Original

Serious infections in patients with rheumatoid arthritis treated with tumor necrosis factor alpha antagonist therapy

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Abstract

Aim and objetives: the use of tumour necrosis factor alpha therapy antagonist leads to an increased risk of serious infections in rheumatoid arthritis, with a high morbi-mortality in clinical practice. There are no published data in our country. This study pretends to provide knowledge about the local epidemiology of serious infections and to identify associated risk factors.

Methods: An observational and retrospective study, included 50 patients treated at least for one year between 2006-2012. Demographic characteristics, clinical and epidemiological characteristics of serious infections and risk factors associated with serious infections were described.

Results: A predominant female population in middle age was detected. Just one serious infection was described, corresponding to serious skin and soft tissue infection that resolved with intravenous antibiotics. Risk factors were described including inmunosuppresive therapy with steroids and prior surgery.

Conclusions: Demographic profile of analized population and their clinic profile of serious infectios are similar to others populations. Low incidence of serious infections in this cohort could be related with less comorbilities than others populations, although, this topic needs further investigations to corroborate or discard this observation.

Keywords: rheumatoid arthritis, antiTNF therapy, serious infections, epidemiology, disease-modifying anti rheumatic drugs, prevalence.

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Affiliation of the authors: ¹Rheumatology Department, "Dr. Rafael Ángel Calderón Guardia" Hospital. 2Rheumatology Department, Mexico Hospital, Abbreviations: RA, rheumatoid arthritis; antiTNFa, tumor necrosis factor alpha antagonist; HLA, human leukocyte antigen; TNFa, tumor necrosis factor alpha; MMP, Matrix metalloproteinases; DMARDs. Disease modifying antirheumatic drugs; TB, tuberculosis; DM, diabetes mellitus; COPD, chronic obstructive pulmonary disease. Marsbcr@gmail.com

Rheumatoid arthritis (RA) is the most common autoimmune inflammatory arthropathy worldwide, affecting approximately 1% of the population.¹ In Latin America, the estimated prevalence is close to 0.5%,² Similar to that observed in Spain in 2001.³ In Brazil for 2013 it is estimated that it affected about 1% of the population.⁴ Unfortunately, there are no epidemiological studies in our country.

From the pathophysiological point of view, new pathways are often found involved in the immunologic cascade that characterizes RA. The role of some inflammatory mediators such as antibodies, growth factors, adhesion factors, cytokines such as tumor necrosis factor alpha (TNF α) and matrix metalloproteinase (MMP) have been defined.

TNFα plays a central role in the pathogenesis of RA.^{5,7} It is produced mainly by synovial macrophages and to a lesser extent by lymphocytes in response to proinflammatory stimuli;^{1,8,9} it's expressed as a transmembrane protein activated by a specific MMP (TNF converting enzyme), after which it is converted into a soluble

protein that is oligomerized to form a homotrimer constituting the active form. The effects of TNF α are mediated by two structurally distinct receptors: receptor 1 (TNF R-I) and receptor 2 (TNF R-II).^{8,10,11}

The binding of TNF α to the receptor activates several intracellular signaling pathways, including the activation of transcription factors such as the nuclear factor kappalight-chain-enhancer of activated B-cells (NF κ B), protein kinases such as MAP kinase and proteases such as caspases among others.⁹ In RA, TNF α induces the production of proinflammatory cytokines such as interleukin 1 (IL-1), interleukin 6 (IL-6), chemokines such as interleukin 8, and increases vascular permeability and expression of endothelial adhesion molecules among other effects.¹²

Given its chronic condition, RA requires continued use of various medications. The current therapeutic arsenal is made up of non-steroidal anti-inflammatory drugs, traditional disease modifying anti-rheumatic drugs (DMARDs), steroids, and in recent years biological disease-modifying antirheumatic drugs or simply biological have been introduced which include TNF α antagonists (antiTNF α), therapy anit-CD20 and anti-IL-6 therapy among others,^{1,13} fundamental in the treatment of RA in patients who do not respond clinically to treatment with traditional DMARDs.^{1,7,14}

