Review Article

TNF and TNF Receptor Superfamily Members in HIV infection: New Cellular Targets for Therapy?

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Tumor necrosis factor (TNF) and TNF receptors (TNFR) superfamily members are engaged in diverse cellular phenomena such as cellular proliferation, morphogenesis, apoptosis, inflammation, and immune regulation. Their role in regulating viral infections has been well documented. Viruses have evolved with numerous strategies to interfere with TNF-mediated signaling indicating the importance of TNF and TNFR superfamily in viral pathogenesis. Recent research reports suggest that TNF and TNFRs play an important role in the pathogenesis of HIV. TNFR signaling modulates HIV replication and HIV proteins interfere with TNF/TNFR pathways. Since immune activation and inflammation are the hallmark of HIV infection, the use of TNF inhibitors can have significant impact on HIV disease progression. In this review, we will describe how HIV infection is modulated by signaling mediated through members of TNF and TNFR superfamily and in turn how these latter could be targeted by HIV proteins. Finally, we will discuss the emerging therapeutics options based on modulation of TNF activity that could ultimately lead to the cure of HIV-infected patients.

1. Introduction

The term tumor necrosis factor (TNF) came into existence in 1975 with the work of Carswell and colleagues while studying hemorrhagic necrosis by endotoxin [1]. It was described as a host factor, a glycoprotein induced in response to endotoxin that has the capacity to kill the tumor. As the time progressed, TNF was realized to be rather a member of a superfamily that governs by binding to their receptors. TNF and TNF receptors (TNFR) are growing members of ligand and receptor superfamily that regulate several complex signaling pathways leading to apoptosis, inflammation, cellular differentiation, and antiviral state. The first member of TNF superfamily discovered is TNF-alpha (old name cachectin), a pleiotropic proinflammatory cytokine that plays pivotal role in several pathological conditions due to inflammation and infection [2]. Role of TNF in malignancies and inflammation conditions like arthritis have been reviewed extensively elsewhere [3–5].

Till date TNF superfamily comprises of 19 ligands and 29 receptors [4]. All members are proinflammatory in nature playing diverse roles [4]. Most of the members act like dual edge sword, both beneficial and in adverse role [4, 6, 7]. First two members of TNF ligand (TNFL) superfamily were TNF-alpha and TNF-beta, recognized first at protein level followed by identification of their respective cDNAs, while rest of the members were discovered based on cDNA sequence homology [4, 8, 9]. All members of TNF superfamily and their receptors have been comprehensively reviewed recently [4]. Besides TNF-alpha and TNF-beta, TNFL superfamily include CD40L, CD30L, FasL, TNF-related apoptosis-inducing ligand (TRAIL), lymphotoxin-beta (LT-beta), LIGHT, receptor activator of NF-kappaB ligand (RANKL), 4-1BBL, CD27L, OX40L, TNF-related weak inducer of apoptosis (TWEAK), a proliferation-inducing ligand (APRIL), B-cell activating factor (BAFF), vascular endothelial cell-growth inhibitor (VEGI), ectodysplasin A (EDA)-A1, EDA-A2, and GITRL

[4, 10]. TNFR superfamily includes TNFR1, TNFR2, LT-betaR, OX40, CD27, CD40, CD30, 41-BB (CD137), Fas, TRAILR1 (DR4), TRAILR2 (DR5), TRAILR3, TRAILR4, OPG, RANK, Decoy (DC) R3, TWEAKR, NGFR, Transmembrane Activator and CAML interactor (TACI), BAFFR, LIGHTR (HVEM), DR3, glucocorticoid induced TNF receptor (GITR), EDAR, XEDAR, TROY, RELT, DR6, and B-cell maturation protein (BCMA) [4, 7]. Extracellular domains of TNFR family members have a typical cysteine rich motif. However, intracellular domains show variation contributing to diverse functions [7, 11]. On the basis of presence or absence of 45 amino acid long regions in their intracellular domain called death domain, TNFR members are categorized into two groups [4]. Presence of death domain is critical for the interaction with other proteins leading to cell death. For example, TNFR1 possess this death domain on the other hand, TNFR2 does not have the death domain.

