# Autoantibodies to TNF $\alpha$ in HIV-1 infection: prospects for anti-cytokine vaccine therapy

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**Summary** – Tumor necrosis factor alpha (TNF $\alpha$ ) is a proinflammatory cytokine principally involved in the activation of lymphocytes in response to viral infection. TNF $\alpha$  also stimulates the production of other cytokines, activates NK cells and potentiates cell death and/or lysis in certain models of viral infection. Although TNF $\alpha$  might be expected to be a protective component of an antiviral immune response, several lines of evidence suggest that  $TNF\alpha$  and other virally-induced cytokines actually may contribute to the pathogenesis of HIV infection. Based on the activation of HIV replication in response to TNF $\alpha$ , HIV appears to have evolved to take advantage of host cytokine activation pathways. Antibodies to TNF $\alpha$  are present in the serum of normal individuals as well as in certain autoimmune disorders, and may modulate disease progression in the setting of HIV infection. We examined TNF $\alpha$ -specific antibodies in HIV-infected non-progressors and healthy seronegatives; anti-TNF $\alpha$  antibody levels are significantly higher in GRIV seropositive slow/non-progressors (N = 120, mean = 0.24), compared to seronegative controls (N = 12, mean = 0.11). TNF $\alpha$  antibodies correlated positively with viral load, (P = 0.013, r = 0.282), and CD8+ cell count (P = 0.03, r = 0.258), and inversely with CD4+ cell count (P = 0.003, r = -0.246), percent CD4+ cells (P = 0.008, r = -0.306), and CD4 :CD8 ratio (P = 0.033, r = -0.251). TNF $\alpha$  antibodies also correlated positively with antibodies to peptides corresponding to the CD4 binding site of gp160 (P = 0.001, r = 0.384), the CD4 identity region (P = 0.016, r = 0.29), the V3 loop (P = 0.005, r = 0.34), and the amino terminus of Tat (P = 0.001, r = 0.395); TNF $\alpha$ antibodies also correlated positively with antibodies to Nef protein (P = 0.008, r = 0.302). The production of anti-TNF $\alpha$  antibodies appears to be an adaptive response to HIV infection and suggests the potential utility of modified cytokine vaccines in the treatment of HIV infections as well as AIDS-related and unrelated autoimmune and CNS disorders. © 2001 Éditions scientifiques et médicales Elsevier SAS

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In targeting immune cells for viral infection, HIV has evolved regulatory mechanisms that permit rapid deployment in response to immune activation signals. The same pathways that induce T cell proliferation and macrophage activation are usurped by HIV to activate viral replication [1-6]. TNF $\alpha$  is a potent proinflammatory cytokine integral to antiviral immune responses, which induces apoptosis of virally-infected cells and contributes to cytotoxic T cell-mediated lysis and lymphocyte and monocyte differentiation [7]. Interaction of TNF $\alpha$  with cell surface receptors can transduce a variety of intracellular signals via phospholipase A2, protein kinase C

\*Correspondence and reprints. E-mail address: jayrapp@astro.temple.edu (J. Rappaport). and activation of several transcription factors including NFKB [8]. It is well established that the major enhancer element within the HIV-1 LTR promoter is comprised of two NFKB binding sites responsive to T cell activation and cytokine activation signals [1-6]. These same transcriptional elements are involved in IL-2 and IL-2R gene expression [3-6]. TNFa production results in coordinate activation of certain inflammatory cytokines, including chemokines, via autocrine and paracrine mechanisms [9-12]. TNF $\alpha$ production results in alteration of endothelial cell monocyte/macrophage interactions, leading to invasion of inflammatory cells into various tissues including the central nervous system [12-14]. This process appears to be a critical pathway in the development of HIV-associated neurologic disorders, with increased accumulation of activated monocytes/ macrophages in the CNS compartment.

It is well established that healthy individuals as well as patients with autoimmune disorders produce natural autoantibodies against TNF $\alpha$  [15]. Anti-cytokine autoantibodies (i.e. anti-TNF $\alpha$  and interferon- $\gamma$ autoantibodies) appear to play a regulatory role in the immune response [16], and may serve to counteract the potential deleterious effects of these inflammatory mediators in vivo. In order to evaluate the significance of anti-TNF $\alpha$  autoantibodies in HIV infection, we produced recombinant human TNF $\alpha$ as a fusion protein, assayed serum samples from HIV infected slow/non-progressors and seronegatives in ELISA, and correlated levels of antibodies to TNF $\alpha$ with serologic and hematologic evaluation parameters.