AntiTNFα are drugs that have been shown in multiple controlled clinical studies to be very effective in the treatment of inflammatory diseases such as RA, juvenile idiopathic arthritis, Crohn's disease and psoriatic arthritis.^{5,8,15-19} In RA, they have been shown to delay the progression of joint damage,6,20,21 they improve symptoms and signs, as well as functional status and quality of life.^{6,22}

The therapeutic effects of antiTNFα in RA are mainly grouped into two major mechanisms: downregulates the local and systemic production of inflammatory cytokines; and decreases the activation and migration of lymphocytes at the joint level.^{23,24} This implies a decrease in the levels of IL-1, IL-6, MMP and in the expression of endothelial adhesion molecules.^{25,26} Previous studies have shown a decrease in the synovial macrophage population by induction of apoptosis mediated by the antiTNFα.⁸

In Costa Rica, the use of antiTNFα drugs started in 2005. Currently our social security system has three antiTNFα: Etanercept, Adalimumab and Infliximab for the management of various pathologies in specialties such as Rheumatology, Dermatology and Gastroenterology among others. In Rheumatology the antiTNF α are used for diseases including RA, Spondyloarthropathies and Adult Still's Disease among others.

From its clinical use it became apparent that the inhibition of TNFa is associated with an increase in the incidence of severe infectious processes.^{4,8,27,28} In 2001 the US Food and Drug Administration of the United States (FDA) issued a warning regarding infectious processes associated with antiTNFa.18 Subsequent studies showed an increased risk for severe infections by intracellular microorganisms mainly in the upper respiratory tract, lung and skin; due to pathogens such as tuberculosis (TB) and other mycobacteria, viruses, endemic fungi and some bacteria,^{5,18,19,27,29-32} Because TNFa plays a greater role in the defense against microorganisms (especially intracellular) preventing spread, by activating the formation of granulomas.^{8,13,16,30,32,33} Some of the mechanisms described are related to activation and differentiation of macrophages, as well as the stimulation to the formation of phagosomes.³⁴

Studies for several years have shown a higher incidence of severe infectious diseases requiring hospitalization in patients with RA versus patients without RA, causing significant morbidity and mortality.^{5,7,15,27,35,36} These are defined as those requiring hospitalization for management or for antibiotic therapy, which are potentially fatal and in addition, those that caused death.^{27,37,38} In general, severe infectious processes are considered to be related to immunological alterations of the disease, especially cellular immunity due to a decrease in the number and function of suppressor T lymphocytes and natural killer lymphocytes (NK lymphocytes),³⁶ to the drugs used for its control or a combination of both.^{5,13,27,30,34,37}

Some factors associated with an increased risk of severe infectious processes with antiTNFa therapy in RA patients are: steroid dose, first six months of anti-TNF treatment, recent previous surgery, lymphopenia and comorbidities such as Diabetes Mellitus (DM) and Chronic Obstructive Pulmonary Disease (COPD).^{4,16,18,27,29,38}

Costa Rica lacks clinical studies on patients treated with antiTNFa drugs, including the development of infectious and non-infectious complications due to its prescription. The purpose of the study was to characterize the demographics of the population using antiTNFa therapy from the "Dr. Rafael Angel Calderón Guardia" Hospital, as well as to determine the clinical behavior of the infectious diseases associated with antiTNFa, in order to contribute relevant information about this population and the associated infectious diseases in our country.

Methods

The present observational and retrospective study was performed at the "Dr. Rafael Ángel Calderón Guardia" Hospital of the Costa Rican Social Security, in Costa Rica. Previous authorization from the local bioethics committee (CLOBI-37-09-2014), the database of patients using antiTNFa therapy from the local pharmacotherapy committee of the "Dr. Rafael Ángel Calderón Guardia" Hospital was analyzed and it was compared with the list that in semiannual form was sent of the service of Rheumatology to said committee. From the list the population of patients with RA treated with antiTNFa drugs was identified and treated for at least one year, during the period between 2006 and 2012; the patients were excluded that had a diagnosis other than RA, as well as those with RA treated with antiTNFa intermittently, either because of a pregnancy, therapeutic failure or for a period of less than one year. The demographic characteristics of the population was described, as well as the clinical and epidemiological characteristics of the severe infections developed after the onset of antiTNFa and the risk factors associated with severe infection(s).

