Association of Antigen-Stimulated Release of Tumor Necrosis Factor-Alpha in Whole Blood with Response to Chemotherapy in Patients with Pulmonary Multidrug-Resistant Tuberculosis

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Key Words
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Abstract

Background: We have previously reported that TNF-\alpha levels correlate to total mycobacterial burden in tuberculosis (TB) patients. Objective: To characterize the dynamics of cytokine responses in TB patients during chemotherapy to identify potential surrogate markers for effective treatment. Methods: Following induction by culture filtrate proteins in whole blood, production patterns of TNF-\alpha, IL-10, IFN-\gamma and IL-12 were measured in 23 non-multidrug-resistant (MDR)-TB and 16 MDR-TB patients and in 31 healthy controls. Rates of mycobacterial clearance from the sputum were then measured and compared. Results: Prior to the initiation of chemotherapy, TNF-\alpha and IL-10 levels were significantly higher in TB patients than in healthy controls while IFN-\gamma and IL-12 levels were similar. During chemotherapy, the levels of all 4 cytokines increased. We evaluated these responses separately in patients that did and did not clear their sputum culture at 2 and 6 months. At 2 months, decreases in both IFN-\gamma and IL-12 correlated strongly with a successful early response, while after 6 months of therapy, when half (7/14) of MDR-TB patients were still sputum culture positive, downregulation of TNF-\alpha was uniquely correlated with sputum conversion between the groups. Conclusion: Our findings suggest the possibility that the regulation of TNF-\alpha production in whole blood may be a more specific indicator of sputum conversion at 6 months than IFN-\gamma, IL-12 or IL-10 in MDR-TB patients.

Introduction

\textit{Mycobacterium tuberculosis} (Mtb) is the causative agent of pulmonary tuberculosis (TB). Currently, there are two major problems facing the medical community...
concerning the fight against Mtb. First, the increasing emergence of multidrug-resistant (MDR)-TB has rendered common therapies against the bacteria ineffective. Second, the current testing methodologies for the diagnosis of TB are extremely slow. Because Mtb is a slow-growing organism, it can take up to 8 weeks for an accurate Mtb diagnosis to be made. As a result, there is an increasing need for both novel therapies and rapid diagnostic methods.

After Mtb infection, healthy individuals develop a cellular immune response, which arrests the growth of the microorganism, preventing progression to clinical TB [1, 2]. However, susceptible individuals develop pulmonary TB. Upon inhalation of aerosolized Mtb, the pathogen is engulfed by macrophages. Once inside the phagosome, virulent Mtb inhibits the acidification of the compartment [3]. This inhibition allows the bacilli to reside within the macrophage, sequestered away from the bactericidal and antigen-generating machinery of the cell. As a result, Mtb establishes a safe and long-term resting area for replication, which eventually results in cell lysis and bacterial spread.

Upon stimulation by a pathogen, macrophages engulf the offending particle, and upon its destruction, they present smaller peptide antigens on their surface. These antigens are then recognized by Th1 cells, which in turn secrete various cytokines, including IFN-γ, IL-12 and TNF-α. These cytokines in turn activate resting macrophages, which trigger the immune response. In order to suppress the immune response once an infection has been cleared, and to prevent autoimmune responses to self-antigens, other cytokines downregulate the immune system. IL-10 activates B and Th2 cells while inhibiting Th1 cytokine production. Activated Th2 cells secrete IL-4, which also inhibits Th1 cell activity and inactivates macrophages.

Previously, it has been reported that Mtb stimulation of peripheral blood mononuclear cells (PBMCs) isolated from TB patients resulted in a diminished production and expression of Th1 cytokines, including IFN-γ and IL-2 [4]. Additionally, this same report demonstrated that after 9 months of drug therapy, IFN-γ levels were markedly increased, coinciding with successful treatment. As a result, it has been suggested that different clinical states of Mtb can be correlated with cytokine profiles.

