

Enteropathogenic *Escherichia coli* activates the RhoA signaling pathway via the stimulation of GEF-H1

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Enteropathogenic *Escherichia coli* delivers a subset of effectors into host cells via a type III secretion system, and this step is required for the progression of disease. Here, we show that the type III effectors, EspG and its homolog Orf3, trigger actin stress fiber formation and the destruction of the microtubule networks beneath adherent bacteria. Both effectors were shown to possess the ability to interact with tubulins, and to stimulate microtubule destabilization *in vitro*. A recent study showed that microtubule-bound GEF-H1, a RhoA-specific guanine nucleotide exchange factor, was converted to its active form by microtubule destabilization, and this sequence of events resulted in RhoA stimulation. Indeed, EspG- and Orf3-induced stress fiber formation was inhibited by the expression of dominant-negative forms of GEF-H1 and RhoA, but not of Rac1 and Cdc42, and by treatment with a ROCK inhibitor. These results indicate that the impact of EspG/Orf3 on microtubule networks triggers the activation of the RhoA–ROCK signaling pathway via GEF-H1 activity. This report reveals for the first time that a pathogen can exploit the host factor GEF-H1.

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Introduction

Enteropathogenic *Escherichia coli* (EPEC) remains a major cause of infantile diarrhea worldwide, mostly in the developing world (Donnenberg and Kaper, 1992). Related pathogens, which cause disease by similar mechanisms (DeVinney *et al*, 1999), include enterohemorrhagic *E. coli* O157:H7,

rabbit EPEC, and *Citrobacter rodentium*, which is found in mice. These pathogens induce a characteristic histopathological lesion, referred to as an attaching/effacing (A/E) lesion (Moon *et al*, 1983), which is defined by the intimate attachment of bacteria to the intestinal epithelial surface and the effacement of host cell microvilli, which is involved in the development of diarrhea (Abe *et al*, 1998). Factors responsible for A/E lesion formation are encoded by a 35-kilobase pair region referred to as the locus of enterocyte effacement (LEE) (McDaniel *et al*, 1995), which includes the type III secretion system, the translocated intimin receptor (Tir) (Kenny *et al*, 1997), intimin (Donnenberg *et al*, 1993b), and EPEC-secreted proteins (Esp). We demonstrated the supermolecular structure of the type III secretion system (Sekiya *et al*, 2001), which includes an EscF-needle portion and an EspA-sheath-like structure at the tip of the needle. EscN, a cytoplasmic ATPase, is essential for the secretion of effectors (Jarvis *et al*, 1995), and the presumed pore-forming factors EspB and EspD are delivered to the host plasma membrane, and these factors are a prerequisite for the translocation of effectors into host cells (Ide *et al*, 2001). To date, five effectors, which are translocated into host cells via the type III secretion system, have been identified and characterized in EPEC; these effectors include Tir (Kenny *et al*, 1997), EspF (McNamara *et al*, 2001), EspG (Elliott *et al*, 2001), EspH (Tu *et al*, 2003), and Map (Kenny *et al*, 2002). In addition, the EspG homolog Orf3, which is encoded on the EspC pathogenicity islet within the EPEC genome (distinct from the LEE), is thought to be an effector (Elliott *et al*, 2001). Both EspG and Orf3 effectors share homology with the *Shigella* VirA effector. Although it has been demonstrated that *Shigella* VirA promotes the destabilization of host microtubules and Rac1 activation (Yoshida *et al*, 2002), the contributions of EspG/Orf3 effectors to EPEC virulence remain unclear. Map induces the transient formation of Cdc42-dependent filopodia, while EspH represses filopodium formation and enhances the actin-rich pedestal formation beneath adherent bacteria. A/E lesion formation is mediated by a Tir–intimin interaction (Kenny *et al*, 1997), which is required for the initiation of the disease process (Abe *et al*, 1998). However, an *in vivo* study with adult human volunteers suggested that additional EPEC factor(s) trigger diarrhea, since diarrhea still developed in four of 11 volunteers who ingested an intimin mutant strain that was incapable of A/E lesion formation (Donnenberg *et al*, 1993a). It has been reported that EPEC-induced diarrhea may be involved in a loss of intestinal barrier function, with ion and fluid movement from the intestinal submucosa into the lumen (Hecht, 2001). When polarized epithelial cells were infected with EPEC, redistribution of the tight junction-associated protein occludin was observed, and this redistribution was correlated with the disruption of the tight junction barrier (Yuhan *et al*, 1997; Simonovic *et al*, 2000). Although the precise function of EspF

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in host cells is not yet clear, EspF is known to be required for the disruption of epithelial cell junctions during EPEC infection, but EspF is not involved in the formation of the A/E lesion (McNamara *et al*, 2001). Thus, EPEC effectors have multiple functions in host cells, and their synergistic effects are thought to contribute to EPEC-mediated diarrhea.

In this study, we demonstrated that actin stress fiber formation was induced by EPEC infection in Swiss 3T3 cells. This effect was dependent on the type III secretion system, but not on Tir–intimin interactions. It has been reported that the formation of actin stress fibers is regulated by the RhoA–ROCK signaling pathway (Kimura *et al*, 1996; Leung *et al*, 1996; Hall, 1998). Although the Tir-induced accumulation of actin filaments has been observed during A/E lesion formation, this phenotype was not found to be affected by the addition of the RhoA inhibitor (Ben-Ami *et al*, 1998); moreover, this phenotype was required for the activity of N-WASP (Kalman *et al*, 1999), the Arp2/3 complex, and Nck (Gruenheid *et al*, 2001), indicating that EPEC may exploit different signal transduction pathways in the formation of A/E lesions and actin stress fibers.

To determine whether or not the type III effectors are involved in actin stress fiber formation, various effector mutants were constructed, and an infection assay was carried out. We found that EspG and Orf3 effectors are involved in the induction of actin stress fiber formation in a Tir–intimin interaction-independent manner. We further demonstrated that both effectors induce the disruption of microtubule networks beneath adherent bacteria by the direct association of EspG and Orf3 with tubulins. One of the aims of the present study was to determine how EspG/Orf3 governs the crosstalk between the actin cytoskeleton and microtubules. We revealed here that a newly identified guanine nucleotide exchange factor, GEF-H1 (Ren *et al*, 1998; Krendel *et al*, 2002), is exploited by EspG/Orf3 effectors to induce actin stress fiber formation during EPEC infection. A recent study also showed that endogenous GEF-H1 is localized in the microtubules in its inactive form and it switches to an active form when it is released from microtubules. Indeed, both EspG/Orf3 effectors were revealed to possess the ability to release GEF-H1 from the cytoskeleton to the cytosol as a result of the microtubule destabilization. We report here that the GEF-H1-mediated signaling pathway is triggered by EPEC infection via the activity of EspG/Orf3 effectors.

