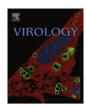
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# Caveolin-1 suppresses Human Immunodeficiency virus-1 replication by inhibiting acetylation of NF- $\kappa$ B

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#### ABSTRACT

Caveolin-1 is an integral membrane protein primarily responsible for the formation of membrane structures known as caveolae. Caveolae are specialized lipid rafts involved in protein trafficking, cholesterol homeostasis, and a number of signaling functions. It has been demonstrated that caveolin-1 suppresses HIV-1 protein expression. We found that co-transfecting cells with HIV-1 and caveolin-1 constructs, results in a marked decrease in the level of HIV-1 transcription relative to cells transfected with HIV-1 DNA alone. Correspondingly, reduction of endogenous caveolin-1 expression by siRNA-mediated silencing resulted in an enhancement of HIV-1 replication. Further, we observed a loss of caveolin-mediated suppression of HIV-1 transcription in promoter studies with reporters containing mutations in the NF- $\kappa$ B binding site. Our analysis of the posttranslational modification status of the p65 subunit of NF- $\kappa$ B demonstrates hypoacetylation of p65 in the presence of caveolin-1. Since hypoacetylated p65 has been shown to inhibit transcription, we conclude that caveolin-1 inhibits HIV-1 transcription through a NF- $\kappa$ B-dependent mechanism.

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#### Introduction

Human Immunodeficiency Virus-1 (HIV-1) is a human retrovirus that infects CD4+ T lymphocytes and myeloid cells. Our studies and those of others have demonstrated that HIV-1 biology is critically dependent upon cholesterol (Chertova et al., 2006; Graham et al., 2003; Liao et al., 2001). HIV infection of T cells enhances expression of genes encoding proteins involved in cholesterol biosynthesis (Zheng et al., 2003; Zheng et al., 2001). Conversely, inhibiting the cholesterol biosynthetic pathways during HIV-1 infection suppresses replication of the virus (del Real et al., 2004). Cholesterol-enriched cellular membrane microdomains known as lipid rafts are the sites for HIV-1 entry and egress (Liao et al., 2001; Nguyen and Hildreth, 2000). The treatment of cells with cyclodextrins has a potent effect on the integrity of lipid rafts (Graham et al., 2003; Liao et al., 2001). Notably, HIV-1 susceptible cells treated with cholesterol-sequestering cyclodextrins are less susceptible to HIV-1 infection (Graham et al., 2003; Liao et al., 2001; Liao et al., 2003). Additionally, disruption of lipid rafts reduces virion infectivity and release from cholesterol-enriched cells (Booth et al., 2006;

Gould et al., 2003; Jouvenet et al., 2006; Liao et al., 2003; Nguyen et al., 2003; Welsch et al., 2007). The dependence on cholesterol for multiple aspects of HIV-1 biology suggests that cellular mechanisms that regulate cholesterol homeostasis may modulate HIV-1 replication (del Real et al., 2004; Graham et al., 2003; Liao et al., 2001; Liao et al., 2003; Mujawar et al., 2006; Waheed et al., 2007).

In addition to cholesterol and glycosphingolipids, lipid rafts are also comprised of proteins (Hooper, 1999; Lajoie and Nabi. One such protein, caveolin-1, is a cholesterol-binding protein that is responsible for the formation of caveolae, specialized membrane structures that represent a distinct class of lipid rafts (Kurzchalia and Parton, 1999). Caveolins are a family of proteins that consist of four isoforms including: caveolin  $-1\alpha$ ,  $-1\beta$ , -2and -3. These proteins are expressed in most nucleated cells, excluding T-lymphocytes (Arakawa et al., 2000; Glenney and Soppet, 1992; Matveev et al., 1999; Razani et al., 2002). Caveolae are 60-100 nm flask-like invaginations in the plasma membrane. Caveolae, like lipid rafts in general, are enriched in cholesterol, sphingolipids, and other raft-specific constituents (Dobrowsky, 2000; Fielding and Fielding, 2000; Frank et al., 2006; Harder and Simons, 1997: Hooper, 1999: Kurzchalia and Parton, 1999: Ouest et al., 2004). Caveolae function as scaffolds for signaling proteins and as vesicles for intracellular cholesterol transport (Arakawa et al., 2000; Fielding and Fielding, 1997; Frank et al., 2006; Ikonen,



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1997). Caveolin-1 has been reported to function as an inhibitor of signaling molecules such as endothelial nitric oxide synthase (eNOS) and is known to modulate processes such as chronic inflammation (Razani et al., 2001; Vargas et al., 2002; Yeh et al., 2004). Several studies have demonstrated that caveolin-1 regulates inflammation through the p38 MAPK and NF- $\kappa$ B pathways (Garrean et al., 2006; Lv et al.; Pontrelli et al., 2006; Wang et al., 2006). Another group reported that caveolin-1 null mice (cav-1 -/-) displayed a significant decrease in NF- $\kappa$ B activity after LPS stimulation, indicating that caveolin-1 plays a critical role in some antimicrobial defense mechanisms (Garrean et al., 2006).

