

# Thrombospondin-1-Induced Vascular Smooth Muscle Cell Chemotaxis: The Role of the Type 3 Repeat and Carboxyl Terminal Domains

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**Abstract** Thrombospondin-1 (TSP-1), an acute phase reactant implicated in vascular disease, is a matricellular glycoprotein with six domains that confer different functions. The authors have shown TSP-1 induces vascular smooth muscle cell (VSMC) chemotaxis via extracellular signal-regulated kinases-1 and -2 (ERK) and p38 kinase (p38) and that a fusion protein of the carboxyl terminal (COOH) and type 3 repeat (T3) domains independently induce VSMC chemotaxis. The purpose of this study was to determine whether COOH-, T3-induced VSMC chemotaxis, or both, is dependent upon ERK or p38 activation. To determine if the T3, COOH, or type 2 repeat domain (T2, control domain not associated with chemotaxis) activate ERK, p38, or both, VSMCs were exposed to each fusion protein (20 µg/ml for 15, 30, 60, or 120 min), serum-free media (SFM, negative control), or TSP-1 (20 µg/ml for 30 min, positive control). Western immunoblotting was performed for activation studies. Using a microchemotaxis chamber, VSMCs pre-incubated in SFM, DMSO (vehicle control), PD98059 (10 µM), or SB202190 (10 µM) were exposed to each domain, TSP-1, or SFM. After 4 h (37°C), migrated VSMCs were recorded as cells/five fields (400 ×) and analyzed by paired *t*-test. ERK was activated by T2, T3, and COOH. However, p38 was activated by T3 and COOH, but not T2. T3 and COOH-induced VSMC chemotaxis were inhibited by PD98059 or SB202190, but more completely by SB202190. The T2 domain had no effect on VSMC chemotaxis. These results suggest activation of the p38 pathway may be more specific than ERK for COOH- and T3-induced VSMC chemotaxis. *J. Cell. Biochem.* 89: 500–506, 2003. © 2003 Wiley-Liss, Inc.

**Key words:** thrombospondin; ERK; p38 kinase; smooth muscle cell; chemotaxis

Vascular smooth muscle cell (VSMC) chemotaxis—a form of directed cell migration along a gradient in response to a soluble chemoattractant—contributes to development of vascular pathologic states, including intimal hyperplasia and atherosclerotic plaque formation. Throm-

bospondin-1 (TSP-1)—a matricellular glycoprotein chemotactic for VSMCs—is not integral to arterial wall structure, but is expressed locally after acute arterial injury, in atherosclerotic plaque, and early intimal hyperplasia [Miano et al., 1993; Yabkowitz et al., 1993; Reed et al., 1995; Roth et al., 1998]. “Matricellular refers to a group of modular, matrix proteins that do not contribute to structural integrity, but whose functions are accomplished by binding to matrix proteins, cell-surface receptors, cytokines, and proteases, all of which interact with the cell surface to exert different effects” [Bornstein, 1995].

The structure of TSP-1, 420 kDa in size, consists of the following six domains: amino terminal or heparin binding (NH<sub>2</sub>), procollagen homology, properdin or type 1 repeat, type 2 repeat (T<sub>2</sub>), type 3 repeat (T<sub>3</sub>), and carboxyl terminal (COOH). Several receptors also have

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been identified and interacted with domains or peptide sequences from TSP-1. An RGD sequence in the T3 binds to the integrin  $\alpha_v\beta_3$  [Lawler et al., 1988]. Two sequences from the carboxyl-terminal domain that contain a VVM motif bind CD47 and regulate activity of the integrins  $\alpha_{IIb}\beta_3$ ,  $\alpha_v\beta_3$ , and  $\alpha_2\beta_1$  in specific cell types [Gao et al., 1996; Chung et al., 1997; Wang and Frazier, 1998]. Differential expression of the receptors for these domains and the varying accessibility of these domains within the intact TSP-1 molecule are potential mechanisms for regulating the effects of TSP-1 on cell behavior. Responses of cells to TSP-1 include adhesion, spreading, chemotaxis, and haptotaxis through cooperative interactions with multiple domains [Roberts et al., 1987]. The focus of this research is on the mechanism of TSP-1-induced VSMC chemotaxis. A fusion protein synthesized to the COOH domain has been determined to be the predominant domain that induces chemotaxis [Nesselroth et al., 2001]. The NH2 and T3 domains also independently induce VSMC chemotaxis.

