A Role for Syndecan-1 in Coupling Fascin Spike Formation by Thrombospondin-1

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Abstract. An important role of cell matrix adhesion receptors is to mediate transmembrane coupling between extracellular matrix attachment, actin reorganization, and cell spreading. Thrombospondin (TSP)-1 is a modulatory component of matrix expressed during development, immune response, or wound repair. Cell adhesion to TSP-1 involves formation of biochemically distinct matrix contacts based on stable fascin spikes. The cell surface adhesion receptors required have not been identified. We report here that antibody clustering of syndecan-1 proteoglycan specifically transduces organization of cortical actin and fascin bundles in several cell types. Transfection of COS-7 cells with syndecan-1 is sufficient to stimulate cell spreading, fascin spike assembly, and extensive protrusive lateral ruffling on TSP-1 or on syndecan-1 antibody. The underlying molecular mechanism depends on glycosaminoglycan (GAG) modification of the syndecan-1 core protein at residues S45 or S47 for cell membrane spreading and on the VC2 region of the cytoplasmic domain for spreading and fascin spike formation. Expression of the VC2 deletion mutant or GAG-negative syndecan-1 showed that syndecan-1 is necessary in spreading and fascin spike formation by C2C12 cells on TSP-1. These results establish a novel role for syndecan-1 protein in coupling a physiological matrix ligand to formation of a specific matrix contact structure.

Key words: cell adhesion • extracellular matrix • proteoglycan • actin • protrusions

Introduction

The integrative effects of extracellular matrix on cell function depend on linkage from the matrix across the plasma membrane to the actin cytoskeleton. Such transmembrane coupling is mediated by numerous cell surface adhesion receptors which have specific matrix ligands (for review see Hynes, 1999). The interactions support cell matrix adhesion and motility and also initiate signaling responses which regulate many aspects of cell behavior (for reviews see Clark and Brugge, 1995; Schwartz et al., 1995; Yamada and Miyamoto, 1995). Thrombospondin (TSP)-1 is a multifunctional glycoprotein component of extracellular matrix that is widely expressed during organogenesis and which is locally upregulated in adult tissues upon platelet activation, inflammatory response, or wound healing. Within the extracellular matrix, TSP-1 binds to other matrix glycoproteins, collagen V, perlecain, certain proteases, or cytokines and is associated with cell surfaces in nonfibrillar patches (for reviews see Bornstein, 1995; Adams, 1997a). TSP-1 functions as a cell adhesion molecule and also regulates cell adhesion to other matrix components (Murphy-Ullrich and Hook, 1989). These properties have led TSP-1 to be considered with tenascins and SPARC as a regulatory component of extracellular matrix (for review see Chiquet-Ehrismann, 1995). To understand the functions of these molecules in different tissue contexts, it is important to establish their cellular mechanisms of action.

This laboratory has shown that cell attachment and spreading on TSP-1 substrata is associated with organization of the cortical actin cytoskeleton to form stable, radial spikes which contain F-actin and the actin-binding protein fascin (Adams, 1995, 1997b). These spike structures and associated lamellae mediate adhesion to TSP-1 and are also needed in cell motile behavior on TSP-1 (Adams, 1997b). Fascin spikes are assembled in the absence of focal contacts, and their assembly is dependent on the maintenance of a pool of the nonphosphorylated, actin-binding form of fascin and the regulated activities of Rac and Cdc42 small GTPases (Adams, 1995; Adams et al., 1999; Adams and Schwartz, 2000). Thus, the interactions of TSP-1 at the plasma membrane transduce signals into cells.
which have highly specific effects on the organization of F-actin and fascin within the cell cortex. To understand the molecular basis for this response, it is of obvious relevance to identify the necessary adhesive receptors.

Each subunit of the TSP-1 homotrimer contains four major domains that are involved in cell attachment to TSP-1: the NH₂-terminal domain, the type 1 repeats, an arginine-glycine-aspartate motif in the last type 3 repeat, and the COOH-terminal globular domain (for review see Adams 1997a). Widespread analyses of many cell types have established that each domain interacts with different cell surface–binding partners. The NH₂-terminal domain has a high affinity for heparin and heparan sulfate (HS)-glycosaminoglycans (GAGs) and can mediate endocytosis of TSP-1 in conjunction with low density lipoprotein receptor–related protein (Sun et al., 1989; Godyna et al., 1995; Mikhailenko et al., 1995). The type 1 repeats contain multiple binding sites for GAGs and a cell type–restricted receptor, CD36 (Asch et al., 1992; Guo et al., 1992a,b; Pancake et al., 1992; Li et al., 1993; Gantt et al., 1997). The arginine-glycine-aspartate site, which is active in a conformation-dependent manner according to the nature of the disulfide pairings within the type 3 repeats, binds αvβ3 integrin on endothelial and smooth muscle cells and to some extent αIβ3β3 on platelets (Lawler et al., 1988; Lawler and Hynes, 1989; Sun et al., 1992). Cell attachment involving other integrins has been reported in individual cell types, for example for α4β1 and α5β1 integrins in activation of adherent T cells, but the binding sites for these interactions have not been mapped precisely (Yabkowitz et al., 1995). Activated αvβ3 integrin binds a site within the NH₂-terminal domain (Krutzsch et al., 1999). Two peptide motifs in the COOH-terminal globular domain act as binding sites for CD47/IAP (Gao et al., 1996).