Number of TNF ligand versus receptor suggests that some of the ligands interact with more than one receptor to achieve their goal [4]. TNF ligands and receptors are mostly expressed by immune cells. However, under certain pathophysiological conditions their presence has been documented in other cell types as well.

2. TNF-Alpha-Mediated Cell Signaling: An Overview

Most extensively studied member of TNF superfamily is TNF-alpha. TNF-alpha is produced in response to pathological conditions like inflammation and infection mainly by activated macrophages and T lymphocytes [4, 7], but also by several cell types including natural killer (NK) cells, mast cells, and fibroblasts. TNF-alpha is synthesized as pro-TNF, a 25 kDa plasma membrane bound protein that is further processed by metalloproteinase called TNF-alpha converting enzyme into a 17 kDa soluble form [12]. Both forms are functional in their trimeric forms via binding to their receptors. Data suggest that plasma membrane associated 25 kDa TNF-alpha form binds to the TNFR2 with high affinity whereas soluble 17 kDa form interacts with TNFR1 with high specificity [13, 14].

TNF-alpha triggers several signaling cascades which include apoptotic pathways, NF-kappaB stimulation, and activation of p38 MAPK, ERK, and JNK [4, 7] (Figure 1). Binding of the ligand TNF-alpha to its receptor TNFR1 leads to the recruitment of a 34 kDa adapter protein called TNFR-associated death domain (TRADD). Latter interacts with the cytopathic death domain of TNFR1 through its own death domain [15] (Figure 1). Further TRADD directly binds to Fas-associated death domain protein (FADD) and activates apoptosis via caspase cascade [16] (Figure 1). On the other hand, TRADD also interacts with TNF receptor associated factor (TRAF) protein 2 followed by sequential recruitment of receptor interacting protein (RIP), TGF-beta-activated kinase 1 (TAK1), and IκB kinase (IKK) complex [7, 17, 18]. IKK phosphorylates IκB resulting in the degradation of IκB. As a result, NF-kappaB translocates into the nucleus to activate transcription of effector molecules including mediators of

inflammation such as chemokines, interleukin-(IL)-6, IL-8, IL-18, and cyclooxygenase-2 [4, 7, 17, 19] (Figure 1).

In addition, TNF-alpha can induce cell proliferation through induction of transcription factor called activator protein-1 (AP-1) by binding to TNFR1 followed by sequential contribution of TRADD, TRAF2-RIP, MEKK1, MKK7 and JNK [4, 7, 20] (Figure 1).

Cell signaling associated with TNFR2 is poorly understood. TNFR2 lacks death domain, despite of that it interacts with TRAF2 through which it can activate transcription factors NF-kappaB and AP-1 (Figure 1). There are several reports where TNFR2 has been reported to be involved in cellular proliferation, apoptosis, and the induction of granulocyte-macrophage colony-stimulating factor (GM-CSF) secretion [21–23].

Significance of TNF-alpha can be evaluated by this fact that several human pathogens have evolved mechanisms to combat TNF-alpha-mediated response against infection [24]. 2013 has been marked as 30 years of discovery of HIV. In this review, we will focus on TNF and TNFR and their family members in context to HIV infection and potentially how to modulate them by TNF inhibitor therapy.

3. Role of TNF and TNFR Superfamily Members in HIV Pathogenesis

3.1. TNFR1 and TNFR2

3.1.1. TNF and HIV Entry. The first and foremost part of any virus life cycle is its entry into permissive cells. TNF-alpha is known to target HIV entry step specifically in macrophages but not in peripheral blood lymphocytes [25]. Notably, in cell culture, TNF-alpha is released by primary macrophages infected with HIV type 1 (HIV-1) or treated with HIV envelope protein gp120 [26]. One of the plausible strategies of inhibition of HIV entry by TNF-alpha may be by downregulating the expression of HIV receptor and coreceptors on cell surface (CD4 and CCR5) that may explain inhibition of HIV entry into permissive cells [27]. In addition, GM-CSF secretion is stimulated by TNF-alpha that in turn can downregulate CCR5 and may inhibit the entry of CCR5-dependent viruses into macrophages [28]. Moreover, it has been reported by Herbein and colleagues that pretreatment of tissue culture-differentiated macrophages (TCDM) with human recombinant TNF-alpha (hrTNF-alpha) resulted in remarkable delay in detection of HIV DNA long terminal repeat (LTR) as a result of strong inhibition of virus entry into these cells. Furthermore, using TNF-R1 and TNF-R2 mutants, they demonstrated that this inhibition was mediated through TNF-R2 not TNF-R1 [25–27].