TSP-1 has been reported to induce VSMC chemotaxis through extracellular signal-regulated protein kinases-1 and -2 (ERK) [Gahtan et al., 1999] and p38 kinase (p38) [Wang et al., 1999]. Whether the effect of the T3 and COOH domains on VSMC chemotaxis is through ERK, p38 activation, or both was examined, based on the observation that specific domains of TSP-1 independently induce chemotaxis.

## METHODS

### VSMC Culture

Bovine aortic VSMCs were isolated and seeded in a culture medium containing serum free medium (SFM, Dulbecco's modified Eagle's medium, Gibco-BRL, Gaithersburg, MD), 10% fetal bovine serum (FBS), and antibiotics ( $1 \times$  penicillin G sodium, streptomycin sulfate, and amphotericin B, Gibco-BRL) in primary culture [Gahtan et al., 1999]. Subconfluent monolayers of VSMCs in early passage (2–5) were used for each experiment.

### Chemoattractants and Inhibitors

Purified TSP-1 from human platelets and fusion proteins to T2, T3, and COOH domains were prepared as described previously [Legrand et al., 1992; Adams and Lawler, 1994]. The

cDNAs for the type two repeats (amino acids 559 to 669), the type three repeats (amino acids 674 to 925), and the C-terminal domain (amino acids 926 to 1152) of TSP-1 were prepared by PCR. The sequence of the PCR products was verified and they were cloned into pGEX vectors (Amersham Pharmacia Biotech, Piscataway, NJ). All sequencing was done by the chain termination method of Sanger [Sanger et al., 1977]. The fusion proteins were expressed and purified using glutathione-sepharose following protocols that were provided by the manufacturer. The concentration of the TSP-1, T2, T3, and COOH proteins was 20  $\mu$ g/ml. This concentration was chosen, because peak chemotaxis occurred by this concentration [Gahtan et al., 1999; Wang et al., 1999].

PD98059 and SB202190 were purchased (Calbiochem, La Jolla, CA). The concentration, 10  $\mu$ M, was used for both inhibitors as this concentration has been previously established for TSP-1 [Gahtan et al., 1999; Wang et al., 1999]. The concentration 10  $\mu$ M was used because it induced inhibition without causing toxic or nonspecific effects of the inhibitors.

### Western Blot Analysis

VSMCs were exposed to SFM, T2 (control domain not associated with chemotaxis), T3, or COOH for 0, 15, 30, 60, and 120 min (20  $\mu$ g/ml) and TSP-1 (20  $\mu$ g/ml, positive control) for 30 min, using similar methods described previously [Gahtan et al., 1999]. VSMCs were washed twice with ice-cold phosphate-buffered saline solution and scraped in lysis buffer containing 50 mmol/L HEPES (pH 7.4), 150 mmol/L NaCl, 10% glycerol, 1% Triton X-100, 1.5 mmol/L  $MgCl_2$ , 1 mmol/L EGTA, 100 mmol/L sodium fluoride, 10 mmol/L sodium orthovanadate, 10  $\mu$ g/ml leupeptin, and 1 mmol/L phenylmethylsulfonylfluoride. Western blot analysis was performed by SDS-PAGE (10% polyacrylamide) and transferred to a nitrocellulose membrane (Amersham, Arlington Heights, IL). Blocking was performed in a 5% non-fat milk-tris-buffer saline tween-20 (TBST) solution. Blots were incubated in a 5% non-fat milk-TBST solution with anti-p-ERK 1/2 (Cell Signaling, Beverly, MA), anti-p-p38 (Cell Signaling), and a secondary antibody tagged with HRP (Cell Signaling). The fold inductions were normalized by reprobing for total ERK 1/2 or total p38 (Cell Signaling) after stripping its respective membrane. Immunoreactivity was

determined by means of enhanced chemiluminescence (Amersham) and quantified with an UN-SCAN-IT densitometer (Silk Scientific, Orem, UT). To verify that the inhibitors used in the chemotaxis assay were specific to ERK (PD98059) and p38 (SB202190), quiescent cells were exposed to either inhibitor for 30 min and then exposed to T3 or COOH (20  $\mu\text{g}/\text{ml}$ ) for 30 min and then processed as described above.

