

Inhibition of the ERK phosphorylation plays a role in terbinafine-induced p21 up-regulation and DNA synthesis inhibition in human vascular endothelial cells

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Abstract

Previously, we showed that terbinafine (TB) induces cell-cycle arrest in cultured human umbilical vein endothelial cells (HUVEC) through an up-regulation of the p21 protein. The aim of this study is to delineate the molecular mechanisms underlying TB-induced increase of p21 protein. RT-PCR analysis demonstrated that the mRNA levels of p21 and p53 were increased in the TB-treated HUVEC. The p21 promoter activity was also increased by TB treatment. Transfection of HUVEC with p53 dominant negative (DN) abolished the TB-induced increases of p21 promoter activity and protein level, suggesting that the TB-induced increase of p21 is p53-dependent. Western blot analysis demonstrated that TB decreased the levels of phosphorylated extracellular signal-regulated kinase (ERK). Over-expression of mitogen-activated protein kinase (MEK)-1, the immediate upstream activator kinase of ERK, abolished the TB-induced increases of p21 and p53 protein and decrease of thymidine incorporation. The ERK inhibitor (PD98059) enhanced the TB-induced inhibition of thymidine incorporation into HUVEC. Taken together, these data suggest that the decrease of ERK activity plays a role in the TB-induced up-regulation of p21 in HUVEC. On the other hand, pretreatment of the cells with geranylgeraniol (GGOH), farnesol (FOH), or Ras inhibitor peptide did not affect the TB-induced decrease of thymidine incorporation. Taken together, our results suggest that TB might cause a decrease of MEK, which in turn up-regulates p53 through the inhibition of ERK phosphorylation, and finally causes an increase of p21 expression and cell-cycle arrest. © 2008 Elsevier Inc. All rights reserved.

Keywords: Terbinafine; p53; p21; ERK; DNA synthesis

Abbreviations: CDK, cyclin-dependent kinase; DEPC, diethyl-pyrocabonate; DMSO, dimethylsulfoxide; DN, dominant negative; DTT, dithiothreitol; ECL, enhanced chemiluminescence; ECs, endothelial cells; ECGS, endothelial cell growth supplement; ERK, extracellular signal-regulated kinase; FBS, fetal bovine serum; FTI, farnesyl transferase inhibitor; FOH, farnesol; GGTI, geranylgeranyl transferase I inhibitor; GGOH, geranylgeraniol; GST, Glutathione-S-Transferase; HUVEC, human umbilical vein endothelial cells; M199, medium 199; MEK, mitogen-activated protein kinase; TB, terbinafine; PMSF, phenylmethyl sulfonyl fluoride; SDS-PAGE, sodium dodecyl sulfate-polyacrylamide gel electrophoresis; HRP, horseradish peroxidase; RT-PCR, reverse transcription-polymerase chain reaction; Hb, hemoglobin.

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Introduction

Angiogenesis, the process of generating new capillary blood vessels from pre-existing ones, is essential for embryogenesis, but happens rarely in the adult with few exceptions (e.g., female reproductive system and wound healing). A pathological imbalance in this process contributes to numerous malignant, ischemic, infectious, inflammatory and immune disorders (Carmeliet, 2005). It is believed that the major steps in angiogenesis include: (a) degradation of basement membrane by proteases; (b) migration of endothelial cells (ECs) into the interstitial space and sprouting; (c) ECs proliferation at the migrating tip; (d) differentiation into new vessels; and (e) the stabilization and maturation of vessels by mediator molecules that recruit mesenchymal cells to vessel walls (Bussolino et al., 1997; Gupta and Qin, 2003). A useful approach for the treatment of pathological angiogenesis might exist in the further development of novel anti-angiogenic drugs.

Angiogenesis is controlled by a number of growth factors. Through their transmembranal receptors, growth factor signals activate signaling pathways leading to an array of responses. Mitogen-activated protein kinase (MEK) and ERK isoforms, which are involved in activating endothelial cell proliferation (Nakagami et al., 2000, 2001), appear to be the principal signaling pathways in growth factor signaling. It has been reported that ERK inhibitor caused cell apoptosis, which was accompanied by increased p53 accumulation (Kim et al., 2002; Kwon et al., 2004). By elevating levels of p21 and decreasing levels of cyclins A and B, wild type p53 can prevent abrogation of arrest (Eastman, 2004).

Terbinafine (TB), an inhibitor of fungal ergosterol biosynthesis, is a newly synthesized oral antimycotic drug (Petranyi et al., 1984). Previous reports have indicated that TB has relative few drug interactions and is safe for clinical uses (Abdel-Rahman and Nahata, 1997). The cream form and oral tablet of TB have been approved for clinical uses in the United States (Gupta and Shear, 1997), and the oral formulation has been on the market in various countries (Gupta et al., 1998). Previously, we demonstrated that TB could inhibit the proliferation and migration of cultured HUVEC, the formation of capillary-like tube, and sprouting of capillaries (Ho et al., 2004). The TB-induced cell-cycle arrest in HUVEC occurred when the cyclin-dependent kinase 2 (CDK2) was inhibited, just as the protein level of p21 was increased. Although the elevation of p21 protein levels has been demonstrated to be responsible for TB-induced cell-cycle arrest, the molecular mechanism underlying the TB-induced increase of p21 expression is still unclear. In the present study, we found that the ERK-mediated pathway may contribute to the TB-induced up-regulation of p53 protein, which is responsible for regulating the p21 protein level in HUVEC.