3.1.2. TNF and HIV Postentry Stages. TNF-alpha can activate HIV-1 in chronically infected T cell lines and promonocytic cell lines through translocation of NF-kappaB to the nucleus followed by activation of HIV LTR [29–33]. However, contradictory findings have been also reported where TNF-alpha has shown to inhibit HIV-1 replication in several cell types including freshly infected peripheral blood monocytes, alveolar macrophages, and TCDM [25, 34, 35]. These findings

FIGURE 1: TNF and TNFR superfamily-mediated cell signaling. Binding of TNF- α to TNFRs results in activation of its receptors followed by recruitment of adaptor proteins (TRADD, FADD, TRAF, and RIP) in sequential manner that activates several signaling cascades leading to the activation of transcription factors NF- κ B, AP-1, and/or caspase cascades. Most of the members of TNF superfamily activate also NF- κ B.

indicate that TNF- α may have contrasting impact on HIV-1 replication in chronically infected cells and cells coming in contact with the virus for the first time [35]. In addition, TNF- α induces several HIV suppressive factors such as RANTES in lymphoid cells [36, 37] and alveolar macrophages [35], MIP-1 α , and MIP-1 β in human fetal microglia cells [38, 39] that may explain the negative role of TNF- α in HIV replication.

3.1.3. TNF and HIV-Induced Apoptosis and Transformation. CD8 $^{+}$ T-cell apoptosis that occurs in HIV pathogenesis could result from the interaction between macrophage-membrane bound TNF- α with TNFR2 present on CD8 $^{+}$ T cells [40]. Additionally, HIV-1 Tat is known to induce the expression of TNF- β in a human B-lymphoblastoid cell line (Raji cells). There is a possibility that HIV-1 Tat protein induces the growth of Kaposi's sarcoma cells via TNF- β induction [41–43].

3.2. CD40. CD40 a member of TNFR superfamily, is a 45–50 kDa integral membrane glycoprotein found on B-cells, monocytes, dendritic cells, endothelial cells, and epithelial cells [44, 45]. Ligand for CD40 is CD40L (CD154), a 33 kDa transmembrane glycoprotein that is mainly expressed by activated B-cells, T-cells, and platelets [45, 46] (Figure 1). CD40-CD40L mediated signaling plays indispensable role in the development of cellular and humoral responses [46]. Membrane dissociated truncated soluble CD40L (sCD40L) is released by activated cells and binds to the CD40 molecule expressed on the target cells to activate it [46, 47].

Interestingly, increased levels of sCD40L in the cerebrospinal fluid and plasma of HIV-infected patients with cognitive impairment have been documented [46, 48]. *In vitro* experiments with recombinant CD40L (rCD40L) and HIV-1 Tat show that they act synergistically to enhance the yield of TNF- α by microglia and monocytes [46]. This enhancement may be contributed by ability of Tat to increase CD40 expression via NF- κ B activation [46, 49]. Furthermore sCD40L interacts with CD40 leading to CD40-mediated signal cascade resulting in activation of NF- κ B in microglia and monocytes [46]. This results in production of high amounts of inflammation mediators such as TNF- α that may explain HIV-associated dementia [46].

Tat is a multifunctional HIV protein. Notably, platelet activation has been documented in HIV-1-infected individuals [50–52]. Tat has been reported to activate platelet to release CD40L via interacting with chemokine receptor CCR3 and beta3 integrin both *in vitro* and *in vivo* [52]. Tat-stimulated CD40L activates B-cells resulting in increase in Ig yield [52]. Thus there is fair possibility that Tat-stimulated CD40L in platelet may contribute to HIV-related thrombocytopenia [52]. In addition, Tat is known to enhance the expression of TNF ligand superfamily members FasL and TRAIL in macrophages and T cells. As a consequence apoptosis can be induced in the bystander cells [53–55].