The multiplicity of cell-binding sites presented by the intact TSP-1 molecule complicates analysis and interpretation of the activities of individual adhesive receptors in transmembrane coupling and organization of the actin cytoskeleton. Yet a knowledge of the mechanism involved is critical in understanding how TSP coordinates the assembly of fascin spikes in preference to other types of matrix contacts. Fascin spikes are also formed by cells adherent to Engelbreth-Holm-Swarm laminin and tenasin C and are localized at the leading edge of migratory cells where they have been functionally implicated in motile behavior (Adams, 1997b; Fischer et al., 1997; Yamashiro-Matsumura et al., 1998). Thus, identification of the coupling mechanism for TSP-1 could prove of general significance in understanding how matrix contacts are coordinated by cells within complex, physiological extracellular matrices.

We hypothesized that ligation and clustering of individual adhesive receptors by specific antibodies could be used to identify roles for TSP-1–binding receptors in fascin spike assembly. We report here that, either by antibody clustering of syndecan-1 or by transfection of syndecan-1 expression constructs, ligation of syndecan-1 specifically coordinates the organization of fascin structures in several cell types. The molecular mechanism of these effects depends on the presence of GAG chains and on the cytoplasmic domain of the core protein. These novel results establish syndecan-1 as a functionally important transducer of fascin spike formation by TSP-1.

Materials and Methods

Cell Lines and Materials

C2C12 myoblastic cells were cultured in DME containing 20% FCS. Mouse embryo fibroblasts (MEF-1) and COS-7 green monkey kidney cells were obtained from American Type Culture Collection and cultured in DME containing 10% FCS. Human lung microvascular endothelial cells (HLMECs) were obtained from CLONTECH Laboratories, Inc. and cultured in the manufacturer’s endothelial growth medium supplemented with 10 ng/ml EGF, 1 μg/ml hydrocortisone, 3 μg/ml bovine brain extract, and 10% FCS (BioWhittaker). Antibodies reactive with mouse syndecan-1 (CD138; rat monoclonal antibody 281-2) (Jalkanen et al., 1985), mouse β1 integrin subunit (CD29; antibody HA2/5) (Kinashi and Springer, 1994), mouse α5 integrin subunit (CD49e; antibody 5H10-27), and mouse β3 integrin subunit (CD61; hamster antibody C9G2) were obtained from BD Pharmingen. N-18 goat antibody to the NH₂-terminus of mouse syndecan-1 was obtained from Santa Cruz Biotechnology, Inc. Rabbit antibody to CD36, reactive with human and mouse CD36, was from Research Diagnostics. Mouse monoclonal antibody 301 to mouse CD47/IAP and mouse hindlimb muscle from neonatal wild-type or CD47-null mice were gifts from Frederick Lindberg (University of Washington, St. Louis, MO) (Lindberg et al., 1996). Primary skeletal myoblasts were prepared from the dissected tissue by sequential trypsinization and replating, and their CD47 status was confirmed by immunostaining. Mouse monoclonal antibodies to human adhesion receptors included antibody SMO to CD36 (Serotec) (Hogg et al., 1984), BRIC 126, B6H12, and 2E11 to CD47 (Biodesign; AMS Biotechnology) (Avent et al., 1988; Brown et al., 1990), monoclonal antibody 13 to β1 integrin (Akiyama et al., 1989), L M609 to αvβ3 integrin (Cheresh, 1987), and antibodies 1D4 and B44 to syndecan-1 (Biodesign) (Dore et al., 1998). Purified nonimmune rat or mouse immunoglobulins were obtained from Sigma–Aldrich.

FACS® Analysis

Single cell suspensions of 10⁶ C2C12 or COS-7 cells were incubated with antibodies to receptor domains diluted in PBS on ice for 30 min. Cells were washed three times in PBS at 4°C, resuspended in 1:50 dilutions of the appropriate FITC–conjugated secondary antibody, and incubated for 30 min on ice. The cells were washed again and passed through a Becton Dickinson FACScan™. 10,000 cells were analyzed per sample.

Extraction of Proteoglycans

Cells were plated at 10³ cells per 90-mm dish and allowed to attach for 2 h. Cells were then lysed in 8 M urea, 0.2% Triton X-100, 10 mM Tris-HCl, pH 8.0, and sonicated on ice, using a Branson model 450 Sonifier in four 10-s bursts at amplitude setting 8 with 10 s cooling time allowed between each burst. The lysates were centrifuged to clarify, and the pellet corresponding to 10⁶ cells was combined with 100 μl packed volume of DEAE-Sepharose (Sigma–Aldrich) which had been prepared by washing in 8 M urea, 0.2% Triton X-100, 10 mM Tris-HCl, pH 8.0. The mixture was rotated end-over-end for 16 h at 4°C. The DEAE-Sepharose beads were washed in TBS and eluted with 1 M NaCl in lysate digestion buffer (100 mM Hepes, pH 6.5, 10 mM CaCl₂, 0.5% CHAPS, 0.2 mg/ml BSA). Proteoglycans were collected as described by Sanderson et al. (1992). Enzymatic cleavage of chondroitin sulfate (CS) and HS-GAG chains was performed by digesting the proteoglycan samples two times with dH₂O followed by incubation with either 50 μg/ml chondroitinase ABC, 50 μg/ml heparitinase II (Seikagaku), or a mixture of both enzymes for 2 h at 37°C. A second digestion of enzyme(s) was made and the reaction mixtures were incubated for an additional 2 h at 37°C. Samples were then concentrated and desalted by centrifugation in Centricon 10 concentrators (Amicon) and combined with SDS-PAGE sample buffer containing 100 mM DTT as a reducing agent. Electrophoresis of samples was performed after reducing conditions as described by Koda et al. (1985), using 4–15% SDS-polyacrylamide gradient gels and Tris/borate buffer containing 40 mM Tris, 60 mM borate acid, 0.8 mM EDTA, 1 mM Na₂SO₄, and 0.1% SDS. Samples were electrophoresed overnight at 60 V onto nitrocellulose (0.22-μm pore size; Bio-Rad Laboratories) and probed for syndecan-1 by incubation with 5 μg/ml monoclonal antibody 281.2.1 for 1 h, followed by ECL detection of alkaline phosphatase–conjugated secondary antibody (Tropix; PerkinElmer) using Hyperfilm ECL (Amersham Pharmacia Biotech).
Transfection and Cytoskeletal Organization Assays