Materials and methods

Cell culture

HUVEC were harvested from human umbilical vein by enzymatic dissociation as described (Moraes and Stastny, 1977). The cells were grown in M199 medium containing 10% fetal bovine serum (FBS), sodium heparin (100 U/mL), endothelial cell growth supplement (ECGS, 0.03 mg mL⁻¹, Biomedical Technologies Inc., MA, USA) and antibiotics in gelatin-coated plates, and incubated at 37 °C in 5% CO₂ in air. Cells from passages 5–9 were used.

Western blot analyses

To determine the protein levels in HUVEC, the total proteins were extracted and Western blot analyses were conducted as previously described. Briefly, HUVEC were cultured in 15 cm petri dishes. After reaching subconfluence, the cells were rendered quiescent and then treated with various concentrations of TB (Novartis Pharmaceuticals Corporation, East Hanover, New Jersey USA) and incubated at 37 °C. At different time points, the cells were washed with phosphate buffer saline (NaCl 0.8%, Na₂HPO₄ 0.144%, KH₂PO₄ 0.024%, KCl 0.02%, pH 7.4), incubated with extraction buffer (Tris 50mM, pH 7.5, NaCl 150mM, PMSF 1mM, NP-40 1%, and 0.1% SDS) on ice, and then centrifuged at 12,000 × g for 30 min. The cell extract was then boiled in a ratio of 3:1 with sample buffer (Tris–HCl 250 mM, pH 6.8, glycerol 40%, dithiothreitol 400 mM, SDS 8% and bromophenol blue 0.2%). Electrophoresis was carried out using 12% SDS-polyacrylamide gel (2 h, 110 V, 40 mA, and 50 µg protein per lane). Separated proteins were transferred to PVDF membranes (1 h, 400 mA), treated with 5% fat-free milk powder (Anchor, Manurewa, NZ) to block the nonspecific IgGs, and incubated for 1h with mouse anti-human antibody specific for p21, p53, p-ERK, ERK or GAPDH (Jackson ImmunoResearch Laboratories, PA, USA). The blots were then incubated with anti-mouse IgG (Jackson ImmunoResearch Laboratories, PA, USA) linked to HRP (1:1000) for 1 h. Subsequently, the blots were developed using the ECL (enhanced chemiluminescence) system (Amersham Biosciences, UK). The bands were quantified by densitometry, using Image Pro-Plus 4.5 Software (Media Cybernetics, MD, USA).

Immunoprecipitation and kinase activity assay

As previously described (Liang et al., 1999), equal amounts of total cellular lysate were immunoprecipitated with anti-JNK1 antibody (2 mg/L) and protein A/G-PLUS agarose for 15 h at 4 °C. Kinase assay was carried out in 45 µL of kinase buffer (40 mmol/L Tris–NaOH pH 7.5, 0.5 mol/L NaCl, 0.1% NP-40, 6 mmol/L EDTA, 6 mmol/L EGTA, 10 mmol/L b-glycerophosphate, 10 mmol/L NaF, 10 mmol/L PNPP, 300 µmol/L sodium orthovanadate, 1 mmol/L benzamide, 2 mmol/L PMSF, 10 g/L aprotinin, 1 g/L leupeptin, and 1 mmol/L DTT) containing 5 mmol/L cold ATP, 10 mCi [³²P] ATP (Amersham Biosciences, UK), and 1 mg GST-c-Jun fusion protein (Santa Cruz Biotechnology, CA, USA) as substrate, and incubated for 20 min at 25 °C. Each sample was mixed with 8 µL of 5 × Laemmli's loading buffer to stop the reaction. The samples were analyzed by 8% SDS-PAGE, and the gel was then dried and subjected to autoradiography.

[³H]thymidine incorporation

The [³H]thymidine incorporation was performed as previously described (Lin et al., 2002). Briefly, HUVEC were applied to 24-well plates in growth medium (M199 plus 10% FBS and ECGS). After the cells had grown to 70–80% confluence, they were rendered quiescent by incubation for 24 h in M199 containing 2% FBS. M199 supplemented with 10% FBS and 0.05% DMSO (control) or TB was added to the cells and the cultures were allowed to incubate for 24 h. During the last 3 h of the incubation without or with TB, [³H]thymidine (Amersham Biosciences, UK) was added at 1 µCi/mL (µCi = 37 kBq). Incorporated [³H]thymidine was extracted in 0.2 N NaOH and measured in a liquid scintillation counter.