In an effort to reduce the amount of time required to accurately diagnose an Mtb infection, some attention is being focused on identifying biomarkers of Mtb infections; examples include molecules that are secreted by the pathogen itself, or host factors derived from immune responses. In this study, we build upon our previous findings that TNF-α, IFN-γ and IL-10 profiles are distinct in non-MDR-TB versus MDR-TB patients [5]. Using sputum clearance as an indicator of positive responses to chemotherapies, we report that patients on anti-Mtb chemotherapy have altered TNF-α and IFN-γ levels that correlate with favorable responses to the drugs. As a result, we propose that these cytokines, particularly TNF-α, could serve as potential biomarkers for response to therapy.

**Patients and Methods**

**Study Subjects**

A total of 39 patients (35 men and 4 women) who were diagnosed with active TB by clinical findings, radiological examination and the demonstration of acid-fast bacilli (AFB) in their sputum at the National Masan TB Hospital (NMTH) were en-
Sputum Culture and Drug Susceptibility Test

Sputum culture positivity for MtB was confirmed both by BacT/Alert and conventional Ogawa media. Briefly, sputum specimens were decontaminated with N-acetyl-l-cysteine/4% NaOH and inoculated into BacT/Alert and Ogawa media. When the Ogawa growth was detected as positive, a drug susceptibility test was carried out on Löwenstein-Jensen plates containing anti-TB drugs. The ‘absolute concentration method’ was used as previously described [6, 7]. The critical concentration for isoniazid and rifampin was 0.2 and 40 μg/ml, respectively. All inoculated slants were incubated at 37°C and growth was checked visually every week. The sputum culture staining for AFB was assessed at the time of admission, and 2 and 6 months after starting proper treatment.

Blood Collection and Cytokine Whole-Blood Assay

Blood samples were drawn into heparinized Vacutainers and processed within 2 h for the cytokine whole-blood assay [5, 8]. Blood was collected at the time of admission, and 2 and 6 months after the start of treatment prescribed by NMTH staff. Blood diluted with RPMI-1640 (1:10) containing penicillin-streptomycin (Gibco) and 10% fetal bovine serum was distributed into 96-well plates and cultured in triplicate in the presence of medium alone or with phytolam gamburtin (PHA) or lipopolysaccharide (LPS; Sigma-Aldrich, St. Louis, Mo., USA) or culture filtrate protein (CFP) at a final concentration of 10 μg/ml. Cultures were maintained in a 5% CO₂ atmosphere at 37°C. Cultured supernatants were harvested 1, 3 and 6 days after culturing, and stored at −70°C until use. Cytokines were measured at day 1 for IL-12 and IL-10, day 3 for TNF-α and day 6 for IFN-γ (optimal time points for each cytokine were determined by preliminary experiments, data not shown).

Measurement of Cytokines by ELISA

For cytokine analysis, two-site capture immunoassays with OptEIA monoclonal antibody sets (BD Pharmingen, Los Angeles, Calif., USA), streptavidin-horseradish peroxidase conjugate and recombinant cytokines as standards were used. Briefly, 96-well plates were coated with capture antibodies to each cytokine (IFN-γ, IL-12, TNF-α and IL-10) according to the manufacturer’s protocol. The concentrations were given in the ELISA set instructions as dilution factor, 1:250. Plates were incubated overnight at 4°C and blocked at room temperature with 5% FBS in PBS for 1 h. The plates were washed three times. Samples were added and incubated for 2 h at room temperature. Working detector (detection antibody + avidin-HRP reagent) was added to each well and the plate was incubated for 1 h at room temperature. After washing 7 times, substrate solution (BD Pharmingen TMB substrate reagent set) was added to each sample. The plates were incubated for 30 min and the reaction was stopped by adding stop solution. The plates were read at 450 nm in an ELISA reader (OPTImax, Sunnyvale, Calif., USA). A standard curve was generated from the known concentration of the standard according to the method provided in the kit. Cytokine levels were calculated from the standard curve by computing absorbance values against the test concentration. Detection limits of the cytokine kits used were as follows: IFN-γ – 5 pg/ml; IL-12 – 7.5 pg/ml; TNF-α – 15 pg/ml, and IL-10 – 7.5 pg/ml.

Statistical Analysis

The Mann-Whitney U test was used for data analysis between groups. Differences were considered significant when p ≤ 0.05.