Worth mentioning, CD40L can be embedded on the surface of HIV-1 virion generated by peripheral blood and through budding from stimulated CD4 $^{+}$ T cells in cell culture as well as in HIV-1 infected patients [56–58]. CD40L-associated virions can induce strong activation of B-cells

and modulate genes including members of TNF superfamily (FAS, A20, TNIP1, CD40, lymphotoxin alpha, and lymphotoxin beta), cytokines, and transcription factors [57, 58]. Additionally, macrophages expressing Nef or activated by CD40L-CD40 receptor interaction release factors (CD23 and soluble ICAM) which makes T cells present in their vicinity susceptible to HIV infection, thus expanding the HIV cellular reservoir [59].

3.3. Lymphotoxin-Beta Receptor (LT-betaR). Another important member of growing TNFR superfamily is lymphotoxin-beta receptor (LT-betaR) that governs the signaling pathways involved in the organogenesis of lymphoid tissue and function of follicular dendritic cells in a manner distinct from TNFR signaling pathways through activation of NF-kappaB [60–65] (Figure 1). LT-betaR stimulation favors HIV-1 replication in monocytes [62]. Additionally, when TNF receptors and LT-betaR are activated by their respective ligands (TNF-alpha and LT-alpha1beta2), an additive effect on HIV-1 replication is observed in U1 cells [62].

3.4. CD27. CD27, a TNFR superfamily member, is a 55 kDa type 1 transmembrane protein that exists in a homodimeric form [66]. Ligand for CD27 is CD27L (CD70), a member of TNF ligand superfamily. CD27 plays an important role in the activation of T cells and infection of T cells by HIV-1 [67, 68]. One critical feature of HIV-1 life cycle is its integration into the host genome using virus-encoded integrase. There are reports describing preference of HIV integration into transcriptionally active genes [68, 69]. A recent study shows that HIV-1 integrates into the coding region of CD27 gene in CD4+ T cells which may disturb CD27 open reading frame and hence can hamper the help response of CD4+ T cells [68]. This can result in inefficient removal of HIV-1 from CD4+ T cells by cytotoxic CD8+ T cells [68].

3.5. CD30. TNFR superfamily member CD30 is present on activated T cells, B-cells, several other transformed lymphocytes, and NK cells [70–72]. CD30 plays role in triggering developmental process in B-cells via CD30-CD30 ligand interaction [72, 73]. Ligation of CD30 with an anti-CD30 monoclonal antibody (functionally equivalent of CD30L) in chronically HIV-1-infected human T cell line ACH-2 has been reported to enhance the HIV gene expression via binding of activated NF-kappaB to the HIV-1 LTR [74, 75]. Investigation of molecular mechanism responsible for the induction of NF-kappaB revealed that NF-kappaB translocation into nucleus was mediated by TRAF2, independent of TNF-alpha/beta (Figure 1).

3.6. Fas. The Fas also known as Apo-1, CD95 and TNFSF6, is a TNFR superfamily member that governs apoptosis when activated by its ligand FasL (Figure 1). Role of Fas-FasL signaling cascade in HIV pathogenesis has been extensively reviewed [55, 76]. Production of Fas and FasL is increased in CD4+ T cells isolated from HIV infected individuals [77, 78]. Increased expression of Fas is observed in B-cells, CD4+, and CD8+ T cells whereas FasL increased expression is associated with macrophages, NK cells, and monocytes [78, 79]. *In vitro*

studies reveal that HIV-infected macrophages can induce apoptosis in Jurkat T cells and in peripheral blood T lymphocytes via FasL that could be one factor responsible for the depletion of lymphocytes during HIV pathogenesis [78, 80].

3.7. 4-1BB (CD137). 4-1BB (CD137) is a TNFR expressed predominately on T cells, NK cells, mast cells, and neutrophils [4]. Ligation of 4-1BB with agonistic monoclonal antibodies has been shown to effectively increase the HIV-1 replication in CD4+ T cells isolated from HIV-1 infected patients. There is a possibility that 4-1BB receptor may be involved in the activation of HIV from latency in CD4+ T cells [81].