### Chemotaxis Assay

A 48-well modified Boyden microchemotaxis chamber (NeuroProbe, Inc., Gaithersburg, MD) was used for chemotaxis assays [Nesselroth et al., 2001]. In the lower chamber, 30  $\mu\text{l}$  of one of the following chemoattractants were placed: T2, T3, COOH domain, GST (fusion protein control), or TSP-1 (20  $\mu\text{g}/\text{ml}$ ). A poretics membrane (8- $\mu\text{m}$  pores) was applied, followed by the gasket and upper chamber. VSMCs (50,000/well) after 30 min exposure to SFM, PD98059 (ERK inhibitor by direct inhibition of MEK1, 10  $\mu\text{M}$ ), SB202190 (p38 inhibitor, 10  $\mu\text{M}$ ), or DMSO (vehicle) were placed into the top wells. Assays were conducted for 4 h at 37°C. The membrane was then removed from the chamber, fixed in 70% ethyl alcohol, stained with hematoxylin, and mounted on a slide. Results were recorded as the total number of cells migrated to the undersurface of the membrane for five high-power fields (400 $\times$ ) per well.

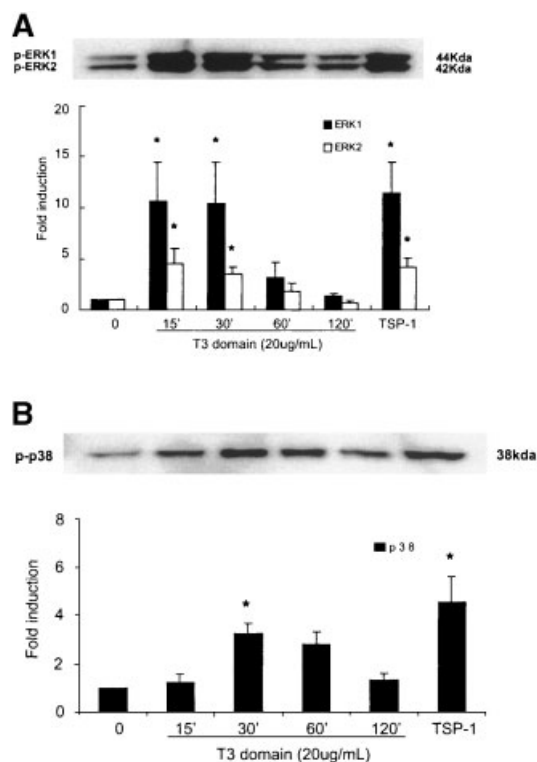
### Statistical Analysis

For the activation studies, differences in densitometry data were assessed by paired *t*-test. For the chemotaxis experiments, the number of migrated VSMCs per five fields under high power (400 $\times$ ) per well was calculated for each experimental group. Each experiment was performed in triplicate. The data were analyzed by paired *t*-test. A *P* value less than 0.05 was considered significant.

## RESULTS

### ERK and p38 Activation Studies

The T3 domain induced ERK activation by 15 min [ERK1,  $10.62 \pm 3.79$  (fold increase compared to SFM, mean  $\pm$  SEM),  $P = 0.042$ ; ERK2,  $4.5 \pm 1.4$ ,  $P = 0.04$ ], to a level similar to that seen after a 30-min exposure to TSP-1 (Fig. 1A). After 30 min, ERK activation declined. Peak activa-