# MATERIALS AND METHODS

# Production of the recombinant tumor necrosis factor $\alpha$

The cDNA of the mature hTNF $\alpha$  was obtained by PCR amplification from the plasmid pORF-hTNF $\alpha$  (Invivogen, San Diego, CA), cloned in the prokaryote expression vector pRSET-A (Invitrogen, San Diego, CA), and confirmed by automated DNA sequencing. This construct permits expression of TNF $\alpha$  as a fusion-protein with an N-terminal polyhistidine-tag, thereby permitting purification by metal chelate affinity chromatography. *Figure 1* illustrates the amino acid sequence of the hTNF $\alpha$  fusion-protein. <sup>1</sup>MRGSHHHHHHGMASMTGGQQMGRDLYDDDDKDRW
<sup>35</sup>GSELEVRSSSRTPSDKPVAHVVANPQAEGQLQWLN
<sup>69</sup>RRANALLANGVELRDNQLVVPSEGLYLIYSQVLFK
<sup>105</sup>GQGCPSTHVLLTHTISRIAVSYQTKVNLLSAIKSPC
<sup>140</sup>QRETPEGAEAKPWYEPIYLGGVFQLEKGDRLSAEI
<sup>175</sup>1<sup>76</sup>NRPDYLDFAESGQVYFGIIAL

## Molecular weight: 22 kDa Estimated pI: 6.14

**Figure 1**. Amino acid sequence of the hTNF $\alpha_r$  fusion protein. The fusion domain containing the polyhistidine tag is in italics.

The plasmid pRSETA-TNF $\alpha$  was transformed into E. coli (BL21-DE3 CodonPlus, Stratagene, San Diego, CA) for protein expression. Bacteria were grown in one liter of 2X LB media. TNFa fusion protein expression was induced at O.D. 0.6-0.8 with 1 mM IPTG (Calbiochem, San Diego, CA), and grown overnight at 37° C and 250 rpm. The culture (O.D. 1.4) was collected by centrifugation. Pellets were resuspended in 50 mL of Buffer A (300 mM NaCl, 50 mM Hepes, pH 7.2, 0.025% Octyl-β-Glucoside and CHAPS). The cells were disrupted by sonication using a Branson Sonifier (3X) for 30 s with 5 min on ice between sonication intervals. Homogenate was centrifuged at 14,000 rpm (30,000 g) for 15 min; supernatant was applied to a 5 mL Talon<sup>®</sup> Metal Affinity Resin column (Clontech, Palo Alto, CA), and packed in an XK16/20 column (Amersham-Pharmacia, Piscataway, NJ). The column was washed with 10 column volumes of Buffer A, followed by 4 column volumes of Buffer A containing 10 mM imidazole, and 6 column volumes of Buffer A containing 50 mM imidazole. Elution was performed with 5 column volumes of Buffer A containing 500 mM imidazole. Chromatography was performed throughout at a flow rate of 1 mL/min using an Acuflow Series III pump (LabAlliance, State College, PA). Affinity purified TNFa was collected in 20 mL fractions and protein containing fractions were pooled. The sample (15 mL; 2.6 mg/mL) was diafiltered under nitrogen with five volumes of Buffer B (300 mM NaCl, 50 mM Hepes, pH 7.2) using an Amicon 8050 stirred cell with a 10,000 NML 45 mm polyethersulfone membrane (Millipore, Bedford, MA) pretreated with 5% polyethylene glycol reacted with Bisphenol A (Sigma-Aldrich, Steinheim, Germany). The final preparation (30 mL; 0.8 mg/mL.) was stored in aliquots at  $-80^{\circ}$  C.

