum drug and ADA levels were determined using Lisa-Tracker Duo (Theradiag, Croissy Beaubourg, France), the ADA-positive samples underwent an inhibition test, and the true-positive samples and 11 ADA-negative samples underwent genetic HLA typing.

All patients signed a written consent to be evaluated for the HLA genotyping. The consents are physically conserved in the Genetic Unit of the Hospital. All patients gave their written informed consent to this retrospective study according to the Declaration of Helsinki and to Italian legislation (Authorization of the Privacy Guarantor No. 9, December

12, 2013). The institutional review board, the Health Director of San Giovanni di Dio Hospital in Florence, reviewed and approved this study and the use of clinical and laboratory data of common clinical practice, with respect to privacy law, for clinical and scientific studies and publications.

Drug levels

Drug levels were determined using Lisa-Tracker Duo, an enzyme-linked immunoassay (ELISA) for the quantitative determination of drug levels in human serum samples. TNF- α was coated onto a polystyrene microtiter plate (six rows of eight wells), and the diluted sample was added to allow binding; after incubation, the unbound proteins were removed by washing. Subsequently, biotinylated antihuman IgG antibodies were added and, after incubation, the unbound antibodies were removed by washing. Horseradish peroxidase-labeled streptavidin was then added, and bound to the complex formed by the biotinylated anti-IgG antibodies. After incubation, the wells were washed once again to eliminate any excess conjugate, and the bound enzyme was revealed by adding a 3,3',5,5'-tetramethylbenzidine (TMB) substrate (the intensity of the color was proportional to the amount of drug). The enzymatic reaction was stopped by adding 0.25 of H₂SO₄, and the optical density was read by means of a spectrophotometer at 450 nm. A calibration range allowed the quantity of drug in each sample to be expressed in micrograms per milliliter.

ADAs

Lisa-Tracker Duo was also used to evaluate ADAs. Depending on the dosage, the drug was coated onto a polystyrene microtiter plate (six rows of eight wells), and the diluted sample was added to the antibody-coated wells to allow binding; after incubation, the unbound proteins were removed by washing. The biotinylated drug was then added and, after incubation, the unbound antibodies were removed by washing. Subsequently, horseradish peroxidase-labeled streptavidin was added and bound to the complexes formed by the biotinylated drugs. After incubation, the wells were washed again to eliminate any excess conjugate, and the bound enzyme was revealed

Table 1 Characteristics of patients at baseline

Disease	Gender (M/F)	Age (years)	Mean duration of disease (years)	DAS28	ASDAS	DAPSA	MTX dose (mg/week)	LEF dose (mg/day)	SSZ dose (mg/day)
RA, 90 patients	12/78	59±6	5±2	4.8±1.2	–	–	12.2±2.1	16.1±1.4	–
AS, 85 patients	74/11	54±4	4±2	–	1.7±0.2	–	–	–	1.5±0.4
PsA, 73 patients	33/40	47±7	3±2	4.2±0.8	–	3.8±0.6	10.2±1.2	15.2±1.3	1.6±0.3

Note: '–' indicates data not available.

Abbreviations: AS, ankylosing spondylitis; ASDAS, Ankylosing Spondylitis Disease Activity Score; DAPSA, Disease Activity in Psoriatic Arthritis; DAS28, Disease Activity Index 28; F, female; LEF, leflunomide; M, male; MTX, methotrexate; PsA, psoriatic arthritis; RA, rheumatoid arthritis; SSZ, sulfasalazine.

by adding a TMB substrate (the intensity of the color was proportional to the amount of ADAs). The enzymatic reaction was stopped by adding 0.25 of H₂SO₄, and the optical density was read by means of a spectrophotometer at 450 nm. A calibration range allowed the quantity of ADAs in each sample to be expressed in nanograms per milliliter.

ADA inhibition

An aliquot of each serum sample was diluted 1:2 in dilution and wash buffer (TDL), and in parallel, another aliquot of the same sample was diluted 1:2 in TDL with the addition of drug at a concentration of 100 µg/mL. All the diluted samples were incubated 1 hour at room temperature, and then vortexed. After this incubation, 100 µL of the standards, diluted positive controls, and diluted serum samples were added to the microplate wells and assayed using the Lisa-Tracker ADA kit as described earlier. After reading the optical density of each well at 450 nm, the percentage of inhibition in the samples in the presence of 100 µg/mL of drug was calculated as follows: inhibition (%) = 100 × 1 – (the result from the drug-spiked sample/result from the unspiked sample). A sample showing >50% inhibition can be considered truly positive for ADAs.