3.8. OX40 (CD134). OX40 (CD134), a member of the TNFR superfamily, is expressed on activated CD4+ T cells and neutrophils. It is crucial for the survival of antigen specific CD4+ T cells [4, 82]. Natural ligand for this receptor is OX40L (also called CD252 and gp34), a member of TNFL superfamily. OX40-gp34 interaction, in HIV-1 (both acutely and chronically) infected T cell lines result in increase in HIV-1 replication, independent of TNF-alpha or TNF-beta production. The increase in viral replication has been shown to mediate by activation of NF-kappaB followed by stimulation of HIV-1 LTR [33]. Recent study suggests that OX40 activation suppresses the CCR5-tropic (R5) HIV-1 infection in PBMCs by generating anti-HIV beta-chemokines [83].

3.9. DR 4 and DR 5. Death receptor (DR), DR4 (also called TRAILR1, TNFRSF10A, and Apo2) and DR5 (also called TRAILR2 and TNFRSF10B) are the fourth and fifth members of TNFR superfamily, respectively. They are expressed in most of the normal as well as transformed cells [4]. They govern their activity by binding to their ligand TRAIL (tumor-necrosis-factor related apoptosis inducing ligand or Apo2L) leading to receptor oligomerization, assembly of death inducing signaling complexes that ultimately govern the activation of caspase pathways [4, 78]. TRAIL is a TNFL superfamily member expressed in NK, T cells, and dendritic cells. In HIV-infected CD4+ T cells, TRAIL is involved in the apoptosis of infected cells by binding with DR4 and DR5 [4, 78]. Plasma level of TRAIL has been found to be high in HIV-1 infected individuals, whereas HIV-1 infected patients undergoing antiretroviral therapy demonstrate decrease in TRAIL levels in plasma and also decreased viral load suggesting crucial role of TRAIL in HIV-1 pathogenesis [84]. *In vitro* testing of recombinant TRAIL (rTRAIL) against HIV-infected peripheral blood lymphocytes and monocyte-derived macrophages isolated from HIV-infected patients results in apoptosis of the target cells. However, rTRAIL shows no effect against target cells isolated from uninfected patients [84]. This raises the possibility of using TRAIL as anti-HIV agent.

4. HIV Proteins Mimicking TNF/TNFR Signaling

Several HIV proteins especially Vpr, Tat, and Nef exhibit molecular mimicry with respect to TNF signaling in HIV-infected cells particularly in macrophages (Figure 2).

Viral protein R (Vpr) is a small (14 kDa) multifunctional virion-associated accessory protein that participates in import of viral preintegration complexes to the nucleus. In addition, Vpr exerts antiapoptotic effect in HIV infected cells on the other hand, it induces apoptosis in the surrounding cells [85]. Besides these functions, Vpr also triggers mitochondrial dysfunction [86, 87]. Although Vpr is dispensable for HIV replication in T cells, it is critical for HIV replication in nondividing cells, for example, macrophages [10] (Figure 2). There are several reports where Vpr has been shown to stimulate HIV-1 growth using serum derived or synthetic Vpr [88, 89]. Virion derived (HIV-1) Vpr activates NF-kappaB in primary T cells, macrophages as well as in promonocytic cell line U937 [90]. Similarly, synthetic Vpr has been shown to activate NF-kappaB, AP-1, and JNK in primary macrophages and U937 cells resulting in increase in viral replication [10, 89] (Figure 2). In addition, recombinant Vpr stimulates HIV production by utilizing toll-like receptor 4 (TLR4) and IL-6 secretion [87]. Synergistic effect of Vpr and Tat leads to enhanced stimulation of HIV-1 LTR [91]. Even Tat alone mimics TNF-alpha. Like TNF-alpha, Tat triggers translocation of NF-kappaB into nucleus and activation of AP-1/cJun by MAPK activation JNK, p38, and ERK1/2 [10, 92] (Figure 2).

In contrast, there are several reports describing the suppression of NF-kappaB activity by Vpr. It has been shown that glucocorticoid receptor and Vpr act in harmony to inhibit NF-kappaB mediated gene expression via a pathway involving the suppression of poly (ADP-ribose) polymerase (PARP)-1 nuclear trafficking in response to TNF-alpha [85, 93].