Results

EPEC infection of mammalian cells induces stress fiber formation

To investigate whether EPEC infection influences the host actin cytoskeleton, serum-starved Swiss 3T3 cells were infected with wild-type (WT) EPEC and the architecture of the actin networks was visualized by F-actin staining with rhodamine-labeled phalloidin. Although EPEC is a human pathogen, we used mouse Swiss 3T3 cells for this experiment, since the dynamics of the actin cytoskeleton are easily observed in these fibroblastic cells. EPEC has the ability to adhere to and form tight microcolonies on tissue culture cells and human intestinal cells (Rosenshine *et al*, 1992); these activities are mediated by a bundle-forming pilus (Giron *et al*, 1991). As expected, the accumulation of F-actin beneath microcolonies of adherent bacteria was also observed in the mouse fibro-

blastic cell line, Swiss 3T3 (Figure 1A, arrows). In addition to this accumulation, actin stress fibers with parallel bundles extending into projections of the cell periphery were prominently induced in the cytoplasm under conditions of EPEC infection (Figure 1A, arrowhead). Although the formation of a pedestal structure, represented by the accumulation of F-actin beneath the adherent bacteria, was not detected in all of the infected cells with the *eae* mutant (defective in intimin production) or the *tir* mutant (defective in Tir production), these cells still developed extensive actin stress fibers (Figure 1B and C). It was of note that stress fiber formation, observed here in EPEC WT-infected cells, was not detected with either the *espA* mutant (defective in the formation of the EspA-sheath in type III machinery) or the *escF* EPEC mutant (defective in the formation of the EscF-needle in type III machinery) (Figure 1B and C). The same was true for the *espB* and *espD* mutants (defective in pore formation on the host plasma membrane) (Figure 1B and C) and the *escN* mutant (defective in secretion of type III-secreted proteins) (Figure 2A and B). Therefore, the results of the present series of experiments strongly suggested that certain unknown type III-secreted effector(s) are involved in EPEC-induced actin stress fiber formation in infected cells.

EspG/Orf3 is involved in stress fiber formation in host cells infected with EPEC

To identify the secreted bacterial protein(s) involved in the formation of stress fibers in host cells, we constructed a series of nonpolar mutants with a defect in the genes *espF*, *orf19* (defective in Map production), *espG*, *espH*, or *orf3*, and we also considered an *espG/orf3* double-knockout mutant (due to the fact that EspG shares 43.5% identity with Orf3; Elliott *et al*, 2001). Swiss 3T3 cells were infected with each of the mutants in order to investigate their ability to elicit actin stress fiber formation. The results demonstrated that the ability of the *espG/orf3* double-knockout mutant to induce stress fiber formation was greatly reduced, in comparison with that of the other type III effector mutants, such as the *espF*, *orf19*, *espG*, *espH*, and *orf3* mutants (Figure 2A and B). Complementation of the *espG/orf3* double-knockout mutant with a plasmid encoding EspG (p99-EspG) or Orf3 (p99-Orf3) restored the ability to form actin stress fibers (Figure 2A and B), indicating that both proteins participated in the formation of stress fibers. To confirm this possibility, pEGFP-EspG or pEGFP-Orf3 (enhanced green fluorescent protein (EGFP)-tagged EspG or Orf3, respectively) was transfected into Swiss 3T3 cells and the cells were observed for the formation of stress fibers. The Swiss 3T3 cells overexpressing EGFP-EspG or EGFP-Orf3, but not EGFP alone, were able to induce actin stress fibers (Figure 2C and D), thereby suggesting the following conclusions: (i) EspG and Orf3 are responsible for the formation of stress fibers, and they possess functional redundancy, and (ii) other effectors are not involved in this series of events.

Although EspG is secreted via the type III machinery (Elliott *et al*, 2001), it remains unclear as to how Orf3 is secreted. To address this issue, the secretion profile of Orf3 was analyzed by immunoblotting using anti-Orf3 antibodies. We detected Orf3 in the culture supernatant prepared from the EPEC WT, but not from the type III machinery mutant, indicating that Orf3 is also secreted via the type III machinery (data not shown).