RT-PCR

Total cellular RNA was extracted from HUVEC with Trizol (Invitrogen Corporation, CA, USA) according to the manufacturer's protocol (Uchima et al., 2004). The RNA pellet was washed in 70% cold ethanol, air-dried and redissolved in 20 µL of diethyl-pyrocyanate (DEPC)-treated water (Arkel and Jackson, 2003). Two microliters was used in a total of 20 µL reaction volume as a template for PCR amplification. PCR was performed under standard conditions in 20 µL; 10 mM Tris, pH 8.3, 40 mM KCl, 1.5 mM MgCl₂, 250 µM dNTP, 10 pM each primer (sense and antisense) and 1 U Taq DNA polymerase (Lee et al., 2006). The experimental conditions were as follows: 95 °C for 1 min; 55 °C for 1 min; and 72 °C for 1 min for 35 cycles. The following primers were used: p53 5'-ATGAGGAGCCGAGT-CAG-3' and 5'-TCAGTCTGAGTCAGGCCCTTC-3'; p21 5'-AGGAGGCGC-CATGTCAGAAC C-3' and 5'-TCCTGTGGGCGGATTAGGGCT-3'; p27 5'-CC-ACGAAGAGTAAACCCGGG-3' and 5'-GTCTGCTCCACAGAACCAGG-3';

GA3PDH 5'-CCACCATGGAGAAGGCTGGGGCTCA-3' and 5'-ATCACGCCA-CAGTTTCCCGGAGGGG-3'. The PCR products were electrophoresed on 1.8% agarose in 1× Tris-acetate-EDTA (TAE) buffer, and stained with ethidium bromide solution.

Preparation of plasmid constructs

p21 promoter. The pWWP-luc plasmid construct, containing promoter region of human p21/WAF1 between positions -2300 and +8 was a gift from Dr. B. Vogelstein (Johns Hopkins University, Baltimore, MD) (el-Deiry et al., 1993). p21 promoter was subcloned into pGL3-basic luciferase reporter vector (Promega Corporation, WI, USA) at the unique HindIII site, and the sequence identity was confirmed by ABI PRISM 377 DNA Analysis System (Perkin-Elmer, MA, USA).

p53 dominant negative (DN) construct. For amplification of the hot-spot mutant (V143A) of p53 (Wong et al., 1999), recombinant PCR was performed. Synthesized cDNA (SuperScript III First-Strand Synthesis System for RT-PCR; Invitrogen Corporation, CA, USA) from HUVEC was used as a template and Phusion high-fidelity DNA polymerase (Finnzymes Oy, Espoo, Finland) with 3' → 5' exonuclease activity was used in subsequent PCR. In the first round PCR, forward (5'-ATGGA-GGAGCCGCAGTCAG-3') and reverse (5'-TCAGTCTGAGTCAGGCCCTTC-3') primers were used to amplify full-length wild type p53 (WT-p53). Reactions were treated in a thermal cycle machine to incubate at 98 °C for 30 s followed by 35 cycles of: 98 °C for 10 s, 66 °C for 30 s, 72 °C for 30 s and a final incubation of 72 °C for 7 min. PCR product of WT-p53 was cleaned up and used for the next PCR. In the second round PCR, two primers set 1 and 2 were used to synthesize fragment A and

fragment B. Primer set 1: forward (5'-CTTAAGCTTGCCGCCATGGAGGAG-3'), and reverse (5'-CACAGATGCGCAGGGCAGGT-3'). Primer set 2: forward (5'-ACCTGCCCTGCGCAGCTGTG-3') and reverse (5'-GTGCTGGATATCTCAGTCTGAGTC-3'). Two-step PCR reactions were applied to synthesize fragment A: a first cycle of 40 s at 98 °C and 72 °C for 1 min followed by 98 °C for 10 s, 72 °C for 1 min for 34 cycles and a final incubation of 72 °C for 7 min. Standard PCR for fragment B was treated at 98 °C for 30 s followed by 35 cycles of: 98 °C for 10 s, 67 °C for 30 s, 72 °C for 30 s and a final incubation of 72 °C for 7 min. For each fragment, the amplified PCR product was gel purified and used as a mixed template for synthesizing p53^{V143A} in the PCR reaction: a first cycle of 40 s at 98 °C, 67 °C for 30 s and 72 °C for 1 min followed by 98 °C for 10 s, 67 °C for 30 s and 72 °C for 1 min for 34 cycles and a final incubation at 72 °C for 7 min. The synthesized PCR product of p53^{V143A} was purified, digested with restriction enzyme HindIII and EcoRV, and cloned into pcDNA3.1+ (Invitrogen Corporation, CA, USA). The p53^{V143A} sequence fidelity was confirmed by ABI PRISM 377 DNA Analysis System (Perkin-Elmer, MA, USA).

Constitutive active MEK-1 construct. MEK-1 cDNA was a gift from Prof. Ruey-Hwa Chen (Institute of Biological Chemistry, Taipei, Taiwan) (Mansour et al., 1994; Chen et al., 2005).

Cell transfection

Transfection was performed using jetPEI-HUVEC transfection reagent (Polyplus Transfection, Bioparc, France) and following the manufacturer's protocol. Briefly, jetPEI-HUVEC/DNA mixture was added drop-wise onto the DMEM-Glutamax medium (Invitrogen Corporation, CA, USA) containing 2% FBS,

