# **TNF**α**-mediated proliferation of vascular smooth muscle cells involves Raf-1-mediated inactivation of Rb and transcription of E2F1-regulated genes**

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Atherosclerosis is characterized by hyperplastic neointima and an inflammatory response with cytokines such as TNF $\alpha$ . TNFα is a pleiotropic cytokine that mediates inflammatory, proliferative, cytostatic and cytotoxic effects in a variety of cell types, including endothelial cells and vascular smooth muscle cells (VSMCs). Interestingly, TNF $\alpha$  has been shown to play two very opposing roles in these cell types; it inhibits proliferation and induces apoptosis in endothelial cells, while it enhances the proliferation and migration of VSMCs. Here we show that TNF $\alpha$  is capable of stimulating proliferation of rat VSMCs as well as human VSMCs in a Raf-1/MAPK-dependent manner. TNFα could increase the expression of E2F-regulated proliferative *cdc6*, *Thymidylate synthase (TS)* and *cdc25A* genes in Aortic smooth muscle cells (AoSMC), as seen by real time PCR assays. There is an activation of the stress-induced kinase, JNK1, in VSMCs upon TNF $\alpha$  stimulation. TNF $\alpha$  was capable of inducing binding of the Raf-1 kinase to Rb, and treatment with the Rb-Raf-1 inhibitor, RRD-251, could prevent TNFαinduced S-phase entry in AoSMCs. In addition, inhibition of Raf-1 or Src kinases using pharmacologic inhibitors could also prevent S-phase entry, while inhibition of JNK was not as effective. These results suggest that inhibiting the Rb-Raf-1 interaction is a potential avenue to prevent VSMC proliferation associated with atherosclerosis.

### **Introduction**

Atherosclerosis is the hardening of the arteries that leads to stroke or heart attack and is characterized by many complex changes to the arteries.<sup>1</sup> Development of atherosclerosis is a stringently regulated and complex process that results from aberrations in endothelial cell and vascular smooth muscle cell (VSMCs) function.<sup>2,3</sup> Endothelial cells form the lining of the blood vessels and the heart, functioning as a barrier by regulating permeability, thrombogenicity and production of growth inhibitory molecules.4 Endothelial cells also respond to mechanical forces. Endothelial cells are contact-inhibited under normal conditions, but when endothelial cells sense an injury such as abrasion of a vessel, they proliferate and migrate, leading to re-endothelialization at sites of injury.4 At the same time, vascular smooth muscle cells proliferate and migrate from the injured arterial wall into the vessel lumen, leading to vessel thickening and occlusion, called restenosis.5,6 Intimal hyperplasia characterized by VSMC proliferation and extracellular matrix (ECM) deposition is a major process contributing to restenosis.7 Atherosclerotic lesions can be blocked if inhibition of VSMCs is effective.<sup>8</sup> Several growth factors and cytokines are capable of stimulating VSMC migration and proliferation, such as platelet-derived growth factor (PDGF), which plays a vital role in the development of restenosis.9-11

PDGF can stimulate VSMC proliferation and migration at sites of stress.10,12 It has been shown that suppression of PDGFR activation can inhibit VSMC proliferation by decreasing activation of its downstream signaling molecules.<sup>13-15</sup> PDGF is a potent mitogen that mediates arterial response and stimulates proliferation and matrix production.<sup>16</sup> PDGF signaling leads to downstream activation of proliferative genes that contribute to atherosclerosis and restenosis.17

Tumor necrosis factor  $\alpha$  (TNF $\alpha$ ) is a pleiotropic inflammatory cytokine known to play a pathophysiological role in vascular atherosclerosis.18-20 It is known to be expressed by vascular smooth muscle cells in atheromatous plaques.<sup>21</sup> By signaling through cell membrane receptors, TNFα can induce mitogenic and proapoptotic effects, activating monocytes, endothelial cells and macrophages.<sup>22</sup> Accelerated atherosclerosis is characterized by hyperplastic neointima and an inflammatory response with cytokines such as  $TNF\alpha$ .<sup>23-25</sup>  $TNF\alpha$  has been shown to play two opposing roles in cardiac pathology, including inhibition of endothelial cell proliferation and enhancement of apoptosis, while stimulating vascular smooth muscle cell proliferation and migration.8,26,27 Although there are conflicting reports on the ability of TNFα to stimulate VSMC proliferation, there is compelling evidence defining the migration stimulating activity of this cytokine.28,29 TNFα, like other chemoattractants, such as PDGF, stimulates VSMC migration through the MAPK pathway.<sup>30</sup>