Cells for transient transfection were plated at 30–40% confluency and transfected with plasmid DNA by use of Superfect reagent (QIAGEN) according to the manufacturer’s instructions. The plasmids used included expression plasmids for enhanced green fluorescent protein (EGFP) (CLONTECH Laboratories, Inc.), EGFP-fascin, wild-type and GAG-addition site mutants of mouse syndecan-1 (Saunders et al., 1989; Langford et al., 1998), and wild-type rat syndecan-2 (Klasa et al., 2000). Syndecan-1 mutants lacking the VC2 regions of the cytoplasmic domain were prepared by PCR-based mutagenesis using as a primer pair the forward primer 5′-TCTGGGAGCATGAGACG and the reverse primer 5′-CTGTAGCCACCCGATGAGCT and Pfu polymerase (New England Biolabs, Inc.). The PCR product was cloned into the pCDNA3 expression vector by TOPO cloning kit (Invitrogen). The 5′ and 3′ junctions of the cDNA were confirmed by DNA sequencing, using the dye-terminator chain-termination method and standard forward and reverse vector primers. Expression of cDNAs was confirmed by in vitro translation using the TNT kit (Promega).

Transiently transfected cells were used in adhesion assays 36–42 h after transfection. Glass coverslips were coated with 50 nM recombinant human TSP-1, prepared using the baculovirus system as described (Adams et al., 1998) or with the appropriate immunogoldogulins overnight at 4°C. To orient antireceptor antibodies, dishes were first coated with 10 μg/ml avidin (Vector Laboratories) and then with 10 μg/ml biotin-conjugated antiamoeglobulin antibodies of the appropriate species specificity (Vector Laboratories or ICN Biomedicals). Surfaces were then coated at 4°C overnight with 40 μg/ml solutions of the antireceptor antibodies. All coverslips were rinsed in TBS and blocked with 2 mg/ml heat-denatured BSA for 1 h. Single cell suspensions, prepared by release with 10 mM EDTA or by trypsinization, were allowed to attach for 1 h at 37°C in serum-free DME. Nonattached cells were removed by rinsing in TBS and the adherent cells were fixed before analysis of cytoskeletal organization by indirect immunofluorescence. For staining with rhodamine-phalloidin, antivinculin monoclonal VIN 11.5 (Sigma-Aldrich), or mouse monoclonal 4G10 to nophluorescence. For staining with rhodamine-phalloidin, antivinculin monoclonal VIN 11.5 (Sigma-Aldrich), or mouse monoclonal 4G10 to phosphotyrosine (Upstate Biotechnology) cells were fixed in 3.7% paraformaldehyde and processed as described previously (Adams, 1995). For staining with monoclonal antibody to fascin (Dako), cells were fixed in absolute methanol and fascin localization was detected with an FITC-conjugated goat anti-mouse secondary antibody (ICN Biomedicals). For quantitation of spread area or length of fascin actin bundles either digital images were processed in ImagePro Plus or tracings from photographic enlargements were scanned into Adobe Photoshop® and then processed. Excel worksheets were used to calculate descriptive statistics and to determine significance using an unpaired two-tailed t test.

Cell Motility Measurements

Nunc slideflasks were coated with antibodies to adhesion receptors as described above or with 50 nM recombinant TSP-1 and blocked with heat-denatured BSA. EDTA-released cells were suspended at 3 × 105 cells/ml and plated for 30 min. Nonadherent cells were removed and the behavior of the attached cells was tracked over the next 1.5 h by time-lapse videomicroscopy as described (Adams, 1997b). Cells were recorded at 10 frames/min and alterations in spread area, ruffling behavior, or cell locomotion were measured from traces. Ruffling behavior was scored as the extension of lateral, substratum-adherent ruffles at cell margins (lateral ruffles). Cell locomotion was scored as the displacement of the cell centroid over time. The extension of short, dynamic projections from the dorsal cell surface (apical projections) was also scored. At least 50 cells were traced for each transfection condition, and five replicate experiments were analyzed in total. For parameters that appeared altered between control or experimental transfections, statistical significance was determined in Microsoft Excel using a two-tailed t test.

Results

Expression of Adhesion Receptors for Thrombospondin-1 by C2C12 Myoblasts

C2C12 skeletal myoblasts attach to TSP-1 through predominant interactions with the type 1 repeats and COOH-terminal domain (Adams and Lawler, 1994). With the aim of identifying the adhesive receptors which couple cell attach-
staining only; Fig. 1 A). However, after trypsin treatment for 2 min at 37°C expression of syndecan-1 was reduced and more heterogeneous (Fig. 1 A, panel Syn-1[T]). As expected, because of the known protease sensitivity of the syndecan-1 extracellular domain, trypsinization for 5 min at 37°C resulted in a complete loss of cell surface syndecan-1 (data not shown; Fitzgerald et al., 2000).