Results

Patient Description and Response to Chemotherapy

Thirty-nine subjects were enrolled in this study (table 1). All subjects were examined for clinical symptoms and underwent standard radiographic assessment. The mean age of the patients was 49.4 ± 2 years (range, 22–87 years). They included 14 newly diagnosed, 6 relapse, 7 chronic, 6 treatment failure, and 6 default cases as defined by the World Health Organization [9]. One case of pleurisy was observed and 11 subjects had concomitant diabetes mellitus, and 1 had hypertension. None of the patients were HIV positive. At the time of initial blood sampling, the patients had been started on chemotherapy for <1 week. Thirty-one Microbacterium bovis BCG-vaccinated, healthy individuals (15 men and 16 women) also participated as controls, and their mean age was 34.6 ± 1.5 years (range, 23–54 years). Of the 39 TB patients, 10 were culture-confirmed non-MDR-TB while 16 were culture-confirmed MDR-TB patients. Thirteen of the initial patients were sputum culture negative at the start of the therapy, and remained so throughout the 6-month study. Although they were culture negative, 12 of 13 of these patients experienced significant clinical improvement with normal first-line chemotherapy during the course of this study while 1 patient did not. For that reason, this non-responsive patient was omitted from further analysis. In total, therefore, 38 patients were considered to have diagnosed as TB and were included in subsequent analysis: 22 had drug-susceptible disease and 16 MDR disease.

Of the 16 sputum-positive drug-susceptible subjects, 9 had achieved sputum conversion after the first 2 months of therapy, and the last subject achieved sputum conversion after 6 months. In total, 100% of the non-MDR-TB subjects were sputum negative at the end of the therapy. Of the 16 MDR-TB subjects, only 2 subjects had achieved sputum conversion after 2 months. After 6 months of treatment, an additional 6 subjects (8 in total) had converted. In total, 50% of the MDR-TB patients had con-
verted to sputum negative by the conclusion of the study (table 1).

**Whole-Blood Cytokine Profiles during Chemotherapy**

To explore whether changes in cytokine responses before, during and after 6 months of drug therapy would provide a surrogate marker for sputum smear and culture, we collected whole blood from healthy volunteers \((n = 31)\) and our cohort of TB patients. This blood was stimulated with CFP from Mtb and 4 cytokines were measured: IFN-\(\gamma\), TNF-\(\alpha\), IL-10 and IL-12 (fig. 1).

At the start of treatment, there was no significant difference in the IFN-\(\gamma\) and IL-12 levels between TB patients and healthy controls. IFN-\(\gamma\) levels increased with therapy by 2 months \((p \leq 0.05)\) and reached a plateau almost threefold higher compared to levels before treatment by 6 months of therapy. IL-12 levels similarly increased after 2 months of therapy \((p \leq 0.01)\) but decreased to levels before therapy by 6 months (fig. 1). In contrast, significantly higher levels of TNF-\(\alpha\) and IL-10 were observed in TB patients than healthy controls at the start of the treatment, and these cytokine levels also increased during...
drug treatment (fig. 1). In general, all 4 cytokines showed a complex dynamic response that appeared to fall into one of two general trends: either a transient increase at 2 months followed by a return to baseline levels at 6 months, or an increase at 2 months followed by a further increase at 6 months.

Cytokine Response Pattern Does Not Correlate with MDR Status But Is Mtb Specific

In an attempt to understand whether these two patterns of response were the result of previous treatment history or were specific to strains of Mtb that had drug resistance, we divided the patients into non-MDR- and MDR-TB groups and examined patterns of cytokine production between these groups, but no significant differences were observed (fig. 2). Likewise, when comparing sputum culture-negative subjects that showed clinical improvement to sputum culture-positive patients, no differences in responses of cytokine profiles were observed (data not shown).