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The apoptosis induced by TNF superfamily requires binding of a ligand to its receptor, leading to oligotrimerization of receptors.31-34 This results in aggregation of death domain-containing proteins allowing recruitment of TRADD (TNF receptor 1-associated death domain protein). TRADD binds FADD (Fas-associated death domain-containing protein) and TRAF-2 (TNF receptor 1-associated protein 2) proteins, which, in turn, leads to activation of procaspase-8 and apoptosis signal-regulating kinase 1 (ASK1), respectively.<sup>32,35-37</sup> TNF $\alpha$  treatment leads to simultaneous activation of the ASK1-JNK/p38 death signal.<sup>36-39</sup> Reports on the effects of TNFα on apoptosis or proliferation in VSMCs are conflicting.<sup>26</sup> Several investigations report that TNFα itself does not induce VSMC proliferation, while other studies suggest TNFα induces proliferation of VSMCs through NFκB-mediated transcription mechanisms.<sup>26,40</sup> Regarding apoptosis, there are also inconsistent reports. Certain studies have shown that  $TNF\alpha$  could induce apoptosis in VSMCs via caspase-3 activation, while others found no pro-apoptotic activity for TNF $\alpha$  in these cells.<sup>28</sup> Further investigations revealed that effects of  $TNF\alpha$  in VSMCs can vary depending on the phenotype of the cells.<sup>26</sup>

Although PDGF and TNFα have very different signaling intermediates, their downstream functions require MAPK activation. It is therefore important to identify the upstream mechanisms contributing to increased proliferation by these two stimuli. Stimulation of VSMCs with PDGF leads to downstream activation of Erk1/2 via the Ras/Raf/MAPK (mitogen-activated protein kinase/extracellular-signal-regulated-kinase) pathway. Activated ERK1/2 rapidly translocates to the nucleus where they target transcription of genes that regulate cell cycle progression, such as *cyclin D1*. 41 Cyclin D1 binds cdk4/6, and together they facilitate S-phase entry through phosphorylation of the retinoblastoma (Rb) protein.<sup>42</sup> Stimulation of VSMCs with  $TNF\alpha$ has been shown to enhance proliferation through ERK1/2,<sup>43</sup> although the exact mechanism is not known.

The retinoblastoma tumor suppressor protein, Rb is a vital regulator of the mammalian cell cylcle, and its inactivation facilitates S-phase entry.<sup>44,45</sup> During cell cycle progression, Rb is inactivated through multiple waves of phosphorylation mediated by kinases associated with D- and E-type cyclins in the  $G<sub>1</sub>$  phase.  $Rb$  controls the  $G_1$ -S boundary by repressing the transcriptional activity of the E2F family of transcription factors.<sup>46</sup> E2Fs transcriptionally regulate the expression of many genes necessary for DNA synthesis and cell cycle progression, such as *cyclins A* and *E*, *cdc2*, *thymidylate synthase (TS)*, *dihydrofolate reductase*, *cdc6* and *cdc25A*. 47-49 However, Rb-E2F interaction also has a pivotal role in regulating apoptosis and the intricacies of Rb-E2F interaction, and its regulation of cell fate is context dependent.<sup>50,51</sup> Cyclins and cyclin-dependent kinases phosphorylate Rb in mid- to late  $G<sub>1</sub>$  phase releasing transcriptionally active E2F1, whereas Raf-1 kinase binds to Rb and phosphorylates Rb in the early  $\mathsf{G}_\text{\tiny{1}}$  phase. This priming phosphorylation by Raf-1 is necessary for subsequent phosphorylation and inactivation of Rb.<sup>52</sup> Our previous studies demonstrated that disruption of Rb/Raf-1 interaction significantly inhibits proliferation, angiogenesis as well as tumor growth in vivo in an Rb-dependent manner.53 The Rb-Raf-1