HIV Nef, the most abundantly expressed HIV accessory protein, is a multifunctional 27 kDa myristoylated cytoplasmic protein expressed in early phase of viral life cycle [94]. Nef helps in the establishment of HIV persistence in infected cells [95] and interferes with several signaling events [10]. Recombinant Nef (rNef) has been shown to induce expression and release of several cytokines mediated by NF-kappaB activation in culture monocyte-derived macrophages (MDMs) [96, 97]. In U937 cells and MDMs, exogenously added rNef triggers NF-kappaB activation resulting in HIV LTR activation [98] (Figure 2). In addition, rNef induce, transcription of several inflammatory genes in response to addition of rNef to MDMs. Convincible, analysis of rNef treated MDMs supernatants revealed induction and release of TNF-alpha and other macrophage inflammatory proteins (MIP-1alpha and MIP-1beta) and IL6 [96]. Moreover, in chronically infected promonocytic cells U1, addition of rNef leads to increase in HIV-1 replication [98]. Furthermore, rNef is able to rapidly and transiently induce phosphorylation of several key-signaling molecules including alpha/beta subunits of Ikappa B kinase, ERK1/2, JNK, and p38 in MDMs [97]. Signaling scenario observed in post-rNef treatment is more or less similar to what is observed in post TNF-alpha treatment [10] suggesting their similar impact on HIV infection at least in mononuclear macrophages (Figure 2).

5. Targeting Members of TNF and TNFR Superfamily in HIV-1 Infection

In this section, we will discuss current status and future potential therapeutic use of TNF based therapies for HIV-1 and HIV-1 related diseases and their pros and cons.

The targeting of TNF signaling has proven the most successful and clinical efficacious therapy at reducing the inflammation in several diseases. Several TNF related inflammatory cytokines and their cognate receptors are now in preclinical or clinical phase of development as a possible target for treating various diseases such as cancer, autoimmune, and inflammatory disorders. HIV infection is characterized by immune activation and inflammation [99]. Therefore, TNF blocking agents and/or TNF inhibitor therapy could be useful to modulate HIV disease.

There are several types of drugs targeting TNF and TNFR superfamily that are being used for therapeutic application [100]. Five different antibodies or receptors based drugs targeting TNF and LTalpha are approved for treating various inflammatory diseases. The chimeric antibody infliximab, TNF targeted drug, was approved in 1998 followed by etanercept [101]. The fully human antibodies adalimumab and golimumab were approved in 2002 and 2009, respectively. Certolizumab pegol, another therapeutic monoclonal antibody with pegylated Fab fragment was approved in 2008. As polyethylene glycol does not cross the placenta, this drug can be administered to pregnant women who have autoimmune diseases and are in need of anti-TNF-alpha therapy [102]. These drugs neutralize the biological activity of TNF-alpha by binding with high affinity to the soluble as well as transmembrane form of TNF-alpha. Thus they prevent the binding of TNF-alpha to their natural receptors. Worth mentioning, drugs adalimumab and infliximab have the potential to lyse the cells involved in inflammation. Besides the availability and affordability, these drugs fall in the class of immune suppressors which may have serious complications such as blood disorders, infections, liver injury, skin lesion, and reactivation of tuberculosis [100].

5.1. Anti-TNF Therapy and HIV-1 Infection. HIV-1 infection induces TNF expression, and high amount of TNF is present in all stages of HIV-1 infection [29, 103, 104]. This elevated level of serum TNF has been associated with increased viral replication and depletion of CD4+ T cells [105, 106]. The treatment of HIV-1 infected patients with thalidomide (a weak TNF inhibitor) reduces serum TNF level that results in lower viral load [107, 108]. Furthermore, it has been shown that LMP-420, a small inhibitor of TNF, suppresses the transcription and biosynthesis of TNF, which ultimately inhibits the replication of HIV-1 [109]. Multiple studies have reported that use of anti-TNF therapy in patients with HIV-1 does not appear to increase the mortality rates [110]. As there is a theoretical risk that immunosuppressive drugs increase the risk of opportunistic infection and progression to HIV-1 disease, several studies have shown that anti-TNF therapy may improve the HIV-1 associated symptoms [111]. Further, etanercept has been used as anti-TNF therapy in HIV-1 infected patients for the treatment of rheumatic disease. In

FIGURE 2: TNF signaling mimicking by HIV proteins. Early expressed HIV proteins Vpr, Tat, and Nef mimic TNF- α as they activate TNF- α governing signaling pathways to enhance HIV replication in infected cells.

most of the cases, the therapy was well tolerated. In addition, no opportunistic infection was observed in HIV-1 infected rheumatic patients unless they had uncontrolled HIV [112, 113]. Similar results were obtained by using other anti-TNF agents such as infliximab and adalimumab in HIV-1 infected patients. These drugs were safe and effective with normal CD4+ T cell counts [114, 115]. Therefore, in the patients whose HIV disease is under control of HAART, the anti-TNF therapy may be helpful for the treatment of autoimmune disease without enhancing plasma viremia [111, 116]. Worth mentioning, above clinical studies are based on a little number of patients; therefore, results must be analyzed based on large cohorts to assert a safe and real benefit to the community.