The syndecan-1 core protein contains multiple sites for addition of HS- and CS-GAGs, and these posttranslational modifications are made with a high degree of cell type specificity (Kokenyesi and Bernfield, 1994; reviewed by Bernfield et al., 1999). To determine the nature of the GAG substitutions in C2C12 cells, we examined the apparent molecular mass of syndecan-1 after extraction of proteoglycans in combination with digestion with GAG lyases specific for either HS or CS. In control extracts, syndecan-1 resolved on immunoblots as bands of apparent molecular mass of ~120 and 97 kD with associated smearing (Fig. 1 B, lane 1). Upon digestion with chondroitinase ABC, the 120-kD band was much reduced, and the majority of syndecan-1 appeared as a broad band in the range of 66–97 kD that comigrated with core protein (Fig. 1 B, lanes 2 and 4). The breadth of this band likely relates to the tendency of syndecan-1 cores to form SDS-resistant oligomers as well as the presence of various HS substitutions. Upon digestion with heparitinase II, the 120-kD band was lost, and syndecan-1 appeared as an extended higher molecular weight smear as well as the ~97-kD species (Fig. 1 B, lane 3). Double digestion with both enzymes resulted in complete conversion of the diffuse higher molecular weight forms into the core protein band (Fig. 1 B, lane 4). Thus, in C2C12 cells, syndecan-1 is expressed as a mixed CS/HS proteoglycan.

**Direct Ligation of Individual TSP-1 Adhesion Receptors Has Distinct Effects on F-Actin Organization**

To dissect the effects of ligation and clustering of individual TSP-1 receptors on cell attachment and actin cytoskeletal organization, the panel of antibodies reactive with the extracellular domains of mouse β1 or β3 integrin subunits, CD47, or syndecan-1 were used as adhesive substrata for C2C12 cells. All the antibodies supported quantitatively

**Table I. Effects of Adhesion Receptor Antibodies on Fascin Distribution in C2C12 Cells**

| Antibody substratum                      | Input cells attached | Attached cells with fascin spikes | Attached cells with localized vinculin |
|------------------------------------------|----------------------|----------------------------------|----------------------------------------|
| Nonimmune IgG                            | 6 ± 3.8              | 0                                | 0                                      |
| Anti-β1 integrin                          | 81 ± 5.3             | 0                                | 78 ± 4.2                               |
| Anti-β3 integrin                          | 77 ± 6.6             | 0                                | 64 ± 6.5                               |
| Anti-CD47                                 | 58 ± 3.7             | 10 ± 5.8                         | 0                                      |
| Anti-syndecan-1, EDTA release             | 72 ± 6.3             | 88 ± 4.3                         | 0                                      |
| Anti-syndecan-1, 2 min treatment with trypsin | 56 ± 4.9             | 68 ± 3.2                         | 0                                      |
| Anti-syndecan-1, 5 min treatment with trypsin | 11 ± 2.3             | 0                                | 0                                      |

Cells were plated for 1 h in serum-free medium on antibody-coated surfaces, then fixed and stained for fascin or vinculin. Results are given as mean ± SEM from five experiments.
increased cell attachment relative to nonimmune rat IgG. C2C12 underwent partial spreading on the β1 integrin subunit antibody and displayed organization of F-actin at cell margins (Fig. 2 and Table I). Cells on the β3 integrin subunit antibody also spread partially and developed concentrations of F-actin at points on the cell surface (Fig. 2). On the CD47 antibody, cells spread more extensively and displayed larger actin-containing ruffles at cell margins (Fig. 2). For all these antibodies, similar results were obtained using either trypsinized or EDTA-disaggregated cells (shown in Fig. 2 for trypsinized cells only).

In contrast, EDTA-disaggregated C2C12 cells spread fully on 281-2 antibody to syndecan-1 and displayed zones of F-actin ruffles and small projections at cell margins (Fig. 2). Cells trypsinized for 2 min, many of which retained cell surface syndecan-1 (Fig. 1 A), showed a less uniform response (Table I and Fig. 3). Extensively trypsinized cells which lacked syndecan-1 attached at only background levels to the antibody and did not spread (data not shown). To establish that these responses were not due to differing affinities or avidities of individual antibody/antigen binding interactions, F-actin organization was examined on surfaces coated with different concentrations of the antibodies to β1 integrin, β3 integrin, CD47, or syndecan-1. The cells spread poorly on any antibody at 5 μg/ml coating concentration. Actin spikes and ruffles were organized by EDTA-dissociated cells on the 281-2 antibody in the concentration range 30–100 μg/ml and were not organized by cells adherent to the β1 or β3 antibodies even when they were coated at 100 μg/ml. Cells adherent on the CD47 antibody formed small F-actin ruffles on surfaces coated with ≥50 μg/ml of the antibody (data not shown).
Effects of Antibody Ligation of Adhesion Receptors on the Distribution of Fascin and other Actin-associated Proteins

To establish whether F-actin organization by antibody ligation of any of the adhesion receptors resulted in localized actin bundling by fascin, the antibody-adherent C2C12 cells were stained for fascin. In the very few cells which attached to rat IgG, fascin was diffuse (not shown). On the β1 or β3 integrin antibodies, fascin was also diffuse and the cell margins did not show fascin projections (Fig. 3 A). On the CD47 antibody, fascin appeared diffuse, and in 90% of the cells staining was uniform throughout the irregular, actin-rich marginal regions of the cells. The occasional cell showed small fascin-containing structures (one such cell is shown in Fig. 3). In striking contrast, fascin localized to radial spike structures in the cortex of EDTA-released cells on the syndecan-1 antibody (Fig. 3). C2C12 cells trypsinized for 2 min were more heterogeneous in spreading and spike formation than the EDTA-treated cells, and ~68% of the cells showed fascin spikes or less well-organized fascin-containing projections (Fig. 3). Quantification of these results confirmed that the syndecan-1 antibody strongly and specifically evoked the formation of fascin spikes (Table I). Because antibodies to different epitopes on CD47 vary in their ability to perturb CD47-mediated functions (Brown et al., 1990), we further examined the role of CD47 in fascin spike formation by comparing the ability of wild-type and CD47-null primary skeletal myoblasts to form fascin spikes on TSP-1. The two cell populations were indistinguishable with regard to adhesion and the organization of fascin spikes (Fig. 3 B).