**Figure 1.** TNFα stimulates proliferation in vascular smooth muscle cells. (A) Rat A10 VSMCs were serum starved and subsequently stimulated with serum, TNFα or PDGF for 18 h, and BrdU incorporation was measured. (B) A similar assay was done using human AoSMCs.

pathway has been shown to play a proliferative role in response to growth factor signaling as well as nicotine stimulation of a variety of cell lines,53-55 but the contribution of this pathway in VSMC proliferation is not known. Here, we show that  $TNF\alpha$  stimulates proliferation of VSMCs by activating Raf-1/MEK/ERK pathway and facilitating Rb-Raf-1 interaction, leading to the expression of E2F-regulated proliferative genes. These studies were conducted on primary human aortic smooth muscle cells (AoSMCs) as well as rat A10 cells, which are immortalized vascular smooth muscle cells. Further, disruption of the Rb-Raf-1 interaction by the small molecule inhibitor RRD-251 prevented TNFα-induced proliferation of AoSMCs, raising the possibility that this could be a viable mechanism to combat certain steps of atherosclerosis.

#### **Results**

**TNF**α **stimulates proliferation of vascular smooth muscle cells.** Since there are conflicting reports on the effects of



**Figure 2.** TNFα activates the Raf/MAPK pathway in VSMCs. (A) Time-course stimulation of AoSMCs results in Raf-1 activation, highest at 1 h, and ERK1/2 activation peaks at 30 min. (B) Activation of ERK1/2 coincides with JNK1 activation from 30 min of TNFα treatment. (C) PDGF and TNFα time-course stimulation shows ERK1/2 and JNK activation occurs simultaneously from the different stimuli.

TNFα-induced proliferation of VSMCs, we examined the effects of serum, TNFα and PDGF on VSMCs by BrdU incorporation assays. Rat A10 cells, which are vascular smooth muscle cells derived from thoracic aorta of embryonic rat, were serum starved for 24 h and subsequently re-stimulated with serum,  $TNF\alpha$ (100 ng/ml) or PDGF (100 ng/ml) for 18 h and S-phase entry was measured using standard BrdU incorporation assays. TNFα could stimulate proliferation in vascular smooth muscle cells to a certain extent (**Fig. 1A**). In the same manner, we examined the effects of  $TNF\alpha$  in primary human aortic vascular smooth muscle cells (AoSMCs). As shown in **Figure 1B**, TNFα could stimulate S-phase entry comparable to PDGF in human VSMCs; serum was used as the positive control.

**TNF**α **activates Raf/MAPK pathway in vascular smooth muscle cells.** TNFα has been shown to induce migration of aortic smooth muscle cells through ERK1/2 activation.<sup>30</sup> We wanted to examine if TNFα treatment had any effects on Raf-1 kinase in these cells. Treatment of aortic smooth muscle cells with TNFα for 10 min, 30 min, 1 h and 2 h led to a shift in Raf-1 migration, indicative of Raf-1 phosphorylation, as seen by western blotting (**Fig. 2A**). Indeed, phosphorylated Raf-1 can be seen by western blotting for serine 338 on Raf-1; activation was highest at 1 h of TNFα treatment (**Fig. 2A**). ERK1/2 activation was seen in response to TNFα and was highest at 30 min of treatment (**Fig. 2A**). The stress-activated protein kinase/Jun-N-terminal kinase (SAPK/JNK) is a member of the MAPK family that is potently and preferentially activated by stresses such as UV irradiation, ceramides and cytokines like  $\text{TNF}\alpha$ .<sup>34</sup> In certain instances, JNK can be activated by growth factors.<sup>56-58</sup> We next examined if TNFα was capable of activating stress kinases in a proliferative scenario. To this end, AoSMCs were serum starved and stimulated with TNFα or PDGF for 30 min (time point when Raf-1 activation and ERK1/2 activation was present). Western blotting