5.2. Costimulatory TNFRs and HIV-1-Specific T Cell Response. Several studies have been carried out to compare the efficacy of different costimulatory TNFR family members for the activation of HIV-1-specific T cells *in vitro*. These costimulatory signaling pathways could be used to activate CD8 T cell responses to HIV *in vivo* (Figure 3) [117]. The OX40L signaling pathway plays an important costimulatory role in DC/T cell interactions. OX40 binding to CD4+ T cells by human OX40L-IgG1 enhances *ex vivo* expansion of HIV-1 specific CTL from HIV-1 infected individuals [118]. This mechanism of CTL expansion was independent of induction of cytokines such as IL-2 or any inhibitory effect on CD4+ T helper cells, but it was associated with a direct effect on proliferation of CD4+ T cells. This mechanism of action of OX40 represents a potentially novel immunotherapeutic strategy that could prevent the persistent HIV-1 infection [118]. Like OX40, the 4-1BB is transiently expressed following TCR ligation [119, 120]. Furthermore, the ligation of 4-1BB/4-1BBL enhances CTL expansion that has both antiviral and

antitumor activity [121]. In addition, dual costimulation by OX40L in combination with 4-1BBL resulted in improved expansion and effector function of CTL over costimulation with individual costimulatory molecules [122]. Furthermore, urelumab is a monoclonal antibody that specifically binds to and activates 4-1BB expressing immune cells and stimulates the CTL response against tumor cells. Although 4-1BB ligand (4-1BBL) could be an important costimulatory molecule for exhausted CD8+ T cells from chronically infected patients, anti-4-1BB therapy is associated with several immunological side effects such as splenomegaly, hepatitis, and several immunological disorders [123, 124]. Additionally, it has been reported that stimulation of 4-1BB in T cells enhances HIV-1 replication [81]. To be an effective therapy, the 4-1BB agonist should induce HIV-1 specific CD8+ T cell response and also should not induce viral replication. This issue may be solved by incorporation of 4-1BBL along with CD8+ T cell epitopes into vaccine vectors. The incorporation of 4-1BBL into a fowlpox vector along with HIV-1 Gag enhances Gag-specific CD8+ T cell responses, suggesting this could be a useful approach in a therapeutic vaccine [125].

5.3. TNF and TNFR Superfamily and HIV-1 Reservoirs. The persistence of HIV-1 reservoirs is a major challenge for complete viral eradication in HIV-1 infected patients [126–128]. To date, resting or memory CD4+ T cells are the most well-characterized HIV-1 reservoirs [128]. During the development of memory CD4+ T cells, the NF- κ B signaling ensures T cell survival during the initial differentiation of effector cells into memory T cells. OX40 along with CD30 induces NF- κ B-dependent expression of antiapoptotic genes such as Bcl-2 and Bcl-xL that in turn play an important role in the survival of memory CD4+ T cells

FIGURE 3: 4-1BB and OX40 signaling enhance anti-HIV immunity, leading to therapeutic effects. Agonists 4-1BB or OX40 specific antibodies can induce enhanced anti-HIV T cell (CD4+ and CD8+ T cell) response in HIV-1 infected patients. The ligation of OX40/4-1BB antibodies activates OX40/4-1BB signaling that results in enhanced CD4+ T cell response through increased cytokine production and increased survival of memory T cells. TRAFs and NF-kappaB signaling has been shown to be important for the generation of memory T cells. Potential antiviral mechanisms of OX40/4-1BB include an increase in CD8+ T cell cytotoxicity through perforins and granzymes and an increase in Fas/FasL mediated HIV-infected cell killing.