#### **Patient samples**

HIV-1-infected Serum from non-progressors (N = 120) were compared with samples from seronegative controls (N = 12) for autoantibodies to TNF $\alpha$ . Slow/non-progressor (S/NP) volunteers were previously enrolled in the genetic resistance to immunodeficiency virus (GRIV) cohort to study genetic and serologic parameters associated with nonprogression. This cohort, established in France, has been useful in identifying genetic polymorphisms and serologic responses associated with non-progression in the setting of HIV infection [17-22]. The inclusion criteria for S/NPs were as follows: seropositive for HIV-1 and asymptomatic for more than eight years, with a CD4+ cell count greater than 500/mm<sup>3</sup> without prior antiretroviral therapy. All seropositives in this group were Caucasians for the purpose of simplifying genetic analysis. Serum samples from HIVnegative individuals were collected at MCP-Hahnemann University (Philadelphia, PA) with informed consent. All seronegatives used in this study were Caucasians for comparison with the GRIV S/NP samples.

#### ELISA assay for TNFα autoantibodies

The recombinant human TNF $\alpha$ , 0.5 µg per well diluted in 100 µL of 50 mM NaHCO<sub>3</sub>, pH 9.0, was immmobilized in Maxisorb<sup>TM</sup> microtiter plates (Nunc, Rochester, NY) by overnight incubation at 4° C. Plates were blocked for three hours at room temperature with 250 µL/well 1X PBS containing 3% IgG-free BSA (Sigma, St. Louis, MO). Plates were washed 6X in 1X PBS with 0.05% Tween 20 using an Ultrawash Plus plate washer (Dynex, Chantilly, VA). Diluted sera (1:500, 50 µL per well) were incubated overnight at 4° C while shaking. Plates were washed as before and Protein G-HRP (50 mL, 1:1000 dilution; Biorad, Hercules, CA) was added and incubated for 2 h at room temperature while shaking. Plates were washed again as before and developed by addition of TMBlue (Intergen, Milford, MA), 50µL per well. The reaction was stopped after 5 min by the addition of 50 µL 2N HCl, and plates were read at  $OD_{\lambda 450}$  on an MRX microplate reader with Revelation software (Dynex, Chantilly, VA).

# Statistical analysis

Statistical analysis, including ANOVA and both parametric and nonparametric correlations, was performed using SPSS 10.0 software for Windows (SPSS Inc., Chicago, IL). Briefly, TNFα antibodies in both GRIV S/NP and seronegative controls were determined to be normally distributed using the Kolmogorov-Smirnov test for normality; ANOVA was performed to test for significant differences between TNFa antibody levels of GRIV S/NP relative to seronegative controls. TNF $\alpha$  antibody levels in GRIV S/NP were correlated with peripheral blood lymphocyte and T cell subset counts, antibody responses to viral proteins including p24, Tat and Nef, and both Tat and gp120-derived peptides. Antibodies to peptide 34, as well as CD4 and CD8 cell counts, and the percent CD4 positive cells were determined to be normally distributed by Kolmogorov-Smirnov. In contrast, viral load, CD4:CD8 ratio, and antibodies to peptides 32, 135, 164, 243, 356, 385, and Tat and Nef fusion proteins were not normally distributed. Correlations between  $TNF\alpha$  antibodies and viral load, CD4 count, CD4 percent, CD4:CD8 ratio, antibodies to peptides 32, 34, 243, 356, and Nef protein were significant using both parametric and nonparametric tests; Pearson's coefficients and P-values are reported in the text. Although the correlation between TNFa antibodies and CD8 cell count only passed the parametric Pearson procedure, it was deemed significant because both sets of data were normally distributed. Similarly, although correlations between antibodies to TNF $\alpha$  and peptides 135 and 385 only passed the non-parametric Kendall and Spearman procedures, they were deemed significant because antibodies to both peptides 135 and 385 were not normally distributed; Spearman's coefficients and *P*-values were reported in the text.

