

ABSTRACT

EMANI, SIRISHA. MOLECULAR CHARACTERIZATION OF T REGULATORY CELLS IN FIV-INFECTION (Under the direction of Dr. Wayne Tompkins)

Naturally occurring CD4⁺CD25⁺ T regulatory cells (Treg) play important roles in maintaining immunologic self-tolerance in addition to controlling the magnitude of anti-microbial immune responses. However, the capacity of these CD4⁺CD25⁺ Treg cells to control immune responses both in vivo and in vitro is not well established. CD4⁺CD25⁺ Treg cell-mediated suppression can control autoimmune diseases; transplantation tolerance and graft versus host disease and, in contrast hinder tumor immunity and immunity to infectious agents. As Treg cells have been reported to be involved in several diseases, this study focused on molecular characteristics that enables them to maintain energy and also resistance to programmed cell death along with the effect of FIV-infection on regulation of the above phenotypic characteristics. Our results show that feline CD4⁺CD25⁺ Treg cells are phenotypically and functionally anergic as indicated by elevated levels of the cyclin dependent kinase inhibitors, CdkI's, (p21^{cip1}, p16^{ink4}, and p27^{kip1}), and resistance to mitogen-induced proliferation compared to their counter parts CD4⁺CD25⁻ T cells. Importantly, CdkI's are constitutively over-expressed only in FIV-infected cats. As expected Treg cells from FIV-infected cats that over-expressed CdkI's expressed low levels of the cyclins (mainly cyclins D) and phosphorylated retinoblastoma protein (pRb) that are responsible for cell cycle progression. We investigated the role of TGFβ signaling and found that TGFβ1 plus ConA stimulation was able to convert CD4⁺CD25⁻ T cells to CD4⁺CD25⁺ T cells with functional and phenotypic characteristics including upregulation of CdkI's and bcl-2. The differential expression of CdkI's and bcl-2 between the two CD4⁺ T cell subsets may be linked to TGFβ-Smad pathway. Consistent with upregulation of CdkI's and bcl-2, we found that although natural and TGFβ1 converted CD4⁺CD25⁺ Treg cells are anergic, they are more resistant to activation induced cell death compared to CD4⁺CD25⁻ T cells functionally which correlated with increased bcl-2 to bax ratio in Treg cells. Thus, the molecular characterization of this unique population of Treg cells may be essential for understanding their role and function for developing effective therapeutics and vaccination especially against chronic infections such as Acquired Immune Deficiency Syndrome (AIDS).

**MOLECULAR CHARACTERIZATION OF T REGULATORY CELLS
IN FIV-INFECTION**

BY

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LIST OF ABBREVIATIONS

- AICD**- Activation Induced Cell Death
- Apaf-1**- Apoptosis protease activating factor-1
- APC**- Allophycocyanin
- CAD**- Caspase Activated DNase
- Cdk**- Cyclin dependent kinase
- CdkI**- Cyclin dependent kinase Inhibitor
- CREB**-c AMP Response Element Binding Protein
- Con-A**- Concavalin-A
- FITC**- Fluorescein Isothiocyanate
- FIV**- Feline Immunodeficiency Virus
- GITR**- Glucocorticoid-Induced TNF Receptor
- rhIL-2**- recombinant Interleukin-2
- LPS**- Lipopolysaccharide
- NF κ B**- Nuclear Factor associated with Kappa light chain of B cells
- PBMC**- Peripheral Blood Mononuclear cells
- PE**- Phycoerythrin
- PI**- Propidium Iodide
- TCR**- T cell receptor
- TGF β** – Transforming Growth Factor-beta
- TNF α** – Tumor Necrosis Factor-alpha
- Treg** – CD4⁺CD25⁺ T regulatory cells

1. INTRODUCTION

CD4⁺CD25⁺ T regulatory cells (Treg) are naturally anergic, arrested in the G0/G1 phase of the cell cycle. Similar to anergic CD4⁺ Th cells, CD4⁺CD25⁺ Treg cells fail to produce IL-2 and proliferate in response to TCR engagement by antigens, but respond to polyclonal stimuli such as IL-2 (Powell, Ragheb et al. 1998; Lechler, Chai et al. 2001) . Naturally occurring CD4⁺CD25⁺ Treg cells, comprising 5-10% of the circulating CD4⁺ T cell population, were originally defined as regulatory T cells that maintain peripheral self-tolerance (Maloy and Powrie 2001; Shevach 2002). More recent studies have demonstrated that Treg cells also play a major role in modulating the magnitude and duration of protective T cell immune responses to pathogens (Read, Malmstrom et al. 2000; Iwashiro, Messer et al. 2001; Belkaid, Piccirillo et al. 2002; Hori, Carvalho et al. 2002; Montagnoli, Bacci et al. 2002; Hisaeda, Maekawa et al. 2004; Belkaid and Rouse 2005). These pathogen-induced Treg cells, similar to natural Treg cells, are anergic and possess potent immunosuppressor function once activated by TCR engagement and/or certain nonspecific stimuli (Belkaid and Rouse 2005).

Treg cells are arrested in the early G0/G1 stage of the cell cycle, suggesting that anergic factors influence early events in the cell cycle. Progression through the G0/G1 to S phase of the cell cycle is regulated by several cyclins and cyclin-dependent kinases (Cdk) whose activities are in turn controlled by their inhibitors, CdkI's. Jackson SK reported that Th1 cell anergy could be linked to increased expression of the cyclin-dependent kinase inhibitors, p21^{*cip1*} and p27^{*kip1*}. Current models propose that p27^{*kip1*} inhibits cyclin-

dependent kinase-2 (cdk2) expression, thus maintaining the cell in a quiescent state (Coats, Whyte et al. 1999; Mohapatra, Agrawal et al. 2001; Mohapatra and Pledger 2001). Down-regulation of p27^{kip1} is associated with entry of cells from G1-S phase and its up-regulation is associated with G₁ growth arrest. p27^{kip1}^{-/-} mice exhibit gigantism with disproportionately enlarged spleen, thymus, and lymph nodes (Fero, Rivkin et al. 1996; Kiyokawa, Kineman et al. 1996; Nakayama, Ishida et al. 1996), suggesting the importance of p27^{kip1} in regulating the homeostatic proliferation of T cells in vivo. On the other hand, reports suggest that p21^{cip1} may either act as an inhibitor of apoptosis (Suzuki, Tsutomi et al. 1999) or a promoter of apoptosis (Peschiaroli, Figliola et al. 2002). T cell development in mice deficient in p21^{cip1} occurs normally. But, p21^{cip1} knockout mice rapidly lose tolerance to nuclear antigens and develop an autoimmune-like disease reminiscent of systemic lupus in humans (Balomenos, Martin-Caballero et al. 2000), suggesting their role in regulating T cell proliferation and homeostasis.

T cell apoptosis also plays a critical role in maintaining immune homeostasis. It has been reported that human CD4⁺CD25⁺ Treg cells are prone to apoptosis which is associated with decreased levels of bcl-2 expression (Taams, Smith et al. 2001). Conversely, others have reported that mouse CD4⁺CD25⁺ Treg cells are resistant to apoptosis, as they do not undergo clonal deletion by viral superantigen *in-vivo* (Papiernik, do Carmo Leite-de-Moraes et al. 1997), and are resistant to Fas-induced apoptosis *in-vitro* (Banz, Pontoux et al. 2002). This urged us to determine if the reported resistance to activation induced cell death (AICD) of feline CD4⁺CD25⁺ Treg cells compared to their counterparts

CD4⁺CD25⁻ T cells in correlation with the expression of anti-apoptotic proteins and pro-apoptotic proteins involved in the mitochondrial apoptotic pathway.

Recently, it has been suggested that CD4⁺CD25⁺ Treg cell homeostasis can also be maintained in the presence of IL-2 and TGFβ1 (Horwitz, Zheng et al. 2004). The development of T regulatory cells (Fontenot, Gavin et al. 2003) and Treg cell suppressor activity (Chen, Jin et al. 2003), is known to be programmed by the transcription factor Foxp3, which is selectively regulated in peripheral immune compartments by TGFβ1, a major growth regulatory cytokine. TGFβ1 induces the expression of p21^{cip1} and p27^{kip1} in various cell types. Depending upon the cell type and the environment TGFβ1 exerts variable effects. Upon stimulation with TGFβ1 lymphocytes exhibit either a G₁ arrest or apoptosis. TGFβ1 is expressed in the G₁ phase of the cell cycle in growth-arrested cells (Howell, Sun et al. 1993) and binding of TGFβ1 to its receptor activates a wide range of pathways during cell cycle progression. TGFβ1 is constitutively expressed at elevated levels on CD4⁺CD25⁺ Treg cells, that are arrested in the G₀/G₁ phase of the cell cycle, compared to CD4⁺CD25⁻ T cells (Maggi, Cosmi et al. 2005). TGFβ1 by itself or along with other superfamily members, such as BMPs, and activins, can induce p21^{cip1} transcription, an event that is thought to link cell growth inhibition, and/or apoptosis depending upon the target cell type (Zauberman, Oren et al. 1997; Ishisaki, Yamato et al. 1998; Jernvall, Aberg et al. 1998; Ghosh-Choudhury, Ghosh-Choudhury et al. 2000; Franzen and Heldin 2001). It has been a subject of debate whether TGFβ1-induced p21^{cip1} expression is linked to inhibition of cell proliferation (Hu et al., 1998; Cheng et al., 2001). Transcriptional induction of p21^{cip1} responsiveness to TGFβ1 could be mediated

either by SMAD pathway or by Ras/mitogen activated protein kinase (MAPK) pathway (Kivinen, Tsubari et al. 1999; Pardali, Kurisaki et al. 2000; Pouliot and Labrie 2002). The p21^{cip1} promoter possesses binding site/s for transcription factors of the Sp1 family along with Smads (Moustakas, Pardali et al. 2002) which is targeted by both Smad- and Ras/MAPK-mediated pathways. TGFβ1 induces phosphorylation of cyclic AMP response element binding proteins (CREB) via MAPK pathway. IL-2Rα gene possesses binding sites for CREB/ATF, AP-1 and Smad3. It has also recently been reported that TGFβ1 along with anti-CD3 and anti-CD28 stimulation could induce IL-2Rα expression (CD25). Also the anti-apoptotic gene bcl-2 processes a CREB binding site on its promoter region (Wilson, Mochon et al. 1996). Therefore, TGFβ1 along with TCR stimulation could induce the expression of CD25, bcl-2, as well as the cell cycling protein inhibitors such as p21^{cip1}. Thus, the TGFβ1 pathway may be a useful mechanism to explain the phenomenon of cell growth inhibition, yet programmed cell survival in Treg cells.

2. BACKGROUND AND SIGNIFICANCE

A. Generation of CD4⁺CD25⁺ Tregulatory cells

Although the existence of a separate subset of T cells specifically involved in suppression of immune responses was proposed 30 yrs ago, it was only over the past 5-10 years that a distinct lineage of CD4⁺CD25⁺ T regulatory cell population or suppressor cells was identified. The function of the thymus, in addition to regulatory T cell differentiation and selection, is to maintain immunological self-tolerance by producing CD4⁺CD25⁺ Treg cells capable of preventing autoimmune diseases (Apostolou, Sarukhan et al. 2002). Clonal deletion, one of the major mechanisms for distinguishing self/non-self by the immune system, can establish tolerance to self-reactive B and T lymphocytes during their development in the thymus. However, the deletion mechanism is considered to be leaky, as it allows some of the self-reactive lymphocytes to escape clonal deletion and exit to the periphery. For the past few years it has become evident that Treg cells actively suppress self-reactive T cells in the periphery, avoiding the onset of harmful autoimmune diseases.

It has been demonstrated that the Treg cells acquire their regulatory activity through the thymic selection process and are released in the periphery as functionally committed CD4⁺CD25⁺ Treg cells (Apostolou, Sarukhan et al. 2002). Apart from their generation in the thymus from a defined lineage, recent studies have shown that these CD4⁺CD25⁺

Treg cells can also be activated in the periphery in response to self antigens or pathogen products (Thorstenson and Khoruts 2001; Zhang, Izikson et al. 2001; Hauet-Broere, Unger et al. 2003) or can be converted from normal CD4⁺CD25⁻ T cell subsets (Chen, Jin et al. 2003) via the TGFβ1 signaling pathway .

B. Phenotypic characteristics of Treg cells

CD4⁺CD25⁺ Treg cells constitute 5-10% of the peripheral CD4⁺ T cells in mice, rats, cats, and humans (Maloy and Powrie 2001; Shevach 2002; Vahlenkamp, Bull et al. 2004). Several distinct subsets of T regulatory/suppressor cells had been described by the expression of surface markers and/or cytokine production profiles. The thymus-derived, naturally occurring CD4⁺CD25⁺ Treg cells, TGFβ1 producing Th3 cells (Fukaura, Kent et al. 1996) and IL-10 secreting Tr1 cells (McGuirk and Mills 2002) appear to be phenotypically different, yet functionally similar, as they suppress the proliferation and IL-2 production by stimulated T cells (Vahlenkamp, Tompkins et al. 2004). Although the CD25 (interleukin-2 receptor chain alpha) molecule is expressed on all conventional activated T cells, and not all T regulatory cells express CD25, to date CD25 is still used as a marker to characterize and isolate T regulatory cells. Several other cell surface markers and intracellular markers had been reported to be constitutively expressed by thymus-derived naturally occurring Treg cells, which slightly differ in murine and human species. Murine thymus-derived Treg cells constitutively co-express CD25⁺, CD122⁺ CD69⁺, CD44⁺, CD45RB^{low}, GITR⁺, CD103⁺(alphaE-integrin), CD134⁺(OX-40), CD54⁺(ICAM), whereas human thymus-derived Treg cells express CD25^{high}, CD122⁺,

HLD-DR⁺, CD45RO⁺, CD95^{high}, CD45RB^{low}, CD38^{low}, CD62^{low}, and GITR⁺ phenotype (Asano, Toda et al. 1996). Recent studies on human and murine Treg cells have used additional markers such as CD62L-selectin along with CD25 to differentiate activated T cells from Treg cells. Activated T cells down regulate CD62L-selectin (Mobley and Dailey 1992), whereas Treg cells constitutively express elevated levels of CD62L-selectin (McHugh, Whitters et al. 2002). The development of T regulatory cells (Fontenot, Gavin et al. 2003) and T cell suppressor activity (Curotto de Lafaille, Muriglan et al. 2001), is known to be programmed by the transcription factor Foxp3, which is selectively expressed in CD4⁺CD25⁺ Treg cells. While Foxp3 is perhaps, the most selective marker for Treg cells, its usefulness for detection and isolation of Treg cells is somewhat limited by its intracellular location. The other markers associated with CD4⁺CD25⁺ Treg cells are CTLA-4 and glucocorticoid –induced TNF receptor (GITR), that are induced by Foxp3 (Hori and Sakaguchi 2004). Recent data suggest that cell surface expressed TGFβ may be a useful marker for activated CD4⁺CD25⁺ Treg cells (Chen and Wahl 2003).

C. Functional characteristics of Treg cells

Apart from the use of the above described phenotypic markers, Treg cells can be best identified using their functional properties. A classical feature of CD4⁺CD25⁺ Treg cells is that they are anergic upon TCR mediated-stimulation in vitro, failing to produce IL-2 and undergo proliferation (Lechler, Chai et al. 2001). Some studies have reported that T cell anergy could be overcome by culturing Treg cells in the presence of IL-2 or by addition of anti-CD28 antibody that induces endogenous IL-2 production (Thornton and

Shevach 1998).

The other major functional characteristic of Treg cells is their ability to suppress proliferation and cytokine production by other activated T cells. T cell suppression by Treg cells is cell contact-dependent and occurs when Treg cells are activated via their T cell receptors. Several recent studies have also shown that TCR ligation is not always required for Treg cell immunosuppression of other T cells (Caramalho, Lopes-Carvalho et al. 2003). Treg cells express several Toll-like receptors (TLRs) including TLR-4, a receptor for LPS which can activate murine (Caramalho, Lopes-Carvalho et al. 2003) and feline Treg cells (Vahlenkamp, Bull et al. 2004). Also Vahlenkamp et al. (Vahlenkamp, Bull et al. 2004) reported that potent Treg cell suppressor function could be induced by ConA or rhIL-2 stimulation.

D. Activation of Treg cells in the peripheral immune compartment

CD4⁺CD25⁺ T regulatory cells possess immune suppressive activity in vitro, and can suppress the proliferation and cytokine production of other T cells, both CD4⁺ and CD8⁺, via an undefined cell contact-dependent mechanism (Takahashi, Kuniyasu et al. 1998; Thornton and Shevach 1998). On the other hand, CD4⁺CD25⁺ Treg cells stimulated with antigen presenting dendritic cells or IL-2 lose their anergic phenotype and can proliferate comparable to CD4⁺CD25⁻ T cells in-vivo and in-vitro while retaining the suppressor function (Yamazaki, Iyoda et al. 2003) Some authors reported that CTLA-4 is constitutively expressed on Treg cells and mediates T cell suppression via inhibition of

IL-2 expression (Krummel and Allison 1996; Read, Malmstrom et al. 2000). Others reported that CTLA-4 is not essential for suppression of other lymphocytes, as Treg-mediated suppression exists even in CTLA-4^{-/-} mice. On the other hand, CTLA-4 engagement upregulates TGF β and TGF β RII surface expression on Treg cells, which enhances their suppressor function (Chen and Wahl 2002). In contrast, glucocorticoid-induced TNF receptor (GITR), which is also predominantly expressed on CD4⁺CD25⁺ Treg cells, counterbalances their suppressor activity once it has been engaged by its ligand (McHugh, Whitters et al. 2002; Shimizu, Yamazaki et al. 2002).

It has also been reported in rodents that ligation of GITR with soluble GITR ligand (sGITR-L) can induce GITR-dependent NF κ B activation, thereby up-regulating IL-2 and thus blocking in-vitro suppression mediated by Treg cells (Ji, Liao et al. 2004). Also, engagement of GITR on Treg cells with its ligand antagonizes TGF β signal transduction by inhibiting phosphorylation of Smad2/3, downstream signaling molecules of TGF β -signaling pathway, to block suppression (Chen, Jin et al. 2003).

It has been well established that TGF β 1 is involved in down-regulation of T cell-mediated responses and in controlling autoimmunity (Letterio and Roberts 1998). TGF β 1 produced by Treg cells, bound to its receptor, expressed on other activated T cells could induce suppression of other T cells in a cell contact-dependent mechanism (Gorelik and Flavell 2000; Leveen, Larsson et al. 2002). In support of this, TGF β RII deficient mice exhibit uncontrolled T cell proliferation (Nakamura, Kitani et al. 2001; Zhang, Izikson et al. 2001). Furthermore, TLR-4 triggering of dendritic cells (DCs) has been reported to

induce IL-6, and IL-1 production, enhance Treg proliferation, and reduce the suppressive activity of CD4⁺CD25⁺ Tregs via GITR-dependent pathway (Pasare and Medzhitov 2003; Yamazaki, Iyoda et al. 2003). Th cell activation requires engagement of dendritic cells co-stimulatory molecules along with TLR-induced cytokines down-regulate Treg cell-mediated suppression during infection (Pasare and Medzhitov 2003). This could be one of the multiple mechanisms in which activation of cells of the innate immune system can override the suppressor activity of CD4⁺CD25⁺ Tregs, thus permitting adaptive immune responses to antimicrobial Ags to supersede the suppressive response to self Ags.

E. Involvement of T regulatory cells in auto-immune and infectious diseases

The CD4⁺CD25⁺ T regulatory cell population has a broad repertoire of receptors that recognize various self and non-self pathogen-associated antigens enabling Treg cells to regulate a broad range of antigen-specific responses. In the case of autoimmune diseases, it has been shown in several rodent models that depletion of CD4⁺CD25⁺ Treg cells leads to severe autoimmunity disorders, and passive transfer of these cells prevents autoimmune disease (Salomon, Lenschow et al. 2000; Stephens and Mason 2000). Treg cells have now been known to regulate a wide range of diseases that includes organ-specific autoimmune diseases (including gastritis, thyroiditis and insulin-dependent diabetes) (Salomon, Lenschow et al. 2000; Stephens and Mason 2000; Furtado, Olivares-Villagomez et al. 2001), inflammatory bowel disease (Read, Malmstrom et al. 2000), inflammatory lung disease (Hori, Carvalho et al. 2002), allograft rejection (Hara, Kingsley et al. 2001; Taylor, Noelle et al. 2001; Graca, Thompson et al. 2002), graft

versus host disease (Hoffmann, Ermann et al. 2002; Taylor, Lees et al. 2002), and allergy (Curotto de Lafaille, Muriglan et al. 2001).

In addition to regulating autoimmune and inflammatory diseases, Treg cells regulate sterilizing immunity to infectious organisms such as protozoa (Belkaid, Piccirillo et al. 2002), and fungi (Netea, Suttmuller et al. 2004), bacteria (Raghavan, Suri-Payer et al. 2004), and viruses (Aandahl, Michaelsson et al. 2004). It has also been shown that Treg cells have the ability to regulate innate immune natural killer T cells to prevent *Helicobacter hepaticus*-triggered colitis in T cell-deficient mice models (Maloy, Salaun et al. 2003). The other major role of Treg cells is the suppression of pathogen-induced inflammatory responses. These pathogen-specific Treg cells can prevent immunopathology induced by infections, but in contrast, can also increase the pathogen load and prolong persistence by suppressing protective immune responses. It has been recently suggested that the decreased pathogen induced responses lead to prolong persistence of infection which could serve as a mechanism for the maintenance of pathogen specific immunological memory. Depletion of Treg cells from mice infected with *Leishmania major* or *Pneumocystis carini* initiated an effective adaptive response but resulted in severe inflammation (Belkaid 2003). Similar immune responses were seen in $CD4^+CD25^+$ Treg cells depleted mice with malarial parasites but with hindered memory immune responses (Aseffa, Gumy et al. 2002; Belkaid, Piccirillo et al. 2002; Hori, Carvalho et al. 2002; Hisaeda, Maekawa et al. 2004). Recent studies also demonstrated that removal of Treg cells from the PBMC of HIV-infected individuals resulted in an effective antigen specific response as measured by cytokine production by antigen

stimulated CD4⁺ and CD8⁺ effector T cells (Aandahl, Michaelsson et al. 2004; Kinter, Hennessey et al. 2004). As Treg cells are known to be involved in several diseases, the phenotypic and genotypic characterization, and understanding their role and function may be useful for devising effective therapeutics and vaccination, especially against chronic infections such as AIDS.

F. Role of IL-2 in T regulatory cell function and homeostasis

According to the classical model of T cell activation, productive stimulation of T cells requires two major signals. The first is the engagement of MHC on APC with the TCR, and the second is the interaction of B7 on the APC with the T cell co-stimulatory receptor CD28 (Azuma, Yssel et al. 1993; Powell, Ragheb et al. 1998). In a normal cell, B7 receptors on APC initially engage the CD28 receptor on CD4⁺ T cells transducing a signal that upregulates IL-2 production, an essential cytokine for transit out of the G0/G1 stage of the cell cycle, cell proliferation and the initiation of an immune response (Krummel and Allison 1995; Krummel and Allison 1996). Following APC activation T cells upregulate CTLA-4 on their surface and upon engagement of B7, transduces a signal for the down-regulation of IL-2 and termination of the immune response. If the T cells are stimulated only through the TCR, these T cells also fail to produce IL-2 enters G0/G1 arrested state of long-lasting, partial or total unresponsiveness termed anergy (Karandikar, Vanderlugt et al. 1996; Bluestone 1997; Scheipers and Reiser 1998; Walunas and Bluestone 1998). Anergic T cells possess increased expression of B7 and CTLA-4 molecules upon mitogen stimulation, yet fail to produce IL-2 and proliferate

upon re-stimulation with antigen-competent APC, but respond to polyclonal stimuli such as IL-2 (Chai and Lechler 1997; Powell, Ragheb et al. 1998; Lechler, Chai et al. 2001). The signaling pathway that is involved in IL-2 down-regulation in anergic cells is not well established.