We extended the analysis of receptor ligation by antibodies to additional cell types which show different degrees of spreading on TSP-1 substrata. MEF-1 mouse fi-
broblasts spread and form fascin spikes and filopodia on TSP-1 (Fig. 4 A). Cells remained rounded with diffuse fascin when plated on the anti–mouse β1, β3, or CD47 antibodies (shown for β1 only, Fig. 4 A). When plated on the 281-2 antibody to mouse syndecan-1, MEF-1 formed broad circumferential lamellae containing radial F-actin ribs which corresponded to sites where fascin and actin were bundled (Fig. 4 A). HLMECs express fascin and syndecan-1 but as reported for endothelial cells from other blood vessel sources, do not spread on TSP-1 (Fig. 4 B; Lawler et al., 1988; Murphy-Ullrich and Hook, 1989; Mertens et al., 1992; Adams, J.C., unpublished data). The cells show specific enrichment of F-actin and fascin structures at cell margins when attached to TSP-1 (Fig. 4 B). HLMECs attached to surfaces coated with antibodies to human β1, β3, CD47, or syndecan-1 and showed most microfilament organization on the β1 antibody. However, fascin was completely diffuse in these cells and was not enriched at cell margins (Fig. 4 B; data not shown). On either of two antibodies to human syndecan-1, HLMECs formed short marginal structures which contained F-actin and fascin (shown for BB4 antibody only, Fig. 4 B). We also examined fascin localization in HLMECs plated on surfaces coated with SMO antibody to human CD36 (Hogg et al., 1984). The cells remained rounded and smooth edged and did not show specific localization of fascin (data not shown). Neither the BRIC 126, B6H12, nor 2E11 antibodies to human CD47 caused fascin–actin bundling (Fig. 4 B; data not shown). Thus, in several cell types from different tissue sources, ligation of syndecan-1 specifically promoted the organization of cortical fascin independently of the degree of cell spreading.

To further evaluate effects on the actin-based cytoskeletal organization and compare these with the effects of...
TSP-1, we stained C2C12 cells on antibody substrata for components of focal contacts. Whereas cells on the β1 or β3 antibodies assembled vinculin into small, focal contact-like structures (shown for the β1 antibody only, Fig. 3), vinculin was diffuse in the cells adherent on the CD47 or syndecan-1 antibodies. We also found that cells adherent on the syndecan-1 antibody showed very low and diffuse staining for phosphotyrosine (data not shown). Thus, antibody ligation of syndecan-1 specifically induces the organization of actin and fascin bundles in the absence of focal contact assembly. Furthermore, these results demonstrate that the antiintegrin antibodies used in these experiments had appropriate integrin-activating activities in that they induced cytoskeletal coupling resulting in the focal localization of vinculin (Table I).

Syndecan-1 Is Sufficient for Fascin Spike Formation

To establish whether syndecan-1 is sufficient to mediate assembly of fascin and actin-containing spikes, we examined the effects of transient expression of mouse syndecan-1 in a heterologous cell system, green monkey COS-7 cells. COS-7 cells contain appropriate endogenous coupling mechanisms for spike formation because they express fascin and undergo limited spike formation when attached to TSP-1 (Adams, 1997b). FACS® analysis established that the 281-2 antibody to mouse syndecan-1 did not stain the surface of live COS-7 cells. COS-7 cells did not attach over background to surfaces coated with syndecan-1 antibody (data not shown). COS-7 cells transiently cotransfected with control, empty expression plasmid, and EFGP-fascin or expression plasmid encoding syndecan-1

Figure 6. GAG modifications of syndecan-1 and regions of the cytoplasmic domain are required for fascin spike formation on TSP-1. COS-7 cells were transfected with the indicated syndecan-1 mutants and EGFP-fascin expression plasmids, and 48 h later EDTA-released cells were plated on 50 nM TSP-1 in serum-free medium for 1 h and then fixed and stained for F-actin. Results shown are representative of three experiments, and ≈100 transfected cells were scored per experiment. Bar, 16 μm.
spreading involved a threefold increase in cell area (statistically significant at $P = 0.0001$; see Fig. 7 A). To examine the specificity of this effect, the syndecan-1 transfectants were plated on another matrix ligand, fibronectin. No increase in cell area or alteration in major F-actin structures was observed (Fig. 5; data not shown). Thus, the effects of syndecan-1 depend on its ligation by specific ligands and are not simply a consequence of the surface expression of syndecan-1.