#### **RESULTS AND DISCUSSION**

The T cell depletion characteristic in AIDS may be due in part to TNF $\alpha$  dysregulation; TNF $\alpha$  secretion contributes to wasting and T cell anergy, and can provide a pro-apoptotic signal for both CD4+ and CD8+ T cells. TNF $\alpha$  also has been implicated in HIV encephalophathy and HIV-associated progressive



**Figure 2**. Expression and purification of hTNF $\alpha$  fusion protein under native conditions using metal chelate affinity chromatography. A) Coomassie stain of 14% SDS-PAGE of 5 mL Talon column load, flow-through, and 10 mM and 50 mM Imidazole washes. B) Coomassie stain of 14% SDS-PAGE of 500 mM Imidazole elution. The main band is a monomer of approximately 22 kDa; the less intense band of approximately 45–55 kDa is presumably the TNF $\alpha$  trimer. C) DAB-stained Western blot of 14% SDS-PAGE of load, flow-through, 50 mM Imidazole wash, and 500 mM Imidazole elution using anti-HisG-HRP monoclonal antibody at 1:5000 dilution (Invitrogen, San Diego, CA).

multifocal leukoencephalopathy, as well as other CNS disorders [12-14, 23-27]. In HIV infection, TNF $\alpha$  production is elevated as demonstrated by leukocyte TNF $\alpha$  mRNA levels, as well as membranebound and circulating forms of TNF $\alpha$  [28-32]. Circulating soluble TNF $\alpha$  receptor is also elevated and most likely antagonizes the effects of circulating and membrane-bound TNF $\alpha$  [28, 30, 33, 34]. The importance of activation of TNF $\alpha$  synthesis in the pathogenesis of HIV infection has been further emphasized by studies demonstrating reduced TNF $\alpha$ production with successful application of highly aggressive antiretroviral therapy (HAART) [35-38].

HIV infection increases TNF $\alpha$  gene expression by interaction of the envelope glycoprotein with CD4 receptors and coreceptors [39-41], as well as by Tatmediated activation [14, 42, 43]. TNF $\alpha$  in turn exerts complex effects on T cells, such as stimulating proliferation, and inducing or preventing apoptosis [44]. HIV-induced TNF $\alpha$  dysregulation is involved in apoptosis of CD4+ cells through a process apparently enhanced by TNF $\alpha$ -mediated upregulation of Fas ligand [45-47]. TNF $\alpha$  activation is also involved in CD8+ cell apoptosis in AIDS through interaction of membrane-bound TNFa on macrophages and TNF receptors on CD8+ cells [48]. TNF $\alpha$  also appears to contribute to polyclonal B cell proliferation in HIV infection [49, 50]. Although TNFa increases HIV replication at the level of transcription,  $TNF\alpha$  may actually suppress CCR5 tropic virus infection via induction of  $\beta$  chemokines and inhibition of CCR5 gene expression, as demonstrated with the monocytetropic NSI virus [9]. In contrast, TNFα upregulates the CXCR4 receptor on T cells, increasing the susceptibility to infection with cytopathic, syncytiainducing (SI) viruses [51]. In view of the role of SI variants in the pathogenesis of HIV infection, TNF $\alpha$  may play a critical role in viral evolution toward pathogenic variants. Based on the functional properties of TNF $\alpha$  in facilitating HIV viral replication, evolution of pathogenic variants and dysregulation of the immune system, autoantibodies to TNF $\alpha$  may be an advantageous host response to viral infection and chronic TNF $\alpha$  upregulation.

In order to study the levels of TNF $\alpha$  autoantibodies, we expressed and purified recombinant human TNF $\alpha$  for use in ELISA for autoantibodies in serum from HIV-positive volunteers. TNF $\alpha$  was expressed in soluble form in E. coli after overnight induction with IPTG. From one liter of culture, 24 mg of recombinant protein (rhTNF $\alpha$ ) was purified by affinity chromatography using Talon resin (Qiagen, Valencia, CA). Figure 2 illustrates the protein profiles of the extract and purified fractions obtained during the purification process. A major band of 22 kDa was visualized in agreement with the predicted migration of the hTNFa fusion protein. The slower-migrating band presumably represents the TNF $\alpha$  trimer (45–55 kDa); the trimer represents the active form of this protein in solution [52]. Antigenicity of the unmodified fusion protein was confirmed by Western blot (figure 2), and in ELISA (data not shown) using goat anti-TNFa antibody (R & D Systems, Minneapolis, MN).