for JNK activation using an antibody that recognizes phosphorylated Thr183/Tyr185 residues revealed TNFα- and PDGFactivated JNK in aortic smooth muscle cells (**Fig. 2B**). It was next examined whether TNFα and PDGF activate downstream kinases following similar kinetics. AoSMCs were rendered quiescent by serum starvation and stimulated with TNFα or PDGF for 10 min, 30 min or 2 h. Western blotting using antibodies to pERK1/2 as well as pJNK showed that these kinases were activated at 10 min of stimulation with either agent. The levels of total JNK or ERK1/2 remained mostly unchanged (**Fig. 2C**). It thus appears that  $TNF\alpha$  and PDGF induce ERK1/2 as well as JNK activation following similar kinetics.

**TNF**α**-induced AoSMC proliferation is abrogated by targeting upstream activators of Raf-1.** Since TNFα led to activation of Raf-1 in vascular smooth muscle cells, and since Raf-1 plays a vital role in cell proliferation, we examined if vascular smooth muscle cell proliferation could be inhibited via targeting Raf-1 or kinases that activate Raf-1. Both Src and PKC kinases are known to activate Raf-1 in response to growth factor signaling.<sup>59</sup> To evaluate the importance of these kinases in TNFα-induced proliferation, the Src inhibitor PP2 and the PKC inhibitor Ro-31–8220 were used in BrdU incorporation assays. AoSMCs were serum starved and subsequently stimulated with PDGF or TNFα in the presence or absence of the aforementioned inhibitors for 18 h. BrdU incorporation assays revealed that both inhibition of Src and PKC could efficiently block TNFα- or PDGF-induced S-phase entry (**Fig. 3A**). Targeting upstream of Raf-1 or Raf-1 itself using the multikinase inhibitor BAY-43–9006 could completely inhibit S-phase entry induced by PDGF or TNFα (**Fig. 3B**). Next we examined if inhibition of downstream activation of JNK could prevent S-phase entry induced by PDGF or TNFα. Inhibition of downstream JNK activation did not significantly block proliferation, suggesting that inhibition of Raf-1 function



**Figure 3.** Targeting Raf-1 activation blocks AoSMC proliferation. (A) Src inhibitor (PP2) and PKC inhibitor (Ro-31–8220) block TNFα- and PDGF-induced proliferation. (B) Multi-kinase inhibitor BAY-43–9006, which targets Raf-1, can completely inhibit PDGF- and TNFα-induced proliferation, while the JNK inhibitor (SP600125) does not significantly affect TNFα-induced proliferation. (C) MEK inhibitor PD98059 could block TNFα- and PDGF-induced S-phase entry as seen by BrdU incorporation assays. (D) Raf-1 depletion using Raf-1 siRNAs could inhibit TNFα- and PDGF-induced AoSMC proliferation.

is a viable way to block TNFα-induced proliferation of AoSMCs (**Fig. 3B**). Further, we examined whether MEK inhibitor PD98059 could block PDGF- and TNFα-induced proliferation of AoSMCs. BrdU incorporation assays revealed that inhibition of MEK could block TNFα- and PDGF-induced cell proliferation (**Fig. 3C**).

To confirm the results obtained by targeting Raf-1 by chemical inhibitors, we employed siRNA transfections. AoSMCs were transfected with siRNAs targeting Raf-1, subsequently serum starved and induced with TNFα or PDGF for 18 h. BrdU incorporation assays showed that Raf-1 depletion could abrogate TNFα- and PDGF-induced AoSMC proliferation compared with a non-targeting control siRNA (**Fig. 3D and Sup. Fig. 1**).