The clinical and microbiological data of the severe and non-severe infectious diseases were identified in the patients that fulfilled the inclusion criteria, but only those related to severe infectious diseases were analyzed. It was defined as a severe infection in the cohort, such as that characterized by requiring hospitalization for its management or prescription of antimicrobial therapy, potentially fatal or causing death.

Based on this information, the clinical records were reviewed and the following variables were studied: gender; current age in years; age at the time of diagnosis of RA in years; age at the time of initiation of antiTNFa in years; number of antiTNFa drugs used; current antiTNFa therapy, use of DMARDs or immunosuppressants at the start of antiTNFa therapy; positivity of rheumatoid factor; interval between initiation of antiTNFa therapy and the development of severe infection in months; location of severe infection; isolated microorganism(s); antimicrobial therapy(s) used; dose of antimicrobial(s) used; duration of antimicrobial therapy prescribed in days; comorbidities associated with initiation of antiTNFa therapy including: DM, lymphopenia less than 1500 / mm3, recent surgery in the last 12 months and COPD; patients with doses of oral steroids less than 10 mg/day at the start of antiTNFa therapy; patients with doses of oral steroids greater than 10mg/day at the start of antiTNF α therapy; presence of latent TB. The required data were

collected from the patients' clinical records using a "Data Collection Sheet" prepared for this purpose and approved by the Local Bioethics Committee.

For the analysis and interpretation of the obtained data, measures were used such as average, median, frequency among others. The SPSS program for Windows 14.0 was used as a statistical analysis tool.

Results

As of June 2014, a total of 117 people had received antiTNF α therapy for at least one year, of which only 51 patients were diagnosed with RA. We found 50 files that met the inclusion criteria, 45 corresponded to women and 5 to men. The average age of the 2014 cohort was 52.9 years (SD: ± 10.9); the minimum corresponds to 35 years and the maximum 78. The average age at the time of diagnosis of RA in the population was 38.6 years (SD: ± 10.9) and at the beginning of antiTNF α therapy, 47.9 years (SD: ± 10.6).

Regarding therapy at the time of analysis, 78% of the patients used etanercept and 22% adalimumab. The use of DMARDs or immunosuppressants at the start of antiTNF α therapy averaged 1.44, with methotrexate being the most prescribed. Other parameters related to the treatment are described in Table 1.

Table 1. Characteristics of treatment in patients with RA in biological therapy in the CGH from 2006 to 2012	
Characteristic	Value
AntiTNFα treatment: number (%) One antiTNFα received Two antiTNFα received	(92%) 4 (8%)
Current AntiTNFa treatment: number (%) Etanercept Adalimumab	39 (78%) 11 (22%)
Patients with DMARDs and/or immuno- suppressants at the start of antiTNFα: number (%) One DMARD and/or immunosuppressant Two DMARDs and/or immunosuppressant Three DMARDs and/or immunosuppressant Four DMARDs and/or immunosuppressant	30 (60%) 18 (36%) 2 (4%) 0 (0%)
DMARD type and/or immuno- suppressant prescribed at the start of antiTNFa: number (%) Methotrexate Leflunomide Hydroxychloroquine Cyclosporin A Gold salts Sulfasalazine Azathioprine	36 (72%) 16 (32%) 12 (24%) 3 (6%) 2 (4%) 1 (2%) 1 (2%)

The determination of the rheumatoid factor was found in 49 clinical files, of which 37 were seropositive (75%); the remaining percentage was seronegative.

In the population analyzed, a severe infection was documented, corresponding to a skin and soft tissue infection at the right thigh level. The episode occurred 8 months after the start of antiTNFa; no microbial isolation was achieved in the various cultures performed including blood culture and secretion culture. An adequate clinical response was obtained following intravenous antibiotic treatment with oxacillin at doses of 2 grams every 6 hours and Clindamycin at doses of 600 mg every 8 hours for a period of 7 days, reaching hospital discharge of the patient.

In the analyzed population, risk factors for the development of severe infections were found: 2 patients with DM, 4 patients with a history of recent surgery and 1 patient with lymphopenia at the start of antiTNF α therapy.