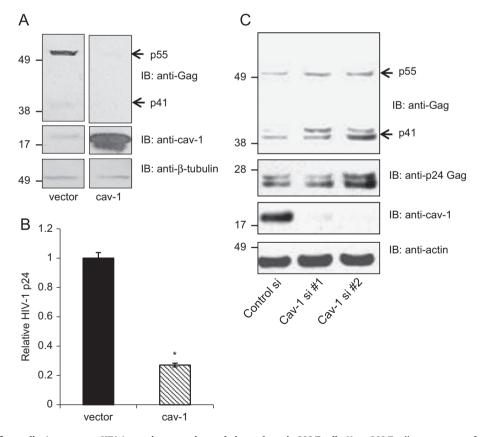
Ectopic expression of caveolin-1 has been shown to suppress the replication of HIV-1 (Llano et al., 2002). However, the molecular mechanism of this inhibition of the HIV-1 life cycle by caveolin-1 remains largely undefined (Carter et al., 2010). Inhibitory effects of caveolin-1 appeared to be HIV-1 specific, as caveolin had little or no effect on other classes of viruses (Llano et al., 2002). Notably, caveolin-1 did not suppress virus species that replicated in the cytoplasm. In another study, HIV-1 infection of macrophages appeared to correlate with an induction of caveolin-1 expression in a HIV-Tat dependent manner (Lin et al., 2010). HIV-1 infection has been shown to lead to induction of a number of signal transduction pathways, some of which involve caveolin-1 (Bennasser et al., 2001; Cheng et al., 2009; Yang et al.).

In the present study, we determined that the inhibitory effect of caveolin-1 on HIV-1 replication is mediated through modulation of the NF- $\kappa$ B-dependent inflammatory pathway. Specifically, we demonstrate that caveolin-1 inhibition of HIV-1-LTR promoter activity is mediated via suppression of NF- $\kappa$ B acetylation.

#### Results

#### Caveolin-1 expression suppresses HIV-1 production

Studies have shown that expression of caveolin-1 creates a restrictive cellular environment for HIV-1 replication (Lin et al., 2010; Llano et al., 2002). More recently, it has been shown that HIV-1 infection induces caveolin-1 expression in macrophages (Lin et al., 2010). These findings have intriguing implications and they may in part explain the persistence of HIV-1 infection in monocytic cells in the absence of the cytopathology observed in lymphocytes. We set out to determine the step of the HIV-1 lifecycle impacted by caveolin-1. We first verified that caveolin-1 expression decreased virus production as previously reported. To that end, 293 T cells were transfected with caveolin-1 or empty vector along with HIV-1 molecular clone, pYU-2. Forty-eight hours post-transfection, virus production was measured using Western blot analysis and HIV-1 p24 ELISA. We observed a significant reduction in cell-associated virus proteins (Fig. 1A) and virus present in cell culture supernatant from cells overexpressing caveolin-1 (Fig. 1B). These observations confirmed that caveolin expression was associated with lower HIV replication. To further explore this relationship, 293 T cells were sequentially transfected, first with control or caveolin-1 specific siRNAs and then with HIV-1 molecular clones. Cells transfected with caveolin-1 siRNA exhibited a reduction in caveolin-1 expression when compared to control cells (Fig. 1C). This reduction of caveleolin-1 expression was sufficient to enhance the level of cell-associated viral proteins when compared to control cells (Fig. 1C). This effect



**Fig. 1. Overexpression of caveolin-1 suppresses HIV-1 protein expression and virus release in 293 T cells**. Here, 293 T cells were co-transfected with HIV-1 molecular clone pYU-2 and caveolin-1 expression construct or vector control. Cells and supernatants were harvested 48 h post-transfection. (A) HIV-1 Gag protein expression was analyzed by Western blot analysis with an anti-Gag mAb (upper panels). We confirmed expression of caveolin-1 by probing with a polyclonal antibody (middle panels) and used  $\beta$ -tubulin as a loading control (bottom panel). (B) Virus particle production was assessed by performing HIV-1 p24 ELISA using supernatants from transfected cells. Graph represents 3 independent experiments done in replicates with S.E.M., *p* value < 0.05. (C) 293 T cells were first transfected with one of 2 independent caveolin-1 specific or control siRNAs, then 48 h later transfected with pYU-2. Protein expression was analyzed h.

was demonstrated with two independent siRNAs. These data indicate that caveolin-1 expression restricts HIV-1 viral replication in 293 T cells as previously reported.

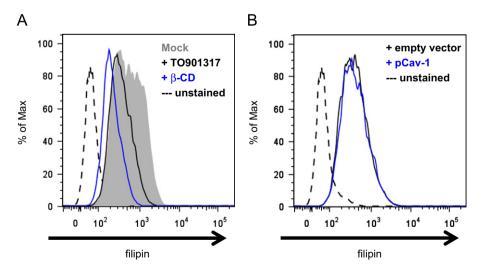
## Caveolin-mediated HIV-1 suppression is independent of modulation of cellular cholesterol

Caveolin-1 proteins bind and transport cholesterol within cells (Fielding and Fielding, 1997; Ikonen and Parton, 2000). Caveolae regulate factors that are involved cholesterol homeostasis, such as SR-BI and ABCA1 (Bist et al., 1997: Frank et al., 2006: Lin et al., 2010: Lin et al., 2009: Wong et al., 2006: Yeh et al., 2004), and some of these affect HIV-1 replication (Morrow et al., 2010: Mujawar et al., 2006; Mujawar et al., 2010). Therefore, we asked whether caveolin-mediated inhibition of HIV-1 involved disruptions in cholesterol homeostasis, as measured by changes in total cellular free cholesterol. To test this we first transfected 293 T cells with caveolin-1 or control vectors. Untransfected cells treated with cholesterol efflux inducer, LXR agonist TO-901317, or cholesterol sequestering agent  $\beta$ -cyclodextrin, served as positive controls for the assay. Cells were harvested two days post transfection. The cells were then fixed and stained with filipin to label total free cholesterol and subsequently analyzed by flow cytometry. As expected, treatment of cells with TO-901317 or  $\beta$ -cyclodextrin reduced the free cholesterol content of cells (Fig. 2A). However, cells overexpressing caveolin-1 had no detectable difference in free cholesterol content when compared to controls (Fig. 2B). These observations were recapitulated in both membrane fraction analyses of transfected cells, as well as fluorescent microscopy (data not shown). These results indicate that caveolin-1 does not cause significant changes in cellular cholesterol localization or content in our model system.