**GAG Substitutions and the Syndecan-1 Cytoplasmic Domain Have Distinct Roles in the Mechanism of Syndecan-1-mediated Cell Spreading and Fascin Spike Organization on Thrombospondin-1**

The extracellular domain of syndecan-1 proteoglycan contains three conserved sites for GAG substitution near the NH$_2$ terminus. These are sites of addition for HS-GAGs in many cell types but can also be substituted by CS-GAGs (Kokenyesi and Bernfield, 1994). HS-GAG chains have well-established functional roles in the activities of syndecans, particularly in binding growth factors such as FGF (for reviews see Rapraeger and Ott, 1998; Bernfield et al., 1999). To determine whether GAGs also contribute to cytoskeletal coupling by syndecan-1, COS-7 cells were transiently cotransfected with EGFP-fascin and a syndecan-1 expression construct in which the three sites of GAG addition, serines 37, 45, and 47, had been mutated to alanines, syn-1/TGM (Langford et al., 1998). Strikingly, these cells spread poorly on TSP-1 compared with the wild-type syndecan-1 transfectants, and lamellipodia were not observed. Instead, the cells formed actin- and fascin-containing projections that appeared long in comparison to the wild-type syndecan-1 transfectants (Fig. 6). These parameters were quantitated by image analysis measurements. The mean cell area of syn-1/TGM transfectants was not significantly increased compared with vector control transfectants. However, the number of spikes per cell and their length were strongly increased (significant at $P = 0.001$; Fig. 7, A and B). Compared with cells expressing wild-type syndecan-1, the number of spikes per cell was somewhat increased (significant at $P = 0.04$), and the mean length of spikes was increased from 8.5 to 13.5 $\mu$m (significant at $P = 0.01$; Fig. 7 B).

COS-7 cells were next transfected with constructs in which the GAG addition sites were individually mutated. Expression of the syn-1/ S37A mutant caused increased cell spreading and spike formation in a similar manner to wild-type syndecan-1 in cells adherent on TSP-1 (not shown). Expression of either the syn-1/S45A or the syn-1/ S47A constructs correlated with enhanced formation of elongated spikes in poorly spread cells, and thus resembled the effects of the syn-1/TGM mutant (Fig. 6). Thus, GAG substitution at either S45 or S47 was needed in lamellar spreading on TSP-1 but was not required for the organization of filopodial-like fascin-containing projections.

To establish whether the cytoplasmic domain of the syndecan-1 core protein was critical for the formation of fascin spikes, COS-7 cells were transfected with a cytoplasmic deletion mutant of syndecan-1. All syndecans have short, 34-residue cytoplasmic domains that contain highly conserved membrane-proximal and COOH-terminal sequences, the C1 and C2 regions, respectively, separated by variable (V) regions which are unique to each family member. The C1 region is important for membrane trans-
location of syndecan-1 (Miettinen et al., 1994). Expression of syndecan-1 in which the V and C2 regions of the cytoplasmic domain had been deleted, syn-1/ΔVC2, did not promote either fascin spikes or lamellar cell spreading on TSP-1 (Fig. 6). Expression of the equivalent deletion mutant lacking the three GAG substitution sites, syn-1/TGM/ΔVC2, had similar effects (Fig. 6).

We also compared the effects of syndecan-2, a syndecan family member that is expressed by many mesenchymal cells, on cytoskeletal organization by cells adherent to TSP-1. The extracellular domain of syndecan-2 contains three analogous GAG addition sites, but the polypeptide sequence of the extracellular domain and of the cytoplasmic V region are unrelated to that of syndecan-1 (for review see Bernfield et al., 1999). Expression of syndecan-2 in COS-7 cells stimulated cell spreading on TSP-1 by 50% (significant at P = 0.04). However, the spread cells were completely smooth edged and lacked fascin spikes or cortical lamellipodia (Fig. 6 and Fig. 7, A and B). Cumulatively, these results demonstrate that the syndecan-1 cytoplasmic domain in combination with effects from GAG chains are jointly needed in coupling extracellular TSP-1 to membrane spreading and fascin cytoskeletal organization.

To determine whether these effects were specific to TSP-1 ligation of syndecan-1, we examined the responses of the transfectants to plating on syndecan antibody or on
fibronectin. On the syndecan-1 antibody, the syn-1/TGM transfectants spread and organized fascin–actin bundles, whereas the syn-1/ΔVC2 transfectants remained round as they did on TSP-1 (Fig. 8 A). Thus, even upon direct, monospecific ligation of the core protein the V and C2 cytoplasmic regions are needed to transduce cytoskeletal organization. To further confirm that the effects of syndecan-1 expression cells resulted from specific ligation of syndecan-1 and not as an immediate consequence of syndecan overexpression, we examined cell behavior on fibronectin. No alterations in cell spreading or fascin distribution were apparent in cells transfected with syn-1/TGM, syn-1/ΔVC2, or syndecan-2 (Fig. 8 A). These cDNAs used were all translatable with equivalent efficiencies (Fig. 8 B), and both Syn1/TGM and Syn1/ΔVC2 proteins were expressed at comparable levels to the wild-type syndecan-1 as determined by FACS® analysis or immunoblot (Fig. 8 B; data not shown). Thus, the differing effects of the various syndecan-1 proteins on fascin organization most likely result from ligation of syndecan-1 by appropriate specific ligands.

To determine the functional role of endogenous syndecan-1 in TSP-1-initiated formation of fascin spikes in C2C12 cells, we first examined the effect of prolonged trypsinization on fascin spike organization by C2C12 cells on TSP-1. Whereas EDTA-released cells spread and formed regions of fascin spikes, <20% of cells trypsinized for 5 min attached, and these remained completely round (Fig. 9, A and B). These cells lack cell surface syndecan-1 (Table I). To examine the specific role of syndecan-1, the syn-1/ΔVC2 or syn-1/TGM constructs were transiently transfected into C2C12 cells. The syn-1/ΔVC2 molecule acted as an effective dominant negative, in that, transfected cells which attached to TSP-1 did not spread or form fascin spikes (Fig. 9 C). Cells expressing the syn-1/TGM construct were altered in morphology on TSP-1, yet formed fascin projections (Fig. 9 D). Thus, in C2C12 cells as in COS-7 cells, GAG substitution of syndecan-1 is important for cell spreading on TSP-1 but does not appear to be required in fascin spike organization.