Regarding the prescription of steroids at the start of antiTNF α therapy, 8 patients had no prescription. Of the 42 patients on steroids, 24 had doses lower than 10 mg/day and 15 with oral doses equal or greater than 10 mg/day. In 3 cases, parenteral steroids were used (methylprednisolone was used at 250mg daily) for clinical control of the disease, for a period of 3 days. The mean oral steroid dose in the cohort was 6.92 mg/day.

Studies of latent TB were positive in 5 patients of the analyzed population. In all of the cases they received chemoprophylaxis from the Infectology department of the "Dr. Rafael Ángel Calderón Guardia" Hospital prior to the start of antiTNF α therapy.

Discussion

The demographic profile of the cohort analyzed is similar to other international studies, with a predominantly midlife female population. Some classic examples are the CORRONA (Consortium of Rheumatology Researches of North America), ReACT (Research in Active Rheumatoid Arthritis trial) and TEMPO (Trial of etanercept and methotrexate with radiographic Patient Outcomes) studies.³⁹⁻⁴²

The period between the onset of RA and the prescription of antiTNF α therapy in the population analyzed is similar to other cohorts with an average of about 10 years.³⁹⁻⁴² In our country, antiTNF α have been

prescribed since 2005, so that patients initially treated with this therapy, correspond to those with a long-standing disease with no initial clinical response to DMARDs prior to that year. The prescription of antiTNF α therapy is limited to patients with a persistent active disease based on the DAS28 (Disease Activity Index of 28 joints), CDAI (Clinical Disease Activity Index) or SDAI (Simple Disease Activity Index) scores, which do not respond to different choices of DMARDs, recommended in national and international institutional treatment guidelines⁴³, so that, to date the antiTNF α do not represent an option for patients with early RA in our social security system.

Similar to other biological records such as the french RATIO (Research Axed on Tolerance of Biotherapies), most patients only received one antiTNF α , in 4 cases it was necessary to change the initial therapy, either by primary or secondary failure. The antiTNF α drugs have shown through multiple studies their effectiveness in the clinical control of RA refractory to the various DMARDs, similar to the one described in this cohort.^{8,16,20,21} Unlike the french RATIO registry where Etanercept was the least prescribed antiTNF α , in our analysis it corresponds to the most prescribed therapy, which obeys reasons of an institutional nature.

Regarding the use of **DMARDs** or immunosuppressants, most patients used only one DMARD (especially methotrexate and less leflunomide) at the start of antiTNFa therapy, similarly to other records such as CORRONA.³⁹ Predominant monotherapy is a striking feature, since it is common in the country to prescribe various combination therapies, including Odell (methotrexate, hydroxychloroquine and sulfazalazine), or combinations of methotrexate with leflunomide among others for the management of patients with persistent active disease, prior to considering the use of biological therapy. The predominant monotherapy in the cohort ,could translate an inability of the various combination regimens of DMARD to achieve prolonged clinical remission or minimal clinical activity of RA, or in turn represent intolerance to the various side effects of these. Such an observation may require further investigation in order to determine possible causes. On the other hand, the use of immunosuppressants in the analyzed population, not considered as current therapeutic options in the various international guidelines for RA, such as gold salts and Cyclosporin A (some associated with adverse effects such as proteinuria, severe cytopenias or arterial hypertension),⁴³ because these therapies they were included in the institutional guidelines in previous years, before considering the use of biological therapy.

The immunological profile of patients is characterized by the positivity of rheumatoid factor in the majority. Currently the seropositivity is considered one of the factors that confer a worse prognosis for RA, as well as extra-articular involvement, presence of radiological erosions or frank functional limitation and the degree of clinical activity based on DAS, DAS28, among others; in which case an early aggressive therapy should be considered.⁴³

The clinical profile of the only severe infection documented consistent with the first reports of severe infectious processes related to antiTNF α , the analysis of the German record of biological RABBIT (Rheumatoid Arthritis: Beobachtung der Biologika Therapie) in 2005 and the English record BSRBR (British Society for Rheumatology Biologics Register) in 2006 significant increases were demonstrated for skin and soft tissue infections related to antiTNF α therapy.^{4,5,7}