In order to verify that the cytokine responses seen were in fact specific to the Mtb antigens present in CFP, we stimulated the same blood samples with non-TB-related mitogens to activate the aforementioned cytokines. For these purposes, PHA was used to stimulate IFN-γ and IL-12, while LPS was used for TNF-α and IL-12 stimulation (fig. 3). No changes in IFN-γ levels were observed between these groups during therapy. Similar results were obtained with TNF-α and IL-12 when comparing CFP and mitogen stimulation. Interestingly, PHA-in-
duced IL-10 production showed a reverse correlation with PHA induction (fig. 3).

**Effect of Sputum Clearance Rate on Cytokine Production Patterns**

In an effort to see if the cytokine production patterns correlated with sputum conversion, we obtained sputum cultures from each subject at the time of blood drawing for cytokine assays and used these results to identify the sputum-positive or sputum-negative status at each time point (fig. 4). Prior to the start of therapy, IFN-γ levels were overall lower in the 12 sputum culture-negative subjects compared to the culture-positive subjects, a trend that continued when comparing groups of patients who had achieved a culture-negative status compared to those who remained culture positive at each time point throughout the study. At the end of the treatment, those who remained culture positive had dramatically increased amounts of IFN-γ compared to levels both before the therapy and after 2 months even compared to those who had converted at 6 months. A similar pattern to IFN-γ levels was seen with IL-12. Lower levels of IL-12 were released in the patients who had converted to culture negative than those who failed to at every time point. This result was statistically significant \( p < 0.05 \) after 6 months of therapy (fig. 4).

TNF-α and IL-10 showed somewhat different patterns than IFN-γ and IL-12 in this cohort. Prior to therapy, sputum-positive and sputum-negative subjects had similar levels of TNF-α and IL-12. Although statistically sim-
ilar, both groups doubled their levels at 2 months. However, a dramatic change in TNF-α was seen at 6 months, when the sputum-negative population returned to cytokine levels before the therapy while remaining elevated in the sputum-positive population (p < 0.001). Strikingly, IL-10, a cytokine which is involved in helping to turn off the immune response, was relatively unchanged by the presence or absence of Mtb in the sputum of subjects before, during or after the therapy.

### Discussion

Cytokine production patterns in TB patients are presumed to vary according to the disease state, and we have previously reported that TNF-α was associated with MDR-TB and mycobacterial load in sputum [5]. However, our previous study was cross-sectional, limiting our ability to evaluate these correlations as markers to potentially predict outcome. In this longitudinal study, we have extended this observation by comparing mycobacterial

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**Fig. 4.** Comparison of cytokine production patterns between culture-positive (black) and -negative (empty) patients. Sputum cultures were performed at each time point and cytokine production was compared between culture-positive (black) and -negative (empty) patients. Only those patients who were sputum culture positive at the previous time point were reevaluated at the next time point of therapy. Therefore, only 26 sputum-positive patients were evaluated at 2 months and only 15 sputum-positive patients were evaluated at 6 months. Each value was calculated as the mean of triplicate cultures after deducting medium alone value. a p ≤ 0.05 and c p ≤ 0.001.
clearance during chemotherapy with TNF-α levels produced by stimulated whole blood in the same patients over time.

Prior to starting the therapy, TNF-α and IL-10 levels were significantly higher in TB patients than in healthy controls. These results are in agreement with other published results for IL-10 [10, 11] and for TNF-α [12–14]. In contrast, IFN-γ and IL-12 levels were not different between healthy controls and TB patients. Many other studies report that the production of IFN-γ is depressed in TB patients compared to healthy individuals [15–17]. However, conflicting reports have demonstrated no differences in IFN-γ mRNA expression and the number of IFN-γ-producing cells between TB patients and healthy controls [18, 19]. This discrepancy may be explained by recent studies reporting that different disease states could induce different IFN-γ responses [17, 20, 21]. In our patient cohort, during anti-TB chemotherapy, the levels of IFN-γ increased significantly after 2 months and much more after 6 months in TB patients, consistent with other reports [4, 8]. IL-12 also increased after 2 months of therapy, but after 6 months of therapy decreased again to levels before therapy, whereas IFN-γ showed the highest production levels at 6 months. Although IL-12 is known to enhance IFN-γ production [2, 22], this observation suggests IFN-γ production is not dependent on IL-12 regulation, as already shown in another study [23, 24]. The decrease in IL-12 production after 6 months of therapy may be related to the increase in the anti-inflammatory cytokine IL-10. IL-10 is known to be produced by macrophages and T cells during Mtb infection and possesses macrophage-deactivating properties including downregulating IL-12 production [25]. Because IL-10 acts as an antagonist of TNF-α and IFN-γ, it appears contradictory that IFN-γ and TNF-α would increase concomitantly with IL-10. However, it is plausible to consider that IL-10 is functioning solely to downregulate the immune system. As drug treatment clears Mtb from the host, the need for IFN-γ and TNF-α production decreases, an event regulated by IL-10. It is also in agreement with the drop in IL-12 to levels before the treatment, further indicating a reduction in the immune response.