Syndecan-1–mediated Cytoskeletal Coupling Stimulates Protrusive Cortical Membrane Ruffling

In addition to their role in cell adhesion to TSP-1, fascin spikes support cell motility on TSP-1 matrix (Adams, 1997a,b; Adams and Schwartz, 2000). We undertook time-lapse experiments to examine the motile behavior of the syndecan-1–transfected COS-7 cells on TSP-1 or on syndecan-1 antibody-coated surfaces. Cells were scored for two aspects of motility: protrusive lateral ruffling activity and locomotion over the substratum. When attached to TSP-1, few of the vector transfectant cells showed extension of spikes in contact with the substratum, there was little ruffling activity at lateral cell margins, and all cells were non-locomotory (Fig. 10). In marked contrast, the syndecan-1 transfectant cells showed greatly increased ruffling at their lateral margins (Fig. 10; significant at \( P > 0.001 \)). These lateral ruffles appeared as dynamic phase dark regions that protruded rapidly along the cell margin or moved centripetally across the spread lamellae. These were also apparent in syndecan-1 transfectants adherent on syndecan-1 antibody (data not shown). Cells expressing Syn1/TGM also showed highly significant increases in lateral ruffling (Fig. 10). In contrast, cells transfected with the syn-1/ΔVC2 construct did not show lateral ruffles and thus appeared similar to vector transfectants (data not shown). To examine the specificity of the ruffling response, we compared the motility of cells transfected with syndecan-2. These cells showed no lateral ruffling. Lateral ruffling thus specifically correlated with the formation of fascin spikes (Fig. 7 B and Fig. 10). The alterations in lateral ruffling occurred in the absence of any change in extension of apical projections, and thus related specifically to adhesion-dependent motility and not to a general change in membrane dynamics (Fig. 10). Interestingly, despite the large increases in cell spreading and lateral ruffling activity, syndecan-1 transfectants did not translocate over the substratum. For adhesion to TSP-1, this result was obtained using coating concentrations of 25 or 50 nM. For adhesion to the syndecan-1 antibody, this result was confirmed at a range of antibody-coating concentrations from 5 to 50 \( \mu \)g/ml. For all concentrations that supported cell spreading, no stimulation of cell locomotion was observed (data not shown). Thus, under these experimental conditions, the most dramatic effect of syndecan-1 ligation on motile behavior is the stimulation of active ruffling protrusions at lateral cell margins (Fig. 10).

Discussion

Our results demonstrate a novel role for syndecan-1 in coupling the organization of fascin spikes in response to a physiological extracellular ligand, thrombospondin-1. We show by several independent approaches that syndecan-1...
has a crucial role in this process. First, antibody ligation of syndecan specifically causes the formation of cortical actin- and fascin-containing bundles in several cell types. Second, overexpression of syndecan-1 in a heterologous cell type is sufficient to cause dramatic enhancement of cell spreading and the formation of fascin spikes in response to TSP-1. Third, treatments which deplete C2C12 cells of endogenous syndecan-1 inhibit cell spreading and fascin spike formation on TSP-1 substrata. Fourth, expression of a cytoplasmic domain deletion mutant of syndecan-1 had dominant negative effects on cell spreading on TSP-1. With regard to the molecular mechanism of this process, we demonstrate that there are distinct requirements for GAG substitution and the COOH-terminal region of the cytoplasmic domain in transducing cell spreading and fascin spike formation.

Syndecan-1 and thrombospondin-1 are coexpressed in many tissues, including skeletal muscle, during development (Corless et al., 1992). TSP-1 is known as a heparin-binding protein, and each subunit of the trimer contains multiple GAG-binding motifs within the NH2-terminal domain and type 1 repeats (for review see Adams, 1997a). Dependent on the tissue source and structural characteristics of the GAGs, TSP-1 binds both HS- and CS-GAGs (Pancake et al., 1992; Herndon et al., 1999). GAG-dependent cell attachment or adhesion to TSP-1 has been reported in many cell types (for example, Roberts et al., 1987; Kaeberg et al., 1989; Adams and Lawler, 1993; Gantt et al., 1997; Wilson et al., 1999). Disruption of focal contacts by soluble TSP-1 and the endocytosis of TSP-1 by low density lipoprotein receptor–related protein are also heparin-inhibitable processes (Murphy-Ullrich and Hook, 1989; Godyna et al., 1995; Mikhailenko et al., 1995; Chen et al., 1996). However, the core proteins involved in GAG-mediated adhesion have remained unknown. By affinity chromatography, syndecan-1 was identified as the major thrombospondin-binding HS proteoglycan in mammalian epithelial cell extracts (Sun et al., 1989). Our data now provide evidence for a role of syndecan-1 in cell spreading and major cytoskeletal responses to TSP-1.

Fascin spikes identify a specific matrix contact structure formed during cell spreading on TSP-1 which has functional roles in cell adhesion and motility on matrix substrata (Adams, 1995, 1997b). Interestingly, fascin spikes are also formed upon cell adhesion to tenascin-C and laminin-1, both of which are also ligands for syndecan-1 (Adams, 1997b; Fischer et al., 1997; for review see Bernfield et al., 1999). Although syndecan-1 has a peripheral distribution in freshly spreading cells, it is not present in focal contacts and colocalizes with actin microfilaments when clustered and in long-term adherent cells (Rapraeger et al., 1986; Carey et al., 1994a). Antibody-mediated clustering of syndecan-1 also promotes actin reorganization in preadherent Schwann cells and Raji cells (Carey et al., 1994b; Lebakken and Rapraeger, 1996). Our results link the binding of syndecan-1 by a physiological extracellular matrix ligand, TSP-1, to a specific effect on the actin cytoskeleton, that is, the formation of fascin spikes and ribs in adherent lamellipodia. Furthermore, this effect of syndecan-1 ligation compared with the effects of ligation of CD36, CD47, β1, or β3 integrins adhesive receptors implies a central role for syndecan-1 in extracellular coordination of actin bundling by fascin.