The purified rhTNF $\alpha$  was used in ELISA to determine the level of autoantibodies from HIV-infected

S/NPs and controls. As shown in *figure 3*, the level of autoantibodies to TNF $\alpha$  was significantly higher (P = 0.005; ANOVA) in the seropositive group (N = 120, mean = 0.244), relative to the seronegative control group (N = 12, mean = 0.114). Among S/NPs, autoantibodies to TNFa positively correlated with viral load (P = 0.013, Pearson = 0.282), and with CD8+ cell number (P = 0.03, Pearson = 0.258). Autoantibodies to TNF $\alpha$  correlated inversely with CD4+ cell number (P = 0.003, Pearson = -0.246), percent CD4+ (P = 0.008, Pearson = -0.306). No significant correlation with age, sex, or total lymphocyte count was observed. Correlation of anti-TNF $\alpha$ antibodies with other antibody responses to Tat and gp160 peptides are shown in table I [53-59]. TNFa antibodies positively correlated with antibodies to gp160 (a.a. 418–444) peptide 32 (P = 0.0001, Pearson = 0.467), and peptide 34 (P = 0.001, Pearson = 0.384), both corresponding to the CD4 binding site. Correlations were also observed with antibodies to peptide 243 containing the 'SLWDQ' CD4 identity region of gp120 (P = 0.016, Pearson = 0.29), and peptide 356 containing a reiterated SLWDQ sequence (P = 0.0001, Pearson = 0.429), as well as peptide 135 (P = 0.005, Spearman = 0.34), encoding the V3 loop (a.a 290-323). There was no correlation with antibodies to several other regions of gp160 (peptides 163, 12, and 23), as illustrated in table I. However, as shown in table I, a correlation between TNFa antibodies and anti-Nef antibodies was observed (P = 0.008, Pearson = 0.302). There was no correlation between  $TNF\alpha$  antibodies and antibodies to recombinant N-terminal polyhistidine fusion Tat protein. Interestingly, a correlation was observed with antibodies to the amino-terminal region of Tat (P = 0.001, Spearman = 0.395), using peptide 385 (a.a. 1-15 of Tat).

Anti-TNF $\alpha$  antibodies may be a natural response to excess TNF $\alpha$  generated as a result of HIV infection. These autoantibodies may serve to mollify the harmful effects of TNF $\alpha$  continually produced by virally-infected cells. The effect of TNF $\alpha$  in activating HIV replication at the level of transcription is well established via the NFKB pathway [2, 59, 60]. It appears that in contrast to the usual antiviral function of TNF $\alpha$ , this cytokine pathway is activated by viral infection and utilized to promote viral infection and injury to the immune system and CNS. The release of soluble TNF receptors in circulation is evi-



**Figure 3**. Box-plot of results from ELISA for serum antibodies (IgG) to TNF $\alpha$ . Anti-TNF $\alpha$  antibodies are significantly elevated (P = 0.005, ANOVA), in GRIV S/NPs (N = 120, mean = 0.24), compared to seronegative controls (N = 12, mean = 0.11). The mean OD of each group is indicated in the box-plot with a cross; the numerical value is also given. Dark horizontal lines connected by a vertical line within each box represent the standard error of the mean for each group. The upper and lower boundaries of the boxes represent the 75th and 25th percentiles, respectively. Whiskers represent the range of values observed in each group. Outliers are indicated with circles.

dent in HIV infection and may provide one means of reducing the harmful effects of TNF $\alpha$  activation. Here we demonstrate the presence of autoantibodies to TNF $\alpha$  in HIV-infected S/NPs, which may provide an additional adaptive response to TNF $\alpha$  dysregulation.

In view of the involvement of TNF $\alpha$  in virus replication and pathogenesis, the development of a vaccine designed to elicit antibodies to TNF $\alpha$  may help restore immune function and at the same time reduce viral replication. Aside from the potential utility of a vaccine to TNF $\alpha$  in HIV infection, this approach may have utility in a variety of clinical disorders where cytokine imbalance contributes substantially to the disease process. For example, in hairy cell leukemia, TNF $\alpha$  may be important for hairy cell leukemia growth [61, 62]. TNF $\alpha$  has been implicated in the pathogenesis of Crohn's disease and other inflammatory bowel diseases and is the target of two therapies: etanercept, a soluble TNF receptor linked to the Fc portion of human IgG1 [63], and infliximab, a