**TNF**α **treatment induces E2F regulated genes involved in proliferation.** Since TNFα was functioning similar to a growth factor in stimulating the proliferation of AoSMCs, we examined whether TNFα-induced proliferation was an E2F-dependent

mechanism. To this end, real-time PCR was performed on three E2F-responsive genes from AoSMCs that were serum starved and subsequently stimulated with either  $TNF\alpha$  or PDGF for 18 h. TNFα could induce cdc25A, cdc6 and TS (thymidylate synthase) gene expression 3.5-, 3- and 1.8-fold respectively (**Fig. 4A–C**). These results were confirmed by western blotting, which showed a reduction in cdc 6, cdc 25A and TS protein levels when Raf-1 as well as E2F1 were depleted by siRNA transfection, confirming their role in the normal expression of cdc 6, cdc 25A and TS (**Fig. 4E and F**). Next, we examined if E2F1 was, in fact, present on these proliferative promoters in response to TNFα. Chromatin immunoprecipitation assays were conducted for this purpose. Treatment of quiescent AoSMCs with TNFα or PDGF for 18 h led to an increase in the association of E2F1 on the cdc25A, cdc6 and TS promoters with a concomitant dissociation of Rb (**Fig. 4D**). In quiescent cells, we consistently observed a faint band for E2F1, whereas a significantly higher amount of



**Figure 4.** TNFα and PDGF induce E2F regulated genes in AoSMCs. (A) Treatment with TNFα and PDGF for 18 h led to 3.5- and 4-fold increase, respectively, in *cdc25A* gene expression in real-time PCR assays. (B) Treatment with TNFα and PDGF for 18 h led to 3.5- and 7-fold increase, respectively in *cdc6* gene expression in real-time PCR assays. (C) Treatment with TNFα and PDGF for 18 h led to 1.8 and 3.8 fold increase, respectively in *TS* gene expression in real time PCR assays. (D) Treatment with TNFα or PDGF led to an increase in E2F1 and dissociation of Rb on the proliferative promoters *cdc25A*, *cdc6* and *TS* in ChIP assays; c-fos was used as the negative control. (E) Western blots for cdc 6, cdc 25A and TS after Raf-1 and E2F1 depletion. (F) Densitometric analysis of the bands to quantify the western blot data.

Rb was found to be associated with these promoters in serumstarved cells. The c-fos promoter was used as a negative control. These results suggest that TNFα can induce changes in the promoter association of E2F1 as well as Rb in AoSMCs comparable to a growth factor.

**TNF**α**-induced AoSMC proliferation involves Rb-Raf-1 interaction.** Our previous studies have shown the importance of the Rb-Raf-1 interaction in mediating proliferation in a wide array of cell lines.53,60 Since Raf-1 activation is evident in response to TNFα-induced proliferation in AoSMCs, we examined if Raf-1-Rb interaction is involved in mediating these effects. Treatment with the Rb-Raf-1 inhibitor RRD-251 in the presence of TNFα or PDGF for 2 h could efficiently reduce Raf-1 levels in both AoSMCs and rat A10 cells (**Fig. 5A and B**). Next, we examined if TNFα stimulation of AoSMCs could induce the Rb-Raf-1

interaction; this was done by IP-WB analysis. Treatment with TNFα and PDGF for 2 h led to an increase in Raf-1 bound to Rb; in addition, there was less E2F1 associated in the TNFα- and PDGF-stimulated complexes (**Fig. 5C**). The Rb-Raf-1 inhibitor RRD-251 could effectively inhibit TNFα-induced Rb-Raf-1 interaction as revealed by immunoprecipitation followed by western blot assays (**Fig. 5D**). These results suggest that treatment of AoSMCs with TNFα leads to an increased association of the Raf-1 kinase with Rb, comparable to growth factor stimulation of cancer cell lines, and that RRD-251 can effectively inhibit this interaction in smooth muscle cells.

We next examined if RRD-251 could prevent serum-, TNFαor PDGF-induced proliferation in AoSMCs. AoSMCs were serum starved and subsequently stimulated with serum, TNFα or PDGF in the presence or absence of 20 μM RRD-251. In



**Figure 5.** RRD-251 inhibits Rb-Raf-1 interaction in AoSMCs. (A and B) Treatment with TNFα or PDGF in the presence of RRD-251 inhibits Raf-1 levels in AoSMCs (A) and A10s (B). (C) TNF $\alpha$  and PDGF treatment induced Rb-Raf-1 binding in AoSMCs. (D) RRD-251 inhibits TNFα-induced Rb-Raf-1 interaction.