IFN-γ is also involved in stimulating TNF-α secretion [26]. This study shows the induction of TNF-α increased as IFN-γ production increased during the course of chemotherapy. Mitogen-induced cytokines showed different profiles from CFP-induced cytokines. No changes in levels were observed in PHA-induced IFN-γ production with therapy but LPS-induced IL-12 and TNF-α production levels correlated with those of CFP-induced levels. In contrast, PHA-induced IL-10 showed an inverse correlation with CFP-induced IL-10 production. The molecular and kinetic basis for this difference in response is not clear, however, successful therapy may have repressed the production of this immunosuppressive cytokine by non-specific stimulation. Our findings suggest that chemotherapy may alter the specific cytokine production profiles.

In order to investigate if mycobacterial clearance induced by chemotherapy is mirrored by alterations in the cytokine response, serial sputum culture and cytokine measurements during therapy were examined. Almost all non-MDR-TB patients (9/10) who were initially sputum culture positive became sputum culture negative after the first 2 months of therapy, whereas only 2 MDR-TB patients obtained sputum negativity. At this time point, both IFN-γ and IL-12 production were decreased more in sputum-negative patients than in sputum-positive patients, indicating the normalization of the production of these 2 cytokines. However, no differences between patient groups were observed in TNF-α and IL-10 levels. By 6 months after the therapy, 6 further MDR-TB patients had become sputum culture negative, and IFN-γ and IL-12 production also decreased in these sputum-negative patients compared to the persistently sputum-positive patients. Notably, IL-10 levels remained unchanged between the two groups. However, there was a marked reduction in TNF-α to levels similar to those seen in the group of sputum-negative patients before the therapy. These findings suggest the potential role of TNF-α as a marker for successful therapy in MDR-TB.

It has already been demonstrated that some MDR-TB patients produce lower levels of TNF-α but not IFN-γ compared with non-MDR-TB patients in PBMCs [27] or in whole blood [5]. Also, antigenic stimulation of peripheral blood monocytes from MDR-TB induces different trends of TNF-α and IL-12 production compared to those from non-MDR-TB patients [27]. The production of nitric oxide from peripheral blood monocytes was depressed in MDR-TB patients and correlated with TNF-α production [28]. Conversely, sputum TNF-α levels decrease following initiation of treatment [29] and a positive correlation was observed between mycobacterial load and TNF-α levels in sputum by 12 weeks of treatment [29]. While this may reflect variation in the patient populations studied it is worth at least considering that this may reflect differential antigenicity of the TB strains that are more prone to develop into MDR disease. It is worth pointing out that at least one virulence factor of Beijing strains (the phenolic glycolipid) has been shown to have potent immunomodulatory properties [30].
TNF-α appears to be the most dynamic of the 4 cytokines investigated here, a finding that should support further evaluation of the response of this cytokine in TB patients. Our results suggest CFP-induced TNF-α levels in whole blood are closely related to the clearance of Mtb in MDR-TB patients independent of IL-10 regulation and could be a useful indicator for monitoring response to therapy in MDR-TB patients. However, the sample size in the present study is too small to make a definitive conclusion and this hypothesis will have to be validated in larger patient cohorts. These studies also suggest the possibility that an immune signature that correlates with sputum conversion may exist that would provide an early indicator for discontinuation of therapy, a significant problem for MDR-TB where treatment courses often exceed 2 years.

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