Table I. Synthetic HIV polypeptides used to detect antibody titers in the GRIV non-progressor population.										
HXB2 position	Peptide number	Amino acid sequence	Source	Presumptive role	Correlation TNFa autoantibodies					
					Pearson		Kendall		Spearman	
					Coeff.	P-value	Coeff.	P-value	Coeff.	P-value
gp160 <sub>294-321</sub>	12	INCTRPNYNKRKRIHGPGRAFYTTK	MN isolate [53]	Env V <sub>3</sub> loop frag- ment	0.14	0.259	0.075	0.369	0.116	0.348
gp160 <sub>584-604</sub>	23	ERYLKDQQLLGIWGCSGKLIC	BRU isolate [54]	C-terminal frag- ment of gp120	0.14	0.256	0.071	0.391	0.115	0.349
gp160 <sub>418-444</sub>	32	CRIKQIINMWQGVGKAMYAPPIEGQIN	Z6 isolate [55]	CD4 binding site of gp120	0.467	0.0001	0.312	0.0001	0.438	0.0001
gp160 <sub>418-444</sub>	34	CRIKQIINMWQEVGKAMYAPPISGQIR	BH8 isolate [56]	CD4 binding site of gp120	0.384	0.001	0.263	0.002	0.391	0.007
gp160 <sub>290-323</sub>	135	FLLAVFCTRPNYNKRKIHIGPGRAFYT- TKNIIG	MN isolate	Env V <sub>3</sub> loop frag- ment	0.148	0.234	0.24	0.004	0.34	0.005
gp160 <sub>837-856</sub>	163	CRAIRHIPRRIRQGLERILL	BRU isolate	C-terminal frag- ment of gp41	0.143	0.244	0.081	0.33	0.096	0.437
gp160 <sub>110-125</sub>	243	SLWDQSLKPCVKLTPL	MN isolate	CD4 identity of gp120 [57]	0.29	0.016	0.221	0.008	0.315	0.009
gp160 <sub>110-125</sub> (reiterated)	356	SLWDQSLWDQSLWDQ	MN isolate	Reiterated SLWDQ frag- ment of gp160 CD4 identity	0.429	0.0001	0.0367	0.0001	0.547	0.0001
gp160 <sub>289-305</sub>	362	NQSVEINCTRPNNNTRK	BRU isolate	Fas identity of gp120	0.119	0.335	0.085	0.307	0.116	0.345
Tat <sub>1-15</sub>	385	MEPVDPRLEPWKHPG	MN isolate	N-terminal frag- ment of transacti- vating regulatory protein	0.157	0.2	0.283	0.001	0.395	0.001
Tat		Fusion protein	III B	Transactivator	0.006	0.96	0.012	0.881	0.018	0.879
Nef	Fusion protein		pSVL4-3	CD4 downregu- lation and patho- genesis	0.302	0.008	0.245	0.002	0.344	0.002

The use of anti-cytokine vaccines in the form of 'kinoids' has been previously proposed [69, 70]. This strategy has provided encouraging results in a murine model of experimental cachexia and type II collageninduced arthritis using a chimeric TNF $\alpha$  vaccine containing immunodominant T<sub>h</sub> epitopes [71]. The application of TNF $\alpha$  vaccine strategies may provide a therapeutic approach to the treatment of HIV infection as well as a wide variety of clinical disorders.

#### CONCLUSIONS

We produced and purified a recombinant  $TNF\alpha$ fusion protein to detect autoantibodies to  $TNF\alpha$  in HIV-infected slow/non-progressors (S/NPs). Anti-TNF $\alpha$  antibodies are significantly elevated in HIVinfected non-progressors, who are naïve to antiretroviral therapy, compared to seronegatives. Because TNF $\alpha$  activates viral replication and induces immune dysregulation, the ability of the host to interfere with chronic TNFa upregulation may be important for prevention of disease progression. Vaccine strategies designed to boost these natural anti-cytokine autoantibodies comprise important therapeutic venues for AIDS treatment. Further studies will be required to determine if anti-TNF $\alpha$  antibodies are indeed a beneficial host response and to evaluate the concept of a TNF $\alpha$ -targeted vaccine.

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