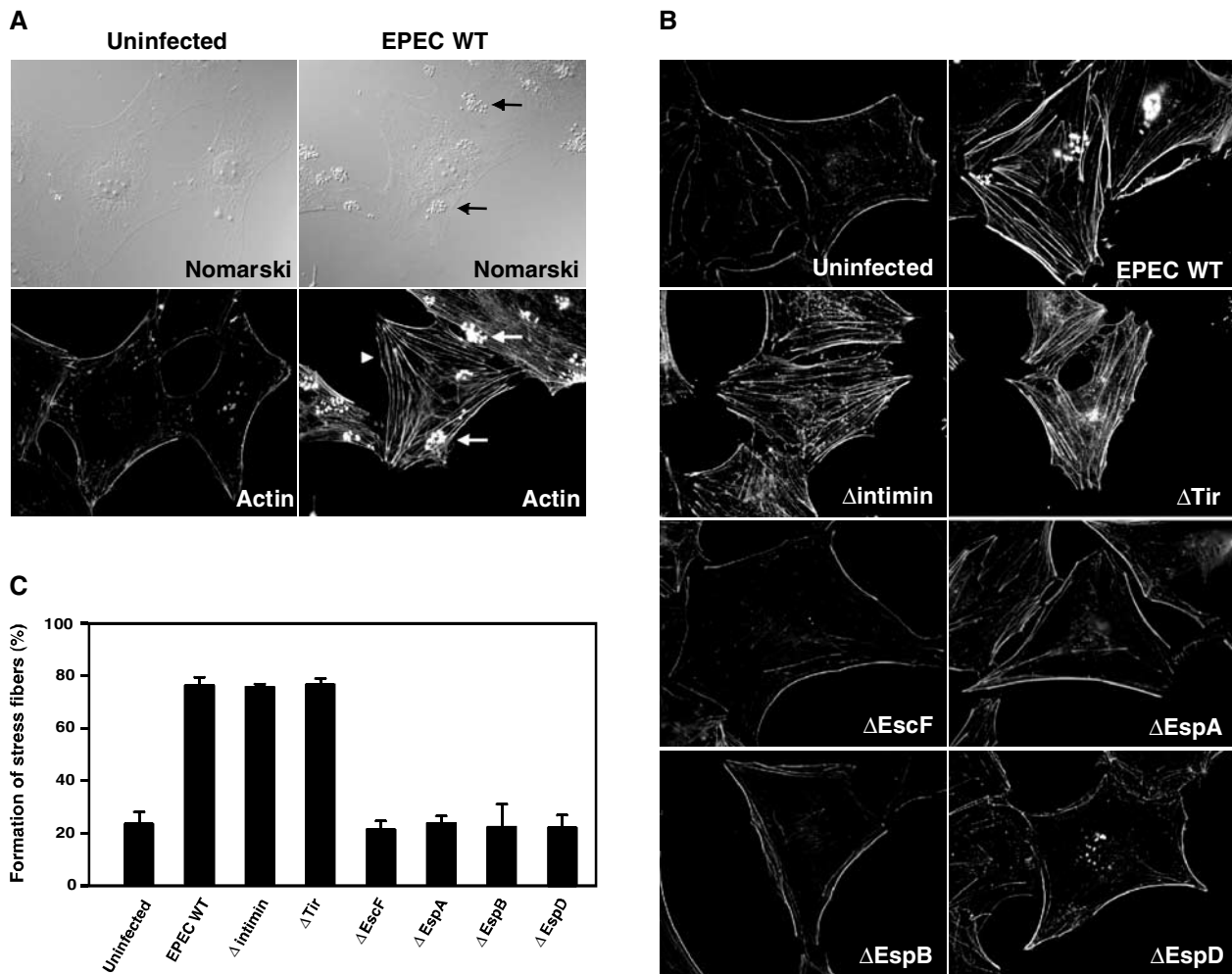


Figure 1 The host cytoskeletal rearrangement as a result of EPEC infection. (A) Swiss 3T3 cells were infected with EPEC WT for 3 h, and cells were fixed and stained with rhodamine-labeled phalloidin. The formation of pedestal structures beneath EPEC microcolonies (arrows) and actin stress fibers (arrowhead) was observed in the cells infected with EPEC. (B) Swiss 3T3 cells were infected with EPEC WT or its mutants for 3 h, and the cells were fixed and visualized as in (A). All fluorescence images were taken with the same exposure time using a CCD camera. (C) The percentage of Swiss 3T3 cells showing marked formation of actin stress fibers is indicated. Swiss 3T3 cells were infected with EPEC WT or its mutants for 3 h, and the cells were fixed and visualized as in (A). Percentages were based on counts of at least 150 cells, and the values are the means \pm standard deviation from three independent experiments.

EspG/Orf3 interaction with tubulins and stimulation of microtubule destruction

EspG and Orf3 shared a striking homology not only with each other but also with the *Shigella* VirA effector (Elliott *et al*, 2001). VirA is secreted via the type III machinery, and is required for promoting the uptake of bacteria by epithelial cells. Since VirA has the ability to interact with tubulins and trigger microtubule destabilization, this finally leads to Rac1 activation (Yoshida *et al*, 2002). Interestingly, cloned EPEC *espG* and *orf3* were able to individually rescue the ability of the *Shigella virA* mutant to invade host cells, implying that both EspG and Orf3 shared functions similar to those of VirA (Elliott *et al*, 2001). To investigate the effects of both EspG and Orf3 on host microtubule networks, Swiss 3T3 cells were infected for 3 h with EPEC WT or the *espG/orf3* double mutant, and the reorganization of both F-actin and the microtubule structures was observed by immunofluorescence microscopy. As shown in Figure 3A, the local microtubule structures beneath the EPEC microcolonies were destabilized by infection with EPEC WT (Figure 3A, arrows),

but not with the *espG/orf3* mutant. These events were reproducibly detected in EPEC WT-infected HeLa and Cos-7 cells, and also with the introduction of cloned *espG* or *orf3* (pEGFP-EspG, pEGFP-Orf3) into Swiss 3T3 cells (data not shown). In addition, the time course of microtubule destabilization after EPEC WT or *espG/orf3* mutant infection was measured, and the percentage of destabilization beneath EPEC microcolonies was calculated (Figure 3B). Destabilization of the local microtubule structures was greatly increased as a result of WT infection. In contrast, no such effect was detected in infection with the *espG/orf3* mutant. Like EPEC WT, the *espG* or *orf3* single mutant still possessed the ability to destabilize the microtubule networks, and complementation of the *espG/orf3* double-knockout mutant with a cloned *espG* or *orf3* restored the ability to destabilize microtubule networks due to functional redundancy (Figure 3C and D). Collectively, both EPEC effectors, EspG and Orf3, are involved in the induction of actin stress fiber formation and the destabilization of microtubule networks *in vivo*.