response to all three stimuli, RRD-251 was capable of inhibiting S-phase entry in AoSMCs (**Fig. 6A**). It is known that binding of Raf-1 to Rb facilitates the complete inactivation of Rb by phosphorylation, resulting in its dissociation from E2F and proliferative promoters, leading to their expression; disrupting this interaction in epithelial cells led to the retention of Rb on E2F1 regulated proliferative promoters, leading to their repression and cell cycle arrest. Given this background, we examined the role of proliferative E2F family members E2F1, E2F2 and E2F3 in TNFα- or PDGF- induced proliferation of AoSMCs. AoSMCs were transfected with E2F1, E2F2, E2F3 or control siRNAs and subsequently induced with  $TNF\alpha$  or PDGF. We observed that E2F1 and E2F3 depletion resulted in a significant reduction in S-phase entry as seen by BrdU incorporation assays, whereas E2F2 had a marginal effect (**Fig. 6D and Fig. S2**). We further examined the effect of RRD-251 in regulating the expression of E2F target genes, such as *cdc6*, *TS* and *cdc 25A* in response to TNFα and PDGF treatment of AoSMCs. Treatment of AoSMC

cells with RRD-251 in the presence of TNF $\alpha$  or PDGF for 18 h could efficiently inhibit the expression of cdc 6, TS and cdc 25A as seen by RT-PCR (**Fig. 6B and C**). These results suggest that inhibiting Rb-Raf-1 interaction leads to the repression of E2F1-regulated proliferative promoters, as in the case of growth factor stimulation of epithelial cell lines, and this is a possible mechanism by which RRD-251 is bringing about the growth arrest of AoSMCs. These observations also raise the possibility that targeting this interaction might be a viable alternative to prevent atherosclerosis.

## **Discussion**

The dynamics of endothelial and vascular smooth muscle cells play a predominant role in the progression of atherosclerosis and restenosis.5,10 Migration, proliferation and differentiation of endothelial cells as well as VSMCs are important pathological responses that contribute to the development of vascular lesions.<sup>61</sup> The "switch" of VSMCs from the quiescent phenotype to the proliferative and migratory phenotype is a vital event in the pathogenesis of atherosclerosis and restenosis post-angioplasty.10,17 Therefore, VSMC proliferation and migration both serve as suitable targets for drug therapy in vascular proliferative disorders.<sup>1,2</sup>

This study provides evidence that TNFα and PDGF evoke similar signaling mechanisms that contribute to VSMC proliferation. Although they are not

equally efficacious in activating these pathways,  $TNF\alpha$  is capable of activating proliferative signaling pathways in the smooth muscle cells, leading to S-phase entry and cell cycle progression. The proliferative response also increased the expression of E2F-regulated genes cdc6, TS and cdc25A; further,  $TNF\alpha$  stimulation led to an increase in the association of E2F1 on these proliferative promoters, with a dissociation of Rb. Interestingly, in response to TNFα stimulation, Raf/MAPK activation occurred, and this coincided with an activation of the stress kinase JNK1. Inhibition of TNFα- or PDGF-induced cell proliferation was observed when pharmacologic inhibitors targeting Raf-1 or upstream regulators of Raf-1 like Src or PKC were used; inhibiting JNK had no effect. This suggests that JNK activation most likely is not responsible for the proliferative responses seen with TNF $\alpha$ , even though it is activated in these cells.

Studies from our lab have shown that upon  $TNF\alpha$  treatment, the kinase ASK1 mediates Rb inactivation as an initial signaling event in Ramos and Jurkat cells.<sup>54</sup> These studies had shown **Figure 6.** Inhibition of Rb-Raf interaction prevents proliferation and expression of E2F1 target proliferative genes in AoSMC. (A) Treatment with RRD-251 inhibits AoSMC proliferation induced by serum, TNF α and PDGF. (B and C) Treatment with RRD-251 inhibits expression of E2F target genes cdc 6, TS and cdc 25A, as seen by RT-PCR. (D) Depletion of E2F1 and E2F3 using siRNAs resulted in significant inhibition of  $\mathsf{TNF}\alpha$  and PDGF induced AoSMC proliferation, whereas E2F2 depletion had only a marginal effect.

that inactivation of Rb by ASK1 kinase was necessary for TNF α to induce apoptosis; further, a mutant ASK1 kinase that was unable to bind to Rb was impaired in its ability to facilitate TNF α-mediated apoptosis in Jurkat and 3T3 cells.