IL-2 signaling is essential for Treg cell development in the thymus, homeostasis and function. On one hand, IL-2 potentially stimulates T cell growth and proliferation, and in the absence of IL-2, Treg cells cannot proliferate or survive in the thymus or in the periphery (Curotto de Lafaille, Lino et al. 2004; Bayer, Yu et al. 2005; Setoguchi, Hori et al. 2005). In contrast, IL-2 is necessary for the maintenance of homeostasis and tolerance to self (Malek and Bayer 2004; Nelson 2004). In the absence of IL-2 or its receptor, a lymphoproliferative auto-immune syndrome was reported in several rodent models (Sadlack, Merz et al. 1993; Suzuki, Kundig et al. 1995; Willerford, Chen et al. 1995) suggesting that IL-2/IL-2R is essential for Treg homeostasis and function.

It has been reported that CD4⁺CD25⁺ Treg cells from mice, humans, and cats constitutively express Foxp3 (Fantini, Becker et al. 2004). Foxp3, a member of fork-head transcription factors has been known to be important for the development and function of Treg cells. Foxp3 has the ability to inhibit IL-2, IL-4, and IFN γ production by direct interaction with the Rel family transcription factors, nuclear factor of activated T cells (NFAT) and NF κ B, which are essential for target gene expression. Foxp3 may also directly bind to IL-2 promoter and block IL-2 transcription. Thus, while it appears that Treg cells are programmed for low levels of IL-2 production, some IL-2 is required for

their survival. How this balance is maintained is not known. However recent data showing a consensus binding element in the IL-2 α gene for Smad4 may provide some insight. Kim et al. (Kim, Kim et al. 2005) identified a novel positive regulatory region in the IL-2 α gene that mediates TGF β 1 induction of this gene via Smad3 binding. Thus, TGF β 1 signaling in Treg cells could constitutively sustain IL-2 α up-regulation and the high affinity IL-2 receptor allowing the survival of Treg cells in the presence of low levels of IL-2.

G. Role of Cyclin dependent kinase Inhibitors (CdkI's) in Treg cell anergy

Anergic cells are blocked at the G0/G1 phase of the cell cycle and this blockade can be induced by cyclin dependent kinase inhibitors (CdkI's). Mitogen-dependent progression through the G1 phase and initiation of DNA synthesis (S phase) during cell cycle is regulated by several cyclin dependent kinases whose activities are in turn controlled by CdkI's. During G1/S phase progression, cyclin D (D1-D3 in most cells and only D2-D3 in lymphocytes) act in mid G1 phase followed by cyclin E and cyclin A in G1/S boundary. Cyclins A and B are involved during S and G2/M phases (Nurse 1994; Morgan 1995; Jackman and Pines 1997; Reed 1997). Cyclin dependent kinases (Cdk's) associate with cyclins and get phosphorylated upon activation. In response to mitogen signals, Cyclin D is induced and associates with Cdk4 and Cdk6 (Sherr 1993), whereas cyclin A associates with Cdk2. The activity of cyclin-Cdk complexes is repressed by CdkI's, which constrain entry into S phase (Hirama and Koeffler 1995; Reed 1997). CdkI's that govern these events have been divided into two groups based on their structure and Cdk

targets. The first class includes INK4 proteins (p16, p15, and p18) and consists of multiple ankyrin repeats, which bind to only cdk4 and cdk6 and are involved in the inhibition of the catalytic subunits of cdk4 and cdk6 (Serrano, Hannon et al. 1993; Guan, Jenkins et al. 1994; Hannon and Beach 1994). The second class of inhibitors includes p21^{cip1}, p27^{kip1}, p53 and p57^{kip2}, which contain characteristic motifs within their amino terminal moieties that enable them to bind both cyclins and CDK subunits (Harper, Adami et al. 1993; Toyoshima and Hunter 1994; Lee, Reynisdottir et al. 1995). Action of these CdkI's can affect the activation of several cyclins and Cdk's that are involved in G1/S transition by inhibiting retinoblastoma protein (Rb) phosphorylation. In normal cells the complexes of D cyclins, Cdk4 and Cdk6 phosphorylates the retinoblastoma protein (pRb), causing the release of E2F transcription factors which activates transcription of cyclin E and cyclin A and other cell cycle related genes. Cyclin E binds Cdk2, promoting progression to the S phase (Sherr 1996; Funk and Galloway 1998). p27^{kip1} and p21^{cip1} are well studied CdkI's that regulate anergy. In resting T lymphocytes, p27^{kip1} levels are high and p21^{cip1} levels are low (Jackson, DeLoose et al. 2001). p27^{kip1} is typically up-regulated via growth factor signaling, whereas p21^{cip1} is induced via p53 in response to DNA damage (Philipp-Staheli, Payne et al. 2001; Wolfrum, Walz et al. 2004). As p21^{cip1} is capable of binding with essentially all the cyclin-cyclin-dependent kinase complexes, it plays a major role in regulating T cell cycling and could be a principle factor in maintaining anergy. In support of this, p21^{cip1} knockout mice rapidly lose tolerance to nuclear antigens and develop a lymphoproliferative autoimmune-like disease reminiscent of systemic lupus in humans (Balomenos, Martin-Caballero et al. 2000). To date, contradictory results have been reported by several groups regarding the

effect of p27^{kip1} and p21^{cip1} in cell cycle progression. Jackson S.K.(Jackson, DeLoose et al. 2001), stated that induction of anergy in Th1 cells is associated with increased levels of CdkI's (p27^{kip1}, and p21^{cip1}) in response to antigenic stimuli but not IL-2. Wolfrain et al. utilized p21^{cip1-/-} and p27^{kip1-/-} knockout mice to show that p21^{cip1} and p27^{kip1} act in synergy to maintain TGFβ induced G0/G1 cell cycle arrest. In contrast to the above studies, Funk J.O (Funk and Galloway 1998), reported that p27^{kip1} and p21^{cip1} are not required for induction and maintenance of T cell anergy.

TGFβ1 has also been shown to induce G0/G1 cell cycle-arrest via transcriptional activation of p15^{ink4}, p21^{cip1} and p27^{kip1}, and post-transcriptional stabilization of p21^{cip1} expression in a number of cells, including T and B lymphocytes (Kee, Rivera et al. 2001; Massague 2004; Wolfrain, Walz et al. 2004). Although there is little information on how TGFβ1 activates CdkI's expression, recent studies demonstrate that the p21^{cip1} promoter contains consensus elements capable of binding SMAD4, a critical downstream element in the TGFβ1 signaling pathway (Yamato, Hashimoto et al. 2001). Cell surface expressed TGFβ1 has also been associated with activated CD4⁺CD25⁺ Treg cells, although it's role in Treg function is controversial (Maloy and Powrie 2001; Shevach 2002).

H. TGFβ-signaling and CD4⁺CD25⁺ Treg cell homeostasis and function

Historically CD4⁺CD25⁺ Treg cells have been defined as a thymus derived distinct CD4⁺ cell lineage with unique anergic and immunosuppressive properties. Recently, it

has been demonstrated that CD4⁺CD25⁺ Treg cells can also be generated ex-vivo by culturing naïve peripheral CD4⁺CD25⁻ T cells in the presence of mitogen and TGFβ (Horwitz, Zheng et al. 2004). Stimulation of TCR, IL-2R and TGFβR can trigger Foxp3 expression in CD4⁺CD25⁻ T cell precursors and these Foxp3 expressing cells can act like conventional T regulatory cells (Fu, Zhang et al. 2004) . In contrast, Foxp3 expression and the suppressive function are prevented upon co-stimulation with CD28 in an IL-4 dependent manner (Fu, Zhang et al. 2004) .

TGFβ is a potent inhibitor of growth in a variety of cells including lymphoid cells (Zhang, Izikson et al. 2001). TGFβ1 is expressed in the G₁ phase of the cell cycle in growth-arrested cells (Chen, Jin et al. 2003) and binding of TGFβ1 to its receptor activates a wide range of pathways during cell-cycle progression. The inhibition of growth modulated by TGFβ could be via the suppression of the activity of cyclin-dependent kinases which leads to arrest of the cell cycle at G₀/G₁ phase (Alexandrow and Moses 1997). The inhibitory mechanism by which TGFβ1 modulates cyclin dependent kinases activity involves upregulation of Cyclin-dependent kinase inhibitors (p21^{cip1}, p27^{kip1}, and p15^{ink}) (Iavarone and Massague 1997).

TGFβ superfamily of cytokines contains more than forty proteins including TGFβ1, TGFβ2, and TGFβ3 (Shi, Wang et al. 1998). TGFβRI/II complex formation is induced by the binding of TGFβ ligands that in turn phosphorylates receptor-activated Smads 2/3 (R-Smads). R-Smads form complexes with Smad4 which allows their translocation into the nucleus where the complex interacts with Smad-binding elements (SBE) (Shi, Wang

et al. 1998). R-Smad complex promotes the transcription of several target genes depending upon the combinations of Smad complex interactions with co-activators and inhibitors of gene transcription (Shi and Massague 2003). As of this date there are seven TGF β I, four TGF β RII, and nine Smad 1-9 proteins involved in the TGF β / TGF β R signaling pathway.

TGF β 1 ligands bind specifically to a characteristic combination of TGF β RI and TGF β RII, which in turn activates a specific subgroup of R-Smad. Smad7 and Smad6 negatively regulate TGF β signaling pathway (Shi and Massague 2003). Smad7 can inhibit TGF β signaling either by preventing the interaction Smad2/3 with TGF β 1 receptor or by interacting with the ubiquitin ligases from the smurf family, which irreversibly terminates the signaling pathway (Bonni, Wang et al. 2001). Smad7 is induced via TGF β -Smad2/3/4 complex along with AP1 and Sp1 transcription factors, providing an auto-regulatory loop for TGF β 1 / TGF β R signaling (Brodin, Ahgren et al. 2000). On the other hand, as mentioned earlier, Treg cells express Foxp3, and TGF β 1 induces Foxp3 expression which in turn down-regulates Smad7. Foxp3-mediated down-regulation of Smad7 induces a positive auto-regulatory loop of TGF β -Smad signaling in naturally occurring and TGF β 1-induced Treg cells in human and murine models (Fantini, Becker et al. 2004). This may be one of the mechanisms by which Treg cells could maintain energy by up-regulation of CdkI's via the positive loop of TGF β - Smad signaling pathway.

As mentioned earlier, it has been reported that CD4⁺CD25⁺ Treg cells express elevated levels of activation molecules such as CD80, CD86, and CTLA-4 compared to their counterparts CD4⁺CD25⁻ T cells (Vahlenkamp, Bull et al. 2004). CTLA-4 in the presence of TCR ligation has been shown to produce TGFβ1 by purified naïve CD4⁺ T cells or T cell clones, which may lead to immune suppression (Jonuleit, Schmitt et al. 2001). TGFβ1 is constitutively expressed at elevated levels on CD4⁺CD25⁺ Treg cells (Chen, Jin et al. 2003; Nakamura, Kitani et al. 2004), that are arrested in the G0/G1 phase of the cell cycle, compared to CD4⁺CD25⁻ T cells. TGFβ1 is also known to up-regulate and activate Smad2/Smad3/Smad4 transcription factors (ten Dijke and Hill 2004), which are in turn responsible for the expression of CdkI's that regulate cell cycle progression at the G1 phase of the cell cycle (Zhang and Lin 1997; Moustakas and Kardassis 1998; Ho, de Guise et al. 2004). However, TGFβ1-dependent expression of CdkI's in Treg cells is not yet determined. TGFβ1 by itself or along with other superfamily members, such as BMP's, and activins, could induce p21^{cip1} transcription, an event that is thought to link cell growth inhibition, and/or apoptosis depending upon the target cell type (Zauberman, Oren et al. 1997; Ishisaki, Yamato et al. 1998; Jernvall, Aberg et al. 1998; Ghosh-Choudhury, Ghosh-Choudhury et al. 2000; Franzen and Heldin 2001). It is not clear if TGFβ1-induced p21^{cip1} expression is linked to cell proliferation. Transcriptional induction of p21^{cip1} responsiveness to TGFβ1 could be mediated either by the SMAD pathway or by the Ras/mitogen activated protein kinase (MAPK) pathway (Kivinen, Tsubari et al. 1999; Pardali, Kurisaki et al. 2000; Pouliot and Labrie 2002). p21^{cip1} promoter possess binding site/s for transcription factors of the Sp1 family along with Smad's (Moustakas, Pardali et al. 2002) which is targeted by both Smad- and

Ras/MAPK-mediated pathways. Smad complex enhances the binding affinity of Sp1 transcription factor, by direct interaction with the p21^{cip1} promoter DNA along with physical interaction with Sp1 (Pardali, Kurisaki et al. 2000). It has also been demonstrated that TGFβ1 activates Jun-family members that could also induce p21^{cip1} expression by binding to the GC rich motifs on the proximal promoter, by physical interaction with Sp1 transcription factor. TGFβ1 induces phosphorylation of cyclic AMP response element binding proteins (CREB) via the MAPK pathway. IL-2R alpha gene possesses binding sites for CREB/ATF, AP1 and Smad3 transcription factors. TGFβ1 along with anti-CD3 and CD28 stimulation could induce IL-2Ralpha expression (CD25). On the other hand, anti-apoptotic gene bcl-2 can be activated transcriptionally by the transcription factor cAMP-response element binding protein (CREB) through a CRE site in the 5'-flanking region (Wilson, Mochon et al. 1996). Therefore, the TGFβ1 along with TCR stimulation in Treg cells could induce the expression of CD25, bcl-2, and also cell CdkI's such as p21^{cip1}. Thus, TGFβ1 pathway may be a useful mechanism to explain the phenomenon of cell growth inhibition yet programmed cell survival in Treg cells.

I. Regulation of apoptotic and anti-apoptotic proteins

The fate of the cell in terms of apoptosis vs. cell survival programming may also depend upon the stage in which IL-2 is down regulated. If IL-2 is deprived at G0/G1 to S phase transition of the cell cycle, which is solely dependent on IL-2, then this, may result in cell survival program (anergy) instead of apoptosis (Krummel and Allison 1996). In contrast, if the cell exited the G1 phase and is at the S or G2 phase of the cell cycle, which also

depends on IL-2, deprivation of IL-2 at this stage of the cell cycle may lead to apoptosis (Krummel and Allison 1996). As CD4⁺CD25⁺ Treg cells are arrested in the G0/G1 phase of the cell cycle, IL-2 may serve as a core component to understand the intracellular molecular mechanisms regulating anergic state and suppressor function of this unique CD4⁺ T cell population.

The well known apoptotic pathways reported in mammalian cells are the death receptor (TNF- α R/TNF- α , Fas/FasL) pathway and the mitochondrial pathway (Bcl-family members). The death receptor pathway is activated when extracellular apoptotic ligands bind their receptors on the cells that receive extrinsic apoptotic signals. Upon receiving a death signal, intra-cellular death domains are activated which in turn activate pro-caspase 8 and/or caspase10, promoting an activation signal for the recruitment of caspase 3 (Boatright and Salvesen 2003). Caspase 3 activates caspase activating DNase (CAD) by cleaving its inhibitor (ICAD) followed by ubiquitination of DNA (Yang, Liu et al. 1997). This death receptor pathway is important in maintaining the homeostasis of the immune system. On the other hand, the mitochondrial pathway is activated via the induction of intrinsic apoptotic signals. During this process mitochondria are permeabilized by pro-apoptotic proteins of the Bcl-family such as bax, and bid, releasing cytochrome C and other mitochondrial proteins (eg. proteases Apaf-1) (Gupta 2003). Cytochrome C binds to apoptotic protease-activating factor (Apaf-1) leading to the activation of caspase-9 and the intrinsic death pathway (Yang, Liu et al. 1997). Anti-apoptotic proteins such as Bcl-2 and Bcl-xL prevent pore formation, thus blocking apoptosis. Therefore, these pathways could be modulated by extracellular signaling cascades via TGF β 1 and intracellular

signaling cascades via MAP kinases.

The fate of the cell exposed to apoptotic stimulus is determined by the balance between the two groups of Bcl-family members. Expression of bcl-2 is known to be an important event in the regulation of cell survival (Adams and Cory 1998). The apparent contradictory responses of different cell types to TGF β 1 signaling may depend upon the relative expression of bcl-2 and bax, as the ratio of bcl-2 to bax in a cell appears to determine survival or apoptosis (Xiao, Oppenlander et al. 2001). A case in point is the different responses of the precursor CD34⁺CD38⁻ and the committed CD34⁺CD38⁺ cell to TGF β 1 signaling. The CD34⁺CD38⁻ precursor with relatively low levels of bcl-2 respond to TGF β 1 by undergoing apoptosis, whereas the committed CD34⁺CD38⁺ with higher levels of bcl-2 are not sensitive to TGF β 1-induced apoptosis (Xiao, Oppenlander et al. 2001).

TGF β , an immune regulatory cytokine that plays a major role in homeostasis, has been reported to be involved in the inhibition of Fas or TNF α R mediated cell death (Schlapbach, Spanaus et al. 2000). Recent reports suggested that TGF β 1 induces the expression of Flice-inhibitory protein (FLIP) that could associate with Fas activated death domains, thus preventing the activation of the caspases mediated pathway (Irmeler, Thome et al. 1997). FLIP is also known to be involved in the activation of transcription factors like NF κ B and ERK inducing cell survival signals (Kataoka, Budd et al. 2000). TGF β 1 can also counteract the cytolytic effect caused by TNF α . TGF β 1 in combination with TNF α has been reported to reduce Fas expression, induce NF κ B activation, and

upregulate anti-apoptotic protein bcl-XL in hepatic stellate cells (Saile, Matthes et al. 2001). On the other hand, in several cancer cell lines, TGF β 1 along with TNF α induced apoptosis via the down regulation of anti-apoptotic protein, bcl-2 (Schuster, Bender et al. 2003). TGF β 1 could induce bcl-2 expression via MAP kinase pathway. As mentioned earlier TGF β 1 induces the expression of CREB transcription factor, and bcl-2 promoter possesses binding sites for CREB (Wilson, Mochon et al. 1996). Thus the fate of the cell, programmed to die or survive, may depend upon TGF β 1 and TNF α - mediated regulation of different Bcl-family members.

J. Regulation of FIV replication in CD4⁺CD25⁺ Treg cells in Feline model

Feline Immunodeficiency Virus (FIV), similar to Human Immunodeficiency Virus (HIV), is a lentivirus that belongs to the Retroviridae family that cause's chronic and progressive acquired immunodeficiency syndrome in domestic cats. FIV has been classified into five clades (clades A-E) similar to HIV with at least nine clades (clades A-D, F-H, J and K). Similar to HIV infection, the immunopathogenesis of FIV infection in cats is characterized by CD4⁺ T cell depletion, reduction in CD4/CD8 ratios, decreased responsiveness to T cell mitogens, and lymph node T cell apoptosis (Lawrence, Callanan et al. 1992; Guiot, Rigal et al. 1997; Tompkins, Bull et al. 2002). The anti-FIV cytotoxic activities, mediated by CD8⁺ T cells that recognize whole virus, FIV gag and env proteins, are initiated during the early stages of infection and persist at the asymptomatic phase even with the loss of CD4⁺ T cells similar to HIV-infection. Although the clinical stages of FIV and HIV infections are well understood, the mechanisms leading to the

reduction of CD4⁺ T cell numbers and functions in these two viral infections are not effectively established. Therefore, FIV model in cats can serve as an efficient small animal AIDS model for humans. Thus, the FIV model may help in exploring the mechanisms involved in regulating FIV-replication by Treg cells, which may be comparable to HIV regulation by T regulatory cells in human subjects.

It has been reported that feline CD4⁺CD25⁺ Treg cells possess the major characteristics of murine and human Treg cells in that they are anergic to mitogenic stimulation, and relatively resistant to activation induced cell death (Vahlenkamp, Bull et al. 2004). FIV-infection results in chronic activation of Treg cells (Joshi, Vahlenkamp et al. 2004; Vahlenkamp, Bull et al. 2004). It has been reported that only activated Treg cells can non-specifically suppress the proliferation of other T cells. Treg cells from FIV-infected cats are endogenously activated and can suppress T cell proliferation without further activation in vitro (Vahlenkamp, Bull et al. 2004).

The ability of CD4⁺CD25⁺ Treg cells to be anergic, yet programmed to cell survival as opposed to apoptosis are unique features that may be essential long-term immunosurveillance and maintenance of peripheral self-tolerance, which may depend on altered CdkI's and anti-apoptotic proteins (bcl-2) expression. Because of this long-term apoptosis resistant phenotypic nature, CD4⁺CD25⁺ Treg cells can serve as a stable reservoir for productive FIV/HIV infection. It has been shown that FIV preferentially replicates in CD4⁺CD25⁺ Treg cells compared to CD4⁺CD25⁻ T cells (Joshi, Vahlenkamp et al. 2004). Interestingly, the transcription factors like NFκB and AP1 involved in the

regulation of IL-2 gene expression are also known to regulate transcription of HIV/FIV genes (Joshi, Garg et al. 2005). HIV/FIV long terminal repeats also possess binding sites for several transcriptional factors like AP1, ATF, CEBP, and Sp1 (Joshi, Garg et al. 2005). The activation of the transcription factors such as AP1, CEBP, and ATF were demonstrated to correlate with FIV-transcription (Joshi, Garg et al. 2005). As mentioned earlier, TGF β 1 signaling has been known to be involved in regulating several of these transcriptional factors such as Sp1, ATF, and NF κ B. Therefore, TGF β 1 signaling may be one of the mechanisms which enable potent suppressor Treg cells to maintain anergy, down-regulate IL-2, yet be programmed for survival, and also serve as excellent reservoirs for FIV/HIV replication.

K. Based on the above literature, the research was focused on the following specific aims.

Specific Aim 1 To determine if feline Treg cells are functionally anergic. If so, what are the phenotypic molecules (mainly the CdkI's) that could be responsible for the maintenance of anergic state in CD4⁺CD25⁺ Treg cells, compared to CD4⁺CD25⁻ T cells?

Specific Aim 2 To determine the activation status of Treg cells from FIV-infected cats compared to the control cats using phenotypic surface markers such as CD62L-selectin.

Specific Aim 3 To determine if Treg cells are more resistant to activation induced apoptosis compared to their counterparts CD4⁺CD25⁻ T cells. If so, how is the expression profile of pro-apoptotic and anti-apoptotic proteins in CD4⁺CD25⁺ Treg cells compared to CD4⁺CD25⁻ T cells?

Specific Aim 4 To determine the surface expression profile of TGFβRII on the two purified CD4⁺ T cell subsets.

Specific Aim 5 To determine if TGFβ converted feline CD4⁺CD25⁻ T cells are functionally and phenotypically anergic similar to that of naturally occurring CD4⁺CD25⁺ Treg cells.