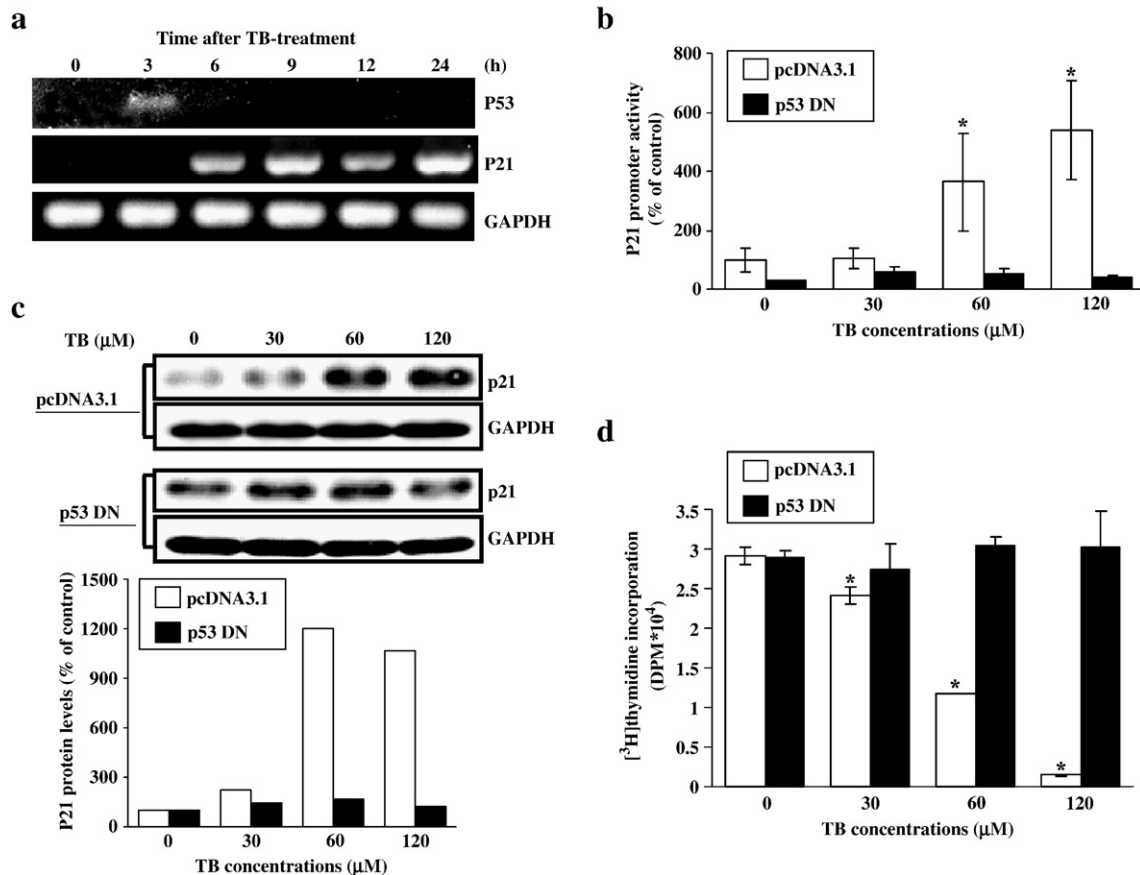


Fig. 1. Involvement of p53 in TB-induced up-regulation of p21 in HUVEC. (a) TB increases the levels of p21 and p53 mRNA in HUVEC. The increased p53 mRNA level was observed at 3 h after TB treatment, whereas the increased p21 mRNA was observed at 6 h after treatment. RT-PCR products of GAPDH were used as an internal control. (b) Transfection of the cells with p53DN abolished the TB-induced increases of p21 promoter activity. Four samples were analyzed in each group, and the values shown represent the means ± s.e.mean. **p* < 0.05 different from control (cells transfected with pcDNA 3.1 without TB treatment). (c) Transfection of the cells with p53DN abolished the TB-induced increases of p21 protein level. The bottom panel shows the quantified results of p21 protein levels, which were adjusted with GAPDH protein level and expressed as percentage of control. (d) Transfection of the cells with p53DN abolished the TB-induced decreases of thymidine incorporation into HUVEC. Four samples were analyzed in each group, and values represent the means of percent of control ± s.e.mean. **p* < 0.05 different from control. TB: terbinafine; p53DN, p53 dominant negative DNA.

mixed gently, and incubated in a humidified 37 °C incubator for 4 h. The transfected cells were incubated in fresh HUVEC growth medium for additional 24 h. M199 supplemented with 10% FBS and TB was added to the cells and the cultures were allowed to incubate for the indicated hours. In these experiments, cells transfected with empty vector were served as a control.

Luciferase assay

After terbinafine treatment, the cells were washed twice with ice-cold phosphate buffer saline and lysed by lysis reagent (Promega Corporation, WI, USA). Luciferase activities were recorded in a 20/20^l luminometer (Turner Biosystems, CA, USA) using the dual luciferase assay kit (Promega Corporation, WI, USA) according to the manufacturer’s instructions. Each experiment was performed at least three times.

Matrigel angiogenesis assay

The Matrigel angiogenesis assay was performed as previously described with minor modifications (Passaniti et al., 1992). A 500 µL portion of a liquid mixture of Matrigel (350 µL, BD Biosciences, CA, USA), M199 medium (150 µL) and VEGF (50 µg/L) (Inoki et al., 2002) with or without terbinafine

was injected subcutaneously into 7-week-old C57/BL6 mice. Matrigel rapidly formed a single, solid gel plug. The plugs allowed VEGF to stimulate angiogenesis. On the 7th day, mice were euthanized, and the Matrigel plugs were excised. Vascularization of Matrigel plugs was quantified by measuring hemoglobin (Hb) content using Drabkin reagent (Sigma-Aldrich, MO, USA) according to the instructions of the manufacturer.