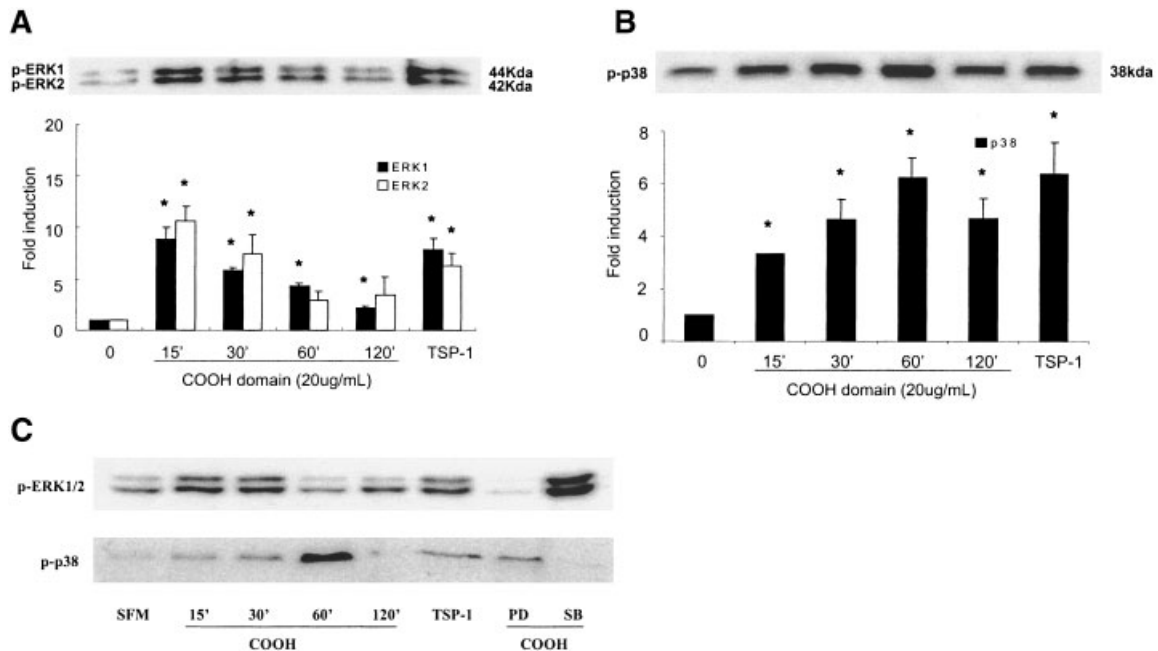


**Fig. 1.** Time course of ERK and p38 activation after T3 exposure. Total cell lysates were analyzed by Western blotting by use of antibodies against phospho ERK1/ERK2 or p38. Representative Western blots and a summary of the densitometry data for three separate experiments are shown. **A:** Activation of ERK is seen at 15 and 30 min of exposure; **(B)** peak activation of p38 is seen at 30 min.  $*P < 0.05$ .

tion of p38 was seen at 30 min of treatment with T3 ( $3.25 \pm 0.42$ ,  $P = 0.006$ ; Fig. 1B).

COOH induced activation of ERK occurred after 15 min of exposure (ERK1,  $8.87 \pm 1.08$ ,  $P = 0.009$ ; ERK2,  $10.61 \pm 1.5$ ,  $P = 0.028$  Fig. 2A). Activation continued at 30 min, and declined thereafter with activation still present at 60 and 120 min for ERK1. Activation of p38 occurred by 15 min of COOH exposure and was sustained during subsequent intervals examined (30, 60, and 120 min; at 60 min,  $6.25 \pm 0.73$ ,  $P = 0.0007$ ; Fig. 2B). PD98059 inhibited ERK activation and SB202190 inhibited p38 activation of COOH (Fig. 2C).

The T2 domain induced ERK activation by 15 min ( $10.9 \pm 4.6$ ,  $P = 0.034$ ; Fig. 3A). Activation declined thereafter and was not significant at the other time points studied (30, 60, and 120 min). No activation of p38 by T2 was noted (at 15 min,  $1.41 \pm 0.36$ ,  $P = 0.17$ ; Fig. 3B). Isolated GST protein had no effect on ERK or p38 activation (data not shown).



**Fig. 2.** Time course of ERK and p38 activation after COOH exposure. Total cell lysates were analyzed by Western blotting by use of antibodies against phospho ERK1/ERK2 or p38. Representative Western blots and a summary of the densitometry data are shown ( $n = 3$ ). **A:** Peak activation of ERK is seen by 15 min of exposure; **(B)** peak activation of p38 is seen by 60 min. **C:** The

inhibitors PD98059 and SB202190 were confirmed to be specific for ERK and p38, respectively. PD98059, but not SB202190, inhibited COOH-induced ERK activation. Similarly, SB202190, but not PD98059, inhibited COOH-induced p38 activation. \* $P < 0.05$ .