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3. CHAPTER-1

**IMMUNE SUPPRESSOR FUNCTION OF CD4⁺CD25⁺ T regulatory
CELLS IN FIV-INFECTED CATS IS ASSOCIATED WITH OVER-
EXPRESSION OF CELL SURFACE TGFβ1 AND TGFβRII AND UP-
REGULATION OF CYCLIN DEPENDENT KINASE INHIBITORS**

ABSTRACT

Cell cycling from G0/G1 to the S phase is regulated by a family of cyclin-dependent kinases (Cdk) and their inhibitors (CdkI's). CD4⁺CD25⁺ T regulatory cells (Treg) are arrested in the G0/G1 phase of the cell cycle and fail to produce IL-2 and proliferate in response to TCR engagement. The expression profile of the Cdk and CdkI genes in CD4⁺CD25⁺ Treg cells and the role they play in maintaining the anergic state has not been studied. We previously reported that CD4⁺CD25⁺ Treg cells from FIV-infected but not uninfected cats are activated in-vivo and possess potent immuno-suppressor function in the absence of additional in-vitro stimulation. In this study we assessed the expression profile of the CdkI's in Treg cells from FIV-infected and uninfected control cats. We demonstrated that the Cdk inhibitors p21^{cip1}, p27^{kip1}, and p16^{ink4} are up-regulated in CD4⁺CD25⁺ Treg cells compared to CD4⁺CD25⁻ T cells from FIV-infected cats. No significant difference in CdkI expression was detected in unstimulated CD4⁺CD25⁺ and CD4⁺CD25⁻ T cells from uninfected cats. Upon stimulation with ConA or LPS alone, p21^{cip1} expression was enhanced in CD4⁺CD25⁺ Treg cells but not in CD4⁺CD25⁻ T cells from uninfected cats, and the presence of rhIL-2 blocked this effect. Up-regulation of CdkI's in CD4⁺CD25⁺ Treg cells from FIV-infected cats is associated with increased expression of TGFβ1 and TGFβRII on the cell surface. We also demonstrated that feline CD4⁺CD25⁺ Treg cells from FIV-infected cats but not control cats are able to suppress the proliferation of activated CD4⁺CD25⁻ T cells. Taken together, these findings suggest that feline CD4⁺CD25⁺ Treg cells from FIV-infected cats, in contrast to CD4⁺CD25⁺ Treg cells from control cats are functionally and phenotypically activated in vivo, and that this activation may be regulated by the TGFβ1/TGFβR signaling pathway.

Keywords: CD4⁺CD25⁺ T regulatory cells; TGFβ1; Cdk; CdkI; FIV

INTRODUCTION

According to the classical model of T cell activation, productive stimulation of T cells requires two major signals. The first is the engagement of MHC-bound peptide on APC with the TCR and the second is the interaction of B7 on the APC with the T cell co-stimulatory receptor CD28 (Greenfield, Nguyen et al. 1998; Harris and Ronchese 1999). Under normal conditions, B7 receptors on APC initially engage the CD28 receptor on CD4⁺ T cells transducing a signal that up-regulates IL-2 production, an essential cytokine for transit out of the G0/G1 stage of the cell cycle, induction of cell proliferation and the initiation of an immune response (Powell, Ragheb et al. 1998). If T cells are stimulated only through the TCR in the absence of co-stimulation, this partial activation leads to a state of long-lasting, partial or total unresponsiveness, termed anergy (Powell, Ragheb et al. 1998). In addition to CD4⁺ T cells that can be rendered anergic by TCR engagement in the absence of B7-CD28 co-stimulation, a second population of CD4⁺ T cells, the CD4⁺CD25⁺ Treg cells appear to be naturally anergic, arrested in the G0/G1 phase of the cell cycle. Similar to anergic CD4⁺ T cells, CD4⁺CD25⁺ Treg cells fail to produce IL-2 and proliferate in response to TCR engagement by antigens, but respond to polyclonal stimuli such as IL-2 (Powell, Ragheb et al. 1998; Lechler, Chai et al. 2001). Naturally occurring CD4⁺CD25⁺ Treg cells, comprising 5-10% of the circulating CD4⁺ T cell population, were originally defined as regulatory T cells that maintain peripheral self-

tolerance (Maloy and Powrie 2001; Shevach 2002). More recent studies have demonstrated that Treg cells also play a major role in modulating the magnitude and duration of protective T cell responses to pathogens (Read, Malmstrom et al. 2000; Iwashiro, Messer et al. 2001; Belkaid, Piccirillo et al. 2002; Hori, Carvalho et al. 2002; Montagnoli, Bacci et al. 2002; Hisaeda, Maekawa et al. 2004; Belkaid and Rouse 2005). These pathogen-induced Treg cells, similar to natural T cells, are anergic and possess potent immunosuppressor function once activated by TCR engagement and/or certain nonspecific stimuli (Belkaid and Rouse 2005). While anergic T cells, including pathogen-induced $CD4^+CD25^+$ Treg cells, are known to be arrested at the G0/G1 phase of the cell cycle, there is little information on what receptors or intracellular signaling proteins maintain IL-2 suppression and the anergic state of Treg cells.

Progression through the G0/G1 to S phase of the cell cycle is regulated by several cyclins and cyclin-dependent kinases (Cdk) whose activities are in turn controlled by their respective inhibitors (CdkI's). During G1/S phase progression, cyclin D (D1-D3 in most cells and only D2-D3 in lymphocytes) are activated in mid G1 phase followed by cyclin E and cyclin A in G1/S boundary (Morgan 1995; Reed 1997; Sherr and Roberts 1999). In response to mitogen signals, Cyclin D is induced and associates with Cdk4 and Cdk6 (Sherr and Roberts 1999), whereas cyclin A associates with Cdk2. The activity of cyclin-Cdk complexes is repressed by CdkI's, which constrain entry into S phase (Reed 1997). The CdkI's that govern these events have been divided into two groups based on their structure and Cdk targets. The first class includes INK4 proteins (p16, p15, & p18) and consists of multiple ankyrin repeats, which bind to only Cdk4 and Cdk6 (Sherr and

Roberts 1999). The second class of inhibitors includes p21^{cip1}, p27^{kip1}, p53 and p57, which contain characteristic motifs within their amino terminal moieties that enable them to bind both cyclins and Cdk subunits (Sherr and Roberts 1999). As p21^{cip1} is capable of binding with essentially all the cyclin-cyclin-dependent kinase complexes, it plays a major role in regulating T cell cycling and could be a principle factor in maintaining anergy. In support of this, p21^{cip1} knockout mice rapidly lose tolerance to nuclear antigens and develop a lymphoproliferative autoimmune-like disease reminiscent of systemic lupus in humans (Balomenos, Martin-Caballero et al. 2000).

While numerous factors are involved in induction of cell-cycle arrest, cytokines, most notably TGFβ1 may play a major role in inducing cell growth arrest in immune cells by up-regulation of CdkI's (Dennler, Goumans et al. 2002; Gartel and Tyner 2002). TGFβ1 has been shown to induce G0/G1 cell cycle-arrest via transcriptional activation of p15^{ink4}, p21^{cip1} and p27^{kip1} expression in a number of cells including T and B lymphocytes (Kee, Rivera et al. 2001; Massague 2004; Wolfrain, Walz et al. 2004). Although there is little information on how TGFβ1 activates CdkI's expression, recent studies demonstrate that the p21^{cip1} promoter contains sequences capable of binding SMAD4, a critical downstream element in the TGFβ1 signaling pathway (Yamato, Hashimoto et al. 2001). Cell surface expressed TGFβ1 has also been associated with activated CD4⁺CD25⁺ Treg cells, although it's role in Treg function is controversial (Maloy and Powrie 2001; Shevach 2002).

Recent studies have demonstrated that feline CD4⁺CD25⁺ Treg cells possess all the functional characteristics of Treg cells described in humans and rodents in that they are anergic, arrested at the G0/G1 phase of the cell cycle, and possess potent immunosuppressive activity when activated (Vahlenkamp, Tompkins et al. 2004). Moreover, we recently reported that CD4⁺CD25⁺ Treg cells from FIV-infected cats, in contrast to Treg cells from uninfected control cats, are activated in-vivo and fully armed for potent contact-dependent immunosuppressor function (Vahlenkamp, Tompkins et al. 2004). To determine what intracellular factors might regulate CD4⁺CD25⁺ Treg cell anergy and suppressive function and what affect FIV-infection of cats might have on these factors, this study focused on the expression profiles of the CdkI's in CD4⁺CD25⁺ and CD4⁺CD25⁻ T cell subsets in FIV-infected and uninfected cats. We demonstrate that in-vivo activated, anergic feline CD4⁺CD25⁺ Treg cells from FIV-infected cats are functionally and phenotypically activated, as indicated by the enhanced expression of CdkI's and potent suppressor function. Moreover, Treg cells from FIV-infected cats over-express TGFβ1 and TGFβRII on their surface, suggesting that this signaling pathway in addition to up-regulating CdkI's and maintaining the anergic state of Treg cells, may play a role in the immunosuppressive function of these cells.

MATERIALS AND METHODS

Animals and Viral Infection

The cats used for this study were specific pathogen free (SPF) obtained from Liberty Labs (Liberty Corners, NJ) or Cedar River Laboratory (Mason City, IA) and housed at the Laboratory Animal Resource Facility at the College of Veterinary Medicine, North Carolina State University. Cats were inoculated with the NCSU1 isolate of FIV, a pathogenic clade A virus (English, Nelson et al. 1994), as described by Bucci et al (Bucci, English et al. 1998). The samples were collected from FIV⁺ cats infected for more than 3yrs and from uninfected control cats, 4 to 6 years of age. FIV-infected cats were housed separately from uninfected control cats. FIV-infection was confirmed by immunoblot analysis and provirus detection by PCR using primers specific for the FIV-p24 encoding sequence.

Cell Collection

Whole blood was collected by jugular venipuncture into EDTA Vacutainer tubes (Becton-Dickinson, Franklin Lakes, NJ) and PBMC isolated by Percoll (Sigma-Aldrich, St. Louis, MO) density gradient centrifugation as previously described (Tompkins, Ogilvie et al. 1987). LN cells were obtained from LN biopsies and single-cell suspensions were prepared by gently injecting sterile PBS into the tissue using 18G needle repeatedly,

until the cells were released from the tissue. Cell counts and viability were determined by trypan blue dye exclusion. Viability was always >90%.

CD4⁺ T cell subset separation

To obtain single lymphocyte subsets, CD4⁺CD25⁺ and CD4⁺CD25⁻ T cells derived from PBMC and pooled LN cells were either enriched by biomagnetic bead depletion and selection or by fluorescent activated cell sorting (FACS) (MoFlo; DakoCytomation, Fort Collins, CO). Biomagnetic bead separation was performed using goat anti-mouse IgG-coated beads (Dynabeads M-450; Dynal, Great Neck, NY) as described by Bucci et al. (Bucci, English et al. 1998). PBMC and LN were depleted of B cells with anti-CD21 mAb (Sertec, Raleigh, NC)-coated beads, and CD8⁺ cells were depleted with anti-CD8 (mAb 3.357 (Tompkins, Gebhard et al. 1990))-coated beads. The CD4⁺ cell population was then enriched for CD25⁺ cells by positive selection using anti-feline CD25 mAb (9F23; kindly provided by K. Ohno (University of Tokyo, Tokyo, Japan)). The enriched CD4⁺CD25⁺ or CD4⁺CD25⁻ T cells were analyzed for purity by flow cytometric analysis and were greater than 90% pure. The purity of the FACS sorted CD4⁺CD25⁺ and CD4⁺CD25⁻ T cells were greater than 95%. In both cases, viability of the purified CD4⁺ T cell subsets was determined by trypan blue dye exclusion and was always >90%. All assays were based on viable cell numbers.

Flow cytometry analysis

At least 5×10^5 PBMC or purified cell populations were stained for surface expression of CD4 and CD25 markers using APC-conjugated anti-CD4 (mAb 30A (Tompkins, Gebhard et al. 1990)), FITC-conjugated anti-CD25 mAb (mAb 9F23), and PE-conjugated anti-CD62L selectin (BD-pharmagen). Monoclonal antibodies for TGF β 1 (mouse anti-human, mAb 240 clone 9016) and TGF β RII (mouse anti-human FAB2HIP clone 25508) were obtained from R and D. Systems. Data were acquired on a FACS caliber flow cytometer (BD Biosciences, Mountain View, CA). Lymphocytes were gated based on forward vs. side scatter, and 20,000 gated events were acquired and stored list-mode fashion for analysis using Cell Quest software.

T cell stimulation assays

T cell subsets were cultured in RPMI 1640 medium containing 10% heat-inactivated FBS, 1% penicillin-streptomycin, 1% sodium bicarbonate, 1% sodium pyruvate, 1% L-glutamine, and 1 mM HEPES buffer. For ConA stimulation, 1×10^5 viable cells/well were cultured in a round-bottom 96-well plate in the presence or absence of 5 μ g/ml Con A and/or 100 U/ml recombinant human IL-2 (rhIL-2) (National Institutes of Health AIDS Research and Reagent Program, Rockville, MD). All assays were run in triplicate. After 2 days of incubation, cells were pulsed with 1 μ Ci of [3 H]TdR/well and harvested 16hrs later using a Filtermake Harvester (Packard Bioscience, Meriden, CT). [3 H]TdR uptake as a measure of proliferation was determined in cpm using a Top Count NXT Microplate scintillation counter (Packard Bioscience).

Co-culture immunosuppression assay

Purified CD4⁺CD25⁻ T target cells (10⁶ cells/ml) were stimulated for 3hrs with 5 µg/ml ConA, washed twice in RPMI 1640, and plated at 5 x 10⁴ viable cells/well in 96-well plates. Varying numbers of freshly isolated autologous CD4⁺CD25⁺ Treg effector cells were added to the target cells to yield E:T ratios ranging from 0.125:1 to 1:1. The assays were run in triplicate. Effector and target cells were cocultured for 3 days, then pulsed with 1µCi of [³H]TdR/well, and harvested 16hrs later as described above. Percent inhibition of proliferation was determined based on proliferation of CD4⁺CD25⁻ T target cells alone and calculated as follows: percent inhibition = (cpm of target cells alone - cpm of coculture) / cpm of target cells alone.

PCR Analysis

Total RNA was isolated from 1x10⁶-2x10⁶ CD4⁺ T cell subsets using RNA extraction kit (Qiagen) and cDNA was prepared using random oligo(dT) primers (reverse transcription system kit from Promega). p21^{cip1} message expression was detected using feline specific p21^{cip1} primer sequences: 5' GGAGCGATGGAACCTTCGACTTTGT 3' (forward) and 5' GCCGGCGTTTGGATGATAGAAAT 3' (reverse). The expression level of GAPDH was determined to ensure equal amounts of RNA in the samples.

Immunoblotting

Equal numbers of purified CD4⁺ T cell subsets (1x10⁶ – 2x10⁶ cells) were lysed using NP-40 lyses buffer for 16-24hrs at 4C. The concentrations of the proteins were

determined using a spectrophotometer (Biorad Smartspec 3000), 30-50µg of proteins were then resolved on 4-12% SDS-polyacrylamide gels (Novex Inc.). The separated proteins were transferred onto nitrocellulose and immunoblotted with specific mAbs to p21^{cip1}, p27^{kip1}, and p16^{ink4} (Santa Cruz biotechnology Inc) followed by HRP-conjugated secondary antibodies (Santa Cruz Biotechnology Inc). Immunodetection was performed by ECL (Pierce) using HyperFilm ECL. To normalize the result, some blots were stripped with the Western Blot stripping buffer (Pierce) and reprobed with anti-actin mAb (Santa Cruz Biotechnology inc.). Specific antibody conjugated blots were analyzed for specific proteins using Lumicapture analyst camera detection system (Lumi- Imager).

Statistical Analysis

The Mann-Whitney U test was used to assess significance of difference between the medians in determining the CD62L-selectin expression, TGFβ , TGFβRII on the purified CD4⁺ T cell subsets in FIV-infected and control cats. p-values < 0. 05 were considered significant.

RESULTS

Anergic phenotype and suppressor function of CD4⁺CD25⁺ Treg cells from FIV-infected and control cats

We reported previously that while CD4⁺CD25⁺ Treg cells from both FIV-infected and control cats were anergic to immune stimulation, only those from FIV-infected cats

displayed strong suppressor function in the absence of in vitro mitogenic stimulation (Vahlenkamp, Tompkins et al. 2004). To confirm that the CD4⁺CD25⁺ Treg cells from the cats used in this study display this pattern of anergy and suppressor function, CD4⁺CD25⁺ and CD4⁺CD25⁻ T cell subsets were purified by flow cytometry sorting of PBMC from FIV-infected and control cats and assessed for anergy and suppressor function. **Figure 1** shows a typical response of the CD4⁺ T cell subsets in these assays. **Figure 1A** shows the CD4⁺CD25⁻ T cells from FIV-infected and control cats proliferate efficiently in response to ConA stimulation, whereas CD4⁺CD25⁺ Treg cells are anergic to ConA stimulation but are responsive to ConA plus rhIL-2 stimulation. In the case of immune suppressor function, CD4⁺CD25⁺ Treg cells from FIV-infected cats demonstrate a strong dose-dependent anti-proliferative effect against ConA-stimulated CD4⁺CD25⁻ T cells, whereas CD4⁺CD25⁺ Treg cells from control cats suppressed the CD4⁺CD25⁻ T cell proliferative response only at the highest E:T ratio (**Figure 1B**) as has been previously reported for murine and feline Treg cells (Caramalho, Lopes-Carvalho et al. 2003; Vahlenkamp, Tompkins et al. 2004). Thus, the CD4⁺CD25⁺ Treg cells from FIV-infected cats and control cats to be used in these studies displayed the predicted functional characteristics, suggesting that they are functionally activated in FIV-infected cats.

CdkI genes are differentially expressed in PBMC from FIV-infected and control cats

Progression through G0/G1 to S phase of the cell cycle is regulated by cyclins and cyclin-dependent kinases (Cdk), whose activities in turn are regulated by their specific inhibitors, CdkI's. As Treg cells are anergic and arrested in the G0/G1 phase of the cell

cycle (Vahlenkamp, Tompkins et al. 2004), we analyzed the expression of some of the CdkI's in CD4⁺CD25⁺ and CD4⁺CD25⁻ T cells to determine their possible role in maintaining energy. We initially examined the expression of the CdkI's, p21^{cip1}, p27^{kip1}, and p16^{ink4} in total PBMC isolated from FIV-infected and control cats. Whole cell lysates of unstimulated PBMC from control and FIV-infected cats were separated by PAGE, transferred onto nitrocellulose and immunoblotted with specific mAbs to the CdkI's p21^{cip1}, p27^{kip1}, and p16^{ink4}. As shown in **Figure 2**, enhanced expression of p21^{cip1}, p27^{kip1}, and p16^{ink4} proteins was observed in PBMC from FIV-infected cats compared to control cats.

CD4⁺CD25⁺ Treg cells from FIV-infected but not control cats preferentially over-express CdkI proteins and mRNA compared to CD4⁺CD25⁻ T cells

To determine whether CD4⁺CD25⁺ or CD4⁺CD25⁻ T cell subsets are contributing to the differential CdkI expression observed in the total PBMC population of FIV-infected cats and uninfected cats (**Figure 1**), highly purified CD4⁺CD25⁺ and CD4⁺CD25⁻ T cell subsets were assessed for CdkI, cyclin, and retinoblastoma protein expression by immunoblotting. As p21^{cip1} expression is reportedly transcriptionally regulated (Harr, Graves et al. 2005), we also examined p21^{cip1} mRNA expression in the two CD4⁺ T cell subsets from PBMC of FIV-infected and control cats. As shown in **Figure 3A**, there is a high level of constitutive p21^{cip1}, p27^{kip1}, and p16^{ink4} protein expression in CD4⁺CD25⁺ Treg cells compared to CD4⁺CD25⁻ T cells from FIV-infected cats. In contrast, there is no discernable difference in the protein expression of CdkI's in CD4⁺CD25⁺ and CD4⁺CD25⁻ T cells from control cats (data shown for only p21^{cip1}, **Figure 3B**). p21^{cip1}

mRNA was also elevated in CD4⁺CD25⁺ Treg cells as compared to CD4⁺CD25⁻ T cells from FIV-infected cats but not control cats (**Figure 3C**). Thus, p21^{*cip1*}, p16^{*ink4*} and to lesser extent, p27^{*kip1*} are over-expressed in unstimulated CD4⁺CD25⁺ Treg cells from FIV-infected cats. In contrast, cyclin D (**Figure 3D**), and phosphorylated retinoblastoma protein (pRb) (**Figure 3D**) were expressed at elevated levels in CD4⁺CD25⁻ T cells compared to CD4⁺CD25⁺ Treg cells. This suggests that only CD4⁺CD25⁺ Treg cells but not CD4⁺CD25⁻ T cells are arrested in the G0/G1 phase of the cell cycle.

Mitogenic activation of CD4⁺CD25⁺ Treg cells from control cats induces p21^{*cip1*} protein expression

As observed by others with murine Treg cells (Caramalho, Lopes-Carvalho et al. 2003), we reported that feline CD4⁺CD25⁺ Treg cells from FIV-infected cats could be induced to strong suppressor function by mitogen ± IL-2 or by IL-2 itself (Vahlenkamp, Tompkins et al. 2004). To assess the p21^{*cip1*} response to mitogenic stimulation CD4⁺CD25⁺ and CD4⁺CD25⁻ T cells from FIV-infected and control cats were stimulated with ConA ± rhIL-2 and assayed for p21^{*cip1*} protein by western blot analysis. ConA stimulation of CD4⁺CD25⁺ or CD4⁺CD25⁻ T cells from FIV-infected cats had no effect on p21^{*cip1*} expression, whereas p21^{*cip1*} was markedly up-regulated in ConA-stimulated CD4⁺CD25⁺ Treg cells, but not in ConA-stimulated CD4⁺CD25⁻ T cells from control cats (**Figure 4A**). ConA induction of p21^{*cip1*} in CD4⁺CD25⁺ Treg cells from control cats was over-ridden by addition of rhIL2 (**Figure 1A**). As we have reported previously that LPS stimulation of CD4⁺CD25⁺ Treg cells from control cats induced immunosuppressor function, we examined the effect of LPS on p21^{*cip1*} expression. Stimulation with LPS

increased p21^{cip1} expression in CD4⁺CD25⁺ Treg cells from control cats, which was overridden by the addition of rhIL-2 (**Figure 4B**). LPS had little effect on either CD4⁺ T cell subsets from FIV-infected cats (**Figure 4B**). Thus, there is a correlation between previously reported mitogen-induced Treg immuno-suppressor function and induction of p21^{cip1} expression. Interestingly, IL-2 down-regulates both the induced and basal levels of p21^{cip1} expression in CD4⁺CD25⁺ Treg cells, which is consistent with its ability to over-ride anergy (Vahlenkamp, Tompkins et al. 2004).

TGFβ1 is up-regulated on the surface of CD4⁺CD25⁺ Treg cells from FIV-infected cats

CD4⁺CD25⁺ Treg cells from humans and mice secrete high levels of TGFβ1 and express TGFβ1 on their surface when appropriately stimulated (Nakamura, K, JEM, Annunziato 2001; Annku, F, JEM, 2002; Cosmi L, Blood, 2003). In view of our observation that CdkI's are up-regulated in Treg cells from FIV-infected cats and the fact that TGFβ1 upregulates expression of CdkI's and induces G0/G1 cell cycle arrest (Dannlers), we examined the expression of TGFβ1 on the surface of PBMC CD4⁺ T cells from control and FIV-infected cats by flow cytometry. **Figure 5A** shows a significant increase in numbers of CD4⁺CD25⁺ Treg cells expressing cell surface TGFβ1 (39.9%) in FIV-infected cats as compared to TGFβ1 positive CD4⁺CD25⁺ Treg cells (10.4%) from control cats. There is no significant difference in expression of TGFβ1 on CD4⁺CD25⁻ T cells from FIV-infected (16.1%) and control cats (11.4%). Analysis of 6 asymptomatic FIV-infected cats and six age-matched control cats revealed a consistent pattern of increased numbers of CD4⁺CD25⁺ TGFβ1 Treg cells in the FIV-infected cats as

compared to controls (**Figure 5B**). These data support the CdkI expression data and the immunosuppressor function data suggesting that CD4⁺CD25⁺ Treg cells from FIV-infected cats are activated in vivo.

TGFβRII is up-regulated on the cell surface of CD4⁺CD25⁺ Treg cells from FIV-infected cats.