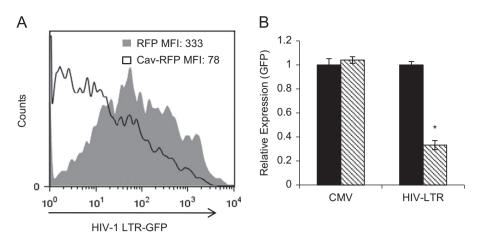
#### Transcription of HIV-1 is suppressed in the presence of caveolin-1

While it was known that manipulations of cholesterol content in HIV-1 susceptible cells have a major effect on virus replication (Liao et al., 2001; Liao et al., 2003), the effect of caveolin-1 on HIV-1 is still largely uncharacterized. Because HIV-1 replication involves a number of essential steps, we thought it necessary to determine at which

replication steps caveolin-1 inhibited HIV-1. Since caveolin-1 is a membrane protein with roles in signaling events, we wanted to determine whether the protein affected early steps in virus replication like transcription, or late steps such as assembly and budding. It has been shown that HIV-1 Gag expression alone is sufficient to generate virus-like particles (VLP) that can be released from cells (Haynes et al., 1991; Rasmussen et al., 1990; Wang et al., 1990). We utilized a cytomegalovirus (CMV) promoter-driven Gag-GFP plasmid to examine VLP release from cells overexpressing caveolin-1. Cells and cell supernatants were collected 48-hours post-transfection in order to measure the HIV-1 VLP production by p24 ELISA. The total amount of HIV-1 Gag in the system was measured. (i.e. intracellular Gag and Gag present in cell supernatants), and those values were used to calculate the virus particle release efficiency (Figure S1). Virus particle release efficiency is the percentage of HIV-1 Gag released from the cell relative to the total amount of Gag produced in the system. No differences in release efficiency were observed between experimental conditions. This indicates that overexpression of caveolin-1 did not alter the ability of virus to exit cells. We next determined the effect of caveolin overexpression on HIV-LTR transcription. To test this we co-transfected caveolin or control plasmids with an HIV-LTR GFP-reporter construct. Flow cytometric analysis of transfected cells revealed a substantial reduction in HIV-LTR activity in cells that expressed caveolin (Fig. 3A). Next we decided to test the specificity of this caveolin-dependent transcriptional suppression by using GFP reporters with either LTR or CMV promoters. The CMV promoter is constitutively active in mammalian cells, and therefore an adequate control for global changes in transcriptional activity in a cell (Fitzsimons et al., 2002). On the other hand, the HIV-LTR is highly regulated and often requires cellular activation before maximal activity can be detected (Cheng et al., 2009: Huang et al., 1994: Kinoshita et al., 1998: Malcolm et al., 2007; Marzio et al., 2002; Sadowski and Mitchell, 2005; Verhoef et al., 1999). Each viral promoter construct was co-transfected with caveolin-1 into 293T cells which were harvested 48 h later for flow cytometric analysis. The fluorescent signals were quantified and plotted in graphs based on the relative mean fluorescence intensity (MFI). In cells transfected with CMV-Gag-GFP there were no differences between caveolin-1-transfected and control cells (Fig. 3B). In contrast, caveolin-1 expression resulted in a substantial



**Fig. 2. Caveolin-1 overexpression in 293 T cells does not cause global changes in free cholesterol content.** 293 T cells transfected with caveolin-1 or empty vector DNA were stained with fluorescent cholesterol-binding reagent filipin at 48 h post-transfection to assess total free cholesterol in the cells. (A) As a positive control for the assay, untransfected cells were treated with conditions that stimulated cholesterol efflux (5 uM TO901317 for 48 h)or a cholesterol sequestering agent (1% Beta-Cyclodextrin for 1 h), prior to filipin staining as described in Methods. Cellular free cholesterol content was determined by flow cytometry and graphed in a histogram as indicated. Dotted lines are control cells that were not stained. (B) Caveolin-1 transfected or control cells were stained with filipin to determine the cellular cholesterol levels of transfected cells. Cells. Data for filipin-stained cell populations are plotted as histograms.



**Fig. 3. Caveolin-1 suppresses HIV-1 promoter activity.** (A) 293 T cells were transfected with caveolin-1-RFP expression construct (filled histogram) or RFP control vector (open histogram), along with HIV-1-LTR GFP reporter. HIV-1 LTR GFP signal was measured by flow cytometry 48 h post transfection and data was plotted in a histogram, where MFI of samples are indicated. (B) 293 T cells were co-transfected with caveolin-1 (stripped) or vector control (black) and either Cytomegalovirus (CMV)—or HIV-1 LTR-promoter driven Gag-GFP plasmids. Cells were harvested 48 h post-transfection and GFP-MFI signal from the individual promoters was analyzed by flow cytometry. Graphs represent 3 independent experiments done in replicates with S.E.M., *p* value < 0.05.

decrease in GFP signal in the LTR-GFP cells. Using HIV-1 specific antibodies in FACS analysis, we were able to observe inhibition of HIV-1 protein expression up to 70% in caveolin-1 transfected cells (data not shown).