Several cell types show quantitatively high attachment to TSP-1 but do not undergo spreading (for example, Asch et al., 1991; Adams and Lawler, 1993). This property does not show a simple correlation with recognition of a particular adhesive domain of TSP-1 (Adams and Lawler, 1993) nor does it correlate in all cell types with a lack of fascin expression (Adams, 1997a,b). We examined HLMECs as a model for a fascin-positive cell type that does not spread on TSP-1 yet is functionally responsive to TSP-1 (Good et al., 1990). Interestingly, these cells did not spread on anti-syndecan antibodies, although cortical enrichments of actin and fascin were apparent on both TSP-1– or antibody-coated surfaces. Several possible mechanisms could underlie this effect. A certain level of syndecan-1 expression could be necessary to promote cell spreading and cytoskeletal organization on TSP-1. However, although the microvascular endothelial cells express lower levels of syndecan-1 than do epidermal keratinocytes, as determined by FACS analysis, epidermal keratinocytes also do not spread on TSP-1 (Adams and Lawler, 1993; Adams, J.C., unpublished data). In C2C12 cells, overexpression of either dominant negative or constitutively active forms of Cdc42 or Rac small GTPases correlates with cell rounding on TSP-1 (Adams and Schwartz, 2000). An interesting speculation is that the constitutive lack of spreading of certain cell types on TSP-1 could result from differences in the sizes of the active pools of these signaling mediators.

Syndecans have principally been characterized as coreceptors with FGF receptors or integrin-adhesive receptors (for reviews see Bernfield et al., 1999; Woods and Couchman, 2000). In particular, syndecan-4 binds the major heparin-binding site of fibronectin and acts coordinately with integrins such as α5β1 in the assembly of focal adhesions (Woods et al., 1993; Couchman and Woods, 1999; Longley et al., 1999; Saoncella et al., 1999). Ligation of α5β1 integrin leads to activation of protein kinase C. By binding and activating protein kinase C, syndecan-4 appears to potentiate this signal for focal adhesion assembly (Vuori and Rosslati, 1993; Oh et al., 1997a,b). Fibronectin supports only limited and transient formation of fascin spikes and ruffles during the initial stages of cell spreading because α5β1-dependent activation of protein kinase C promotes
phosphorylation of fascin and thereby downregulates its actin-binding activity (Adams, 1995; Adams et al., 1999). Thus, the molecular process we have defined for syndecan-1 as a transducer of fascin spike formation appears distinct from the mechanisms by which syndecan-4 participates in focal contact assembly.

In examining the mechanism by which syndecan-1 mediates this effect on fascin, we have uncovered a dualistic activity of syndecan-1. Whereas the effect of direct antibody ligation of the syndecan-1 extracellular domain or transfection of wild-type syndecan-1 stimulated both lamellipodial cell spreading and associated formation of fascin- and actin-containing spikes and ribs, GAG modifications at specific sites on the extracellular domain were needed to support lamellipodial spreading on TSP-1. Cells expressing GAG-negative syndecan-1 or loss-of-function mutants at the S45 or S47 attachment sites showed florid formation of fascin spikes on TSP-1 and poor membrane spreading. This effect was separable from a necessity for the VC2 portion of the cytoplasmic domain in both fascin spike formation and cell spreading.

As a model to explain these results, we propose two separate activities of the extracellular domain: (a) an interaction of TSP-1 with the core protein that is transduced through the cytoplasmic domain, which principally couples spike formation and is permissive for cell spreading; and (b) a GAG-mediated receptor-clustering effect which acts to promote cell membrane spreading. Although a protein–protein interaction of syndecan-1 with TSP-1 has not been reported, a separate role for the core protein would not be entirely unprecedented within the syndecan family. A region in the extracellular domain of syndecan-1 is required for cytoskeleton association but not detergent insolubility (Lebakken and Rapraeger, 1996; Zimmermann and David, 1999). The extracellular domain of syndecan-4 presented as a recombinant, monomeric, uncleaved protein specifically supports fibroblast attachment and binds cell surfaces with nanomolar affinity (McFall and Rapraeger, 1997, 1998). There are several reports of effects on syndecan-1 on cell morphology which do not require the cytoplasmic domain, and thus a separable role of the GAG chains appears plausible (Lebakken and Rapraeger, 1996; Liu et al., 1998). Effects on F-actin distribution that depend on the cytoplasmic domain have also been reported (Carey et al., 1996).

Members of the PDZ family of proteins are known binding partners for the C2 COOH-terminal region of syndecans (Grootjans et al., 1997; for reviews see Rapraeger and Ott, 1998; Bernfield et al., 1999; Zimmerman and David, 1999). However, transfection of syndecan-2, which has similarly positioned GAG attachment sites and shares the common C1 and C2 cytoplasmic regions with syndecan-1, promoted cell spreading but did not lead to organization of fascin spikes on TSP-1. The V region of syndecan-4 uniquely binds and activates protein kinase C (Oh et al., 1997a; for reviews see Rapraeger and Ott, 1998; Woods and Couchman, 1998). The V regions of other syndecans might have other unique binding partners. Our results define a novel and specific requirement for syndecan-1 in coupling between TSP-1 and fascin spike formation. Further studies will be directed towards defining the molecular interactions responsible for the recruitment of fascin and actin into these cortical structures.

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