There is considerable data on the expression of TGFβ1 on activated Treg cells. However there is no information on possible changes in TGFβ1 receptor expression. To assess TGFβRII expression on Treg cells, we performed 3-color flow cytometry on purified CD4⁺CD25⁺ and CD4⁺CD25⁻ T cell subsets from FIV-infected and control cats. Similar to observations of increased TGFβ1 positive CD4⁺CD25⁺ Treg cells from FIV-infected cats compared to control cats (**Figure 5**), TGFβRII was preferentially expressed on the surface of CD4⁺CD25⁺ Treg cells from FIV-infected (19%) compared to control (4%) cats (**Figure 6A**). A very small fraction of CD4⁺CD25⁻ T cells from both FIV-infected and control cats expressed cell surface TGFβRII, although significantly more CD4⁺CD25⁻ T cells from FIV-infected cats expressed TGFβRII than those from control cats (**Figure 6A**). Flow cytometric analysis of 6 asymptomatic FIV-infected cats and 6 age-matched controls confirmed that significantly more CD4⁺CD25⁺ and CD4⁺CD25⁻ T cells from FIV-infected cats expressed cell surface TGFβRII (**Figure 6B**) as compared to their counterparts from control cats. There was no significant difference in TGFβRII expression between CD4⁺CD25^{bright} and CD4⁺CD25^{dim} from FIV-infected cats (**Figure 6C**). Whether these numbers of TGFβRII positive cells accurately reflect the actual numbers of TGFβRII positive T cells, particularly in the FIV-infected cats is not clear

from these data, as the high level of cell surface TGF β 1 expression could block antibody binding to the TGF β RII. However, 3- and 4-color flow cytometry analysis revealed that 80-90% of CD4⁺CD25⁺TGF β 1⁺ T cells also expressed TGF β RII (data not shown) suggesting that the TGF β RII and its ligand are co-expressed on Treg cells.

L-Selectin expression in CD4⁺CD25⁺ vs. CD4⁺CD25⁻ T cells from peripheral blood of FIV-infected and control cats

As shown in **Figures 2, 3, and 4**, we were able to consistently detect p21^{cip1} in unstimulated CD4⁺CD25⁻ T cells from control cats. However, we were not able to detect this protein in CD4⁺CD25⁻ T cells from FIV-infected cats. As CdkI's generally accumulate in G0/G1 resting cells but are down regulated following activation (Balomenos, D, Imm Today, 2000), we speculated that the differential expression of p21^{cip1} in CD4⁺CD25⁻ T cells between FIV-infected and control cats may be due to the activation state of the cells. Loss of L-selectin (CD62L) expression on lymphocytes is a useful phenotypic marker for T cell activation (Gebhart). Therefore, we analyzed the CD4⁺ T cell subsets from FIV-infected and control cats for the expression of CD62L-selectin. **Figure 7A** shows a typical 3-color analysis for the expression of CD62L on CD4⁺CD25⁺ and CD4⁺CD25⁻ T cells from a FIV-infected and control cats. Greater than 80% of the CD4⁺ cells, whether or not they expressed CD25, were CD62L⁺ in the control cat. In contrast, only 41% of the CD4⁺CD25⁻ T cells and 15% of the CD4⁺CD25⁺ Treg cells from the FIV-infected cat expressed CD62L. Analysis of data from a number of cats consistently demonstrate a difference in CD62L expression on CD4⁺ subsets

between FIV-infected and normal cats (**Figure 7B**), indicating a large percent of the cells in both CD4⁺ subsets are activated in FIV-infected cats. As reported in murine and human models CD4⁺CD25^{bright} cells express elevated levels of CD62L-selectin in FIV-infected cats but not control cats compared to CD4⁺CD25^{dim} T cells (**Figure 7C**). The failure to detect p21^{cip1} in CD4⁺CD25⁻ T cells from FIV-infected cats as opposed to CD4⁺CD25⁻ T cells from control cats may be the result of T cell activation in FIV-infected cats.

DISCUSSION

Treg cells are characterized by G0/G1 cell cycle arrest and are anergic to immune activation, as manifested by their inability to produce IL-2 and exit the G0/G1 stage of cell cycle in response to antigen-specific TCR engagement (Powell, Ragheb et al. 1998). Another major characteristic of CD4⁺CD25⁺ Treg cells described in human and rodent species is their ability, once activated through their TCR, to suppress proliferative responses of other antigen- or mitogen-stimulated CD4⁺ or CD8⁺ T cells (Thornton and Shevach 1998; Jackson, DeLoose et al. 2002). We recently described CD4⁺CD25⁺ Treg cells in the cat possessed all the major phenotypic and functional characteristic of Treg cells described in other species i.e. they are G0/G1 cell cycle arrested, IL-2 deficient and anergic to mitogenic stimulation, and when activated capable of suppressing IL-2 production and proliferation of activated T cells (Vahlenkamp, Tompkins et al. 2004).

The hallmark of naturally occurring Treg cells, including those in normal cats is that they require activation by antigen-TCR engagement or a potent mitogen before they can display immunosuppressive function (Thornton and Shevach 1998). In marked contrast to natural Treg cells, CD4⁺CD25⁺ Treg cells from cats chronically infected with the feline AIDS lentivirus (FIV) are activated in-vivo and displayed strong T cell immunosuppressive function in the absence of in-vitro stimulation (Vahlenkamp, Tompkins et al. 2004). In this study we confirmed that CD4⁺CD25⁺ Treg cells from FIV-infected cats, as opposed to normal cats, are functionally immunosuppressive and that this activation phenotype is associated with up-regulation of the Cdk inhibitors p21^{*cip1*}, p27^{*kip1*}, and p16^{*ink4*} and over-expression of TGFβ1 and TGFβRII on their surface.

Induction and maintenance of T cell anergy due to inappropriate TCR signaling has been associated with over-expression of the cyclin-dependent kinase inhibitors p21^{*cip1*} and p27^{*kip1*} (Jackson, DeLoose et al. 2002; Wolfrum, Walz et al. 2004). A direct relationship between p27^{*kip1*} over-expression and immunological unresponsiveness was demonstrated in p27^{*kip1*}-transfected murine and human CD4⁺ T cells (Boussiotis VA, 2000). As CD4⁺CD25⁺ Treg cells, unlike Th cells are naturally anergic, it was of interest to determine if the CdkI's were constitutively up-regulated in these cells or required an induction signal for expression.

In this study we demonstrated that the CdkI's, p21^{*cip1*}, p16^{*ink4*}, and p27^{*kip1*} proteins are expressed at elevated levels in the total PBMC of FIV-infected cats compared to control cats. Upon further examination of CD4⁺ T cell subsets, we observed an increase in

p21^{*cip1*}, p27^{*kip1*}, and p16^{*ink4*} expression in the CD4⁺CD25⁺ Treg cells compared to CD4⁺CD25⁻ T cells. Interestingly, up-regulation of CdkI's in CD4⁺CD25⁺ Treg cells was restricted to those cells from FIV-infected cats.

As these cats are FIV-infected and chronically antigenemic, we speculated that activation of CdkI's in CD4⁺CD25⁺ Treg cells may be due to immune activation. This is consistent with the observation that CD4⁺CD25⁺ Treg cells from FIV- infected cats but not control cats are functionally immunosuppressive and express enhanced levels of cell surface B7.1 and B7.2 in the absence of in vitro stimulation (Vahlenkamp, Bull et al. 2004; Vahlenkamp, Tompkins et al. 2004). We extend this observation in this study by demonstrating that CD4⁺CD25⁺ Treg cells from FIV-infected cats that displayed increased expression of CdkI's also possessed strong immunosuppressive function in the absence of in-vitro stimulation. Others have demonstrated in-vivo activation of immunosuppressive Treg cells in a number of chronic infections, such as HIV, *leishmania* and malaria (Belkaid 2003; Hisaeda, Maekawa et al. 2004; Tsunemi, Iwasaki et al. 2005); Belkaid and Rouse, *Nat Imm*, 2005), but have not examined the CdkI expression profile. While we have not identified the CD4⁺CD25⁺ Treg cell activation factors in chronically infected cats, a number of studies have demonstrated that naïve or natural Treg cells could be induced to suppressor activity by antigenic or mitogenic stimulation (51). We also reported that CD4⁺CD25⁺ Treg cells from naïve cats could be induced to strong suppressor function by LPS and IL-2 (Vahlenkamp, Tompkins et al. 2004). In this study, we also demonstrated that p21^{*cip1*} expression in CD4⁺CD25⁺ Treg cells from control cats was markedly enhanced by ConA or LPS stimulation. As

expected, these mitogen did not induce p21^{cip1} expression in CD4⁺CD25⁻ T cells from either FIV-infected or control cats. Consistent with the observation that exogenous IL-2 is able to override the anergic state of feline CD4⁺CD25⁺ Treg cells in vitro (Vahlenkamp, Tompkins et al. 2004), we observed that rhIL-2 reduced the mitogen-induced, as well as basal levels of p21^{cip1} expression in T cells.

Over-expression of CdkI's in CD4⁺CD25⁺ Treg cells from FIV-infected cats could result from up-regulation of the TGFβ1 signaling pathway. In many cell types, TGFβ1 causes a G0/G1 cell-cycle arrest by inhibiting Cdk activity via activation of p15^{ink4b} and p21^{cip1} expression (Denmler S). A number of studies have demonstrated that TGFβ1 is up-regulated on the surface of CD4⁺CD25⁺ Treg cells following treatment with appropriate activation stimuli (Nakumura N, Annumziato F, Cosmi; (Hoffmann, Ermann et al. 2002), while others have demonstrated that treatment of CD4⁺CD25⁻ T cells with TCR stimuli in the presence of TGFβ1 converted them to phenotypic and functional Treg cells (Chen W, JEM 198; 2003; Fantini MC, JI, 2004). As the TGFβ1 signaling pathway also is capable of up-regulating p21^{cip1} and p27^{kip1} expression and inducing G0/G1 cell cycle arrest, we examined Treg cell from FIV-infected cats for cell surface TGFβ1 expression. Our data clearly demonstrates that the numbers of CD4⁺CD25⁺ Treg cells expressing TGFβ1 on their surface is increased in FIV-infected cats compared to control cats. Importantly, we were also able to demonstrate that TGFβRII is also up-regulated on the surface of CD4⁺CD25⁺ Treg cells from FIV-infected cats, but not control cats. To the best of our knowledge this is the first report of up-regulation of TGFβRII on the plasma membrane of activated CD4⁺CD25⁺ Treg cells. These data provide a direct link between

TGF β 1/TGF β RII expression, activation of CdkI's and Treg suppressor function. As the biological half-life of TGF β 1-family ligands is short and they usually act in an autocrine manner, it is possible that phenotypic and functional activation of Treg cells in FIV-infected cats is the result of autocrine TGF β 1/TGF β R signaling. This is consistent with the observation that activated Treg cells secrete high levels of biological active TGF β 1 (Nakumar, Maloy and Powers). This is also supported by 3- and 4-color flow cytometry analysis revealing that TGF β 1 and TGF β RII were co-expressed on the subset of CD4⁺CD25⁺ Treg cells. Preliminary data analyzing downstream elements of the TGF β 1/TGF β R signaling pathway shows that SMAD2 is phosphorylated and Rb protein is hypophosphorylated in Treg cells from FIV-infected cats suggesting that the pathway is functional (S. Emani, preliminary data). Whether the TGF β 1/TGF β R signaling pathway and up-regulation of CdkI's play a role in the immunosuppressor function of Treg cells remains to be determined. However, it is interesting that TGF β 1 stimulation of CD4⁺CD25⁻ T cells in the presence of TCR engagement converts them to CD25⁺TGF β 1⁺Foxp3⁺ T cells with potent immunosuppressive properties (Chen, Fantini). We have also demonstrated that feline CD4⁺CD25⁻ T cells could be converted to CD4⁺CD25⁺TGF β 1⁺Foxp3⁺ immunosuppressive Treg cells by stimulation with ConA plus TGF β 1 (C. Petty unpublished data). While TGF β 1 signaling may regulate Treg suppressor activity, it is unlikely to directly relate to up-regulation of CdkI's as we have observed that IL-2 treatment of Treg cells reduces both the basal level and induced p21^{cip1} expression, yet has been shown to substantially increase both feline (Vahlenkamp, Tompkins et al. 2004) and murine (Fu, Zhang et al. 2004) Treg suppressor function.

Whether cell surface expressed TGF β 1 plays a direct role in mediating T cell suppression was not evaluated in this study. The role of TGF β 1 immune suppression by rodent and human Treg cells is controversial. While Nakamura et al. reported that both cell bound and soluble TGF β 1 could mediate T cell immune suppression other have reported that soluble TGF β 1 does not mediate suppression (Takasaki T, Int Imm 1998, Thornton AM, JEM 1998). Additional studies are in progress to determine the possible role of TGF β 1 in mediating feline CD4⁺CD25⁺ Treg immunosuppressor function.

An additional observation of interest from this study is that, while we could consistently measure p21^{*cip1*} in unstimulated CD4⁺CD25⁻ T cells from control cats, it was unusual to detect this protein in CD4⁺CD25⁻ T cells from FIV-infected cats. This could be due to the activation state of the CD4⁺ T cells from FIV-infected cats with long-term asymptomatic infection. It is well established that resting or naïve Th cells constitutively express p27^{*kip1*} protein which is rapidly degraded following activation (Kwon TK, 1997; Appleman LJ, Ji, 2000). Analysis of CD62L-selectin expression on CD4⁺CD25⁻ T cells from naïve cats used in this study indicated that the majority were phenotypically naïve as indicated by high expression of cell surface CD62L-selectin. In contrast, the majority of CD4⁺CD25⁻ T cells from FIV– infected cats that did not express p21^{*cip1*} also did not express cell surface CD62L-selectin, indicating that they are activated. A similar state of chronic T cell immune hyperactivation has been described previously in FIV infected cats (Vahlenkamp, Tompkins et al. 2004) and in HIV–infected patients with established infections (Eggena, Barugahare et al. 2005). Whether and how this state of T cell immune hyperactivation would affect p21^{*cip1*} expression remains to be determined.

Although, this study did not address the factors responsible for activating Treg cells in FIV-infected cats, there are a number of possibilities. While natural autoreactive Treg cells are thought to be activated by TCR engagement by their cognate antigen (Maloy and Powrie), pathogen-induced Treg cells can be activated by a number of mechanisms including antigen-specific TCR engagement as has been suggested for a number of pathogens (Price, Schaumburg et al. 2005). $CD4^+CD25^+$ Treg cells also express a number of TLR's and could be activated by pathogen associated molecular pathogens (PAMPS) such as LPS, as has been reported for murine and feline Treg (Caramalho, Lopes-Carvalho et al. 2003; Vahlenkamp, Tompkins et al. 2004). The cytokine microenvironment is another possibility, as IL-2 has been shown to be a potent inducer of Treg immunosuppressor function (Caramalho, Lopes-Carvalho et al. 2003; Vahlenkamp, Tompkins et al. 2004) and a number of cytokine including $TGF\beta_1$, $IFN\gamma$, and IL-6 have been reported to up-regulate $p21^{cip1}$ expression in other cell types (Gartel and Tyner 2002). Direct interactions between FIV and Treg cells is another possibility, as FIV has been shown to preferentially establish a productive infection in $CD4^+CD25^+$ Treg cells, although the infection does not appear to be cytolytic (Joshi, Garg et al. 2005). HIV has similarly been shown to preferentially establish productive infection in $CD4^+CD25^+$ Treg cells as opposed to $CD4^+CD25^-$ T cells (Borvak, Chou et al. 1995). Studies are currently in progress to explore the possible mechanisms of $CD4^+CD25^+$ Treg cells activation in FIV-infected cats.

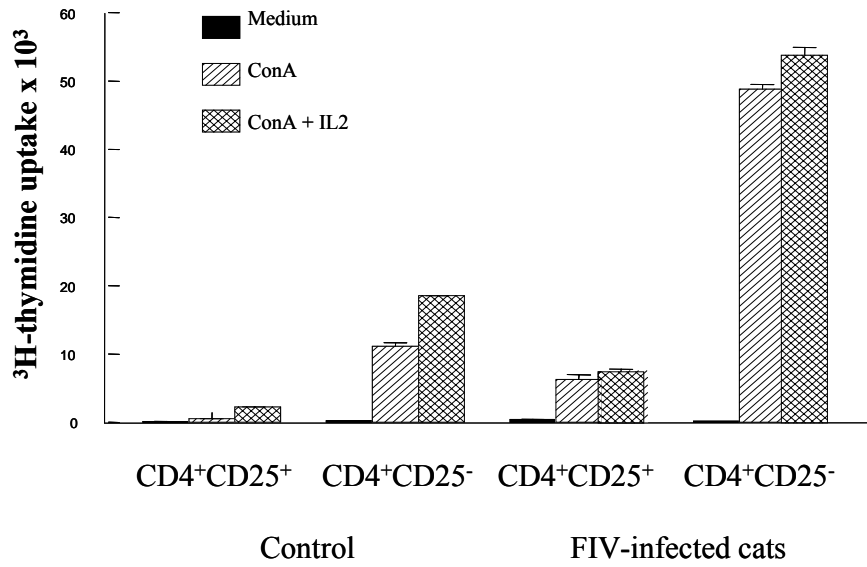
In conclusion, while CD4⁺CD25⁺ Treg cells from both FIV-infected and naïve cats are functionally anergic, as manifested by their inability to produce IL-2 and proliferate upon mitogenic stimulation, only those from FIV-infected cats are activated for strong suppressor function in-vivo. Functional activation of Treg cells from FIV-infected cats correlates with up-regulation of TGFβ1 and TGFβRII on their surface and over expression of the CdkI's, p21^{cip1}, p27^{kip1} and p15^{ink4}. As TGFβ1/TGFβR signaling is known to induce G0/G1 cell-cycle arrest by up-regulation of CdkI's, it can be speculated that the other phenotypic and functional responses observed in Treg cells from FIV-infected cats derive from the TGFβ1/TGFβR signaling pathway. Further, molecular characterization of CD4⁺CD25⁺ Treg cells from FIV chronically infected cats may be helpful in providing experimental basis for understanding the immunopathogenic potential of activated Treg cells and developing immunotherapeutic strategies in treatment for chronic diseases such as HIV/FIV infection.

ACKNOWLEDGEMENTS

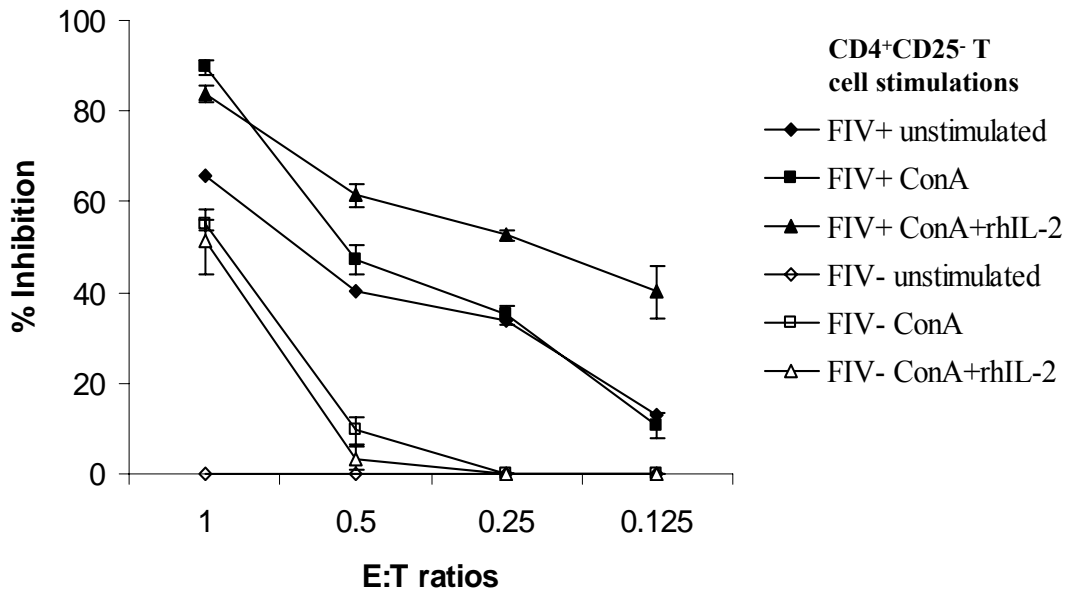
We thank Janet Dow and Debra Anderson for their technical assistance. We thank Dr. Koishi Ohno for providing the anti-CD25 antibody. We also thank NIH AIDS Research and Reference reagent program for providing rHuIL-2. The present study was supported by National Institute of Health grants AI38177, AI 43858, and AI058691.

Figure 1. Suppressor activity and proliferative response of purified CD4⁺ T cell subsets from PBMC of FIV-infected cats to Con A and/or rhIL-2 stimulation. (A) Mo-flow purified CD4⁺ T cell subsets (2×10^5) were incubated with or without ConA (5 μ g/ml) and/or rhIL-2 (100 U/ml) for 48hrs, pulsed with [³H]TdR, and harvested 16hrs later. Bars represent the mean \pm SEM of triplicates. (B) Suppressor activity of freshly isolated CD4⁺CD25⁺ Treg cells on autologous CD4⁺CD25⁻ T cells from FIV-infected cats and control cats. FACS-purified CD4⁺CD25⁻ T target cells were either unstimulated or stimulated with ConA (5 μ g/ml) +/- rhIL-2 (100 U/ml) for 3hrs, then cocultured for 3 days with autologous CD4⁺CD25⁺ Treg cells and were pulsed with [³H]TdR 16hrs prior to the harvest. A constant number of CD4⁺CD25⁻ T target cells (5×10^4 /well) were cultured with varying numbers of CD4⁺CD25⁺ Treg effector cells. Percent inhibition of proliferation was calculated based on proliferation of CD4⁺CD25⁻ T target cells alone ((cpm of target cells alone – cpm of coculture) / cpm of target cells alone). Bars represent mean +/- SEM of triplicate wells for each condition.

A



B



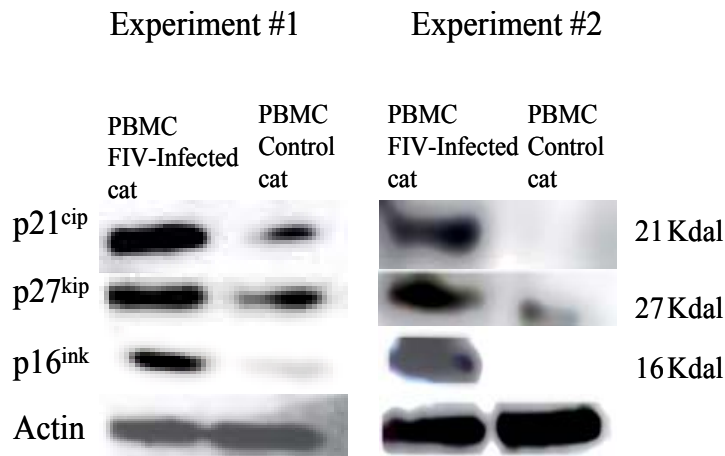


Figure 2. PBMC from FIV-infected cats express elevated levels of CdkI (p21^{cip1}, p27^{kip1}, and p16^{ink4}) proteins compared to control cats. Fresh PBMC (2x10⁶ cells) isolated from FIV-infected and control cats were lysed using NP-40 lysis buffer and separated on SDS-PAGE followed by immunoblotting with specific CdkI primary antibodies and HRP-conjugated goat anti-mouse IgG1 secondary antibody and detected by chemiluminescence. The blots were stripped using stripping buffer and reprobed with actin primary antibody and HRP-conjugated goat anti-mouse IgG1 secondary antibody followed by chemiluminescence detection. These data are representative of gels from two separate experiments, each with an FIV-infected and control cats.