### T3- and COOH-Induced VSMC Chemotaxis Is Functionally Dependent on ERK and p38

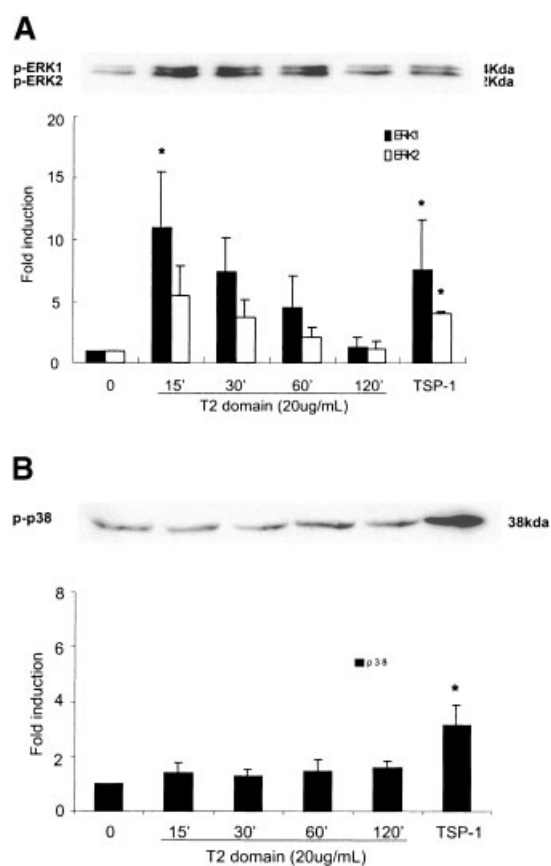
A significant increase in VSMC chemotaxis was observed in response to T3 ( $106 \pm 4.6$ ,  $P = 0.004$ ), COOH ( $146 \pm 3.7$ ,  $P = 0.002$ ), or TSP-1 ( $162 \pm 4.2$ ,  $P = 0.0015$ ), in comparison to SFM ( $21.3 \pm 1.8$ ; Fig. 4A). Both PD98059 and SB202190 independently inhibited T3-, COOH-, and TSP-1-induced VSMC chemotaxis (T3,  $P = 0.01$  and  $0.002$ ; COOH,  $P = 0.0004$  and  $0.0003$ ; TSP-1,  $P = 0.00005$  and  $0.00004$ ). SB202190 inhibited chemotaxis more than PD98059 for each chemoattractant studied (T3,  $P = 0.005$ ; COOH,  $P = 0.021$ ; TSP-1,  $P = 0.014$ ). When making comparisons to the negative control, chemotaxis was still significant in the PD98059 treated groups (T3,  $P = 0.008$ ; COOH,  $P = 0.005$ ; TSP-1,  $P = 0.002$ ); however, chemotaxis was abolished by SB202190 (T3,  $P = 0.11$ ; COOH,  $P = 0.17$ ; TSP-1,  $P = 0.053$ ). The degree of chemotaxis was less for T3 than COOH ( $P = 0.029$ ). Similarly, the degree of chemotaxis was less for COOH than for TSP-1 ( $P = 0.042$ ). DMSO had no intrinsic effect on T3 and COOH-induced VSMC chemotaxis. T2 ( $P = 0.097$ ) and isolated GST protein did not

induce chemotaxis (Fig. 4B). Morphologically, neither the inhibitors nor GST induced any obvious toxic effects in the cells.