Statistics

All data were expressed as the mean value±s.e.mean. Comparisons were subjected to one way analysis of variance (ANOVA) followed by Fisher’s least significant difference test. Significance was accepted at $p < 0.05$.

Results

p53-dependence of TB-induced p21 up-regulation

Previously, we demonstrated that TB exerts an anti-proliferative effect in HUVEC through induction of p21 expression,

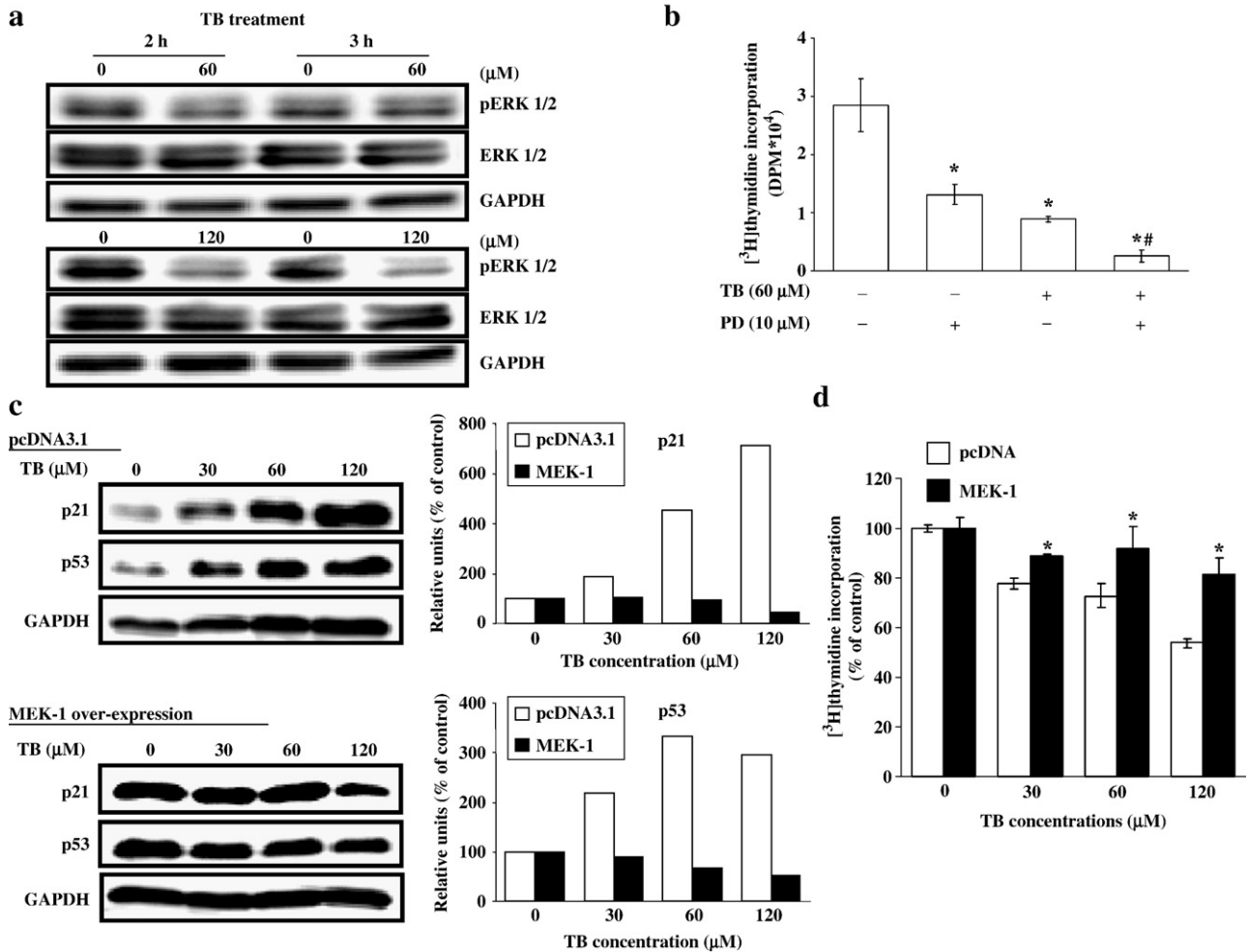


Fig. 2. Involvement of ERK in the TB-induced increases of p53 and p21 protein. After treatment with TB for 2–3 h, the cells were processed for total protein extraction and Western blot analysis. (a) Treatment of the cells with TB (60 and 120 µM) for 2 or 3 h decreased the levels of phosphorylated ERK protein. (b) The TB-induced decrease of thymidine incorporation was enhanced by pretreatment of the cells with PD98059 (an ERK inhibitor). HUVEC were pre-treated with PD98059 for 1 h followed by TB (60 µM) for 24 h. [³H]thymidine incorporation was done to examine the DNA synthesis. Four samples were analyzed in each group, and values represent the means±s.e.mean. * $p < 0.05$ different from control. # $p < 0.05$ different from TB-treated. (c) Over-expression of MEK-1 abolished the increase of p21 and p53 expression induced by TB. The right panels showed the quantified results of p21 and p53 protein levels, which were adjusted with GAPDH protein level and expressed by percentage of control. (d) Over-expression of MEK-1 abolished the TB-induced decrease of thymidine incorporation. HUVEC were transiently transfected with control vector (pcDNA 3.1) or vector coding for constitutive active MEK-1 followed by TB treatment for 24 h. Four samples were analyzed in each group, and values represent the means±s.e.mean. * $p < 0.05$ different from corresponding control (pcDNA-transfected cells).