HLA genotyping

HLA-A, HLA-B, HLA-C, HLA-DRβ1, and HLA-DQβ1 were genotyped by means of the hybridization of sequence-specific oligonucleotide (SSO) probes immobilized on microspheres and amplified genomic DNA samples, followed by flow fluorescence intensity analysis according to the manufacturer's instructions (One Lambda, Canoga Park, CA, USA). In the validation study cohort, the HLA analysis was restricted to HLA-DRβ1*11 and HLA-DQ-03-05. The presence or absence of the haplotypes was determined using nested sequence-specific primer polymerase chain reaction (PCR)-SSP. In both cohorts, HLA-DRβ1*11 alleles were identified by means of the

group-specific sequencing of exon 2 of the HLA-DRβ1*11-positive samples according to the manufacturer's instructions. HLA was classified using serological techniques to broad antigens and splits, with the classic methodical National Institutes of Health (NIH) in microlymphocytotoxicity according to Terasaki and McClelland,⁸ with the trade plates, antigens for HLA-A, HLA-B, HLA-C, HLA-DR, and HLA-DQ. In 201 donors (50.2%), serological DR and DQ typing was confirmed using molecular biology techniques; HLA-DRB1, HLA-B3, HLA-B4, HLA-B5, and HLA-DQB were biologically typed using TaqI RFLP,⁹ or the locus amplification method specification of the second exon of class II and hybridization with a PCR-oligonucleotides specific sequence.¹⁰

Statistical analysis

The relative risk (RR) was calculated comparing the ADA-positive and ADA-negative sample populations by using Microsoft Excel for Windows, 2010 (Microsoft Corp., Redmond, Washington, USA). Then, by evaluating a 95% acceptable confidence, a confidence interval was calculated by deducting it using the standard error methodology.

Results

After 6 months of treatment, we observed ADAs in 2/26 patients treated with IFX (7.69%), 5/83 patients treated with ADA (6.02%), 5/88 patients treated with ETA (5.68%), 4/32 patients treated with CERT (12.5%), and 1/19 patients treated with GOL (5.26%). After inhibition testing, the frequency of ADAs was 2/26 patients treated with IFX (7.69%); 4/83 treated with ADA (4.81%); 0/88 treated with ETA (0%); 4/32 treated with CERT (12.5%); and 1/19 treated with GOL (5.26%) (Table 2). The frequency of HLA alleles in the 11 ADA-positive examined patients was HLA-DRβ-11 0.636, HLA-DQ-03 0.636, and HLA-DQ-05 0.727; the frequency of HLA alleles in the ADA-negative population was HLA-

Table 2 Characteristics of patients with positivity of ADAs

Patient	Age (years)	Disease	TNF-β inhibitor	DMARD	HLA genotype
1	71	PsA	CERT		A2 B13-51 DR07-11 DQb1 02-03
2	42	AS	IFX		A2-30 B18-27 C02-08 DR03-06 DQ02-05
3	49	AS	ADA		A32-32 B14-50 C03-18 DR03-11 DQ03-03
4	53	AS	ADA		A2-26 B27-38 C02-12 DR14-16 DQ 05-05
5	71	RA	CERT		A01-01 B08-15 C07-07 DR03-11 DQ01-03
6	78	AS	IFX		A01-02 B8-40 C05-07 DR14-14 DQ05-05
7	49	RA	ADA		A01-24 C35-37 C04-06 DR01-11 DQ03-05
8	74	RA	CERT	MTX	A11-24 B35-35 C04-04 DR11-16 DQ03-05
9	68	PsA	GOL		A28-30 B13-35 C06 DR07-19 DQ02-05
10	56	RA	ADA	MTX	A2-13 B35 C04-07 DR04-11 DQ03-05
11	67	RA	CERT	MTX	A1 B8-37 C02-12 DR03-11 DQ03-05

Abbreviations: ADA, adalimumab; ADAs, antidrug antibodies; AS, ankylosing spondylitis; CERT, certolizumab; DMARD, Disease Modifying Anti-Rheumatic Drug; GOL, golimumab; IFX, infliximab; MTX, methotrexate; PsA, psoriatic arthritis; RA, rheumatoid arthritis.

DR β -11 0.182, HLA-DQ-03 0.363, and HLA-DQ-05 0.454. The estimated RRs between the ADA-positive patients and the ADA-negative patients were 2.528 (95% CI 0.336–19.036), HLA-DQ-03 1.750 (95% CI 0.289–10.581), and HLA-DQ-05 2.424 (95% CI 0.308–15.449) (Table 3).