within the small intestine lamina propria [129]. Low levels of NF-kappaB activity by control of the TNF and TNFR superfamily members may contribute to the establishment and maintenance of latent HIV-1 reservoirs in memory CD4+ T cells. Activation induced purging of HIV-1 in patients receiving HAART has long been a proposed mechanism to eradicate the latent HIV pool [130]. TNF-alpha has been used to reactivate HIV-1 from latently infected cells, but it is not the optimal reactivation treatment [131]. In Jurkat based HIV-1 latency model, TNF-alpha consistently activates latent HIV-1 provirus. However, in primary CD4 T cell model, TNF-alpha does not appear too effective to purge HIV-1 from latent reservoirs [132, 133]. Moreover, there are some concerns about the toxicity associated with TNF treatment [78]. Since TNF-alpha is only a weak enhancer of HIV-1 reactivation to purge cellular reservoirs, new approaches have been used to target the epigenetic regulation of HIV gene expression (Figure 4). Many studies have reported that the combination of HDACIs (histone deacetylase inhibitors) with TNF-alpha synergistically reactivates HIV-1 from latency [134, 135]. The HDACIs such as trichostatin A (TSA),

trapoxin (TPX), valproic acid (VPA), and sodium butyrate (NaBut) activate HIV-1 transcription by remodeling nucleosome 1 of HIV-1 promoter [136, 137]. Thus, an ideal anti-AIDS therapy would consist in eliminating the pool of latently cells by inducing forced HIV-1 gene expression by HDACIs and TNF-alpha, while maintaining an effective HAART regimen [131, 136, 138]. Furthermore, HDACIs potentiate TNF-alpha mediated NF-kappaB activation and also delay IκBα cytoplasmic reappearance [139]. Thus, the use of TNF-alpha and HDACIs in the presence of HAART not only could purge HIV-1 for latent reservoirs but also could suppress plasma viremia and formation of further new viral reservoirs. In addition to HDACI, inhibitors of histone methyltransferase (HMTI) have also been recently shown to reactivate HIV-1 from latency in resting CD4+ T cells from HIV-infected HAART-treated patients [140]. Furthermore, JNK inhibitors, such as AS601245, prevent HIV-1 reactivation from latency despite potent NF-kappaB activity [141]. Since stimulation of CD40 and 4-1BB induces JNK activation [142], the use of agonist of 4-1BB or CD40 could lead to the reactivation of the HIV-1 from latency. Combining TNF-alpha treatment

FIGURE 4: Potential TNF-based therapies to reactivate HIV-1 from latency. During HIV-1 latency, nuc-1 nucleosome is hypoacetylated. CTIP-2 interacts with Sp1, switching nuc-1 from transcriptionally active to a repressive state. Furthermore, CTIP-2 recruits HDACs that deacetylate the nuc-1 nucleosome. SUV39H1 adds trimethylation mark onto the histone protein H3. Furthermore, HP1 protein stabilizes the nuc-1 in transcriptionally silent state. The combination of TNF-alpha with HDACIs or HMTIs can disrupt the HIV-1 latency. TNF-alpha and HDACIs can trigger the activation of transcriptional activators, such as NF-kappaB (p50/p65 heterodimer). HDACIs may prevent the formation of heterochromatin, resulting in nuc-1 hyperacetylation and remodeling, thereby alleviating the HIV transcriptional block. The use of TNF, HDACIs, and HMTIs enhances HIV-1 LTR transcription and could participate in the clearance of HIV-1 from cellular reservoirs and ultimately could lead to the cure of HIV-infected patients.

with HDACIs/HMTI could have significant impact on the clearance of HIV-1 from cellular reservoirs and ultimately could lead to the cure of HIV-infected patients [143].

6. Conclusion

TNF ligands and TNF receptors superfamily are the integral part of our immune system. TNF signaling exerts significant impact on HIV life cycle. In turn, HIV encoded proteins also modulate TNF signaling pathways resulting in the survival of HIV-infected cells and killing of bystander cells. Anti-TNF therapy has been successfully used in several inflammatory diseases. Combinatorial therapy involving HAART, anti-TNF therapy, along with use of HDACIs/HMTI might be a viable option for the treatment of HIV infection to reach the ultimate goal, the clearance of HIV-1 from cellular reservoirs.

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