Figure 3. CdkI mRNA and protein expression in the purified CD4⁺ T cell subsets from FIV-infected and control cats. Equal number of purified CD4⁺CD25⁺ and CD4⁺CD25⁻ T cells (2×10^6) were lysed with NP-40 lysis buffer to obtain total cellular protein for the detection of p21^{*cip1*}, p27^{*kip1*}, p16^{*ink4*}, cyclin D, pRb expression. Total cell lysates from two individual infected cats (**A,D**) and one control cat (**B**) were separated by SDS-PAGE and analyzed by immunoblot using specific primary antibodies against CdkI's (p21^{*cip1*}, p27^{*kip1*}, and p16^{*ink4*}), followed by HRP-conjugated goat anti-mouse IgG1 secondary antibody and detected by chemiluminescence. The blots were stripped using stripping buffer and reprobbed with actin primary antibody and HRP-conjugated goat anti-mouse IgG1 secondary antibody followed by chemiluminescence detection. (**C**) mRNA expression of p21^{*cip1*} and GAPDH3 in FIV-infected and control cats.

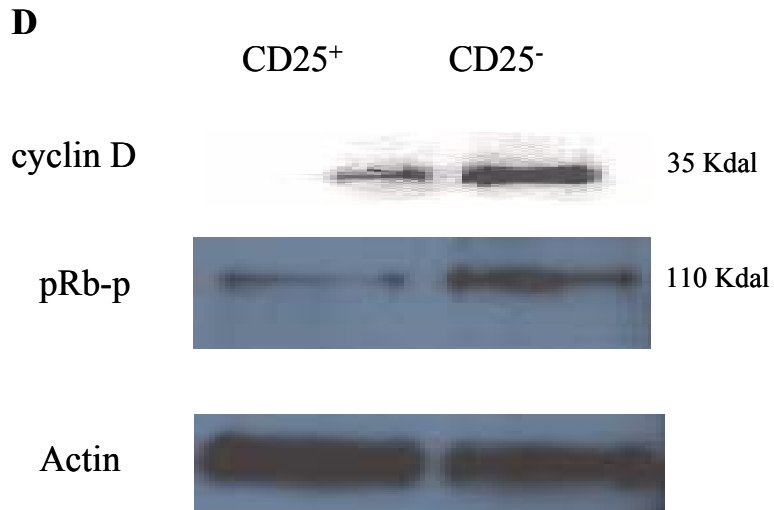
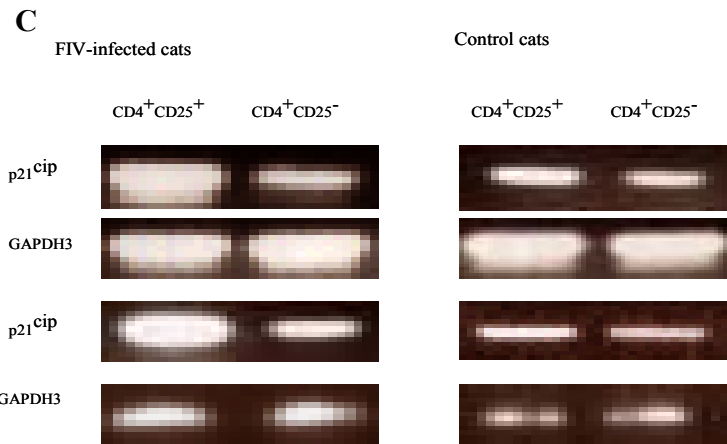
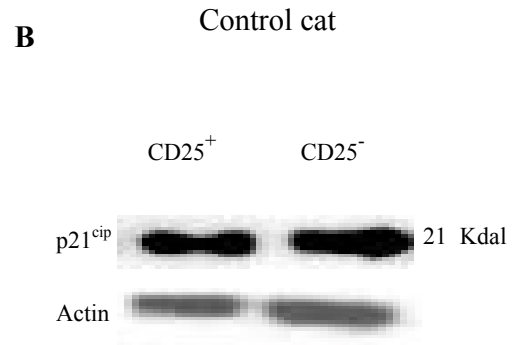
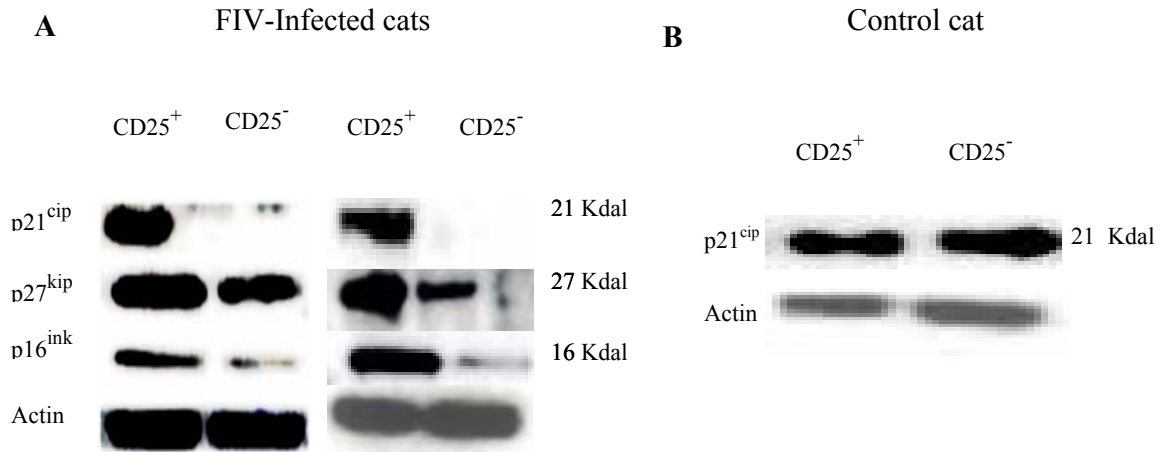
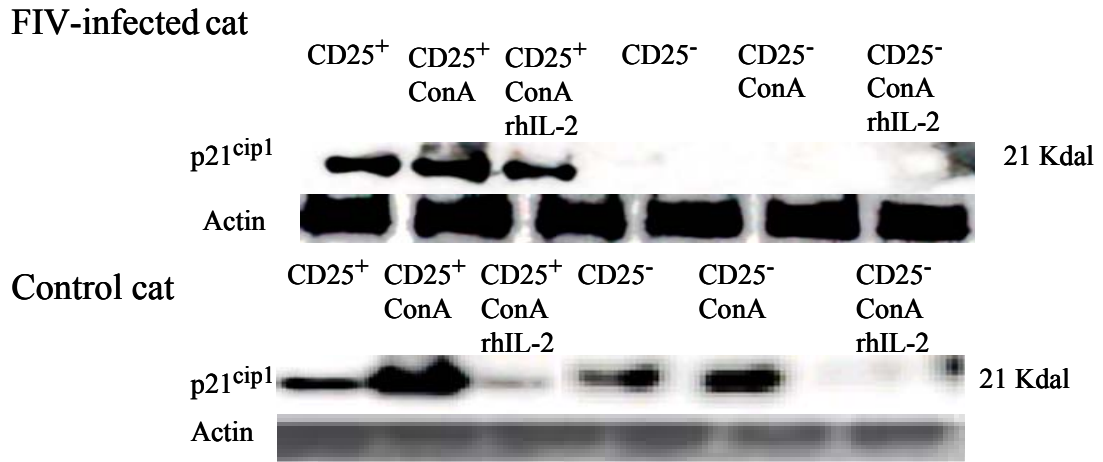


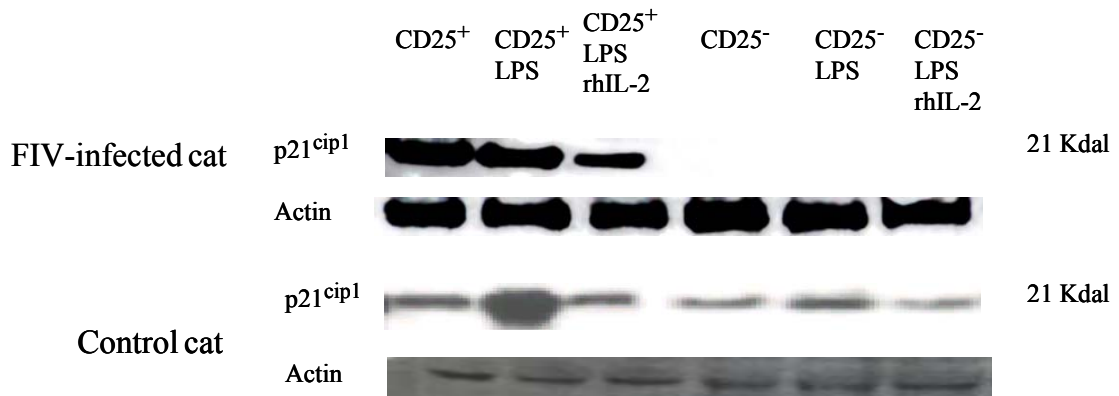
Figure 4. CdkI protein expression in ConA or LPS stimulated CD4⁺ T cell subsets.

Bead purified CD4⁺ T cell subsets were cultured with or without 5µg/ml ConA (**A**) or LPS 10ug/ml (**B**) with or without 100U/ml rhIL-2 from FIV infected cats and uninfected cats cultured for 2 days. Equal numbers (2×10^6) of cultured cells were lysed using NP-40 lysis buffer and separated on SDS-PAGE followed by immunoblotting with specific CdkI primary antibodies and HRP-conjugated goat anti-mouse IgG1 secondary antibody and detected by chemiluminescence. The blots were stripped using stripping buffer and re-probed with anti-actin primary antibody and HRP-conjugated goat anti-mouse IgG1 secondary antibody followed by chemiluminescence detection.

A



B



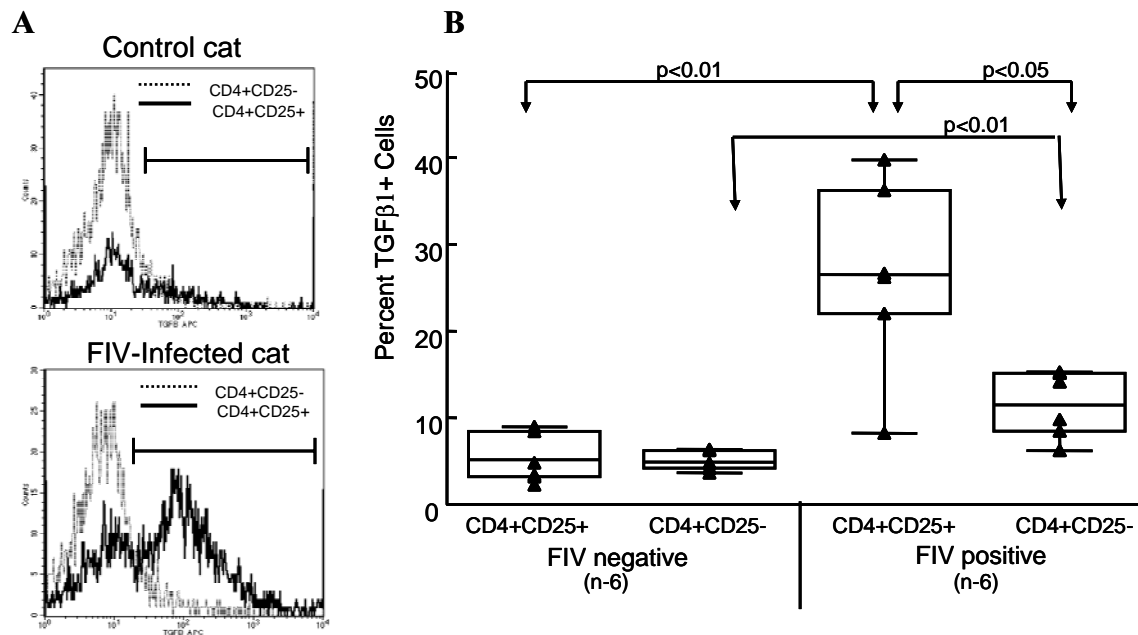
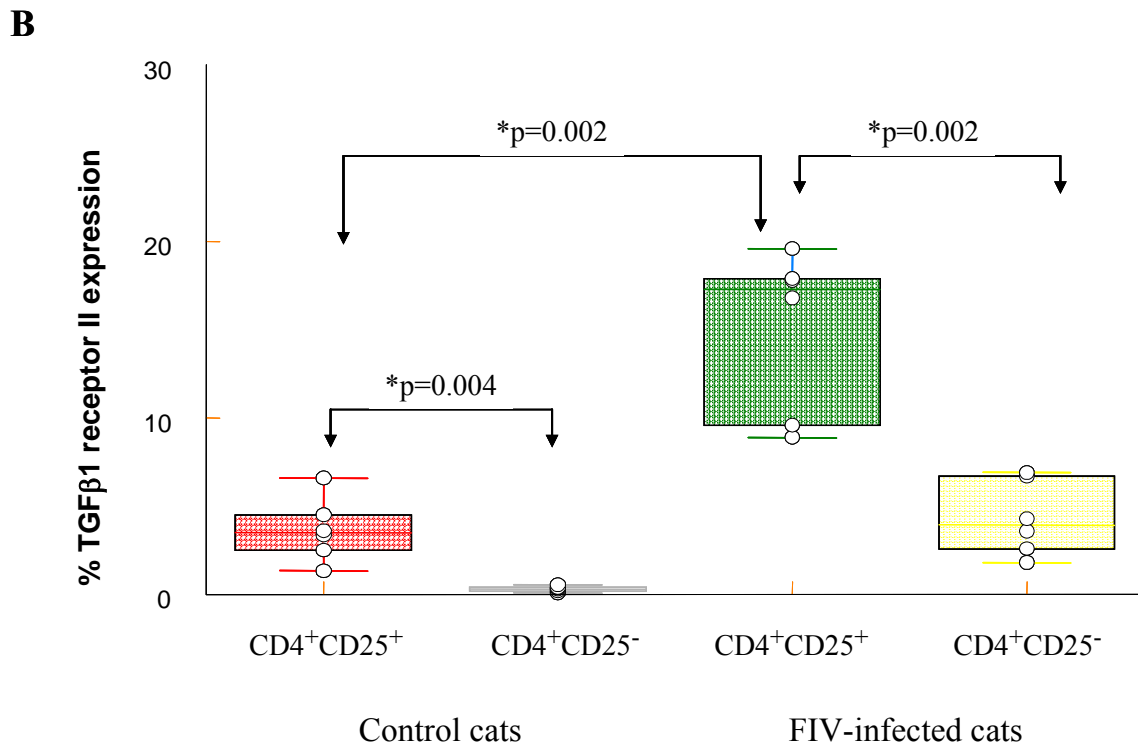
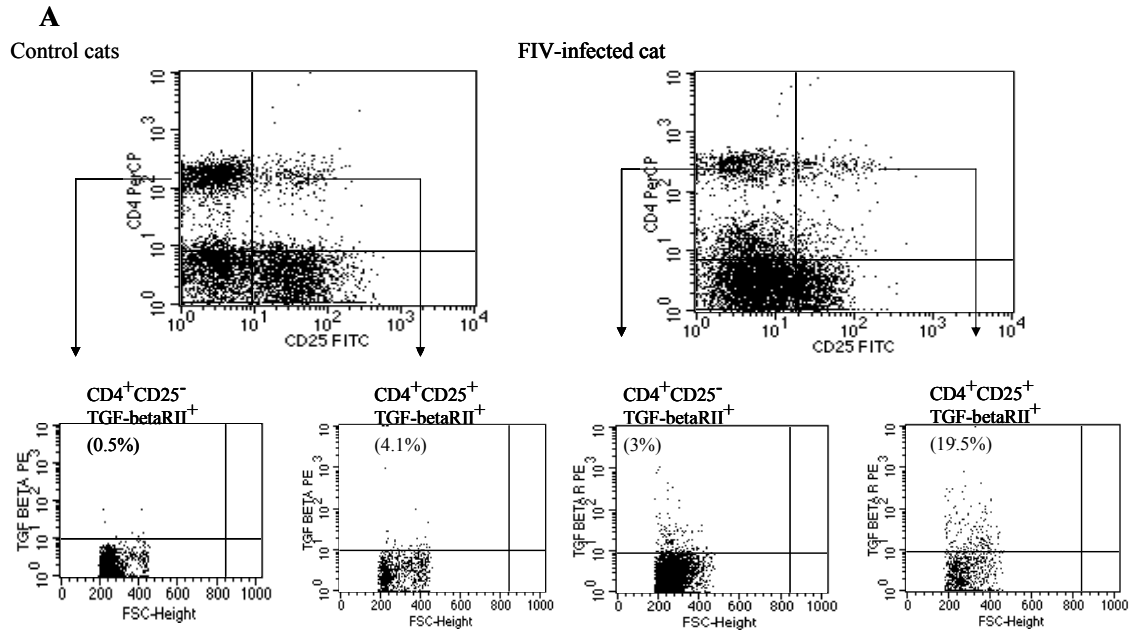


Figure 5. $CD4^+CD25^+$ T cells from FIV-infected cats express surface TGFβ1.

PBMC from FIV-infected and control cats were stained with anti-CD4-PE, anti-CD25-FITC, and anti-TGFβ1-APC. Data were acquired on a FACS Caliber flow cytometer. Lymphocytes were gated based on forward and side scatter and 20,000 gated events acquired for analysis using Cell Quest software. **(A)** Representative flow cytometry histogram of surface TGFβ1 expression on $CD4^+CD25^-$ (dotted line) and $CD4^+CD25^+$ (solid line) T cells from an FIV-infected and control cats. **(B)** Cumulative data for TGFβ1 expression on $CD4^+CD25^+$ and $CD4^+CD25^-$ T cells from FIV-infected (n=6) and control cats (n=6) cats. The middle line in the box whisker plots indicated by an arrow represents the median of the sample. The p-values were calculated by Mann-Whitney U test.



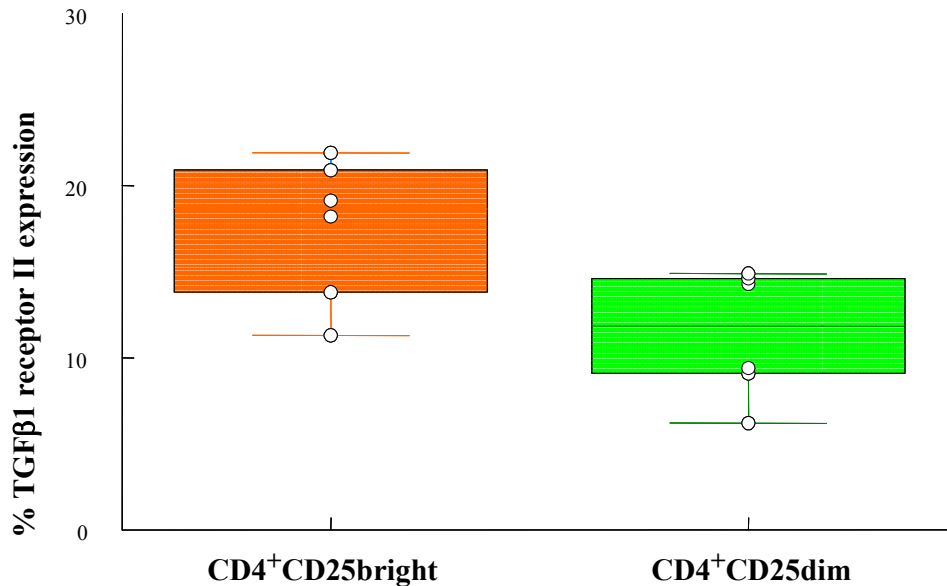
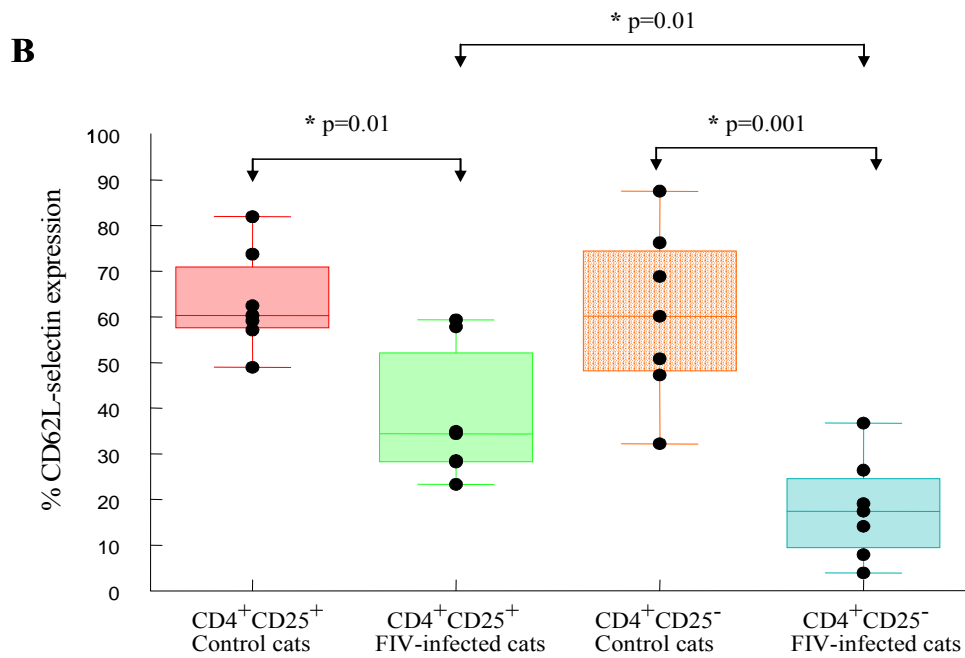
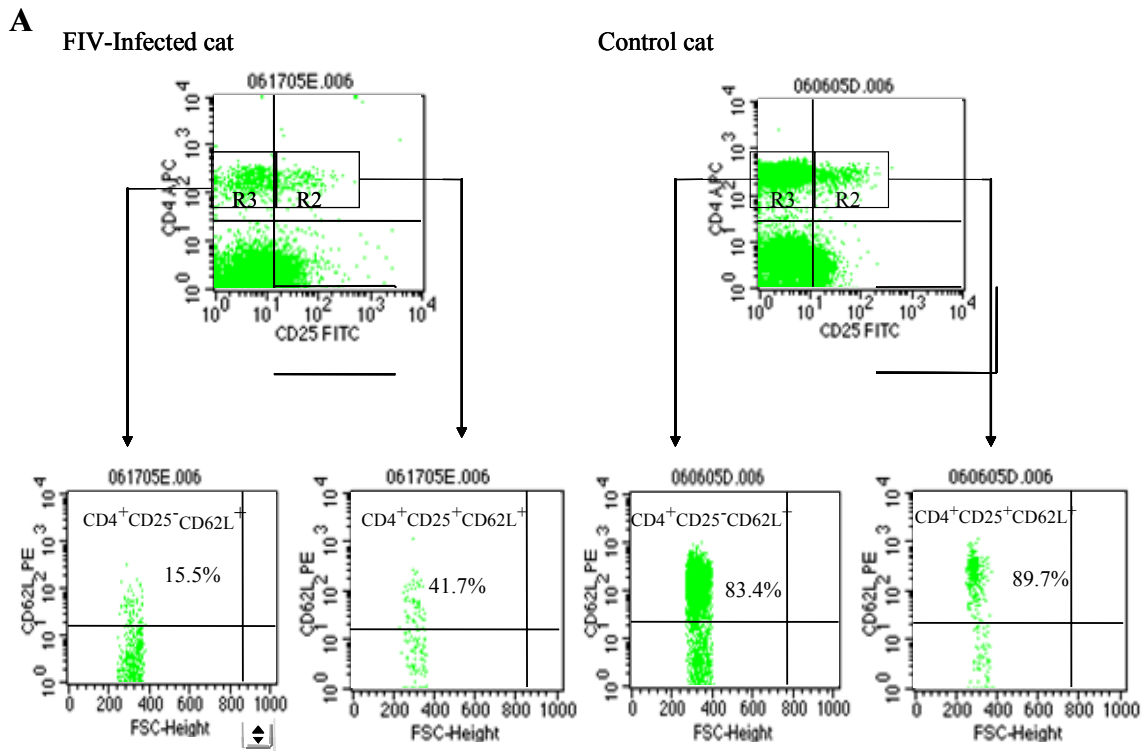
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Figure 6. CD4⁺ T cell subsets express TGFβRII at varying levels in FIV-infected and control cats. (A) Representative dot plots for TGFβRII expression in the two CD4⁺ T cell subsets from FIV-infected and control cats. Mononuclear cells from freshly drawn cat blood were stained with anti-CD4 APC, anti- TGFβRII PE, and anti-CD25 FITC. The cells in this analysis were gated on lymphocytes via their forward and side scatter properties. CD4⁺CD25⁺ and CD4⁺CD25⁻ T cells were analyzed for surface TGFβRII expression in FIV-infected and control cats. **(B)** Cumulative data for TGFβRII expression on CD4⁺CD25⁺ and CD4⁺CD25⁻ T cells from FIV-infected (n=6) and control cats (n=6) cats. **(C)** CD4⁺CD25^{bright} vs CD4⁺CD25^{dim} from FIV-infected (n=6) and control cats (n=6) cats. The middle line in the box whisker plots indicated by an arrow represents the median of the sample. The middle line in the box whisker plots indicated by an arrow represents the median of the sample. p-values were calculated by Mann-Whitney U test.