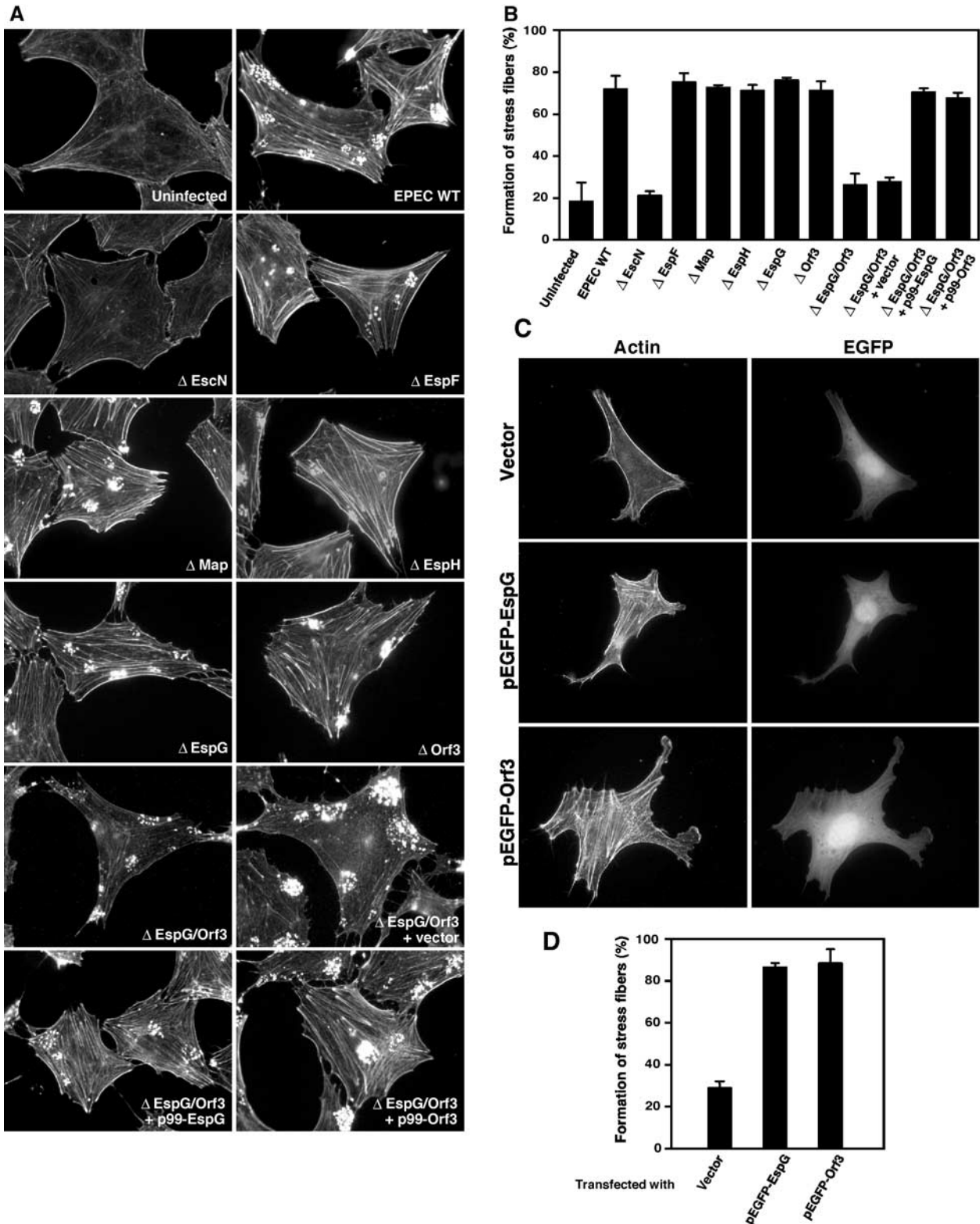


Figure 2 Determination of the EPEC type III effectors required for stress fiber formation. (A) Swiss 3T3 cells were infected with EPEC WT, various type III effector mutants, or an *espG/orf3* mutant complemented with cloned *espG* (p99-EspG), *orf3* (p99-Orf3), or vector alone for 3 h. Cells were fixed and stained with rhodamine-labeled phalloidin in order to detect F-actin. All fluorescence images were taken with the same exposure time. (B) The percentage of Swiss 3T3 cells showing marked formation of actin stress fibers is indicated. Swiss 3T3 cells were infected with EPEC WT or its mutants for 3 h, and the cells were fixed and visualized as in (A). Percentages were based on counts of at least 150 cells, and the values are the means \pm standard deviation from three independent experiments. (C) Swiss 3T3 cells were transfected with the plasmids encoding EGFP-tagged EspG, EGFP-tagged Orf3, and vector alone. Cells were fixed and stained as in (A), and the transfected cells were visualized by EGFP fluorescence. (D) EGFP fluorescence-positive Swiss 3T3 cells were randomly obtained from panel C, and the percentage of cells showing actin stress fiber formation is indicated. Percentages were based on a count of at least 100 cells, and the values are the means \pm standard deviation from three independent experiments.

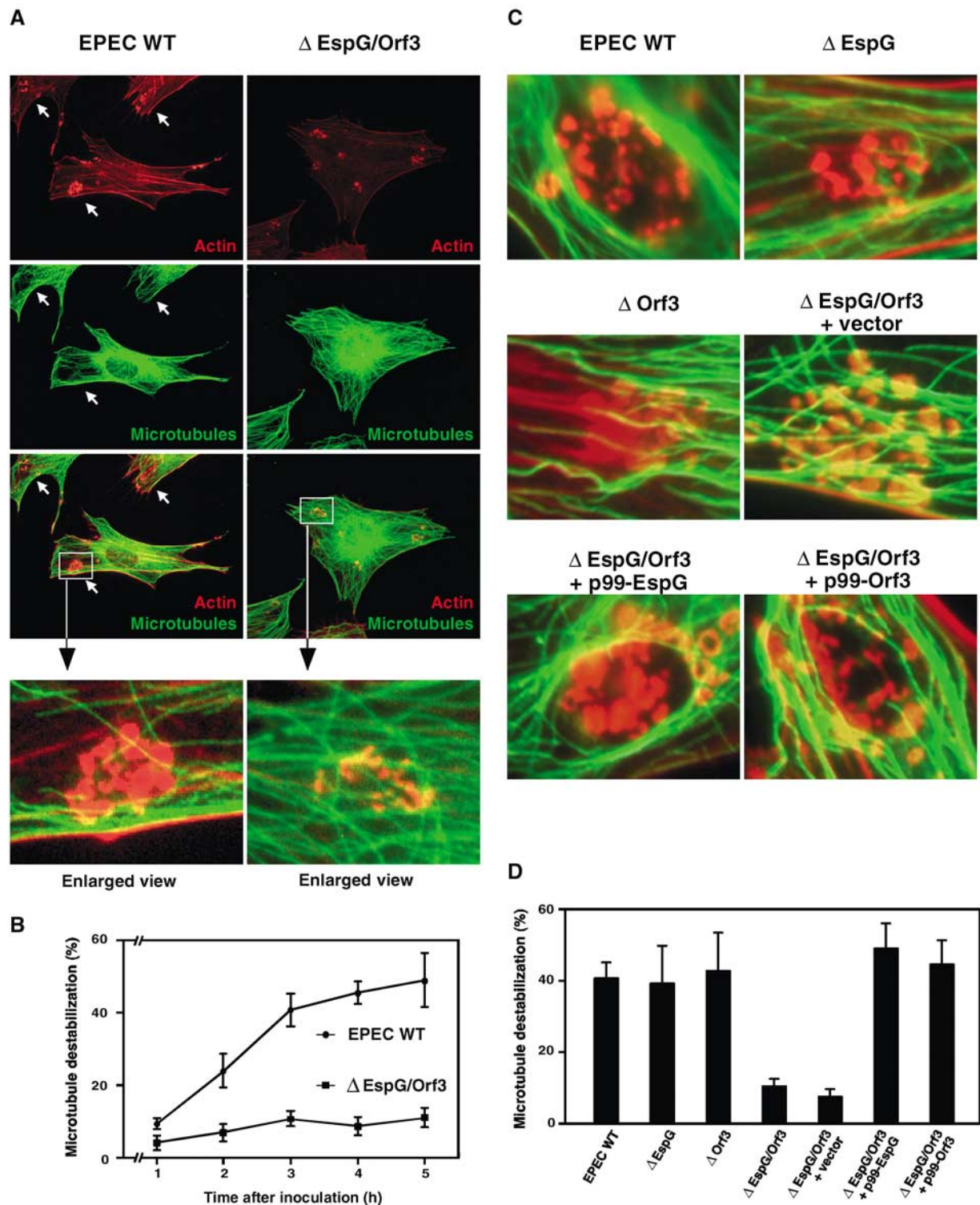


Figure 3 Effect of EPEC infection on microtubule dynamics. (A) Swiss 3T3 cells were infected with EPEC WT and the *espG/orf3* mutant for 3 h, and the cells were fixed and stained with rhodamine-labeled phalloidin and fluorescently labeled anti-tubulin antibodies in order to detect the actin cytoskeleton and microtubule networks, respectively. The arrows indicate disruptions of the microtubule network beneath the adherent EPEC microcolonies. (B) Time course of the local destabilization of microtubule networks after EPEC infection. The percentages of microtubule destabilization beneath the EPEC microcolonies were based on a count of at least 150 microcolonies, and the values given are the means \pm standard deviation from three independent experiments. (C) Swiss 3T3 cells were infected with EPEC WT and its mutants for 3 h and the cells were fixed and visualized as in (A). (D) The percentage of local destabilization of microtubule networks beneath the EPEC microcolonies using various EPEC derivatives. After 3 h infection, percentages were calculated as in (B).