In a similar vein, our recent studies show that TNF α stimulation of human aortic endothelial cells led to sig nificant induction of apoptosis.62 This coincided with Rb phosphorylation, which was mediated by ASK1 kinase. A peptide competitor, which could effectively prevent the binding of ASK1 to Rb could inhibit TNF α-induced apop tosis of endothelial cells. These studies showed that p53 had a limited role in the process. Indeed, the role of p53 in  $TNF\alpha$ -induced apoptosis has been controversial. $63,64$  We also find that TNF α has no effect on p53 expression; however,  $p73\alpha$  levels were found to be upregulated, implicating p73 α as the major contributing factor to apoptosis induced by TNF $\alpha$  in endothelial cells. This induction of p73 $\alpha$ by TNF α was dependent on E2F1 function. Supporting this contention, depletion of E2F1 or ASK1 significantly reduced the ability of TNF α to induce apoptosis in endo thelial cells. Taken together, studies from our lab show that Rb interacts with ASK1 upon apoptotic stimuli, and ASK1 has to overcome Rb function to execute its pro-apoptotic functions, suggesting that Rb acts as a critical connector between apoptotic and proliferative pathways by interact ing with the functionally distinct kinases like Raf-1 and ASK1.54 Thus, the role of Rb phosphorylation by specific kinases is pertinent for directed signaling for apoptotic or proliferative pathways.<sup>65</sup>

The contrasting observation in AoSMCs, where TNF α treatment resulted in a lack of apoptotic response and an increase in proliferation suggests that  $\text{TNF}\alpha$  is involved in multiple pathways depending on the cellular context (**Fig. 7**). We observed activation of Raf-1 and ERK in response to TNF α stimulation, indicative of a proliferative response. It has been shown that a colocalization of TNF α and ERK1/2 occurs, and that ERK 1/2 activation induces the expression of Ets-1, Egr-1 and c-fos in neointimal lesions from rat aortae two weeks post balloon injury.<sup>66</sup> The ChIP and RT-PCR experiments showed recruitment of E2F1 to proliferative promoters, suggesting that E2F1 is a key mediator in the TNF α-induced proliferative or apoptotic pathways in VSMCs or endothelial cells, respectively.

Rb-Raf-1 interaction was found to play a vital role in serum-, PDGF- and TNF α-induced proliferation of VSMCs. Targeting Rb-Raf-1 interaction using RRD-251 could completely inhibit S-phase entry in these cells. Our



lab has previously shown that disrupting the Rb-Raf-1 interaction can prevent endothelial cell adhesion, migration and proliferation.53,54,60 Raf-1 is known to play an essential role in normal wound healing in vivo and migration of keratinocytes and fibroblasts in vitro.<sup>67</sup> This novel function of Raf-1 is independent of its kinase activity and is brought about by the spatial regulation of Rho signaling during migration.68 Apart from its role in cell cycle regulation, emerging evidences also indicate the involvement of Rb-E2F pathway in cell migration and invasion.<sup>69-71</sup> Taken together, the importance of this interaction in both endothelial as well as vascular smooth muscle cell physiology in atherosclerotic lesions needs further evaluation and may provide useful tools in development of therapies for heart disease. The divergent responses of AoSMCs and HAECs (human aortic endothelial cells) to  $TNF\alpha$  thus provide unique therapeutic possibilities: simultaneously targeting the cell cycle of two different cell types within same tissue microenvironment, resulting in opposite and biologically complimentary effects.

#### **Materials and Methods**

**Cell culture.** Human aortic smooth muscle cells were obtained from Lonza (CC-2571) and cultured in smooth muscle basal medium (SmBM) supplemented with growth factors and 5% FBS, according to the manufacturer's instructions (CC-3182). TNFα (Promega) was added at 100 ng/ml. PDGF (Biosource) was added at 100 ng/ml concentration.