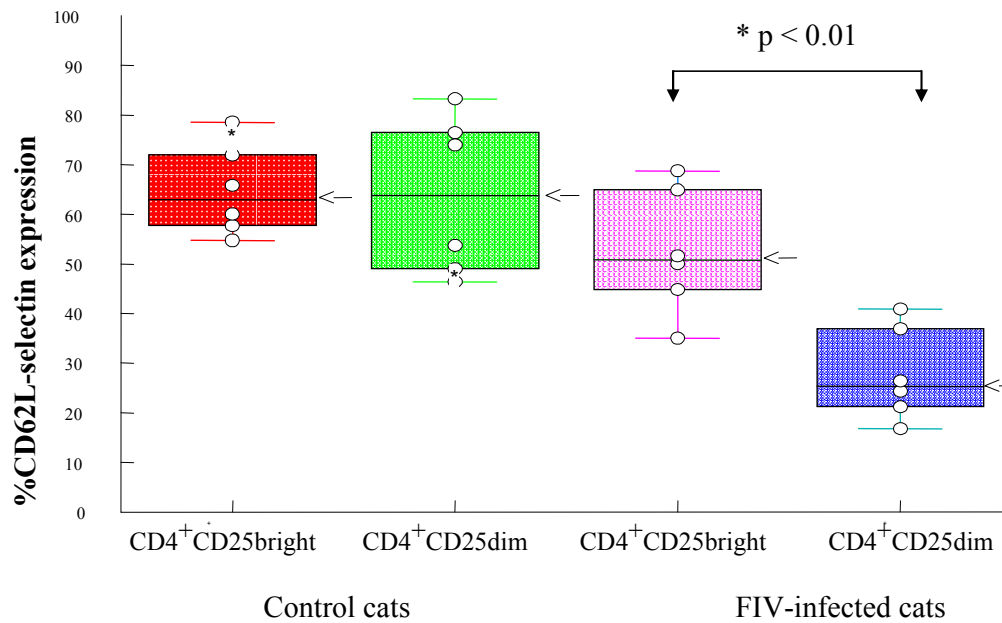
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Figure 7. CD4⁺ T cell subsets express CD62L-selectin at varying levels in FIV-infected and control cats. (A) Representative dot plots for CD62L-selectin expression in the two CD4⁺ T cell subsets from FIV-infected and control cats. Mononuclear cells from freshly drawn cat blood were stained with anti-CD4 APC, anti-CD62L PE, and anti-CD25 FITC. The cells in this analysis were gated on lymphocytes via their forward and side scatter properties. CD4⁺CD25⁺ and CD4⁺CD25⁻ T cells were analyzed for surface CD62L-selectin expression in FIV-infected and control cats. (B) Cumulative data for CD62L-selectin expression on CD4⁺CD25⁺ and CD4⁺CD25⁻ T cells. (C) CD4⁺CD25^{bright} vs CD4⁺CD25^{dim} from FIV-infected (n=7) and control cats (n=7) cats. The middle line in the box whisker plots indicated by an arrow represents the median of the sample. p-values < 0.01 as calculated by Mann-Whitney U test were considered significant.

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4. CHAPTER-2

**TGF β 1/TGF β RII SIGNALING MAY PLAY A KEY ROLE IN THE
MAINTENANCE OF ANERGIC STATE OF CD4⁺CD25⁺ T
REGULATORY CELLS, YET ENABLING THEIR RESISTANCE TO
ACTIVATION INDUCED APOPTOSIS VIA DIFFERENTIAL
EXPRESSION OF SEVERAL G0/G1 CELL CYCLE PROTEINS**

ABSTRACT

CD4⁺CD25⁺ T regulatory (Treg) cells that are known to be anergic, arrested in the G0/G1 phase of the cell cycle, yet programmed for cell survival. However, the molecular mechanism(s) involved in G0/G1 cell cycle arrest and cell survival signaling has not been defined. We previously reported that CD4⁺CD25⁺ Treg cells from chronically FIV-infected cats as opposed to those from control cats are functionally activated in vivo and express high levels of TGFβ1 on their surface and elevated levels of intercellular cyclin-dependent kinase inhibitors (CdkI's). To learn more of the potential role of cell surface expressed TGFβ1 in regulating the phenotypic and functional characteristics of Treg cells, we examine the role of exogenous TGFβ1 in conversion of CD4⁺CD25⁻ T cells to a Treg phenotype. Herein, we report that TGFβ1 in combination with mitogen could convert feline CD4⁺CD25⁻ T cells from peripheral blood to Treg-like cells in vitro. We also demonstrate that, similar to naturally occurring CD4⁺CD25⁺ Treg cells, TGFβ1 in vitro-converted CD4⁺CD25⁻ T cells are anergic and express elevated levels of CdkI, p21^{*cip1*} and anti-apoptotic protein bcl-2 but not pro-apoptotic protein bax, compared to unstimulated controls. Both natural and TGFβ1 converted CD4⁺CD25⁺ Treg cells express high levels of the anti-apoptotic protein bcl-2 and are more resistant to activation-induced cell death than their counterparts CD4⁺CD25⁻ T cells.

Keywords: CD4⁺CD25⁺ T regulatory cells; TGFβ1; CdkI; apoptosis; anergy; bcl-2

INTRODUCTION

Naturally occurring CD4⁺CD25⁺ Treg cells, comprising 5-10% of the circulating CD4⁺ T cell population, were originally defined as regulatory T cells that maintain peripheral self-tolerance (Maloy and Powrie 2001; Shevach 2002). More recent studies have demonstrated that Treg cells also play a major role in modulating the magnitude and duration of protective T cell responses to pathogens (Iwashiro, Messer et al. 2001; Montagnoli, Bacci et al. 2001; Belkaid, Piccirillo et al. 2002; Hori, Carvalho et al. 2002; Hisaeda, Maekawa et al. 2004; Belkaid and Rouse 2005). These pathogen-induced Treg cells, similar to natural Treg cells, are anergic, yet programmed for cell survival, and possess potent immunosuppressor function once activated by TCR engagement and/or certain nonspecific stimuli (Belkaid and Rouse 2005). While anergic T cells, including pathogen-induced CD4⁺CD25⁺ Treg cells, are known to be arrested at the G0/G1 phase of the cell cycle, and thought to be relatively long-lived in the circulation, there is little information on what receptors or intracellular signaling proteins maintain their unique functional state and homeostasis.

We recently demonstrated that feline CD4⁺CD25⁺ Treg cells possess all the functional characteristics of Treg cells described in humans and rodents in that they are anergic, arrested at the G0/G1 phase of the cell cycle, resistant to activation-induced cell death and possess potent immunosuppressive activity when activated (Joshi, Garg et al. 2005). Moreover, we recently reported that CD4⁺CD25⁺ Treg cells from chronically FIV-infected cats, in contrast to Treg cells from uninfected control cats, are activated in vivo

and fully armed for potent contact dependent immunosuppressor function (Vahlenkamp, Tompkins et al. 2004). Despite the activation status of CD4⁺CD25⁺ Treg cells in FIV-infected cats, which may result from the fact that they support a chronic productive FIV infection (Joshi, Vahlenkamp et al. 2004), their numbers remain relatively constant, or may actually increase throughout the course of the disease (Joshi, Vahlenkamp et al. 2004) .

The mechanism(s) regulating the survival and homeostasis of the Treg cell pool in the periphery is poorly understood. However, studies indicate that maintenance of peripheral Treg homeostasis is independent of the thymus. Marie et al. (Marie, Letterio et al. 2005) reported that Foxp3 expression and the size of the Treg cell compartment, as well as suppressor function are dependent on signals induced by TGFβ1 in the periphery and are not thymus dependent. Dieckmann, et al. (Dieckmann, Plottner et al. 2005) demonstrated that activated CD4⁺CD25⁺ Treg cells are capable of inducing a Treg phenotype from activated Th cells in a contact-dependent manner, suggesting that Treg cells may be recruited from the CD4⁺CD25⁻ T cell pool. In support of this, Chen et al. (Chen, Jin et al. 2003) reported that TGFβ1 is capable of converting TCR-stimulated CD4⁺CD25⁻ T cells into Foxp3 positive CD4⁺CD25⁺ Treg cells with potent immunosuppressive function. However, it is possible that the origin of the converted Treg cells in the periphery is not the CD4⁺ Th pool, but rather a small population of CD4⁺CD25⁻Foxp3⁺ cells that have been suggested to be a reservoir of precursor Treg cells (Romagnoli, Tellier et al. 2005). Whatever the source of cells maintaining peripheral Treg homeostasis, data suggest that

Treg cells, once activated, may recruit additional Treg cells from CD4⁺CD25⁻ T cell pool in a TGFβ1-dependent manner.

TGFβ1 is the prototypic member of a large family of structurally related pleiotropic-secreted cytokines that play a pivotal role in the control of differentiation, proliferation and the state of activation of many different cell types, including immune cells (Dennler, Goumans et al. 2002). Because of its dual role in pro-apoptotic and anti-apoptotic signaling, TGFβ1 is fundamental in the maintenance of homeostasis by regulating the balance between cell growth and apoptosis. Evidence from TGFβ1 knockout mice show that TGFβ1 plays a pivotal role in the maintenance of immune cell homeostasis. TGFβ1 null mice that are born, die shortly after weaning as a result of multifocal inflammatory disease with lymphocyte infiltration with multiple organs and autoimmune manifestations (Kulkarni, Huh et al. 1993; Christ, McCartney-Francis et al. 1994; Shull and Doetschman 1994; Yaswen, Kulkarni et al. 1996). TGFβ1 is thought to maintain peripheral Treg homeostasis by its ability to induce cell-cycle arrest as well as promote resistance to AICD. In many cell types, TGFβ1 causes a G0/G1 cell-cycle arrest by inhibiting cyclin-dependent kinase (Cdk) activity via activation of cyclin-dependent kinase inhibitors (CdkI's) (Wolfrain, Walz et al. 2004). We recently reported that CdkI's p21^{cip1}, p27^{kip1}, and p15^{ink4} are upregulated in G0/G1 arrested CD4⁺CD25⁺ Treg cells from FIV-infected cats. Upregulation of CdkI's in CD4⁺CD25⁺ Treg cells correlates with increased cell surface expression of TGFβ1 and TGFβRII (chapter 1). Others have reported increased expression of p21^{cip1} and p27^{kip1} in T cells anergized by TCR engagement in the absence of co-stimulation (Jackson, DeLoose et al. 2002). Wolfrain, et al. (Wolfrain, Walz et al.

2004) reported that TGFβ1 induced G0/G1 cell-cycle arrest of T cells is markedly diminished in p21^{cip1} and p27^{kip1} double knockout mice when T cells are activated under optimal co-stimulation conditions.

TGFβ1 also plays an important role in homeostasis of the immune system via its ability to regulate apoptosis (Sanchez-Capelo 2005). TGFβ1 inhibits or decreases the sensitivity of Fas/FasL and TNFα-induced apoptosis of activated T cells (Wahl, Orenstein et al. 2000). TGFβ1 has also been shown to suppress cytotoxic T cell-induced apoptosis by a contact-dependent mechanism between CD4⁺CD25⁺ Treg cells and cytotoxic T cells (Chen and Wahl 2003). Although the signaling pathways involved in TGFβ1-induced cell survival are not completely understood, recent data suggest that TGFβ1 may inhibit apoptotic signaling by upregulation of the NFκB cell survival signal (Saile, Matthes et al. 2001). Others have reported that TGFβ1 suppresses apoptosis in hepatic stellate cells by activating NFκB and by upregulating bcl-xl, the anti-apoptotic protein of the mitochondrial Fas-mediated pathway (Saile, Matthes et al. 2001). Similarly, TGFβ1 protects cultural embryonic rat neurons from apoptosis by increasing bcl-2 expression (Prehn, Bindokas et al. 1994). In contrast, in oligodendroglial precursor cells, TGFβ1 induces apoptosis which is associated with down-regulation of bcl-xl with no effect on bcl-2 (Schuster, Bender et al. 2003). While it has been reported that human CD4⁺CD25⁺ Treg cells are prone to apoptosis in association with decreased levels of bcl-2 expression (Taams, Smith et al. 2001), mouse CD4⁺CD25⁺ Treg cells are resistant to apoptosis, as they do not undergo clonal deletion when treated with superantigen in vivo (Papiernik, de Moraes et al. 1998), and are resistant to Fas-induced apoptosis in vitro (Banz, Pontoux et

al. 2002). Similar to murine Treg cells, Joshi, et al. (Joshi, Garg et al. 2005) reported that feline CD4⁺CD25⁺ Treg cells were relatively resistant to AICD compared to CD4⁺CD25⁻ T cells.

The apparent contradictory responses of different cell types to TGFβ1 signaling may depend upon the relative expression of bcl-2 and bax, as the ratio of bcl-2 to bax in a cell appears to determine survival or apoptosis (Xiao, Oppenlander et al. 2001). A case in point is the different responses of the precursor CD34⁺CD38⁻ and the committed CD34⁺CD38⁺ cell to TGFβ1 signaling. The CD34⁺CD38⁻ precursor with relatively low levels of bcl-2 respond to TGFβ by undergoing apoptosis, whereas the committed CD34⁺CD38⁺ with higher levels of bcl-2 are not sensitive to TGFβ1-induced apoptosis (Xiao, Oppenlander et al. 2001). Interestingly, bax expression is not changed in either cell in response to TGFβ1 signaling supporting the speculation that the bcl-2/bax ratio determines susceptibility or resistance to apoptosis.

As CD4⁺CD25⁺ are naturally anergic to antigen stimulation, we do not know whether the above discussed cellular responses to TGFβ1 signaling apply. To address some of these questions, this study focused on determining the mechanism(s) regulating cell growth arrest and apoptosis of CD4⁺CD25⁺ Treg cells from FIV-infected and control cats. We also explored the effects of TGFβ1 on expression of cell cycle genes and apoptotic genes in Treg cells. We observed that although Treg cells express elevated levels of CdkI's, they are resistant to apoptosis, which correlates with increased levels of anti-apoptotic proteins (bcl-2) and decreased levels of bax compared to CD4⁺CD25⁻ T cells. We also

report that TGFβ1 plus ConA treatment could convert feline CD4⁺CD25⁻ T cells into immune suppressor cells which possess phenotypic properties of naturally occurring CD4⁺CD25⁺ Treg cells, in that they express cell surface TGFβ1, and elevated levels of p21^{*cip1*}, and anti-apoptotic proteins (bcl-2).

MATERIALS AND METHODS

Animals and viral infection

The cats used for this study were specific pathogen free (SPF) obtained from Liberty Labs (Liberty Corners, NJ) or Cedar River Laboratory (Mason City, IA) and housed at the Laboratory Animal Resource Facility at the College of Veterinary Medicine, North Carolina State University. Cats were inoculated with the NCSU1 isolate of FIV, a pathogenic clade A virus (English, Nelson et al. 1994), as described by Bucci et al (Bucci, English et al. 1998). The samples were collected from FIV⁺ cats infected for more than 3yrs and from uninfected control cats, 4 to 6 years of age. FIV-infected cats were housed separately from uninfected control cats. FIV-infection was confirmed by immunoblot analysis and provirus detection by PCR using primers specific for the FIV-p24 encoding sequence.

Cell collection

Whole blood was collected by jugular venipuncture into EDTA Vacutainer tubes (Becton-Dickinson, Franklin Lakes, NJ) and PBMC isolated by Percoll (Sigma-Aldrich, St. Louis, MO) density gradient centrifugation as previously described (Tompkins,

Ogilvie et al. 1987). LN cells were obtained from LN biopsies and single-cell suspensions were prepared by gently injecting sterile PBS into the tissue using 18G needle repeatedly, until the cells were released from the tissue. Cell counts and viability were determined by trypan blue dye exclusion. Viability was always >90%.

CD4⁺ T cell subset separation

To obtain single lymphocyte subsets, CD4⁺CD25⁺ and CD4⁺CD25⁻ T cells derived from PBMC and pooled LN cells were either enriched by biomagnetic bead depletion and selection or by fluorescent activated cell sorting (FACS) (MoFlo; DakoCytomation, Fort Collins, CO). Biomagnetic bead separation was performed using goat anti-mouse IgG-coated beads (Dynabeads M-450; Dynal, Great Neck, NY) as described by Bucci et al. (Bucci, English et al. 1998). PBMC and LN were depleted of B cells with anti-CD21 mAb (Serotec, Raleigh, NC)-coated beads, and CD8⁺ cells were depleted with anti-CD8 (mAb 3.357 (Tompkins, Gebhard et al. 1990))-coated beads. The CD4⁺ cell population was then enriched for CD25⁺ cells by positive selection using anti-feline CD25 mAb (9F23; kindly provided by K. Ohno (University of Tokyo, Tokyo, Japan)). The enriched CD4⁺CD25⁺ or CD4⁺CD25⁻ T cells were analyzed for purity by flow cytometric analysis and were greater than 90% pure. The purity of the FACS sorted CD4⁺CD25⁺ and CD4⁺CD25⁻ T cells were greater than 95%. In both cases, viability of the purified CD4⁺ subsets was determined by trypan blue dye exclusion and was always >90%. All assays were based on viable cell numbers.

Flow cytometry analysis

At least 5×10^5 PBMC or purified cell populations were stained for surface expression of CD4 and CD25 markers using APC-conjugated anti-CD4 (mAb 30A (Tompkins, Gebhard et al. 1990)), FITC-conjugated anti-CD25 mAb (mAb 9F23), and PE-conjugated anti-CD62L selectin (BD-pharmagen). Monoclonal antibodies for TGF β 1 (mouse anti-human, mAb 240 clone 9016) were obtained from R&D Systems. Data were acquired on a FACS Caliber flow cytometer (BD Biosciences, Mountain View, CA). Lymphocytes were gated based on forward vs side scatter, and 20,000 gated events were acquired and stored list-mode fashion for analysis using Cell Quest software.

Propidium Iodide staining

Purified CD4⁺ T cell subsets from PBMC were cultured with or without ConA (5 μ g/ml) or CD4⁺CD25⁻ T cells stimulated with rhIL-2 (100U/ml) or without ConA (5 μ g/ml) +/- TGF β 1 (10ng/ml) for 24hrs or 48hrs. Propidium Iodide was added to the cultured cells (1×10^5) for 10min incubated on ice followed by a wash in PBS. The cells were then assayed for viability immediately by flow cytometry analysis. For TGF β 1 plus ConA stimulated CD4⁺CD25⁻ T cells, only TGF β 1 positive cells stained positive for PI were determined.

T cell stimulation assays

T cell subsets were cultured in RPMI 1640 medium containing 10% heat-inactivated FBS, 1% penicillin-streptomycin, 1% sodium bicarbonate, 1% sodium pyruvate, 1% L-

glutamine, and 1 mM HEPES buffer. For ConA/TGFβ1 stimulation, 1×10^5 viable cells/well were cultured in a round-bottom 96-well plate in the presence or absence of 5 μg/ml Con-A or 100 U/ml recombinant human IL-2 (rhIL-2) (National Institutes of Health AIDS Research and Reagent Program, Rockville, MD) or 10ng/ml TGFβ1 (R&D Biosciences) . All assays were run in triplicate. For anergy assays CD4⁺CD25⁻ T cells were stimulated for 2 days cells, and for suppressor assays, after 4-5 days of incubation stimulated cells were cocultured for 2 days with autologous ConA stimulated (24hrs) CD4⁺CD25⁻ T cells and were pulsed with 1 μCi of [³H]TdR/well and harvested 16hrs later using a Filtermake Harvester (Packard Bioscience, Meriden, CT). [³H]TdR uptake as a measure of proliferation was determined in counts per minute (cpm) using a Top Count NXT Microplate scintillation counter (Packard Bioscience).

Immunoblotting

Equal numbers of Ficoll separated PBMC or FACS purified CD4⁺ T cell subsets (1×10^6 – 2×10^6 cells) stimulated for 4-5 days with rhIL-2, TGFβ1, ConA or ConA plus TGFβ1 were lysed using NP-40 lyses buffer for 16-24hrs at 4C. The concentrations of the proteins were determined using a spectrophotometer (Biorad Smartspec 3000), 30-50μg of proteins were then resolved on 4-12% SDS-polyacrylamide gels (Novex Inc.). The separated proteins were transferred onto nitrocellulose and immunoblotted with specific mAbs to p21^{cip1}, bcl-2, and bax (Santa Cruz biotechnology Inc) followed by HRP-conjugated secondary antibodies (Santa Cruz Biotechnology Inc). Immunodetection was performed by ECL (Pierce) using HyperFilm ECL. To normalize the result, some blots were stripped with the Western Blot stripping buffer (Pierce) and reprobed with anti-actin

mAb (Santa Cruz Biotechnology Inc.). Specific antibody conjugated blots were analyzed for specific proteins using Lumicapture analyst camera detection system (Lumi- Imager).

Statistical Analysis

The student t-test was used to assess significance of difference between different CD4⁺CD25⁻ T cell stimulations compared to unstimulated controls for detection of activation induced apoptosis in FIV-infected and control cats.

RESULTS

TGFβ1 along with mitogen stimulation can induce anergy in feline CD4⁺CD25⁻ T cells

It has also been recently demonstrated in rodent and human models that apart from naturally occurring CD4⁺CD25⁺ Treg cells generated in the thymus from a defined lineage, CD4⁺CD25⁺ Treg like anergic/suppressor cells may also be induced in the periphery by the action of several immunosuppressive molecules or cytokines like TGFβ1. Also that TGFβ1 along with TCR stimulation can convert naïve CD4⁺CD25⁻ T cells into CD4⁺CD25⁺ anergic/suppressor cells in vitro. To test this hypothesis in a feline model, flow cytometry purified CD4⁺CD25⁻ T cells were stimulated with rhIL-2, or TGFβ1, or TGFβ1 +/- ConA for 2 days and ³H thymidine was added in the last 16hrs to determine anergy. As observed in **figure1**, TGFβ1 along with ConA stimulation of CD4⁺CD25⁻ T cells exhibited significantly reduced TCR-triggered proliferation

compared to CD4⁺CD25⁻ T cells stimulated with rhIL-2 or ConA or TGFβ1 alone. Thus, TGFβ1 plus TCR stimulated CD4⁺CD25⁻ T cells are functionally anergic similar to that of naturally occurring feline CD4⁺CD25⁺ Treg cells (chapter 1).

TGFβ1 along with ConA stimulation induces the expression of cyclin dependent kinase inhibitor (p21^{cip1}) protein.