Discussion

This study shows an association between HLA and genetic factors associated with the occurrence of ADAs in patients with rheumatic diseases. An inadequate response to anti-TNF drugs in RA patients may be due to the development of ADAs.² Previous studies found a correlation between the *IL10* genotype and the production of ADA ADAs.⁴ IL-10, a cytokine that plays a key role in antibody development, is associated with the production of antibodies that inhibit recombinant factor VIII (FVIII) in patients with hemophilia¹¹ and autoantibodies against nicotinic acetylcholine receptors (nAChRs) in patients with myasthenia gravis (MG).¹²

Using high-resolution HLA class I and II typing, we identified three HLA class II alleles associated with the development of antibodies: HLA-DR β -11 (RR 2.528), HLA-DQ-03 (RR 1.750), and HLA-DQ-05 (RR 2.424) in comparison with the ADA-negative population. An association between HLA and the development of ADAs has also been found in other diseases, with HLA-DR β 1*04:01, HLA-DR β 1*04:08, and HLA-DR β 1*16:01 being identified as genetic markers of an increased risk of developing anti-interferon beta (IFN- β) antibodies.¹³ Three studies have shown an association between HLA and ADAs against IFN- β .^{13–15} Smoking is a risk factor for the development of multiple sclerosis (MS), and the risk is greater if smoking is combined with well-known HLA associations. It is also a risk factor for the development of ADAs in IFN- β - and natalizumab-treated MS patients,^{16,17} as well as for the development of RA by means of a mechanism of citrullination.

As in the case of other drugs, the development of ADAs against TNF- α -blocking monoclonal antibodies depends on many factors: environmental and genetic factors, disease

pathophysiology-related variations, and drug preparation and administration. ADAs appear soon after the start of the treatment,¹⁸ and their frequency varies considerably depending on the drug and assay method used.^{19,20} To the best of our knowledge, there are limited published data concerning the T cell epitopes involved in immune responses to anti-TNF agents, although a number of research groups are beginning to investigate this field. One study has identified a T cell epitope associated with the *IGHG1* G1m1 allotype that can induce T cell proliferation in healthy subjects with a different allotype.²¹ As ADA contains this T cell epitope, patients with the *IGHG1* G1m1 allotype may have a reduced propensity to produce ADA ADAs, but this view is contradicted by the results of a study showing increased ADA-induced immunogenicity in patients with this allotype.²²

Some anti-TNF agents may include multivalent antigens that can induce T cell-independent ADA formation as a result of extensive B cell receptor cross-linking (which causes B cell activation regardless of T cell help).²³ A number of studies have shown that concomitant immunosuppression reduces the induction of ADAs, which suggests that this treatment may be sufficient to ensure a good treatment response or at least delay the development of ADAs.^{24,25} As for FVIII treatment, there have been attempts to induce immunological tolerance through high doses of IFX, which is believed to exhaust the immune response.²⁶ The ADA frequencies observed in our study are similar to those reported in the literature.²⁷ The different frequency of the development of ADAs depends on the method used (ELISA, radioimmunoassay [RIA], antigen-binding test, pH-shift anti-idiotypic) the underlying disease (RA, PsA, inflammatory bowel disease, spondyloarthritis) and by the concomitant use of immunosuppressive treatment such as methotrexate.²⁸

It is known that the large individual variations in the treatment responses of patients with RA are influenced by factors such as gender, body mass index, and the degree of inflammation,²⁹ but this is the first study to show an association between HLA and genetic factors associated with the occurrence of ADAs in patients with rheumatic diseases. An important limitation of our study is that HLA-DRB1 and HLA-DQB1 association at the four-digit level is not presented. The number of samples is too small to draw any definite conclusion, and further comparative studies are now required to investigate rheumatic patients who do not develop ADAs vs healthy subjects, and non-ADA patients and healthy subjects.

Disclosure

The authors report no conflicts of interest in this work.

Table 3 HLA frequency in rheumatic patients with and without ADAs

Frequency of HLA-DR β 1-11	Frequency of HLA-DQ-03	Frequency of HLA-DQ-05
Patients with ADAs		
0.636	0.636	0.727
Patients without ADAs		
0.182	0.363	0.454
RR		
2.528	1.750	2.424
95% CI		
0.336–19.036	0.289–10.581	0.308–15.449

Abbreviations: ADAs, antidrug antibodies; RR, relative risk.

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