which in turn inhibits the CDK2 activity (Ho et al., 2004). To examine whether the TB-induced increase of p21 expression is regulated by p53, we performed RT-PCR to examine the expression of p21 and p53 mRNA at different time points after TB treatment (60 μ M). As illustrated in the Fig. 1a, TB-induced increases of p53 and p21 mRNA level in HUVEC began at 3h and 6h, respectively. The RT-PCR products of GAPDH were used as an internal control to monitor amounts of specimen loading. Fig. 1b further demonstrates that TB concentration-dependently increased the p21 promoter activity in HUVEC. Transfection of the cells with p53DN abolished the TB-induced increases of p21 promoter activity (Fig. 1b) and protein level (Fig. 1c) and decrease of thymidine incorporation (Fig. 1d) in HUVEC.

Involvement of ERK in the TB-induced increases of p21 and p53 expression in HUVEC

Since the ERK pathway has been implicated to be involved in the regulation of endothelial cell proliferation, we further examined the levels of total and phosphorylated ERK protein in the TB-treated HUVEC. As illustrated in Fig. 2a, TB de-

creased the levels of phosphorylated ERK in HUVEC. Moreover, treatment of the cells with PD98059, an ERK inhibitor, decreased the DNA synthesis and accentuated the TB-induced inhibition of DNA synthesis in HUVEC (Fig. 2b). To further investigate the involvement of ERK in cell-cycle alternations caused by TB, we over-expressed the MEK-1, the immediate upstream activator kinase of ERK, in HUVEC. Over-expression of MEK-1 abolished the TB-induced increases of p21 and p53 protein (Fig. 2c) and decrease of thymidine incorporation (Fig. 2d).

Isoprenylation might not be involved in the TB-induced growth arrest

To examine whether prenylation was involved in the TB-induced inhibition of proliferation in HUVEC, the cells were pretreated with isoprenoid pyrophosphate precursors (FOH and GGOH) or isoprenoid inhibitors (FTI and GGTI) followed by TB treatment. The results showed that treatment of the cells with isoprenoid inhibitors (Fig. 3a and b) accentuated the TB-induced inhibition of DNA synthesis. However, isoprenoid pyrophosphate precursors (Fig. 3c) did not affect the TB-induced inhibition of