We observed caveolin-1-mediated suppression of transcription from HIV-1 reporters in our system under basal conditions, where no stimulus like LPS or TNF-alpha was administered. We decided to test the ability of caveolin-1 to suppress Tat-activated HIV transcription. The HIV-1 Tat protein is a potent transactivator of HIV transcription and provides a robust increase in LTR activity (Huang et al., 1994; Marzio et al., 2002; Yang et al.), Caveolin-1 was found to suppress both basal and Tat-enhanced transcription in 293 T cells (Fig. 4A). To confirm that caveolin-1 suppresses HIV-1 promoter activity in a physiologically relevant system, we co-transfected THP-1 cells with caveolin-1 expression plasmids and an HIV-1 reporter. The cells were cultured in the absence or presence of TNF-alpha and then we measured the activity of the reporter 12 h post transfection (Fig. 4B). Further, we assessed the capacity of caveolin-1 to suppress HIV-1 transcription when p65 is overexpressed (Fig. 4C). In these experiments, we were able to determine that caveolin-1 was capable of suppressing HIV-1 transcription under all conditions. Correspondingly, we observed an increase in HIV-1 transcription in THP-1 cells that were first transfected with siRNA against caveolin-1 and then infected with a VSV-G pseudotyped HIV-1 reporter virus (Figure S2). Taken together our data strongly implicate caveolin-1 as a transcriptional inhibitor of the HIV-1 promoter

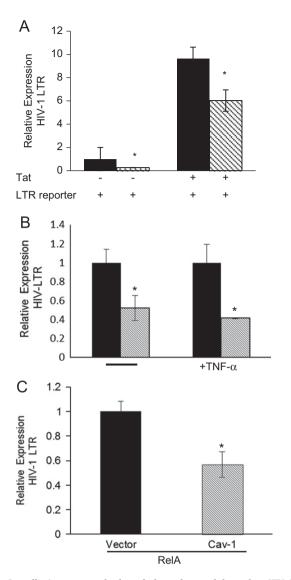
#### Caveolin-1 suppression of HIV-1 replication requires NF-кВ

It is well documented that HIV-1 infection induces a number signaling pathways, including the inflammatory pathway (Herbein and Varin, 2010; Ross et al., 2006). Inflammation is a host defense mechanism against infection and is largely sufficient to create an inhospitable environment for many pathogens. However, this is not the case for HIV-1 infections, which induce inflammation and in fact benefit greatly from the activation of the numerous host transcription factors that are known to mediate the inflammatory response (DeLuca et al., 1999; Montano et al., 1996; Roulston et al., 1993; Verhoef et al., 1999; Westendorp et al., 1995). Caveolin-1, on the other hand, has been shown to suppress inflammatory responses (Chidlow and Sessa; Garrean et al., 2006; Lv et al.; Wang et al., 2006). Since our data indicates that caveolin-1 specifically inhibits HIV-1-LTR transcription, we

hypothesize that NF-KB could mediate caveolin's effect on HIV-1 replication. NF-kB binding sites are present in the HIV-1 LTR, and in cells other than T cells, NF-kB drives the majority of HIV transcription (DeLuca et al., 1999; Montano et al., 1996; Roulston et al., 1993; Westendorp et al., 1995). To determine the contribution NF-κB to inhibition of HIV-1 by caveolin, NF-κB reporters were co-transfected along with caveolin expression constructs into 293 T cells. Fourteen hours post-transfection, the cells were treated with 10 ng/mL of TNF- $\alpha$  for 10 h. After stimulation with TNF- $\alpha$ , cells were harvested and analyzed by microscopy and flow cvtometry. Microscopy showed that the overall level of the NFκB-GFP signal was reduced in caveolin-1 transfected cells (Fig. 5A, top). We noted a difference in the intensity of RFP signal from caveolin-1-RFP transfected cells when compared to RFP vector transfected cells, possibly due to intrinsic differences in protein stability. Flow cytometry data indicated that NF-kB activity was suppressed in cells transfected with caveolin-1 (Fig. 5A, lower). To determine the NF-kB-dependence of caveolin-mediated suppression of HIV-1 transcription, we next transfected cells with either wild-type HIV-LTR (wtLTR) reporters or reporter containing point mutations in the NF-kB binding site (nkLTR). We observed caveolin-1-dependent suppression of transcription only in cells transfected with a wild-type LTR. The mutant HIV-1 reporter (nkLTR) was insensitive to caveolin-1-mediated suppression (Fig. 5B). To further examine the requirement for NF- $\kappa$ B in caveolin-mediated suppression, we next employed specific inhibitors of NF-kB activation in cells transfected with caveolin-1 and HIV-1 molecular clone pNLENG1. Transfected cells were cultured in the absence or presence of 50  $\mu$ g/ml of specific I $\kappa$ B kinase (IKK) inhibitor peptide for 18 h. In cells that received the IkB kinase inhibitor peptide, caveolin-1-mediated suppression was ablated (Fig. 5C). These observations indicate that caveolin suppresses HIV-1 expression not by simply inhibiting NF-κB activity but by a mechanism that requires NF-kB nuclear translocation. Our results also suggest that NF-kB binding to the HIV-1 promoter is a required step for caveolin-1-mediated suppression of HIV-1 transcription.

#### Caveolin-1 decreases the acetylation of NF-*k*B p65

We have demonstrated that caveolin-1-mediated suppression of HIV-1 requires NF- $\kappa$ B translocation, but we have not determined the consequence of that relocation. Our data clearly show that NF- $\kappa$ B-dependent transcription is reduced when caveolin-1



**Fig. 4. Caveolin-1 suppresses both tat-independent and dependent HIV-1 LTR transcription.** Cells were transfected with caveolin-1DNA and transcriptional activity from HIV-1 reporters was measured by flow cytometry. (A) 293 T cells were co-transfected with caveolin-1 (stripped) or empty vector control (black) along with HIV-1 LTR reporter and -/+HIV-1 Tat. HIV-1-LTR-GFP fluorescence was measured 24 h post-transfection by flow cytometry. Displayed is the average of 3 independent experiments. (B) THP-1 monocytic cells were transfected with vector or caveolin along with HIV-1 LTR reporter and treated with 50 ng/ml of TNF-αfor the last 2 h of cell culture before harvest. The GFP signal from the HIV-1-LTR reporter was measured by flow cytometry. Caveolin-transfected samples were normalized to 0.1. (C) THP-1 cells were co-transfected with caveolin-1 expression construct or empty vector control along with ReIA expression construct and HIV-1 LTR GFP reporter. All graphs represent at least 2 independent experiments done in replicates with S.E.M., p value < 0.05