We recently reported that feline CD4⁺CD25⁺ Treg cells are phenotypically anergic as indicated by the elevated levels of expression of cyclin dependent kinase inhibitor proteins (p21^{cip1}, p27^{kip1}, and p16^{ink}) (chapter 1). To determine whether TGFβ1 converted CD4⁺CD25⁻ T cells from control cats (**figure 2A**) and FIV-infected cats (**figure 2B**) are phenotypically anergic, bead purified CD4⁺CD25⁻ T cells were stimulated with or without rhIL-2, or ConA, or TGFβ1 +/- ConA for 4-5 days. Equal numbers of viable cells were lysed and total cellular proteins were separated on SDS-page gels and analyzed by immunoblotting procedures. There was no effect on p21^{cip1} expression in CD4⁺CD25⁻ T cells stimulated with rhIL-2 or ConA or TGFβ1 alone. In contrast, TGFβ1 along with ConA stimulation of CD4⁺CD25⁻ T cells induced elevated levels of p21^{cip1} protein compared to unstimulated controls in FIV-infected and control cats (**figure 2**).

Feline CD4⁺CD25⁻ T cells stimulated with TGFβ1 along with ConA are potent suppressors similar to naturally occurring CD4⁺CD25⁺ Treg cells

To further reconfirm whether the TGFβ–energized feline CD4⁺CD25⁻ T cells were functionally active and able to suppress responder Th cell proliferation (unpublished data), freshly isolated CD4⁺CD25⁻ naive T cells from FIV-infected cats and control cats were cultured with ConA+/- rhIL-2 with or without TGFβ1 for 4-5 days. These stimulated CD4⁺CD25⁻ T cells were cocultured for 48hrs with autologous CD4⁺CD25⁻ T cells cultured with ConA for 24hrs. ³H-thymidine was added in the last 16hrs before harvest. Percent inhibition of CD4⁺CD25⁻ T cell proliferation was calculated.

CD4⁺CD25⁻ T cells stimulated with either rhIL-2 or ConA or TGFβ1 alone from both FIV-infected cats (**figure 3A**) and control cats (**figure 3B**) had any suppressive action. In contrast, TGFβ1 and ConA converted CD4⁺CD25⁻ T cells from both control cats and FIV-infected cats were able to inhibit ConA stimulated CD4⁺CD25⁻ T cell proliferation similar to that of naturally occurring CD4⁺CD25⁺ Treg cells from FIV-infected cats.

TGFβ1 converted CD4⁺CD25⁻ Treg like cells similar to naturally occurring CD4⁺CD25⁺ Treg cells are more resistant to activation induced apoptosis compared to rhIL-2 stimulated CD4⁺CD25⁻ T cells.

Our laboratory previously demonstrated that feline CD4⁺CD25⁺ Treg cells are anergic functionally and phenotypically as indicated by elevated levels of cyclin dependent kinase inhibitors expression compared to CD4⁺CD25⁻ T cells (chapter 1). We also reported that FIV preferentially replicates in CD4⁺CD25⁺ Treg cells compared to CD4⁺CD25⁻ T cell counterparts (Joshi, Vahlenkamp et al. 2004). To determine the potential of the purified CD4⁺ T cell subsets for the resistance to activation induced

apoptosis, CD4⁺CD25⁺ Treg cells and CD4⁺CD25⁻ T cells from FIV-infected and control cats were stimulated with rhIL-2 or TGFβ1, or ConA +/- TGFβ1 for 24hrs or 48hrs. Cell viability was determined using flow cytometry by propidium iodide staining procedures. We demonstrated that Con-A stimulated CD4⁺CD25⁺ Treg cells from either FIV-infected or control cats are significantly more resistant to activation induced apoptosis at 24hrs and even more at 48hrs than CD4⁺CD25⁻ T cells in-vitro (**figure 4A**). Similar to naturally occurring Treg cells, only TGFβ1 along with mitogen stimulation of CD4⁺CD25⁻ T cells induced an anti-apoptotic phenotype compared to the rhIL-2 stimulated controls (**figure 4B**). As expected CD4⁺CD25⁻ T cells were able to proliferate significantly in response to ConA compared to rhIL-2 stimulated controls (**figure 4B**).

Anti-apoptotic but not pro-apoptotic proteins are differentially expressed in the TGFβ1 converted CD4⁺CD25⁻ T cell subsets

To further analyze the molecular bases for the CD4⁺CD25⁺ Treg cells to maintain anergic phenotype yet be resistant to apoptosis compared to CD4⁺CD25⁻ T cells, expression of anti-apoptotic protein (bcl-2) and pro-apoptotic protein (bax) was determined in the two CD4⁺ T cell subsets (**figure 5A**). Bead separated CD4⁺CD25⁺ Treg cells and CD4⁺CD25⁻ T cells isolated from the PBMC of FIV-infected and control cats or CD4⁺CD25⁻ T cells stimulated for 4 days with rhIL-2, TGFβ1, or ConA +/- TGFβ1 were lysed to obtain protein. Total cell proteins were separated on SDS-PAGE gels and further analyzed by western blotting using specific antibodies such as bcl-2 and bax. As shown in **figure 5A** anti-apoptotic protein bcl-2 was expressed at elevated levels in CD4⁺CD25⁺ Treg cells compared to CD4⁺CD25⁻ T cells, and in contrast pro-apoptotic protein bax was

expressed at increased levels in CD4⁺CD25⁻ T cells from both FIV-infected and control cats. Stimulation of CD4⁺CD25⁻ T cells only with TGFβ1 plus ConA induced the expression of anti-apoptotic protein bcl-2, but had no effect on pro-apoptotic protein bax expression (**figure 5B**).

DISCUSSION

Treg cells are characterized by G0/G1 cell cycle arrest and are anergic to immune activation, as manifested by their inability to produce IL-2 and exit the G0/G1 stage of cell cycle in response to antigen-specific TCR engagement (Powell, Ragheb et al. 1998). Another major characteristic of CD4⁺CD25⁺ Treg cells described in human and rodent species is their ability, once activated through their TCR, to suppress proliferative responses of other antigen- or mitogen-stimulated CD4⁺ or CD8⁺ T cells (Thornton and Shevach 1998; Jackson, DeLoose et al. 2001). We recently described CD4⁺CD25⁺ Treg cells in the cat possessed all the major characteristics of phenotypic anergy as indicated by the expression of genes responsible for exiting G0/G1 phase of the cell cycle such as cyclin D, retinoblastoma protein and cyclin dependent kinase inhibitors (chapter 1). We also demonstrated the functional characteristics of Treg cells in cats, similar to those described in other species in that they are G0/G1 cell cycle arrested, IL-2 deficient and anergic to mitogenic stimulation, and when activated capable of suppressing IL-2 production and proliferation of activated T cells (Vahlenkamp, Tompkins et al. 2004).

We recently reported that CD4⁺CD25⁺ Treg cells from chronically FIV-infected cats, in contrast to Treg cells from uninfected control cats, are activated in vivo and fully armed for potent contact dependent immunosuppressor function (Vahlenkamp, Tompkins et al.

2004). In conjunction with functional anergy, in vivo activated CD4⁺CD25⁺ Treg cells from FIV-infected cats expressed elevated levels of CdkI's such as p21^{cip1} compared to CD4⁺CD25⁻ T cells. No significant difference between the two CD4⁺ T cell subsets was observed in control cats (chapter 1). We also reported that activated Treg cells from FIV-infected cats express elevated levels of TGFβ1 and its receptor on their cell surface compared to their counterparts CD4⁺CD25⁻ T cells. We also observed the colocalization of TGFβ1 with its receptor on the cell surface (Tompkins lab unpublished data).

CD4⁺CD25⁺ regulatory T cells (Treg) play an important role in the control of immune responses and the maintenance of immune homeostasis. Although Treg generally inhibit T cell proliferation, the precise regulatory mechanisms of Treg remain unknown, despite reports suggesting possible roles for TGFβ or CTLA-4. In this study, we have provided evidence that feline Treg like cells can be induced/converted from peripheral CD4⁺CD25⁻ naive responder T cells through costimulation of TCR and TGFβ signaling. Several important conclusions can be drawn from the current work. First, TGFβ1 treatment in the presence of TCR stimulation converts naive CD4⁺CD25⁻ responder feline T cells to anergized CD4⁺CD25⁺ Treg like cells that inhibits their proliferation in vitro compared to ConA stimulated CD4⁺CD25⁻ T cells.

The hallmark of naturally occurring Treg cells, including those in normal cats is that they require activation by antigen-TCR engagement or a potent mitogen before they can display immunosuppressive function (Vahlenkamp, Tompkins et al. 2004). TGFβ1 bound to its receptor on Treg cells could induce suppression of other T cells in a cell-contact

dependent manner. We reconfirmed that TGFβ1–converted anergic feline CD4⁺CD25⁻ T cells are potent suppressors similar to professional Treg cells. Functionally, when cocultured with ConA stimulated CD4⁺CD25⁻ T cells, TGFβ1-energized T cells inhibit proliferation and cytokine production of the target T cells similar to naturally occurring CD4⁺CD25⁺ Treg cells.

Induction and maintenance of anergy in feline Treg cells has been associated with over-expression of the cyclin-dependent kinase inhibitors, CdkI's, p21^{*cip1*}, p21^{*kip1*}, and p16^{*ink*} (chapter 1). TGFβ1 causes a G0/G1 cell-cycle arrest by inhibiting cyclin-dependent kinase (Cdk) activity via activation of cyclin-dependent kinase inhibitors (CdkI's) (Kee, Rivera et al. 2001; Massague 2004; Wolfrain, Walz et al. 2004). We recently reported that CdkI's p21^{*cip1*}, p27^{*kip1*}, and p16^{*ink4*} are upregulated in G0/G1 arrested CD4⁺CD25⁺ Treg cells from FIV-infected cats. Upregulation of CdkI's in CD4⁺CD25⁺ Treg cells correlates with increased cell surface expression of TGFβ1 and TGFβRII (chapter 1). In this study we have demonstrated that TGFβ1–converted anergic feline CD4⁺CD25⁻ T cells similar to the professional Treg cells, over-express CdkI, p21^{*cip1*}. Thus, TGFβ1 stimulation along with TCR-engagement can induce phenotypic and functional anergy in CD4⁺CD25⁻ T cells in vitro.

TGFβ1 suppresses apoptosis in cells by activating NFκB and by upregulating Bcl-family members such as bcl-X and bcl-2. CD4⁺CD25⁺ Treg cells exert regulatory effects through a cell-cell contact-dependent manner, and the frequency of Treg cells is crucial to the prevention of inflammation and autoimmunity, as well as induction of tumor antigen

tolerance. It has been shown previously that CD4⁺CD25⁺ Treg cells are potent targets for FIV-infection (Joshi, Vahlenkamp et al. 2004). The activation status of CD4⁺CD25⁺ Treg cells in FIV-infected cats may result from the fact that they support a chronic productive FIV-infection (Joshi, Vahlenkamp et al. 2004). Furthermore, subsets of immune cells are known to respond differently to activation induced apoptosis. These observations prompted us to explore the sensitivity of CD4⁺CD25^{+/-} T cells to activation induced cell death.

A homeostatic phenomenon tends to reduce the peripheral pool of proliferating lymphocytes by inducing apoptosis of activated T cells upon TCR re-engagement. TGFβ1 is important in the maintenance of homeostasis by regulating the balance between cell growth and apoptosis. TGFβ1 is also thought to maintain peripheral Treg homeostasis by its ability to induce cell-cycle arrest as well as promote resistance to AICD. We have shown here that TGFβ1 induced Treg like cells similar to naturally occurring Treg cells are more resistant to activation induced apoptosis compared to ConA stimulated CD4⁺CD25⁻ T cells. This different susceptibility to cell death might be explained by the balance between pro- and anti-apoptotic factors such as FLIP (Irmeler, Thome et al. 1997; Kataoka, Budd et al. 2000), and members of the Bcl-2 family such as bcl-2, and bax (Xiao, Oppenlander et al. 2001) of the mitochondrial Fas-mediated pathway (Saile, Matthes et al. 1999). In this study, we demonstrated that TGFβ1 along with TCR stimulation of CD4⁺CD25⁻ T cells similar to professional Treg cells can induce the expression of anti-apoptotic proteins like bcl-2. In contrast to Treg cells which express low levels of pro-apoptotic proteins, bax. TGFβ1 converted Treg like cells had no effect on bax expression. The expression of elevated levels of bcl-2 correlated with the anti-

apoptotic potential of activated CD4⁺CD25⁺ Treg cells. Therefore, more than the levels of expression of anti- and pro-apoptotic proteins, the ratio between the two may play a critical role in determining the fate of the cell, to die or live, upon activation.

Based on the above observations, we proposed a model that could explain the effects of TGFβ1 signaling in the maintenance of homeostasis by regulating proteins that are essential for the balance between cell growth and apoptosis in Treg cells (**Figure 6**).

TGFβ1 is expressed in the G₁ phase of the cell cycle in growth-arrested cells (Chen, Jin et al. 2003) and binding of TGFβ1 to its receptor activates a wide range of pathways during cell-cycle progression. TGFβRI/II complex formation is induced by the binding of TGFβ ligands that in turn phosphorylates receptor-activated Smads 2/3 (R-Smads). R-Smads form complexes with Smad4 which allows their translocation into the nucleus where the complex interacts with Smad-binding elements (SBE) (Shi, Wang et al. 1998). R-Smad complex promotes the transcription of several target genes depending upon the combinations of Smad complex interactions with co-activators and inhibitors of gene transcription (Shi and Massague 2003) including p21^{*cip1*}, and Foxp3. Foxp3 in turn down regulates the Smad7. Foxp3-mediated down-regulation of Smad7 induces a positive auto-regulatory loop of TGFβ-Smad signaling in naturally occurring and TGFβ1 induced Treg cells in human and murine models (Fantini, Becker et al. 2004). On the other hand Foxp3 inhibits IL-2 transcription. TGFβ1 signaling can also activate MAP-Kinase pathway that induces expression of several transcriptional factors such as ATF, Sp1, and c-Jun. p21^{*cip1*} promoter possesses binding site/s for transcription factors of the Sp1 family along with Smad's (Moustakas, Pardali et al. 2002) which is targeted by both Smad- and

Ras/MAPK-mediated pathways. ATF transcription factor induces phosphorylation of cyclic AMP response element binding proteins (CREB) which in turn targets induction of several genes including bcl-2. Thus, TGF β 1 pathway could be one of the mechanisms by which Treg cells can maintain energy, yet be programmed for cell survival.

In conclusion, TGF β 1 stimulation along with TCR- engagement can induce functional and phenotypic energy in converted Treg like cells, as manifested by their inability to proliferate upon mitogenic stimulation and expression of CdkI, p21^{cip1} respectively. TGF β 1 converted Treg like cells similar to naturally occurring Treg cells are more resistant to activation induced apoptosis compared to mitogen stimulated CD4⁺CD25⁻ T cells, which correlates with the expression of anti apoptotic proteins (bcl-2) and pro-apoptotic proteins (bax). Therefore, TGF β /TGF β R signaling may be one of the mechanisms by which Treg cells could induce G0/G1 cell cycle arrest by up-regulation of CdkI's, yet be resistant to apoptosis by maintaining anti-apoptotic and pro-apoptotic protein ratios for survival.

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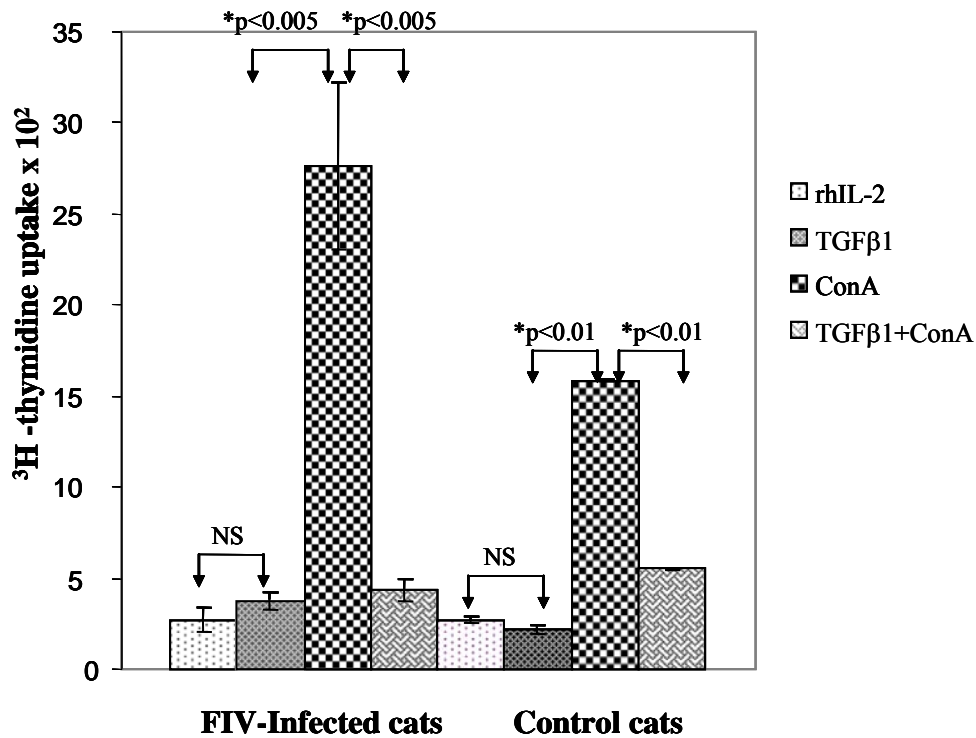


Figure 1. *TGFβ1 along with TCR triggering converts feline CD4⁺CD25⁻ T cells to anergic CD4⁺CD25⁺ Treg like cells in-vitro.* Freshly isolated bead purified CD4⁺CD25⁻ T cells from FIV-infected cats and control cats were stimulated with or without rhIL-2, or ConA, or TGFβ1 +/- ConA for 2 days. ³H-thymidine was added in the last 16hr period to determine cell proliferation. p-values were calculated using student t- test. *p-value < 0.01 was considered significant.

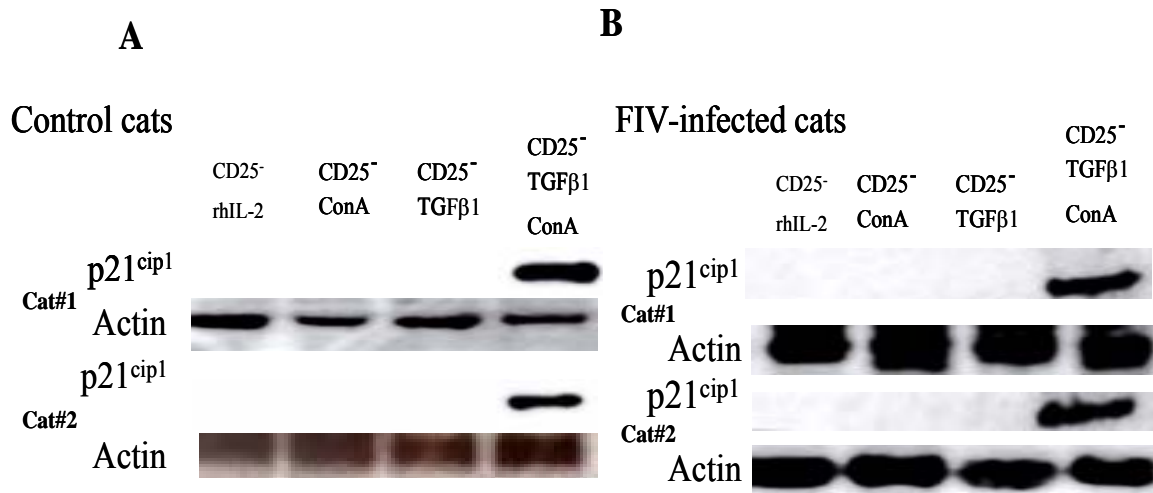


Figure 2. Induction of protein expression of cyclin dependent kinase inhibitor (p21^{cip1}) by TGFβ1. Equal number of purified CD4⁺CD25⁻ T cells (1*10⁶ - 2*10⁶) from control cats (**A**) and FIV-infected cats (**B**) stimulated with or without TGFβ1 (10ng/ml) +/- ConA (5μg/ml) and /or rhIL-2 (100U/ml) for 4-5 days were lysed with NP-40 lysis buffer to obtain total cellular protein for the detection of p21^{cip1}. Total cell lysates were separated by SDS-PAGE and analyzed by immunoblot using specific primary antibodies against CdkI (p21^{cip1}), 21Kdal protein, followed by HRP-conjugated goat anti-mouse IgG1 secondary antibody and detected by chemiluminescence. The blots were stripped using stripping buffer and reprobbed with actin primary antibody and HRP-conjugated goat anti-mouse IgG1 secondary antibody followed by chemiluminescence detection.

Figure 3. Suppressor activity of TGFβ1 converted CD4⁺CD25⁻ T cells on autologous CD4⁺CD25⁻ T cells from FIV-infected cats and control cats. FACS-purified CD4⁺CD25⁻ T target cells from control cats (**A**) and FIV-infected cats (**B**) were either stimulated with rhIL-2 (100U/ml) or ConA (5μg/ml) or TGFβ1 (10ng/ml) +/- ConA for 4-5 days. These stimulated CD4⁺CD25⁻ T cells or freshly isolated CD4⁺CD25⁺ Treg cells from FIV-infected cats were cocultured for 2 days with autologous CD4⁺CD25⁻ T cells cultured for 24hrs in the presence of ConA and were pulsed with [³H]TdR 16hrs prior to the harvest. A constant number of CD4⁺CD25⁻ T target cells (5 x 10⁴/well) were cultured with varying numbers of CD4⁺CD25⁺ Treg cells or the above stimulated CD4⁺CD25⁻ T effector cells. Percent inhibition of proliferation was calculated based on proliferation of CD4⁺CD25⁻ T target cells alone ((cpm of target cells alone – cpm of coculture) / cpm of target cells alone). Bars represent mean +/- SEM of triplicate wells for each condition.