**Lysate preparation, immunoprecipitation and western blotting.** Lysates from cells treated with different agents were prepared by NP-40 lysis as described earlier in reference 52. Polyclonal E2F1, c-Raf, ASK1, cyclin D and E and monoclonal antibodies for cdc 25A, cdc 6 and TS were obtained from Santa Cruz Biotechnology. Monoclonal Rb and Raf-1 were supplied by BD Transduction Laboratories. Polyclonal antibodies to phospho-Rb (807, 811), phospho-Raf-(338), phospho-JNK, phospho ERK 1/2 and phospho Mek1/2 were supplied by Cell Signaling.

**Chromatin immunoprecipitation (ChIP) assay.** ChIP lysates were prepared, and immunoprecipitations were conducted using antibodies against E2F1, Rb and the association with specific promoters detected by PCR as previously described in reference 72. Rabbit anti-mouse secondary antibody was used as the control for all reactions. The sequences of cdc25A, TS, cdc6 and c-fos primers are described in reference 52.

**Real-time PCR.** AoSMC cells were subjected to serum starvation for 24 h, followed by treatment with PDGF or TNF $\alpha$  in presence or absence of RRD-251. Unstimulated serum-starved cells were used as a control. Total RNA was isolated by an RNeasy miniprep kit from QIAGEN following the manufacturer's protocol. One microgram of RNA was DNase treated using RQ1 DNase (Promega), followed by first-strand cDNA synthesis using the iScript cDNA synthesis kit (Bio-Rad). A fraction (1/20) of the final cDNA reaction volume was used in each PCR.73 Primer sequences are as follows: 5'-CTG CCA GCT GTA CCA GAG AT-3' (TS forward primer), 5'-ATG TGC ATC TCC CAA AGT GT-3' (TS reverse primer), 5'-CCC CAT GAT TGT GTT GGT AT-3' (Cdc6 forward primer), 5'-TTC AAC AGC TGT GGC



**Figure 7.** Schematic showing the contrasting roles of TNFα in AoSMCs and HAECs. In HAECs, TNF $\alpha$  stimulation resulted in Rb inactivation by ASK1 leading to elevated E2F1 activation and upregulation of proapoptotic protein p73. resulting in apoptosis.62 Whereas in AoSMCs, TNFα stimulation led to Raf-1-dependent Rb inactivation, elevated E2F1 activity, leading to the expression of proliferative promoters, resulting in increased cell proliferation.

TTA CA-3' (Cdc6 reverse primer), 5'-CTC AAC ACG GGA AAC CTC AC-3' (18S forward primer), and 5'-AAA TCG CTC CAC CAA CTA AGA A-3' (18S reverse primer). Real-time PCR was performed using a Bio-Rad iCycler.

**siRNA transfections.** siRNA oligonucleotides targeting human E2F1, E2F2, E2F3 and Raf-1 were purchased from Santa Cruz biotechnology. Non-target siRNA was used as the control. AoSMCs were transfected either with control siRNA, E2F1, E2F2, E2F3 or Raf-1 siRNA (100 pmol) using oligofectamine (Invitrogen) in accordance with the manufacturer's recommendations. For western blots, cells were harvested 48 h after transfection, and lysates were made. For BrdU incorporation asays, cells were serum starved for 24 h after transfection and induced with TNFα or PDGF for 18 h.

**Proliferation assays.** Bromodeoxyuridine (BrdU) labeling kits were obtained from Roche Biochemicals. Cells were plated in poly-d-lysine coated chamber slides at a density of 10,000 cells per well and rendered quiescent by serum starvation for 24 h. Cells were then re-stimulated with serum,  $TNF\alpha$  or PDGF in the presence or absence of the indicated drugs for 18 h. S-phase cells were visualized by microscopy and quantified by counting three fields in quadruplicate.

#### **Disclosure of Potential Conflicts of Interest**

No potential conflicts of interest were disclosed.

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# **Note**

# Supplemental materials can be found at: www.landesbioscience.com/journals/cc/article/18473

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