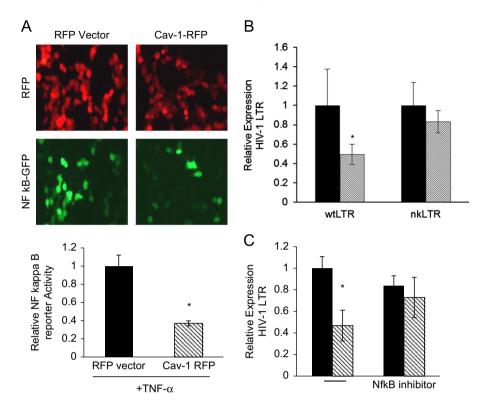
is expressed and that nuclear NF- $\kappa$ B is involved. Interestingly, Chen et al. have described post-translational modifications at K310 that could alter NF- $\kappa$ B transcriptional activity. In their work unacetylated p65 subunit was unable to drive transcription. Nonetheless, up until this point it was unclear how a normally positive modulator of HIV-1 transcription was made to suppress viral replication by caveolin-1. Post-translational modification of p65 is required for optimum transcriptional activity (Chen et al., 2002; Chen et al., 2005). Acetylation is one such modification that has been described as a molecular switch for transcriptional activation, providing an additional layer of NF- $\kappa$ B regulation (Kiernan et al., 2003). Therefore we suspected that caveolin-1 overexpression affected the post-translational modification of the p65, making it less transcriptionally active. To test this possibility in 293 T cells, we first co-expressed caveolin-1 with p65-myc, and then probed for acetylation specifically at K 310. As predicted, we observed a significant decrease in K 310 acetylation (Figure S3). These observations indicated that caveolin-1 suppresses p65 K 310 acetylation.

Intriguingly, Kiernan et al. observed that acetylation at residues 122/123 of p65 has the ability to lower the DNA-binding capacity of NF-kB. Hence, suppression of p65 acetylation could result in increased binding of transcriptionally inert p65 to kB, resulting in reduced expression of NF-kB-responsive genes. In lieu of having available reagents that can be used to probe for acetylation of K 122/123, we assessed global acetylation levels of p65 in the presence of caveolin-1. Here, 293T cells were first co-transfected with p65-myc and caveolin-1 expression constructs or vector control. Thirty-six-hours after transfection, we prepared cell lysates which were subjected to immunoprecipitation of transfected p65 using antibodies directed against the mycepitope in overexpressed p65 or acetylated p65 with pan-specific anti-acetyl-lysine antibodies. Next, we detected the total amount of p65 precipitated in each sample with a pan-specific p65 antibody. We observed a marked reduction in the amount of p65 in the acetyl-lysine immunoprecipitates of samples overexpressing caveolin-1 when compared to vector control (Fig. 6A, lane 8 and 9). However we detected similar amounts of p65 in the samples immunoprecipitated using anti-myc antibody (Fig. 6A, lanes 5 and 6). These observations support the hypothesis that caveolin-1 expression can suppress global acetylation of p65.

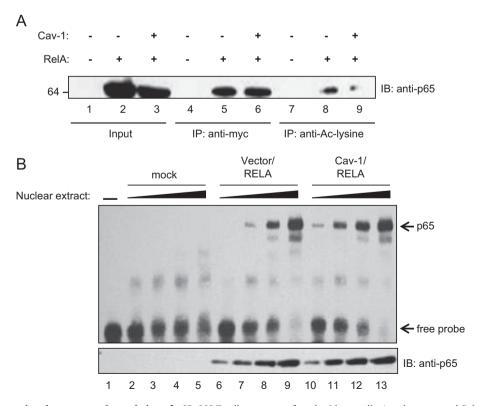
To assess if global hypoacetylation of p65 mediated by caveolin-1 results in differential DNA binding, we performed electrophoretic mobility shift assays (EMSA). For these studies, nuclear extracts were prepared from 293 T cells co-transfected with caveolin-1 and p65-myc expression constructs. We observed a 3.5-fold enhancement in the level of p65 bound to kB-containing probes derived from the HIV-1 LTR when comparing nuclear extracts from caveolin-1 transfected cells to extracts from control cells (Fig. 6B, compare lanes 7 and 11). The shifted species was not detected when using an HIV-1 LTR probe with mutated kB sites (data not shown). This finding was also independently verified by a plate-based transcription factor binding assay (Figure S4). Increased p65 DNA binding was transfection and concentration dependent. These findings indicate that caveolin-1 expression resulted in modifications of NF-kB that not only lowered its transcriptional activity, but also increased the DNAbinding capacity to  $\kappa B$  sites. This occurs via combined hypoacetylation of NF-kB residues K 310 and K 122/123, which regulate transcriptional activity and DNA-binding, respectively (Chen et al., 2002; Kiernan et al., 2003). These observations likely explain the profound effect of caveolin-1 on HIV-1 replication in cells that require NF-kB to drive viral transcription. Our model for caveolin-mediated suppression of HIV-1 showed that caveolin alters the equilibrium between acetylated and unacetylated forms of p65 (Fig. 7). This shift increases the pool of p65 that is transcriptionally inert but capable of DNA binding, thus providing a mechanism for suppressing NF-KB-dependent transcription.