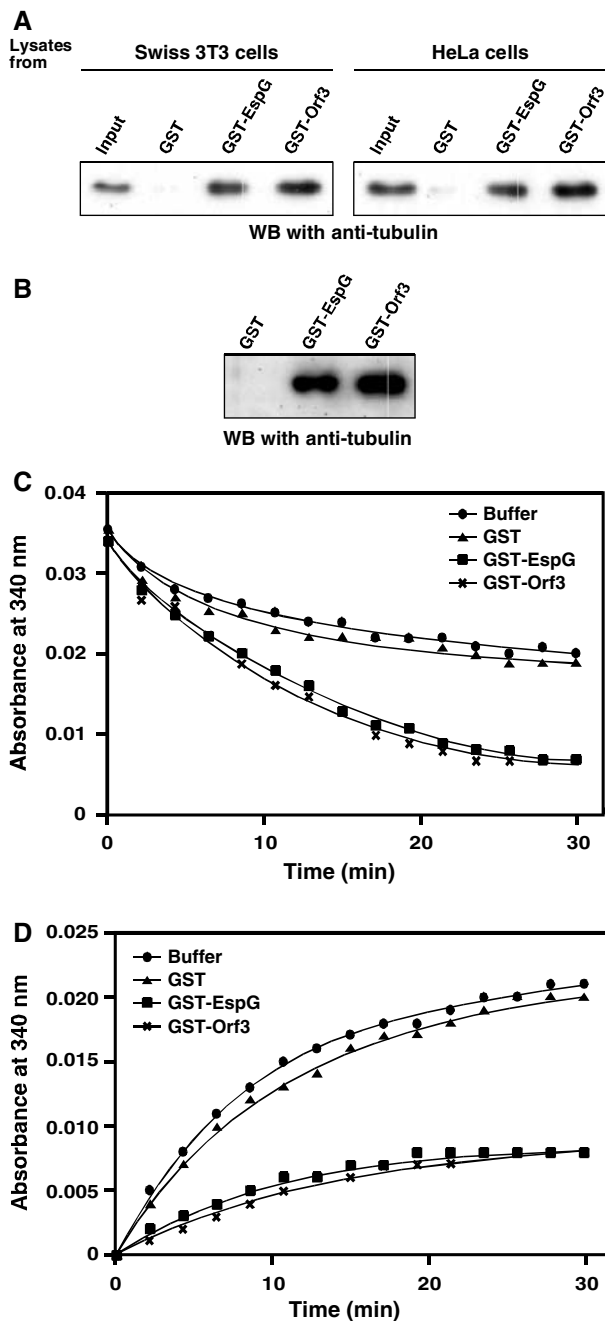


Figure 4 EspG/Orf3 bound to tubulins and induced the destabilization of microtubules *in vitro*. (A) GST fusion proteins bound to glutathione-Sepharose 4B beads were incubated with Swiss 3T3 and HeLa cell lysates for 2 h at 4°C. The beads were washed, and then bound proteins were analyzed by immunoblotting using anti-tubulin antibodies. (B) GST fusion proteins bound to glutathione-Sepharose 4B beads were incubated with purified twice-cycled microtubule proteins for 2 h at 4°C. The beads were washed, and then each sample was subjected to immunoblotting as in (A). (C) EspG/Orf3 induces microtubule destabilization. Purified twice-cycled microtubule proteins (15 μ M) were polymerized at 37°C for 1 h, and then equal volumes of PM-1.3 M buffer alone (circles) or PM-1.3 M buffer each containing 15 μ M of GST alone (triangles), GST-EspG (squares), or GST-Orf3 (crosses) were added. Destabilization of microtubules was monitored spectrophotometrically (absorbance at 340 nm) at 37°C for 30 min. (D) EspG/Orf3 inhibited microtubule polymerization. Purified twice-cycled microtubule proteins (7.5 μ M) were polymerized with PM-1.3 M buffer alone (circles) or PM-1.3 M buffer each containing 7.5 μ M of GST alone (triangles), GST-EspG (squares), and GST-Orf3 (crosses).

Next, using a GST pull-down assay, we investigated GST-EspG and GST-Orf3 for their ability to interact with tubulins contained in Swiss 3T3 or HeLa cell lysates. Analysis of the precipitated proteins with GST-EspG or GST-Orf3 in an immunoblot analysis with anti-tubulin antibodies revealed that the tubulins were co-precipitated with GST-EspG or GST-Orf3, but not with GST alone (Figure 4A). The direct binding of EspG/Orf3 to tubulins was further confirmed by a ligand overlay assay (see Materials and methods; data not shown) and by a pull-down assay with purified twice-cycled microtubule proteins (Yoshida *et al*, 2002) (Figure 4B). Moreover, we considered whether or not EspG/Orf3 would be able to affect either microtubule stabilization or destabilization using twice-cycled microtubule proteins (Figure 4C and D). Twice-cycled microtubule proteins were polymerized at 37°C for 1 h, and the resulting polymerized microtubules (15 μ M) were incubated with an equal volume of GST alone, GST-EspG, or GST-Orf3 (15 μ M each), and the microtubule dynamics were monitored by spectrophotometry (absorbance at 340 nm) (Figure 4C). In the absence of GST-EspG or GST-Orf3, the destabilization of the microtubules was slightly induced after 30 min (Figure 4C, circles), and similar results were observed in an incubation with GST alone (Figure 4C, triangles). In contrast, depolymerization of the microtubules was triggered by the addition of GST-EspG (Figure 4C, squares) or GST-Orf3 (Figure 4C, crosses). Furthermore, microtubule assembly was inhibited by the addition of GST-EspG and GST-Orf3, but not GST alone (Figure 4D). These results clearly demonstrate that EspG/Orf3 has the ability to bind directly to tubulins, and that it promotes the destabilization of microtubules *in vitro*.

EspG/Orf3-induced rearrangement of the actin cytoskeleton is dependent on GEF-H1

Regulation of the actin cytoskeleton by microtubules relies on the activity of the Rho family GTPases (Wittmann and Waterman-Storer, 2001). Microtubule disassembly results in the activation of RhoA, which enhances myosin contractility and actin stress fiber formation (Enomoto, 1996; Liu *et al*, 1998; Ren *et al*, 1999). The newly identified GEF family members, including murine Lfc (Glaven *et al*, 1999), murine p190RhoGEF (van Horck *et al*, 2001), and human GEF-H1 (Lfc homolog) (Krendel *et al*, 2002), are associated with microtubules and function as RhoA/Rac activators. Since the destruction of local microtubule networks and the formation of actin stress fibers were the most pronounced host responses to EPEC infection, we investigated the involvement of GEF-H1 (human) in EspG- or Orf3-induced stress fiber formation in human epithelial HeLa cells, instead of in mouse Swiss 3T3 cells. When HeLa cells were infected with EPEC WT, extensive actin accumulation was detected beneath a substantial number of adherent bacteria. However, actin stress fiber formation was very weak in HeLa cells (data not shown). (In fact, this has been one of the major reasons as to why EPEC-induced stress fiber formation has not been characterized to date.) It is possible that actin may be preferentially supplied for the formation of a pedestal structure, rather than for stress fiber formation in EPEC-infected HeLa cells. To facilitate the detection of pathogen-induced stress fiber formation, including GEF-H1 activation, we created *tir/espG/orf3* triple mutants carrying either p99-EspG or p99-Orf3. Our preliminary data indicated that mutation of the *tir* gene turned off the formation of F-actin aggregation beneath the