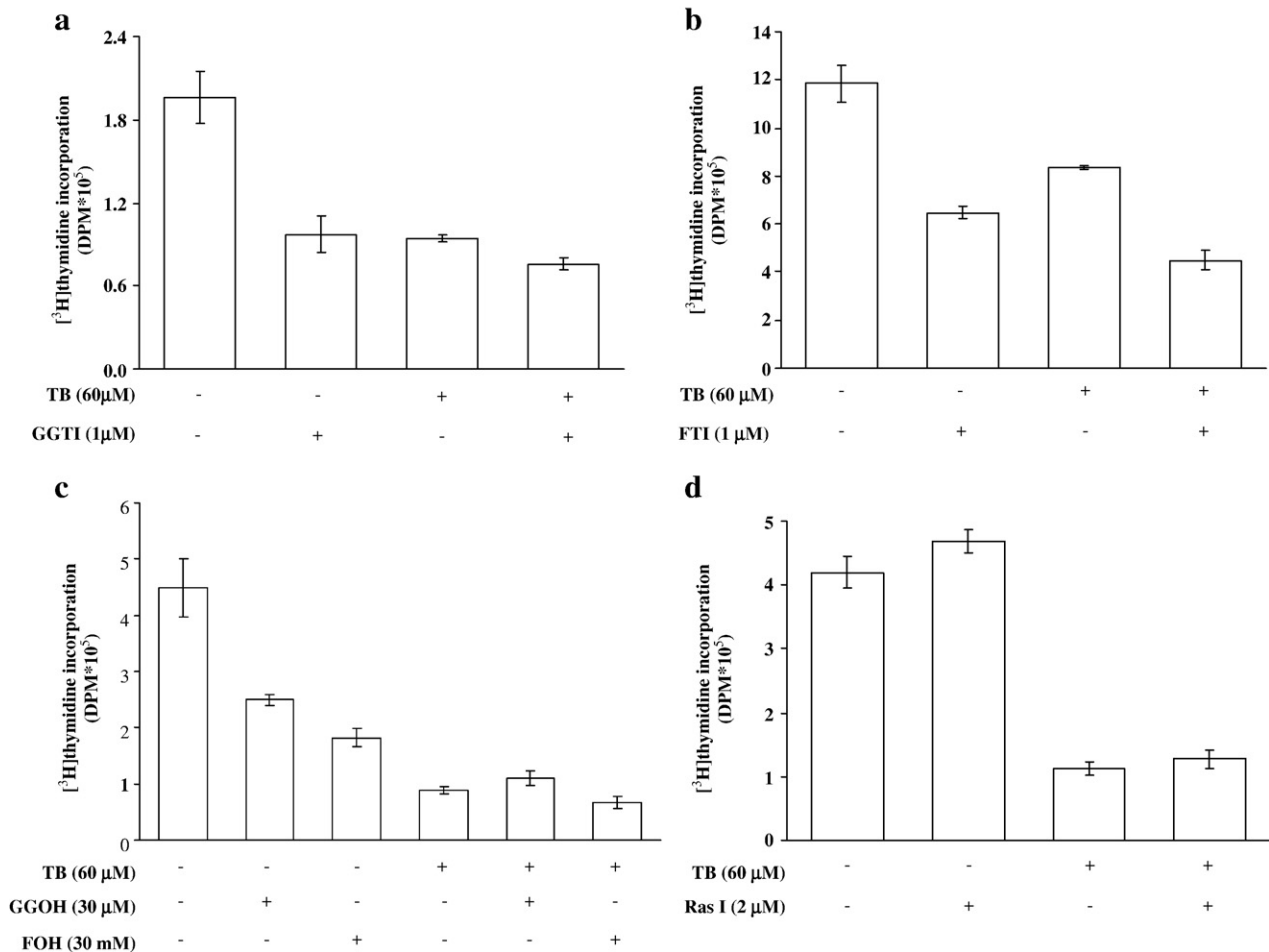


Fig. 3. Ras and protein isoprenylation might not be involved in TB-induced anti-proliferative effects in HUVEC. Pretreatment of the cells with the inhibitors of isoprenoid pyrophosphate, GGTI (a) and FTI (b), accentuated the TB-induced inhibition of DNA synthesis. However, neither the precursors of isoprenoid pyrophosphate, GGOH and FOH (c), nor Ras inhibitor (d) affected the TB (60 μ M)-induced anti-proliferation effects. Four samples were analyzed in each group, and values represent the means \pm s.e.mean. Ras I, Ras inhibitor.

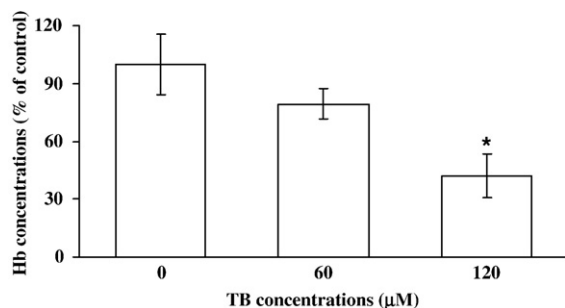


Fig. 4. Effect of TB on angiogenesis *in vivo*. Angiogenesis was assessed by using the Matrigel plug assay. Matrigel mixed with 50×10^{-6} g/L VEGF and phosphate buffer saline or TB (60 or 120 μ M) was injected s.c. into C57BL/6J mice. On the 7th day, the plugs were excised, and Hb levels were measured. The mean Hb concentration of plugs excised from animals treated with vehicle without TB is 100%. Seven or eight samples were analyzed in each group, and values represent the means \pm s.e.mean. * $p < 0.05$ different from control.

DNA synthesis in HUVEC. We also investigated the involvement of Ras in the TB-mediated growth inhibition in HUVEC. Since isoprenylation is important for subcellular distribution and functions of Ras, we further examined the involvement of Ras protein in regulating the HUVEC proliferation. As illustrated in Fig. 3d, Ras inhibitor did not affect the TB-induced inhibition of DNA synthesis in HUVEC.

TB inhibits angiogenesis *in vivo*

Previously, we applied the capillary-like tube formation assay and the chick embryo chorioallantoic membrane (CAM) assay to demonstrate the anti-angiogenic activity of TB (Ho et al., 2004). However, the results from *in vitro* study could not always be replicated in an *in vivo* setting. To confirm the *in vivo* anti-angiogenic effect of TB, we applied the Matrigel angiogenesis assay. As illustrated in Fig. 4, TB (0, 60 and 120 μ M) concentration-dependently decreased Hb content, which represents the blood vessel formation, in the plug.

Discussion

Previously, we demonstrated that TB-induced cell-cycle arrest at the G₀/G₁ phase in HUVEC through an up-regulation of the CDK inhibitor, p21 protein (Ho et al., 2004). The findings of the present study demonstrate the TB-induced p21 and p53 mRNA expression. Transfection of the cells with p53DN abolished the TB-induced increases of promoter activity and protein levels of p21. Treatment of HUVEC with TB decreased the phosphorylated ERK protein levels (Fig. 2a). On the other hand, over-expression of MEK-1 abolished the TB-induced increases of p21 and p53 protein. Apparently, this is the first demonstration that TB induces increases of p21 mRNA and protein in HUVEC through a p53-dependent pathway mediated by inhibition of ERK phosphorylation.