#### Discussion

It has been several years since the first report of caveolin-1dependent inhibition of HIV-1 (Llano et al., 2002). During that time there have been few attempts to identify the mechanism for caveolin-1 suppression of HIV-1. A recent study demonstrated that caveolin-1 interacts with HIV-1 envelope proteins (gp41) and potentially facilitates entry (Hovanessian et al., 2004). A number



**Fig. 5. Caveolin-1-dependentsuppression of HIV-1 transcription is mediated by NF-κB.** (A) 293 T cells were transfected with caveolin-1-RFP or control RFP expression vector with a NF-κB-GFP reporter construct then treated with 50 ng/mL of TNF- $\alpha$  for last 14 h of the experiment. Cells were visualized by fluorescence microscopy (top) 24 h post-transfection and collected and analyzed for NF-κB activity (bottom) by flow cytometry. Relative expression (MFI-GFP) is graphed for RFP-caveolin-1 expression construct (stripped bars) or RFP control vector (black bars) transfected samples. (B) Cells were transfected as in (A) with HIV-1-LTR GFP reporters (wild-type, wtLTR or mutated NF-κB binding site, nkLTR) along with RFP-caveolin-1 expression construct (stripped bars) or RFP control vector (black bars) and 24 h post-transfection reporter activity was analyzed by flow cytometry. (C) 293 T cells were transfected with RFP-caveolin-1 (stripped bars) or RFP control vector (black bars) expression construct with HIV-1-LTR GFP reporters, then treated with lkB kinase inhibitor peptide, SN50 (50 µg/mL) for 18 h. HIV-1-LTR reporter activity was analyzed by flow cytometry. All data shown represents 3 independent experiments. Graph represents three independent experiments done in replicates with S.E.M. *p* value < 0.05.



**Fig. 6. Caveolin-1 overexpression decreases total acetylation of p65.** 293 T cells were transfected with caveolin-1 and myc-tagged RelA expression constructs as indicated. (A) Total cells lysates were immunoprecipitated using either anti-myc or anti-acetyl-lysine antibodies and analyzed by and Western blot analysis. Blots were probed with monoclonal anti-p65 antibodies to determine efficacy of pull-down. (B) Electrophoretic mobility shift assays (EMSA) performed with increasing volumes of nuclear extracts (0.5–4 uL) from caveolin-1 and RelA transfected 293 T cells. DNA binding was determined using biotinylated-HIV-1-LTR oligonucleotides and chemiluminescent detection. Replica nuclear extract samples were analyzed for p65 expression by Western blot (lower panel).

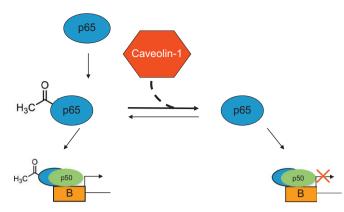


Fig. 7. Model of caveolin-1 mediated suppression of HIV-1 transcription via hypoacetylation of NF- $\kappa$ B p65 subunit. Expression of caveolin results in hypoacetylation of p65 resulting in increased binding to DNA while making the protein transcriptionally inert.

of reports have indicated that caveolin-1 plays some role in modulating viral pathogenesis in vivo (Benferhat et al., 2009a; Benferhat et al., 2009b; Benferhat et al., 2008; Fermin and Garry, 2005; Hovanessian et al., 2004; Huang et al., 2007; Rey-Cuille et al., 2006; Wang et al., 2010). Based on some of our previous studies on the role of cholesterol in HIV-1 biology, we initially suspected that the mechanism of HIV-1 inhibition by caveolin would be linked to its role as a cholesterol transporter. However, data from the current study indicated that caveolin's role in cholesterol homeostasis was not essential to the mechanism by which it inhibits HIV-1 replication. Based on the known role of caveolin in modulation of inflammatory responses, we queried whether altered signal transduction could explain suppression of HIV-1 by the protein. NF- $\kappa$ B is the prototypical initiator of inflammatory processes in cells and is also one of the major transcription factors induced during the course of HIV-1 infection. We hypothesized that caveolin-1 inhibited HIV-1 replication by suppressing NF-kB-induced transcription.

Our results showed that instead of reducing NF- $\kappa$ B association with DNA, caveolin expression increased DNA binding of nuclear NF- $\kappa$ B. Further, mutation of the NF- $\kappa$ B binding site in the HIV-LTR ablated the suppressive effect of caveolin on HIV transcription. These results indicated that caveolin-mediated suppression of HIV-1 required nuclear association of NF-κB and binding to the LTR. While we were completing our studies, Wang et al. published a study confirming that caveolin suppressed HIV-1 transcription through an NF-κB-dependent mechanism (Wang et al., 2011). In that study, the investigators abrogated caveolin-1 mediated suppression of HIV-1 by blocking nuclear translocation of NF-κB with the inhibitor SN50 confirming that NF-κB was necessary for caveolin-1 mediated suppression of HIV-1 replication. We extended our observations and those of the study by Wang et al. by demonstrating that binding of hypoacetylated NF-kB to the HIV-1 promoter is the specific mechanism by which caveolin-1 inhibits HIV-1.

It is possible that cells, such as macrophages, have developed mechanisms to protect themselves from the cytopathic effects of HIV-1 infection. Modulation of caveolin-1 levels during the course of infection may be a host cell response designed to combat HIV-1 replication at the level of viral transcription (Lin et al., 2010). The detailed steps that are involved in the mechanism are still somewhat unclear, but with the evidence presented in this study it is clear that caveolin-1-mediated suppression of HIV-1 requires DNA binding of NF- $\kappa$ B p65. This finding is similar to a phenomenon previously described when unacetylated p65 is present in the nucleus. In the absence of acetylation, NF- $\kappa$ B p65 suppresses transcriptional activity, while showing enhanced DNA binding

capacity. Currently it is unclear how caveolin could affect acetylation of p65. One potential mechanism would be caveolindependent reduction of expression or activity of molecules such as SIRT-1 or p300, which are responsible for acetylation status of NF- $\kappa$ B. Investigating the effect of caveolin-1 on these molecules will be the subject of further studies. Understanding how caveolin modulates factors that regulate NF- $\kappa$ B p65 function may provide important new insights into differences in HIV-1 replication in T cells and macrophages.