EPEC adherent; this finding allowed us to detect the F-actin architecture in epithelial cells such as HeLa cells. As shown in Figure 5A, upon infection of HeLa cells with the *tir/espG/orf3* triple mutant overexpressing EspG, actin stress fibers appeared, a result that was not observed in HeLa cells infected with the triple mutant carrying the vector alone. Importantly, EspG-induced actin cytoskeletal rearrangement was almost completely abolished by the expression of a dominant-negative form of GEF-H1 (GEF-H1 DH^{mut}) in HeLa cells. Indeed, the percentage of cells showing a marked formation of actin stress fibers increased greatly among HeLa cells infected with the *tir/espG/orf3* triple mutant carrying p99-EspG or p99-Orf3 (Figure 5B). To further analyze the direct involvement of GEF-H1 in the EspG/Orf3-mediated actin stress fiber formation, a dominant-negative form of p115-RhoGEF (p115-RGS) (Johnson *et al*, 2003) was also used in this study, since p115-

RhoGEF is well characterized as a RhoA-specific activator. RhoA is activated by the treatment of lysophosphatidic acid (LPA) via p115-RhoGEF activity (Hart *et al*, 1998; Kozasa *et al*, 1998). Indeed, LPA-induced stress fiber formation was inhibited by the expression of p115-RGS (Figure 5B). As expected, the formation of EspG-induced stress fibers was not affected by the expression of p115-RGS (Figure 5A and B). In contrast, GEF-H1 was found to be modulated by the addition of nocodazole, which induces the depolymerization of microtubules. Moreover, the nocodazole-, but not LPA-, induced stress fiber formation was inhibited by the expression of GEF-H1 DH^{mut}. The impact of actin stress fiber formation elicited by EspG or Orf3 in HeLa cells was similar to that elicited via the destruction of microtubule structures by nocodazole treatment, but not by LPA treatment (Figure 5B). The EspG/Orf3-GEF-H1 signaling pathway was further confirmed by a

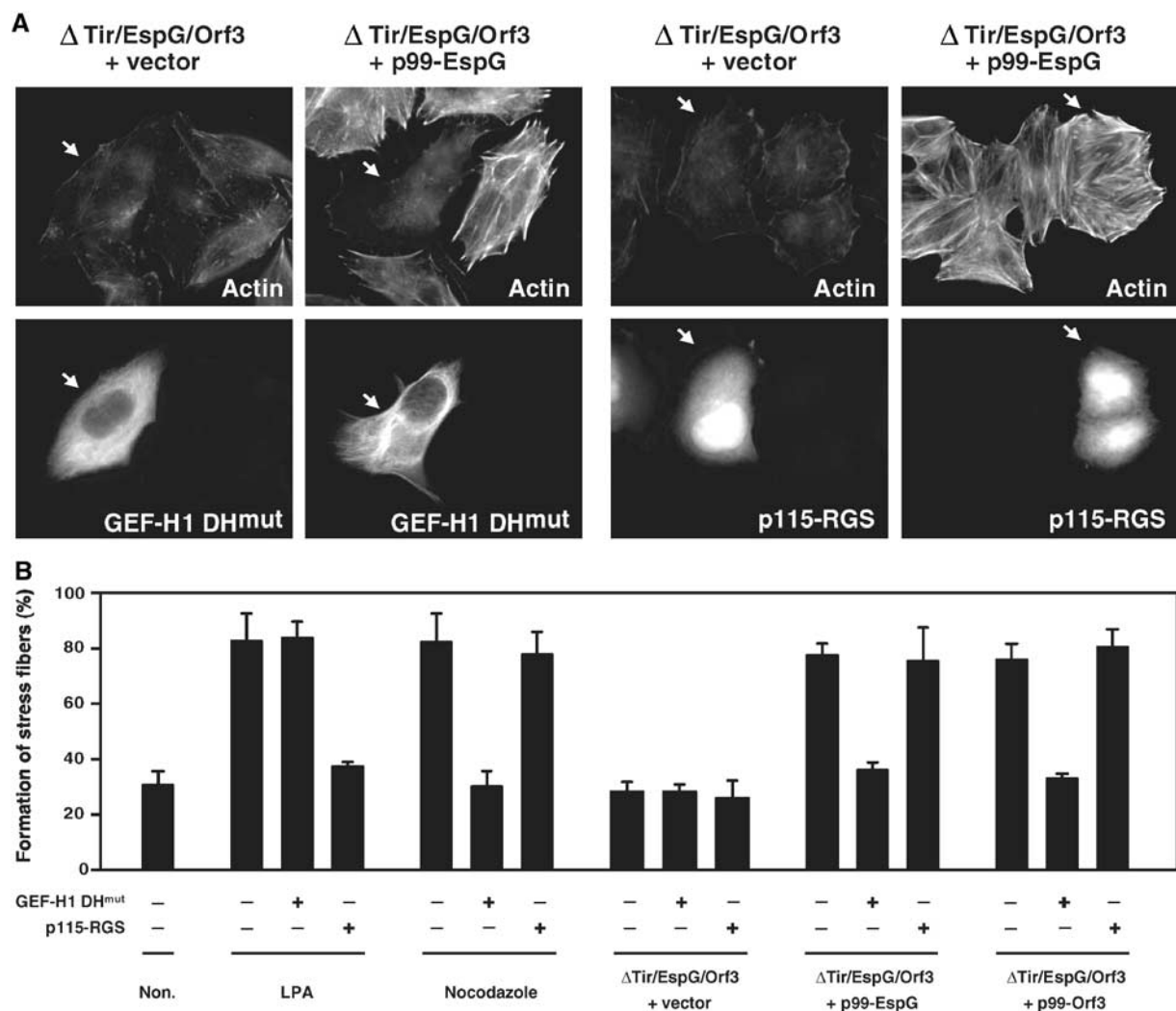


Figure 5 Effect of GEF-H1 on EPEC-induced actin stress fiber formation. (A) HeLa cells were transfected with a plasmid encoding the dominant-negative form of GEF-H1 (GEF-H1 DH^{mut}), containing a Tyr-to-Ala substitution at residue 393 in the DH domain of GEF-H1, or RGS domain of p115-RhoGEF (p115-RGS). Transfected cells were infected with a *tir/espG/orf3* mutant complemented with a cloned *espG* (p99-EspG) or with vector alone for 3 h, at a final concentration of 1 mM of IPTG in order to induce EspG expression. The cells were fixed and stained with rhodamine-labeled phalloidin in order to detect F-actin, and the transfected cells were visualized by EGFP fluorescence. The arrows indicate GEF-H1 DH^{mut} or p115-RGS. (B) HeLa cells were transfected with GEF-H1 DH^{mut} or p115-RGS, and then transfected cells were infected with the EPEC mutant complemented with a cloned *espG* (p99-EspG), *orf3* (p99-Orf3), or vector alone (vector). Otherwise, transfected cells were treated with 1 μ g/ml of LPA for 3 min or 10 μ M nocodazole for 1 h. The percentage of HeLa cells showing marked formation of actin stress fibers is indicated. Percentages were based on counts of at least 75 cells, and the values given are the means \pm standard deviation from three independent experiments.