Severe infection developed within 8 months of initiating antiTNF α therapy, which differs from that described in the literature. The maximum risk has been described in the first 6 months of treatment initiation.4,44 In 2007 Askling and colleagues showed that the risk of infection requiring hospitalization in patients exposed to drug antiTNF α decreases over time, 1.43 (95% confidence interval: 1.18 to 1.73) in the first year to 0.82 (95% confidence interval: 0.62 to 1.08) in the third.¹⁸

Based on the clinical response to the established intravenous antibiotic therapy, it is likely that the cause was a bacterial agent, despite not achieving microbial isolation in the cultures performed. In 2013 an analysis of the BSRBR English registry of skin and soft tissue infections in users of antiTNF α showed gram positive cocci as the main causative agents followed by pseudomonas species. Of 130 severe skin and soft tissue infectious processes, gram positive cocci were isolated in 103 patients, of which 84 corresponded to Staphylococcus aureus, 11 to Streptococcus sp. and 8 cases to negative coagulase Staphylococcus.³⁵

The presence of initiation of antiTNFα therapy of comorbidities such as DM, COPD, as well as lymphopenia and recent surgery have been associated with an increased risk of severe infectious events.^{4,15,16,27-29,34,38} In the present cohort, the prevalence of comorbidities is unusually low compared to studies from other latitudes. Those made in the United States, France and England have prevalences of DM in ranges varying from 6.0% to 18.9%, and COPD between 8.0% and 28.0%.^{16,17,29,33} This lower prevalence of comorbidities does not have a clear explanation and could be a cause of the low incidence of severe infections,

which is why later analyzes are required to corroborate or to discard such observation.

The use of steroids has been identified as a major risk factor in antiTNF α users,^{4,15,16} The percentage of patients receiving steroids (84%) is similar to that of studies such as RATIO, but higher than that reported in the US SABER (Safety Assessment of Biologic Therapy) registry, 60.1%. However, there is a significant difference in relation to high doses of steroids, whereas in the RATIO study up to 50% received oral doses greater than 10mg/day or intravenous boluses and in the SABER study 10.2%, our cohort in 36% received considerable high oral and parenteral steroidal doses.¹⁷

Several studies have demonstrated the importance of TNF α in TB control. In a retrospective cohort study of 112300 Canadian patients with RA, the overall TB rate was 2.2 cases per 1000 people/year (95% confidence interval 2.0 to 2.4). The incidence among patients with antiTNF α therapy was 2.6 per 1,000 people/year (95% confidence interval 1.9 to 3.3).³⁰ In the present study, 5 patients were identified by studies with positive latent TB prior to initiation of antiTNF α therapy. The figure is similar to other Latin American reports, a small Colombian cohort showed a high prevalence of latent TB, higher than that reported in cohorts from other latitudes.^{45,46}

From this investigation it is concluded that the demographic profile of the population using antiTNFa therapy is similar to that described in international studies, as well as the clinical and epidemiological profile of severe infections. The low incidence of severe infections could be related to a lower prevalence of comorbidities in the analyzed cohort, however future prospective national clinical-epidemiological studies are necessary to corroborate or rule out such observation.

The study presents some limitations including a small population to be analyzed, compared to classic RA studies including thousands of patients.

Other epidemiological variables in the analysis were not taken into consideration that could have enriched the results and the discussion, including the presence of sequelae secondary to RA, presence of anemic syndrome or thrombocytosis secondary to the chronic uncontrolled inflammatory process, elevation of the acute phase reactants, decrease in DAS28 after the prescription of antiTNFa therapy, among others, so it is considered a significant weakness not to have obtained a broader epidemiological characterization.

Nor was the primary or secondary failure to antiTNF α therapy adequately characterized, which might

have enriched the analysis, in attempting to establish possible causes. Because of the large difference between the number of patients treated with various antiTNF α therapies, it is impossible to make comparisons.

An important point that limited a broader analysis is the absence of measurements in the clinical file such as the ACR20, ACR50 and ACR70 responses routinely used internationally in clinical trials both to assess the clinical response to antiTNF α therapy as well as therapies with DMARDs.

Preferences

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