#### Materials and methods

#### Cells and constructs

293 T human embryonic kidney cells were cultured in DMEM (GIBCO/Invitrogen, Carlsbad, CA) supplemented with 10% fetal bovine serum, 10 mM L-glutamine and 5 mM Hepes and 100U/ml penicillin and streptomycin (complete DMEM). THP-1 monocytes were cultured in RPMI (GIBCO/Invitrogen, Carlsbad, CA, Carlsbad, CA) supplemented with 10% fetal bovine serum, 10 mM L-glutamine, and 5 mM HEPES (complete RPMI). pYU-2 (obtained from NIH AIDS Research and Reference Reagent Program) is a CCR5tropic HIV-1 molecular clone derived from a human brain viral isolate. Caveolin-1 cDNA was excised from pOTB7 by EcoRI/XhoI digestion and subcloned into pcDNA3.1+ vector, (pCav1). Caveolin-1-myc-RFP and RFP-myc vectors were kind gifts from Dr. Robert Nabi (University British Columbia) were made by subcloning caveolin-1 gene into pRFP-N1 vector (Clontech, Mountain View, CA). NF-KB reporter, Cignal NFKB Reporter (GFP) Kit (CCS-013, SABioscience). Caveolin-1 silencing experiments were performed using Silencer select pre-designed caveolin-1 siRNA, 5'-GCC-GUCUAUUCCAUCUAtt3' (Ambion Inc., Foster City, CA). Wild-type and mutant HIV-1-LTR constructs were cloned into pZsgreen expression vector (cat. no. 632446, Clontech, Mountain View, CA) to generate LTR reporters as previously described (Taylor et al., 2011). The following primers were used to generate HIV-1-LTR and HIV-1-NF-κB mutant LTR reporters: wtLTR-Zsgreen- 5'-TGACATC-GAGCTTGCTACAAGGGACTTTCCGCTGGGGGACTTTCC-3' and nkLTR-Zsgreen 5'-TGACATCGAGCTTGCTACAAGCCACTTTCCGCTGGGGACT-TTCC-3'.

#### Virus stock

Virus preparation and viral infection. Virus stocks were prepared by transfecting 293 T cells with HIV-1 plasmid DNA using Lipofectamine 2000 reagent (Invitrogen). VSV-G envelope-pseudotyped virus was produced as previously reported (Zhou et al., 2005). Culture supernatants were collected 48 h after transfection, centrifuged at  $1000 \times g$  to remove cell debris, filtered through a 0.45-µm-pore-size filter, and concentrated by ultracentrifugation at  $100,000 \times g$  through a cushion of 20% sucrose in PBS. The pelleted virus was resuspended in RPMI with 10% FBS, aliquoted, and stored at  $-80^{\circ}$ C. The viral titer was measured by anti-p24<sup>Gag</sup> ELISA.

#### Transfection

Transfections were carried out by utilizing the Lipofectamine 2000 (Invitrogen, Carlsbad, CA) protocol for plasmid DNA transfection and Oligofectamine (Invitrogen, Carlsbad, CA, Carlsbad, CA) for siRNA transfection per manufacturer's instructions. Preliminary transfections utilized pCav1 and pYU-2. siRNA transfections were done in a sequential manner to maximize the effect of caveolin-1 knockdown on HIV-1 replication. Cells were transfected with siRNA as previously described, and incubated for 24 h. Cells were then transfected with HIV-1 provirus and incubated an additional 48 h before they were harvested. To monitor transfection efficiency, specific siRNAs were co-transfected along with a fluorescent-tagged control siRNA. To determine transfection efficiency of expression plasmids, cells were co-transfected with pMAX vector (Amaxa Biosystems, Braunschweig, Germany) which encodes GFP along with the expression plasmids. The fluorescent signal was measured by analysis on a FACSCalibur (Becton Dickson, Fair View Lakes, NJ) flow cytometer. pNLENG1-EGP is a full length infectious molecular clone of HIV-1 containing enhanced green fluorescent protein(EGFP) between the *env* and *nef* genes. This was kindly provided by D.N. Levy (NYU).

#### ELISA

Virus released from infected cells was quantified using a standard Enzyme-Linked Immunosorbent Assay (ELISA) to measure viral p24 antigen in supernatants (sensitivity 50–200 pg/ml) (Liao et al., 2003). Absorbance was measured on a Fluorstar Optima plate reader (BMG Labtech).

#### Western blot

Protein analysis was accomplished using the NuPAGE gel electrophoresis system (Invitrogen, Carlsbad, CA). Cells were lysed on ice for 30 min in RIPA buffer (50 mM Tris-HCl, pH 7.4, 150 mM NaCl, 1% NP-40, 0.5% sodium deoxycholate, 0.1% SDS, and protease inhibitors (Roche)). Lysates were clarified at 16,000 × g at 4 °C. Protein lysates were resolved on 10% Bis-Tris gels, and then transferred onto nitrocellulose using a semi-dry transfer apparatus (Bio-Rad). Membranes were blocked for 1 h in 5% non-fat milk in PBS, followed by 3 washes in PBST (PBS, 0.5% Tween-20) before probing with primary antibodies. Membranes were then incubated with secondary antibodies conjugated to peroxidase. Chemiluminiscent substrate (ECL, GE Healthcare Life Sciences) was used for detection.