small interfering RNA (siRNA) assay (Figure 6). Two siRNAs homologous to genes coding for GEF-H1 in different regions (GEF-H1-siRNA-I and GEF-H1-siRNA-II; Benais-Pont *et al*, 2003) were used; nonsilencing siRNA was used as a negative control. These siRNAs were introduced into HeLa cells, and the reduction in the transcription of GEF-H1 mRNA was confirmed by RT-PCR (Figure 6A). EspG- and Orf3-induced actin cytoskeletal rearrangement was almost completely abolished by the introduction of GEF-H1-siRNA-I or GEF-H1-siRNA-II into HeLa cells (Figure 6B and C). Collectively, these data clearly demonstrate that GEF-H1 is absolutely required for EspG- and Orf3-induced actin cytoskeletal rearrangement.

EspG/Orf3 induce GEF-H1 release from the cytoskeleton to the cytosol

In a previous study, Krendel *et al* reported that GEF-H1 is localized in the microtubules in its inactive form, and it switches to an active form when it is released from microtubule networks. Hence, to investigate whether GEF-H1 is

released from the cytoskeleton to the cytosol by EPEC infection, both cytoskeletal and cytosolic fractions were separated from the lysates of HeLa cells stably expressing GFP-tagged GEF-H1, and then GFP-GEF-H1 was detected by using anti-GFP antibodies (Figure 7). As was the case with the treatment with nocodazole, the relative amount of GEF-H1 associated with the cytoskeletal components was greatly reduced by infection with EPEC WT. The same results were observed in cells infected with the *espG* or *orf3* mutant (data not shown), but not with the *espG/orf3* mutant, or with the *escF* mutant (defective in the type III secretion system); these findings indicate that both EspG and Orf3 effectors possess the ability to release GEF-H1 from the cytoskeleton to the cytosol, and this effect presumably contributes to switching GEF-H1 to its active form.

EspG and Orf3 stimulate the RhoA-ROCK signaling pathway

Since GEF-H1 specifically promotes nucleotide exchange on RhoA, but not on Rac1 or Cdc42 (Krendel *et al*, 2002), cells

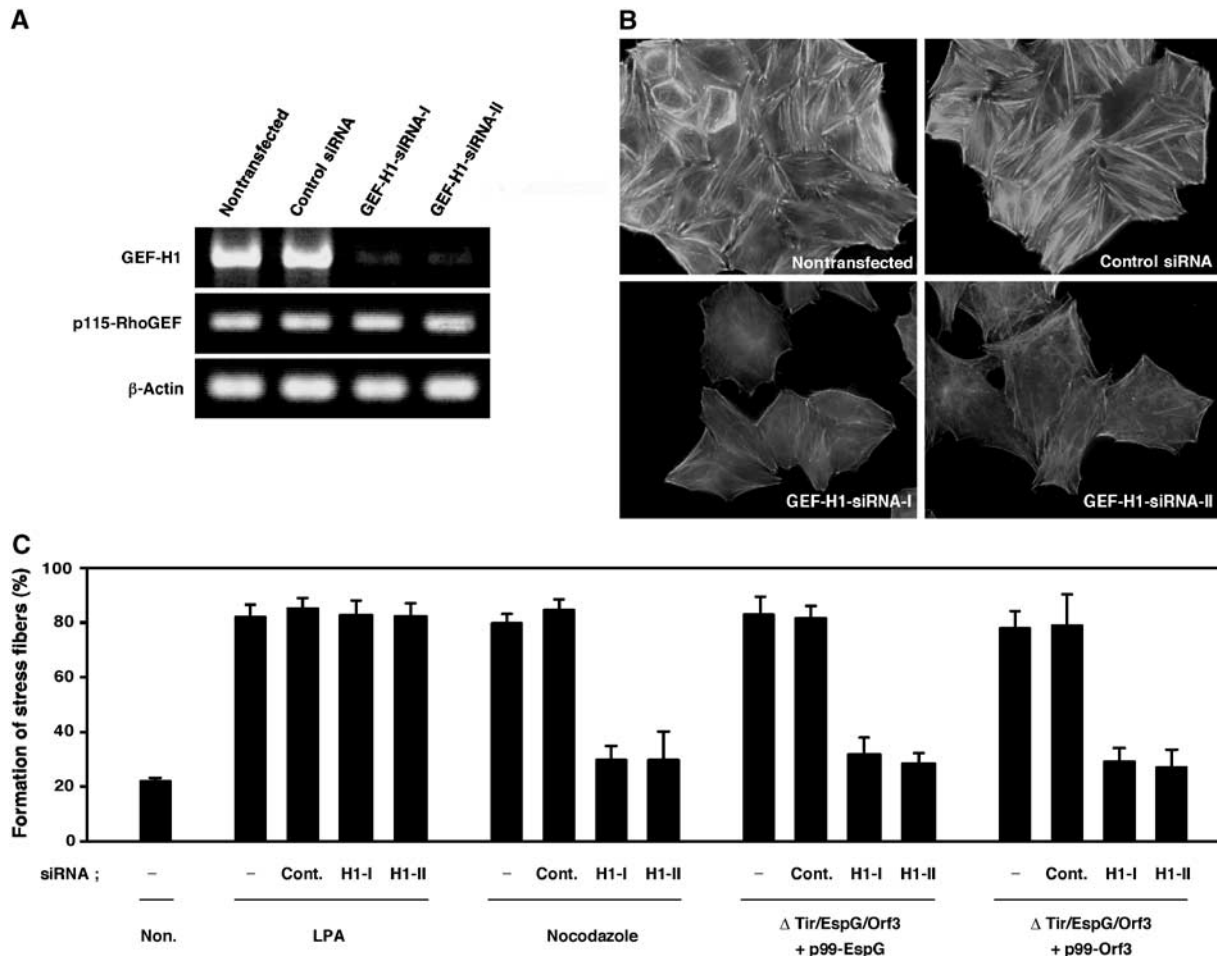


Figure 6 The EspG/Orf3-induced stress fiber formation was inhibited by reduced expression of GEF-H1. (A) Total RNA was purified from HeLa cells transfected with nonsilencing control siRNA, GEF-H1-siRNA-I, or GEF-H1-siRNA-II, and mRNAs for GEF-H1, p115-RhoGEF, and β -actin were analyzed by quantitative RT-PCR. (B) HeLa cells transfected with GEF-H1-siRNA-I, GEF-H1-siRNA-II, or nonsilencing control siRNA were infected with a *tir/espG/orf3* mutant complemented with a cloned *espG* (p99-EspG) for 3 h, at a final concentration of 1 mM of IPTG in order to induce EspG expression. The cells were fixed and stained with rhodamine-labeled phalloidin in order to detect F-actin. (C) The percentage of HeLa cells showing marked formation of actin stress fibers is indicated. HeLa cells were transfected with or without GEF-H1-siRNA-I (H1-I), GEF-H1-siRNA-II (H1-II), or nonsilencing control siRNA (Cont.). Otherwise, the transfected cells were treated with 1 μ g/ml of LPA for 3 min or 10 μ M nocodazole for 1 h. Percentages were based on counts of at least 150 cells, and the values are the means \pm standard deviation from three independent experiments.