### DISCUSSION

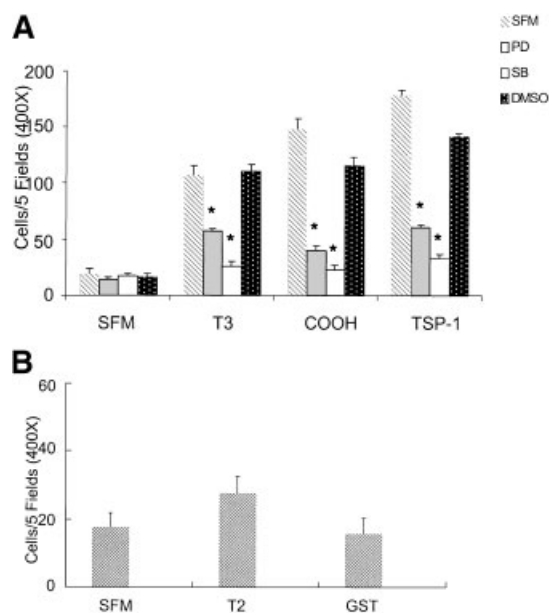
TSP-1 is a matricellular protein because it has the ability to bind both extracellular matrix proteins and cellular receptors, and the many effects of this molecule are the result of these bindings. Multiple, often contradictory, functions have been attributed to TSP-1, varying by both cell type and species. This variability is thought to be due to differences in the extracellular milieu and cell-surface receptors present in various different cell types [Murphy-Ullrich et al., 1993; Hogg, 1994; Gao et al., 1996; Qian et al., 1997]. For human [Patel et al., 1997] and bovine [Nesselroth et al., 2001] VSMCs, TSP-1 has been observed to induce VSMC chemotaxis, relevant to vessel remodeling after acute arterial injury and in the subsequent development of atherosclerotic plaque and intimal hyperplasia.

The cell-signal-transduction events by which TSP-1 induces VSMC chemotaxis have not been well characterized. A functional role for



**Fig. 3.** Time course of ERK and p38 activation after T2 exposure. Total cell lysates were analyzed by Western blotting by use of antibodies against phospho ERK1/ERK2 or p38. Representative Western blots and a summary of the densitometry data are shown ( $n=3$ ). **(A)** Peak activation of ERK is seen after 15 min of exposure; **(B)** T2 does not induce p38 activation. \* $P<0.05$ .

TSP-1-induced VSMC chemotaxis was demonstrated by a number of different integrally related cell signaling pathways, including protein tyrosine kinases, focal adhesion kinase, ERK, p38, p21Ras, phosphatidylinositol 3-kinase, and protein kinase C [Patel et al., 1997; Wang et al., 1999; Willis et al., 1999, 2000]. One set of pathways focused on in the study of TSP-1-induced VSMC chemotaxis, are the mitogen-activated protein kinase (MAPK) pathways. MAPK, a family of serine/threonine protein kinases, consists of three main pathways: ERK, p38, and c-jun-N-terminal kinase (JNK) [Robinson and Cobb, 1997]. The exact mechanism by which these pathways affect cell chemotaxis is not fully understood. Actin, myosin, and microtubules are involved in cytoskeleton remodeling and cell migration. Although less is known about remodeling of



**Fig. 4.** **A:** T3- and COOH-induced VSMC chemotaxis is dependent on ERK and p38. A significant increase in VSMC chemotaxis was observed in response to T3 and COOH versus SFM. Preincubation of VSMCs with PD98059 (10  $\mu$ M) or SB202190 (10  $\mu$ M) inhibited T3- and COOH-induced chemotaxis. \* $P<0.05$  ( $n=3$ ); **(B)** T2 domain did not induce VSMC chemotaxis. No difference in VSMC chemotaxis was observed in response to the T2 domain versus SFM and the GST control fusion protein.

smooth muscle cell myosin and actin, other studies showed that myosin light chain kinase, which phosphorylates and reorganizes cytoskeletal components that facilitate cell movement, also is a substrate for MAPK [Klemke et al., 1997].

Both ERK and p38 have been associated with VSMC migration, as well as cell proliferation, apoptosis, differentiation, cytoskeletal remodeling, and cell cycle function [Bennett and Tonks, 1997; Graf et al., 1997; Hedges et al., 1999]. Since specific inhibitors of ERK (PD98059) and p38 (SB202190) cascades were first described, they have been widely used to investigate their involvement in intracellular signal transduction pathways. PD98059 is a specific inhibitor of the mammalian MEK1/2, which is just upstream of ERK [Dudley et al., 1995]. SB202190 selectively inhibits p38  $\alpha$ -,  $\beta$ -isoforms, but not  $\gamma$ -,  $\delta$ -isoforms [Kumar et al., 1997]. The authors showed previously that TSP-1-activated ERK, and PD98059 inhibited TSP-1-induced VSMC chemotaxis [Gahtan et al., 1999]. Similarly, p38 was activated in TSP-1-stimulated VSMCs, and the p38 inhibitor, SB202190, inhibited

TSP-1-induced VSMC chemotaxis [Wang et al., 1999]. These inhibitors were specific in their ability to selectively block ERK or p38 as demonstrated in our control experiments for ERK and p38 activation (Fig. 2C).