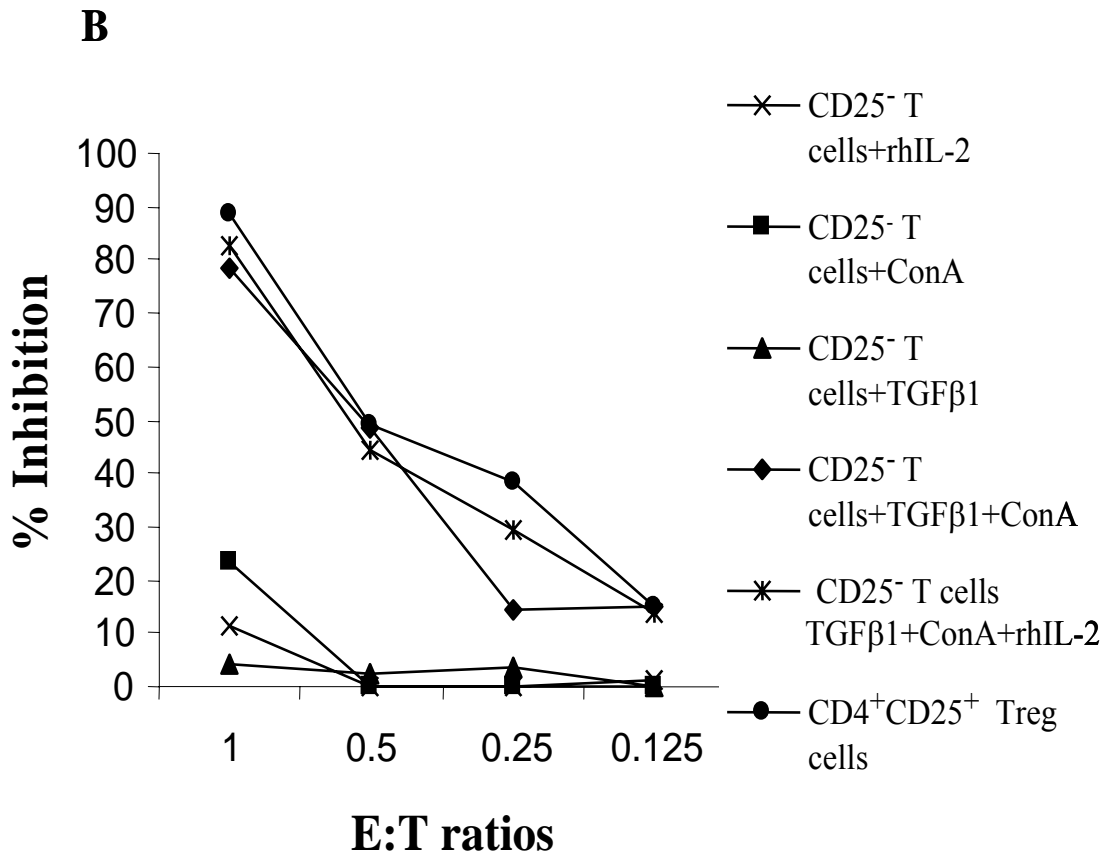
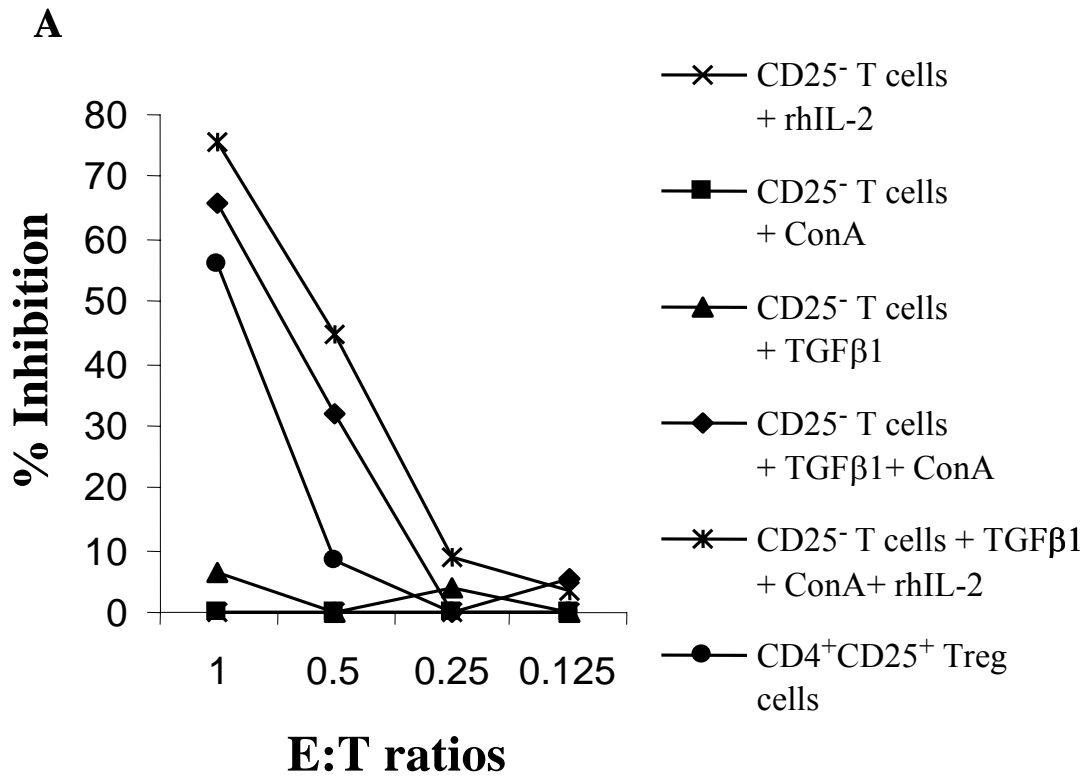
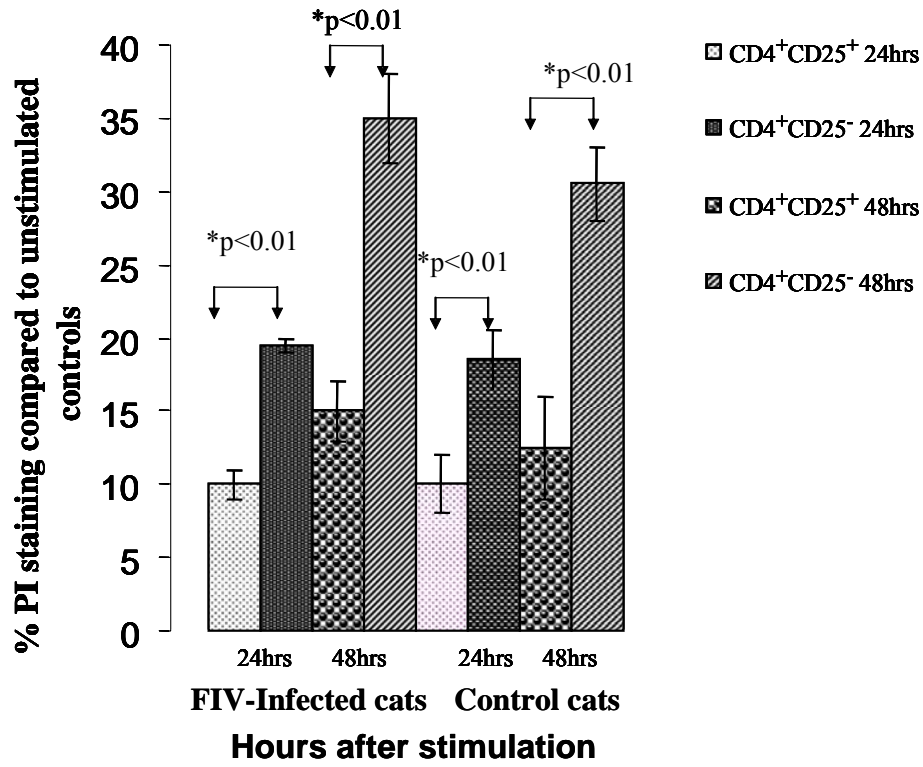


Figure 4. Resistance to activation induced apoptosis. (A) Flow cytometry sorted pure CD4⁺ populations from FIV-infected and control cats were stimulated with or without 5µg/ml Con-A for 24hrs or 48hrs. Propidium Iodide was added to the cells (1*10⁵) at 24hrs or 48hrs time points to analyze cell viability. Percent PI uptake by ConA stimulated cells was calculated by subtracting percent cell death of unstimulated controls of that CD4⁺ T cell subset at that time point. (B) CD4⁺CD25⁻ T cells were stimulated with rhIL-2, or TGFβ1, or ConA +/- TGFβ1 for 2 days. Stimulated (1*10⁵) cells were stained with anti-TGFβ1 antibody and propidium iodide at 48hrs time point to analyze cell viability. % PI positive cells were determined from the total CD4⁺CD25⁻ T cell population for CD4⁺CD25⁻ T cells stimulated with TGFβ1 or rhIL-2, or ConA stimulations (**left**). But for CD4⁺CD25⁻ T cells stimulated with TGFβ1 plus ConA percentage PI staining on only TGFβ1 positive CD4⁺ T cells was determined which relates to percentage cell death of only converted Treg like cells (**right**). *p-value < 0.01 was considered significant.

A



B

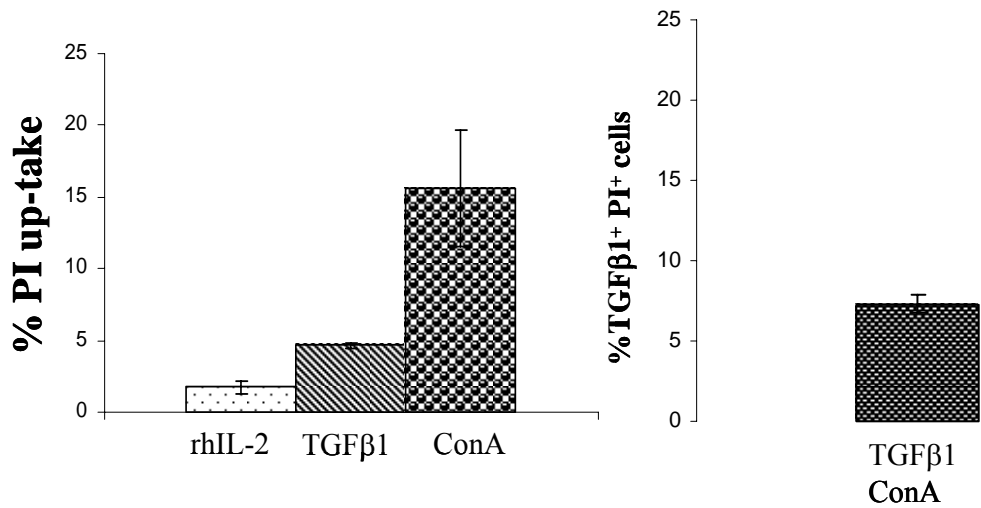
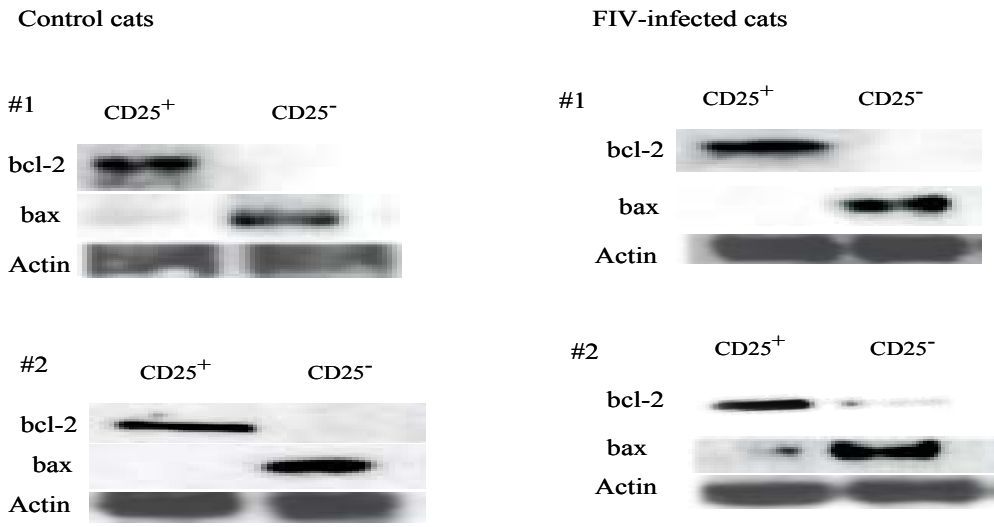


Figure 5. Differential expression of pro-apoptotic and anti-apoptotic proteins in the CD4⁺ T cell subsets. (A-B) Equal number of purified CD4⁺CD25⁺ and CD4⁺CD25⁻ T cells (2x10⁶) (A) or CD4⁺CD25⁻ T cells were stimulated with or without ConA +/- TGFβ1, or TGFβ1, or rhIL-2 for 4 days (B), from control cats (n=2) and FIV-infected cats (n=2) were lysed with NP-40 lysis buffer to obtain total cellular protein for the detection of bcl-2 and bax expression. Total cell lysates from FIV-infected and control cats were separated by SDS-PAGE and analyzed by immunoblot using specific primary antibodies against pro-apoptotic and anti-apoptotic proteins (bax and bcl-2), followed by HRP-conjugated goat anti-mouse IgG1 secondary antibody and detected by chemiluminescence. The blots were stripped using stripping buffer and reprobed with actin primary antibody and HRP-conjugated goat anti-mouse IgG1 secondary antibody followed by chemiluminescence detection.

A



B

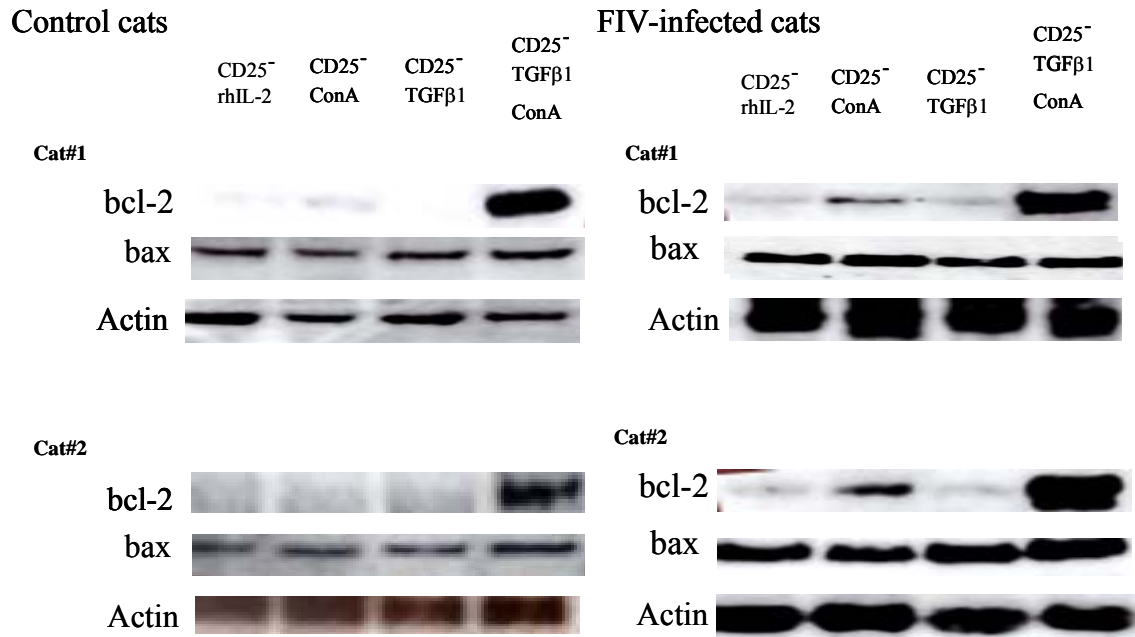
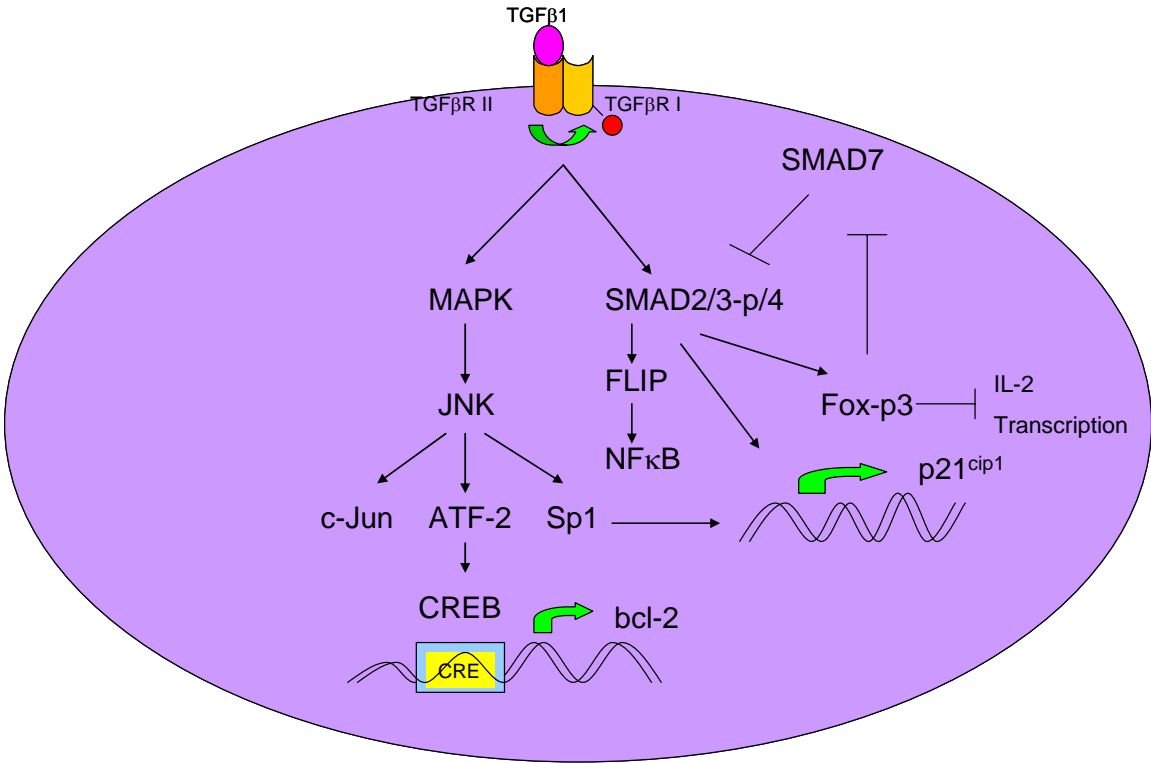


Figure 6. Schematic diagram of TGFβ1/TGFβR signaling pathway in Treg cells

Binding of TGFβ1 to its receptor leads to the formation of TGFβRI/II complex that in turn phosphorylates receptor-activated Smad 2/3 (R-Smads). R-Smads form complexes with Smad4 which allows their translocation into the nucleus where the complex interacts with Smad-binding elements. R-Smad complex promotes the transcription of several target genes including p21^{cip1}, and transcription factors such as Foxp3 and NFκB. Foxp3 in turn down regulates Smad7 which induces a positive auto-regulatory loop of TGFβ-Smad signaling. On the other hand Foxp3 inhibits IL-2 transcription. TGFβ1 signaling can also activate MAP-Kinase pathway that induces expression of several transcriptional factors such as ATF, Sp1, and c-Jun. Transcription factors of the Sp1 family along with Smads can activate p21^{cip1} transcription. ATF transcription factor induces phosphorylation of cyclic AMP response element binding proteins (CREB) which in turn targets induction of several genes including bcl-2.



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5. SUMMARY

- a) Feline CD4⁺CD25⁺ Treg cells are functionally anergic as they do not proliferate upon mitogenic stimulation.
- b) Activated feline CD4⁺CD25⁺ Treg cells are potent suppressors as they can suppress proliferation of mitogen stimulated CD4⁺CD25⁻ T cells.
- c) CdkI's such as p21^{*cip1*}, p27^{*kip1*}, and p16^{*ink4*} are differentially expressed in the total peripheral blood lymphocytes from FIV-infected and control cats.
- d) CD4⁺CD25⁺ Treg cells from FIV-infected cats but not control cats express elevated levels of CdkI proteins compared to CD4⁺CD25⁻ T cells. In contrast cyclin D and phosphorylated retinoblastoma protein (pRb) are expressed at elevated levels in CD4⁺CD25⁻ T cells compared to CD4⁺CD25⁺ Treg cells.
- e) The differential expression of CdkI's in the two purified CD4⁺ T cell subsets was observed even at RNA level.
- f) CD4⁺ T cells from FIV-infected cats are activated in-vivo as detected by reduced levels of CD62L-selectin compared to control cats.
- g) CD4⁺CD25⁺ Treg cells are more activated in FIV-infected cats than in control cats.
- h) Differential expression of the CdkI's between the two CD4⁺ T cell subsets is activation dependent.
- i) TGFβ1 is expressed at significantly elevated levels in CD4⁺CD25⁺ Treg cells from FIV-infected cats compared to autologous CD4⁺CD25⁻ T cells. No significant difference in TGFβ1 expression was observed between the two CD4⁺ T cell subsets from control cats.

- j) TGFβRII is also expressed at significantly elevated levels in CD4⁺CD25⁺ Treg cells from FIV-infected and control cats compared to CD4⁺CD25⁻ T cells.
- k) CD4⁺CD25⁺ Treg cells from FIV-infected express significantly more surface TGFβRII compared to CD4⁺CD25⁺ Treg cells from control cats.
- l) TGFβ1 along with ConA could induce functional anergy in CD4⁺CD25⁻ T cells.
- m) These converted cells are potent suppressors as they can suppress proliferation of other T cells.
- n) In correlation with functional anergy, TGFβ1 plus ConA stimulated CD4⁺CD25⁻ T cells express phenotypic molecules such as CdkI's (p21^{*cip1*}) similar to that of naturally occurring CD4⁺CD25⁺ Treg cells.
- o) Similar to naturally occurring CD4⁺CD25⁺ Treg cells, TGFβ1 plus ConA treated CD4⁺CD25⁻ T cells from FIV-Infected and control cats are more resistant to activation induced apoptosis compared to rhIL-2 stimulated CD4⁺CD25⁻ T cells.
- p) In conjunction with the reduced apoptotic potential, TGFβ1 induced CD4⁺CD25⁻ T cells similar to the professional CD4⁺CD25⁺ Treg cells also express enhanced levels of anti-apoptotic proteins (bcl-2).
- q) Naturally occurring CD4⁺CD25⁺ Treg cells express decreased levels of pro-apoptotic proteins (bax) compared to their counterparts CD4⁺CD25⁻ T cells. Mitogen along with TGFβ1 stimulation of CD4⁺CD25⁻ T cells had no effect on bax expression.

6. CONCLUSIONS

Cell cycling from G0/G1 to the S phase is regulated by a family of cyclin-dependent kinases (Cdk) and their inhibitors (CdkI's). CD4⁺CD25⁺ T regulatory cells (Treg) are arrested in the G0/G1 phase of the cell cycle and fail to produce IL-2 and proliferate in response to TCR engagement. The expression profile of the Cdk and CdkI genes in CD4⁺CD25⁺ Treg cells and the role they play in maintaining the anergic state has not been studied. We previously reported that CD4⁺CD25⁺ Treg cells from FIV-infected but not uninfected cats are activated in-vivo and possess potent immuno-suppressor function in the absence of additional in-vitro stimulation. In this study we assessed the expression profile of the CdkI's in Treg cells from FIV-infected and uninfected control cats.

We report that while CD4⁺CD25⁺ Treg cells from both FIV-infected and naïve cats are functionally anergic, as manifested by their inability to produce IL-2 and proliferate upon mitogenic stimulation, only those from FIV-infected cats are activated for strong suppressor function in vivo. Functional activation of Treg cells from FIV-infected cats correlates with up-regulation of TGFβ1 and TGFβRII on their surface and over expression of the CdkI's, p21^{cip1}, p27^{kip1}, and p16^{ink4}. As TGFβ1/TGFβR signaling is known to induce G0/G1 cell-cycle arrest by up-regulation of CdkI's, it can be speculated that the other phenotypic and functional responses observed in Treg cells from FIV-infected cats derive from over expression of the TGFβ1/TGFβR signaling pathway.

Although CD4⁺CD25⁺ T regulatory (Treg) cells are anergic and arrested in the G0/G1 phase of the cell cycle, they are programmed for cell survival. However, the molecular mechanism(s) involved in G0/G1 cell cycle arrest and cell survival signaling has not been defined. As mentioned earlier CD4⁺CD25⁺ Treg cells from chronically FIV-infected cats as opposed to those from control cats are functionally activated in vivo and express high levels of TGFβ1 on their surface and elevated levels of intercellular cyclin-dependent kinase inhibitors (CdkI's). To learn more of the potential role of cell surface expressed TGFβ1 in regulating the phenotypic and functional characteristics of Treg cells, we examined the role of exogenous TGFβ1 in conversion of CD4⁺CD25⁻ T cells to a Treg phenotype.

Herein we report that TGFβ1 stimulation along with TCR- engagement can induce functional and phenotypic anergy in converted Treg like cells, as manifested by their inability to proliferate upon mitogenic stimulation and expression of CdkI, p21^{cip1} respectively. TGFβ1 converted Treg like cells, similar to naturally occurring Treg cells, are more resistant to activation induced apoptosis compared to mitogen stimulated CD4⁺CD25⁻ T cells, which correlates with the expression of anti apoptotic proteins (bcl-2) and pro-apoptotic proteins (bax). Therefore, TGFβ/TGFβR signaling may be one of the mechanisms by which Treg cells could induce G0/G1 cell cycle arrest by up-regulation of CdkI's, yet be resistant to apoptosis by maintaining anti-apoptotic and pro-apoptotic protein ratios for survival. Further, molecular characterization of CD4⁺CD25⁺ Treg cells from FIV chronically infected cats may be helpful in providing experimental basis for understanding the immunopathogenic potential of activated Treg cells and

developing immunotherapeutic strategies in treatment for chronic diseases such as HIV/AIDS infection.