#### Immunoprecipitation

Cells were first pelleted at  $1000 \times g$  for 5 min and resuspended in NP-40 lysis buffer (50 mM Tris-HCl, pH 7.4, 150 mM NaCl, 1% NP-40, 2 mM EDTA, and protease inhibitors (Roche) then incubated on ice for 30 min. Lysates were clarified by centrifugation at  $16,000 \times g$  for 20 min at 4 °C. Immunoprecipitation were performed using 500 µg of total protein and 10 µg/mL of either antimyc mAb (9E10 clone) or anti-acetyl-lysine polyclonal antibodies. Samples were incubated on ice O/N and immune complexes were recovered by adding protein A/G sepharose for 1 h at 4 °C. Beads were washed twice with lysis buffer and resuspended in loading buffer and prepared for SDS-PAGE.

#### Antibodies

Antibodies used in this study were polyclonal rabbit anticaveolin-1 (BD Transduction Laboratories, cat no. 610059), HIV-1 p24 KC57-FITC and KC57–p24-RD1 (Beckman Coulter, cat nos. 6604665 and 6604667). anti-p65 mAb (Santa Cruz), anti-phospho-p65 (S276; Cell Signaling), anti-I $\kappa$ B $\alpha$  (Cell Signaling), antiphospho-I $\kappa$ B $\alpha$  (14D4; Cell Signal), anti-acetyl-lysine antibody (#9441; Cell Signal). Anti-acetyl-lysine (21623; Abcam), antitopoisomerase (clone H-300; Santa Cruz), and anti-RNA polymerase II (Millipore).

#### Flow cytometry

Cells transfected with cDNA for fluorescent-tagged proteins were pelleted and washed with PBS to remove excess culture medium. Cells were then fixed with 50 ul of 2% paraformaldehyde in PBS for 15 min at room temperature. The cells were then washed using 1 ml of PBS and resuspended in 1ml of FACS buffer (1XPBS, 5% FCS, and 0.1% sodium azide). The cell suspension was analyzed by flow cytometry (FACSCalibur, Becton Dickson) for specific signals associated with expression of tagged proteins. Controls consisted of untransfected cells or cells transfected with empty vector. Protein expression was also assessed by staining cells with directly-conjugated monoclonal antibodies or untagged primary antibodies and directly-conjugated secondary antibodies, followed by flow cytometry analysis.

#### Cholesterol analysis

Cholesterol content of fractions from sucrose gradients was measured using the Amplex Red Cholesterol Oxidase kit (Invitrogen, Carlsbad, CA). Fractions collected at the completion of lipid raft isolation were subjected to the cholesterol oxidase assay as per manufacturer's protocol. The protein concentration of cell lysates determined was by BCA assay (Pierce/THERMO). For filipin staining, transfected cells were grown on 35 mm glass-bottom dishes (MatTek Corporation, Ashland, MA) under normal growth conditions and then fixed in 2% paraformaldehyde in PBS for 15 min. The cells were then incubated on ice with filipin at 50  $\mu$ g/ ml in PBS for 30 min. Analysis of filipin staining was performed using a Nikon TE2000 wide-field microscope (Nikon Instruments, Melville, NY). All images were acquired with a plan fluor  $60 \times 1.3$  NA oil immersion objective equipped with an ultraviolet filter. The digital image acquisition was performed with Nikon Elements Advanced Research Software (Nikon Instruments, Melville, NY).

#### Nuclear binding assays

Transfected cells were lysed in nuclear/cytoplasmic extraction buffer from NE/PER kit (Pierce) per manufacturer's instructions. The protein content of nuclear and cytoplasmic lysates was measured by BCA assay (Pierce). After lysis aliquots were placed at -80 °C until use. An equal amount of protein equivalents were placed into wells containing oligonucleotides with kB binding sites provided with NF-kB p65 Transcription Factor Assay Kit (Pierce/THERMO). Lysates were incubated in the wells for 1 h before washing the plates with provided wash buffer. Next primary antibodies (anti-p65, dilution 1:1000), were added to each well and incubated at room temperature for 1 h, followed by an additional wash step. Finally secondary antibody was added (horseradish peroxidase-conjugated, 1:10000), and incubated for 1 h at room temperature. After a final wash, luminal detection solution was added to the well for 20 min. Chemiluminiscent signal was detected on fluorescent plate reader.

#### Electrophoretic mobility shift assay

Nuclear extracts were obtained as described above. Extracts were thawed and added to binding reactions. Nuclear extract volumes were used between 0.5 and 4.0 ul with 20 fmol of either wild type HIV-1 LTR-biotinylated probe (Biotin-5'-TACAAGG-GACTTTCCGCTGGGGACTTTCCAGGGA-3') or mutant probe (Biotin-5'-TACAAGCCACTTTCCGCTGGCCACTTTCCAGGGA-3') containing point mutations in the NF-κB binding sites in final volume of 20 ul. Reaction mix incubated on ice for 30 min. Pre-run 5% native polyacrylamide in 0.5X TBE at 120 V for 30 min in a

Mini-PROTEAN tetra cell (Bio-Rad). Add 5 ul of 5x loading buffer to each reaction and load onto gel. Run gel at 100 V until dye front runs 2/3 down the gel at 4 °C. Transfer gel on to a nylon membrane in 0.5X TBE for 30 min at 400 mA for 30 min (at constant 25 V). Crosslink membrane with a UV crosslinker. Probes detected as described by LightShift Chemiluminescent EMSA kit protocol (Pierce/THERMO).

#### Statistical analysis

All experiments were reproduced two to four times, and representative experiments are shown. The average of repeat experiments was plotted and the standard deviation of mean was calculated from experimental values. The student *t* test was used to determine statistical significance of the differences.

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#### Appendix A. Supplementary information

Supplementary data associated with this article can be found in the online version at http://dx.doi.org/10.1016/j.virol.2012.05.016.

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