Through the use of fusion proteins to each of the domains of TSP-1, we have shown that T3 and COOH domains are involved in VSMC chemotaxis [Nesselroth et al., 2001]. The COOH domain contains a binding site for the integrin-associated protein (CD47) that is associated with cell attachment, platelet aggregation, cell migration, and VSMC chemotaxis [Klemke et al., 1997]. One previous study shows that a peptide agonist to CD47, 4N1K, independently induced VSMC chemotaxis [Guo et al., 1998]. Consistent with this, we noted that anti-CD47 antibodies inhibited TSP-1-induced VSMC chemotaxis (unpublished data). The authors demonstrated previously that a fusion protein to the T3 domain also independently induced VSMC chemotaxis [Nesselroth et al., 2001]. In addition, TSP-1-induced VSMC chemotaxis can be blocked by RGD (the major binding site for T3)-containing peptides [Patel et al., 1997]. The activity of the RGD containing peptide implies involvement of the  $\alpha_v\beta_3$  integrin, which is associated with development of arterial lesions, including intimal hyperplasia. Since  $\alpha_v\beta_3$  also associates with CD47, it is possible with  $\alpha_v\beta_3$  concomitantly binds to the RGD site on the T3 domain and the CD47 on the COOH domain to stimulate VSMC chemotaxis.

Since the COOH and T3 domains are the primary sites mediating induction of TSP-1-induced VSMC chemotaxis [Nesselroth et al., 2001], we sought to determine whether the COOH and T3 domains independently activated the same cell-signaling pathways as the whole TSP-1 protein. The findings of the current study confirmed that COOH and T3 domains independently activated ERK and p38 in VSMCs. Complete inhibition of either p38 or ERK was able to inhibit VSMC chemotaxis. These findings suggest when complete inhibition of one pathway is present, activation of the other pathway may not be sufficient to induce chemotaxis, or that cross-talk between ERK and p38 is essential. Previous studies have addressed cross-talk between ERK and p38 pathways in a specific and nonspecific manner [Singh et al., 1999; Birkenkamp et al., 2000]. The findings of this study suggest that p38 may be more specific for chemotaxis, as there was

p38 activation by COOH and T3, but not T2. In contrast, each of the three fusion proteins induced a similar degree of ERK activation. In addition, a more sustained degree of p38 activation by COOH in comparison to T3 occurred (see Figs. 1B and 2B). This observation also correlated closely with VSMC chemotaxis, in which chemotaxis was abolished after SB202190 exposure (see Fig. 4A). After inhibition with PD98059, a small degree of VSMC chemotaxis remained, which was significant.

The N-terminal domain of TSP-1 is essential to support attachment of various cultured cells including VSMCs and endothelial cells [Asch et al., 1991]. Others reported that the N-terminal A2.5 binding domain of TSP contains an important binding site for sulfated glycoconjugates and appears to have an important role in tumor cell chemotaxis [Taraboletti et al., 1987]. Our previous study showed that the NH2 domain also is involved in VSMC chemotaxis, but the degree of chemotaxis is much less than that observed for COOH [Nesselroth et al., 2001]. The cell signaling pathways involved in the induction of chemotaxis by NH2 is an area for future study.

In conclusion, this study indicates activation of the ERK and p38 pathways by T3, COOH, or both, and regulates VSMC chemotaxis. Our findings further suggest activation of the p38 pathway may be more specific than ERK for COOH- and T3-induced VSMC chemotaxis. Selective inhibition of T3 or the COOH domain of TSP-1 may be a helpful adjunct in prevention of VSMC chemotaxis, a process associated with development of arterial lesions.

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