It has been suggested that p21 is a transcriptional target of the tumor suppressor gene p53 and might promote p53-dependent cell-cycle arrest or apoptosis (Mendoza-Rodriguez and Cerbon, 2001; Jerry et al., 2002; Yu and Zhang, 2005). However, p21 can also be activated through a p53-independent pathway (Marchetti

et al., 1996). Our previous study in the COLO-205 cells showed that the protein level of p53 and the binding between p53 protein and p53 consensus binding site in p21 promoter DNA probe were increased by TB treatment, and TB treatment caused an increase in the level of p53 protein, which in turn up-regulated the p21 protein, and finally caused cell-cycle arrest (Lee et al., 2003). In the HUVEC, our previous data showed that TB treatment resulted in an increase in the protein levels of p21 and p53 (Ho et al., 2004). In this study, we demonstrated that TB induced an increase in the mRNA levels of p21 and p53 in HUVEC, suggesting that transcriptional regulation is involved in the TB-induced increases in p21 and p53 protein levels. p53 has been identified as a host protein interacting with the SV40 large T (Lane and Crawford, 1979). It is essential for preventing improper cell proliferation and maintaining the genome integrity following genotoxic stress. Following DNA damage, hypoxia, oncogene over-expression, and nucleotide depletion, p53 undergoes extensive post-translational changes which result in the increased binding of p53 to DNA and transcriptional activation of its target genes (Vogelstein et al., 2000; Yu and Zhang, 2005). These targets regulate a number of cellular processes; among which the best studied is a block in the cell-division cycle. The p53 protein directly stimulates the expression of p21. In the present study, RT-PCR results showed that the induction of p53 mRNA expression induced by TB treatment was observed prior to p21 mRNA induction (Fig. 1a). In the p53DN-transfected HUVEC, the increased p21 promoter activity and protein levels induced by TB were abolished (Fig. 1b and c). Taken together, our results suggest that TB increased p21 expression through p53 transactivation on the p21 promoter.

Mitogen-activated protein kinase (MAPK) cascades have been shown to participate in a diverse array of cellular programs, including cell division. In mammalian systems, four MAPK modules, ERK1/2, JNK and p38 MAPK have been identified. Activation of the MAPK pathway has been commonly associated with cell-cycle progression. For example, activation of the oncogenic MEK1 caused transformation of the NIH3T3 cells (Okazaki and Sagata, 1995), and injection of the constitutive active form of MEK1 to the NIH3T3 cells induced DNA replication (Yuen et al., 2001). In response to a wide variety of extracellular stimuli (such as growth factors, cytokines, and environmental stresses), ERK become phosphorylated and activated through a three-kinase signaling cascade (Marshall, 1995). The ERK are vital in regulating diverse cellular activities, including cell proliferation, cell-cycle progression, differentiation, apoptosis, and tumor invasion (Marshall, 1995; Kolch, 2000). ERK inhibitor has been reported to increase p53 accumulation (Kim et al., 2002; Kwon et al., 2004). In the present study, we showed that TB inhibited the phosphorylation of ERK (Fig. 2a), and over-expression of MEK-1 abolished the TB-induced increases of p53 and p21 protein level (Fig. 2c). These findings suggest that ERK inactivation might be involved in the TB-induced increases of p53 and p21 expression.

In the cholesterol synthesis pathway, squalene epoxidase converts squalene to 2,3-oxidosqualene. TB, an inhibitor of squalene epoxidase, might alter the protein prenylation by changing the expression of farnesyl pyrophosphate (FPP) and geranylgeranyl

pyrophosphate (GGPP), which are the upstream molecules of squalene and might regulate protein post-translational modifications. It has been identified that membrane localization of the small G proteins, such as Ras, is critical to their functions. The membrane localization of the small G proteins is dependent on a post-translational modification that results in attachment of hydrophobic prenyl groups that anchor the small G proteins to plasma membranes (Magee and Marshall, 1999). Activation of Ras triggers a phosphorylation cascade that successively activates MEK1/2 and the ERK-MAP kinases, which are translocated to the nucleus and activate transcription factors involved in cell proliferation. Previously, we showed that TB concentration-dependently decreased the protein levels of Ras in HUVEC and Ras inhibitor peptide abolished the FOH-induced prevention of TB-induced migration inhibition in HUVEC (Ho et al., 2006). However, our data from the present study showed that pretreatment of the cells with Ras inhibitor did not significantly affect the TB-induced inhibition of DNA synthesis (Fig. 3d). These data suggest that Ras might be involved in the TB-induced inhibition of endothelial cell migration, but not proliferation.

In the present study, our results showed that pretreatment of the cells with FTI accentuated the TB-induced inhibition of DNA synthesis (Fig. 3b); and this has led us conjecture the involvement of farnesylation in TB-induced growth arrest. Unexpectedly, FOH did not reverse TB-inhibited DNA synthesis. FTIs have been reported to induce hypophosphorylation of retinoblastoma protein (pRb), a key player in the transition from the G1 to S phase (Sepp-Lorenzino and Rosen, 1998; Edamatsu et al., 2000), and increase expression of p21 at the transcriptional level in the p53-independent pathway (Feldkamp et al., 1999). Besides Ras, Rheb, which is required for cell-cycle progression and cell growth, is also modified by farnesylation. We speculate that FTI and TB might inhibit the DNA synthesis of HUVEC through different pathways.

Previously, we showed that TB suppresses proliferation of cultured HUVEC by inhibiting DNA synthesis and arresting cell-cycle activity through up-regulation of p21. In the present study, we further investigate the molecular mechanisms underlying TB-induced up-regulation of p21 in HUVEC. We demonstrate that TB could cause ERK inactivation, which in turn increases p53 protein expression and results in the transactivation of p21 promoter and increase in p21 protein, and finally leads to growth arrest in HUVEC. Demonstration of TB-induced inhibition of angiogenesis *in vivo* together with our previous *in vitro* studies strongly suggests potential applications of TB in anti-angiogenic therapy.

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