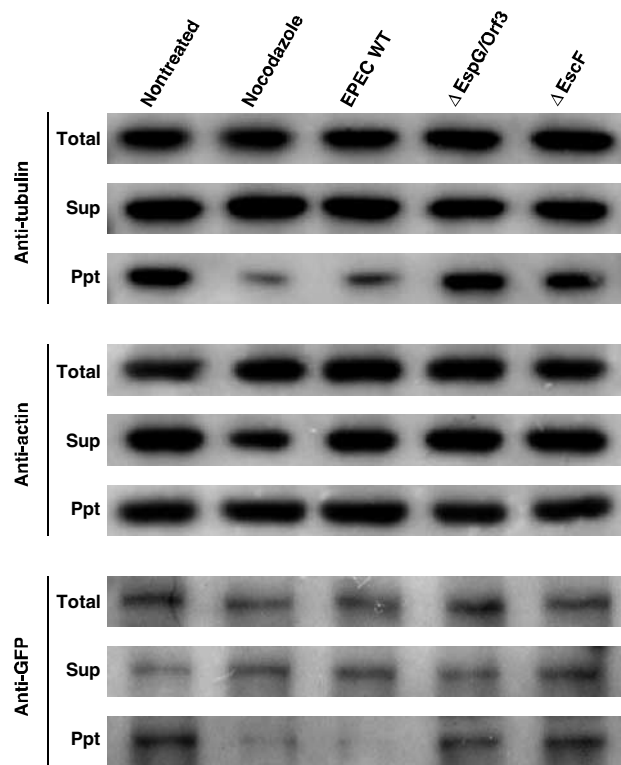


Figure 7 EspG/Orf3 induces GEF-H1 release from the cytoskeleton to the cytosol. HeLa cells stably expressing GFP-GEF-H1 were infected with EPEC WT or its mutants for 3 h. Otherwise, HeLa cells were treated with 10 μ M nocodazole for 1 h as a positive control. After infection, the cells were washed and lysed for the preparation of total protein (Total). The cytosolic fraction (supernatant; Sup) and the cytoskeletal fraction (precipitate; Ppt) were separated by centrifugation. In all, 5% of the total recovered whole-cell lysates or the cytosolic fraction and 20% of the cytoskeletal fraction were subjected to SDS-PAGE. The amount of tubulin, actin, and GEF-H1 was analyzed by Western blotting using anti-actin antibodies, anti-tubulin antibodies, or anti-GFP antibodies, respectively.

transfected with pFLAG-N19 (a RhoA dominant-negative mutant), pFLAG-N17-1 (a Rac1 dominant-negative mutant), and pFLAG-N17-42 (a Cdc42 dominant-negative mutant) were infected with EPEC WT, and the effects of the respective dominant-negative forms on bacterial-induced actin stress fiber formation were investigated by F-actin staining. Figure 8A shows the representative data, which reveal that the formation of EPEC-induced actin stress fibers was blocked when the RhoA dominant-negative form (N19) was expressed but not when the Rac1 (N17-1) or Cdc42 (N17-42) dominant-negative form was expressed. To further confirm these observations, at least 150 FLAG-positive cells with F-actin accumulation underneath the adherent bacteria were randomly chosen from each series, and the percentage of cells showing extensive stress fiber formation was calculated (Figure 8B). The number of cells with extensive actin stress fibers decreased greatly in the EPEC-infected Swiss 3T3 cells overexpressing FLAG-N19 (Figure 8B). This finding was reaffirmed by a pull-down assay using the Rho-binding domain (RBD) of rhotekin, which can bind to active RhoA (GTP-bound form). The level of the active form of RhoA in EPEC WT-infected cells was significantly higher than that in cells infected with the *espG/orf3* double mutant or the *escF*

mutant (Figure 8C). Moreover, the *espG* as well as the *orf3* single mutants still possessed the ability to activate RhoA, and complementation of the *espG/orf3* double-knockout mutant with p99-EspG or p99-Orf3 restored this ability. These findings indicate that RhoA was activated by EPEC WT infection, and that this event was dependent on the delivery of EspG/Orf3 into host cells by the type III secretion system.

Since these results strongly suggest that EspG- and Orf3-induced stress fiber formation is stimulated by activating the RhoA signaling pathway, we treated Swiss 3T3 cells with Y27632, a ROCK inhibitor, and we then analyzed the resulting effects on stress fiber formation. ROCK, a downstream effector of RhoA, is an excellent candidate for mediating RhoA-induced changes in the actin cytoskeleton (Kimura *et al*, 1996; Leung *et al*, 1996; Hall, 1998). As expected, Tir-induced F-actin accumulation beneath the adherent bacteria, caused by EPEC WT infection, was not affected by the addition of Y27632 (Figure 9A and B). However, the formation of actin stress fibers in parallel bundles, induced by EPEC WT and the *tir* mutant, was completely inhibited in the presence of Y27632. Collectively, these results demonstrate that stress fiber formation, when induced by the RhoA–ROCK signaling pathway, proceeds via a mechanism that is independent of the Tir–intimin interaction. This process is instead triggered via the stimulation of GEF-H1 activity, which is elicited by the destruction of host microtubule networks by EspG and Orf3 effectors.

Discussion

We demonstrated in this series of experiments that both EspG and Orf3 can modulate GEF-H1-mediated signaling by the disruption of host microtubules. EspG- or Orf3-induced GEF-H1 modulation might thus contribute to an increase in paracellular permeability, rather than to the disruption of tight junctions. A recent study has shown that GEF-H1 is associated with tight junctions and that it regulates size-selective paracellular permeability by the activation of the RhoA–ROCK signaling pathway (Benais-Pont *et al*, 2003). Furthermore, GEF-H1 was not found to have an effect on overall cell morphology, including tight junction assembly, as was determined by measuring the transepithelial electrical resistance (TER). The disruption of tight junctions has been observed in studies of EPEC WT infection, but not in those of *espF* mutant infection, indicating that the EspF effector is involved in tight junction disruption (McNamara *et al*, 2001). These findings are supported by data from a previous report (Elliott *et al*, 2001). As was the case with the infection of EPEC WT, infection of an *espG/orf3* double mutant equally induced a reduction in TER, indicating that neither EspG nor Orf3 is involved in tight junction disruption (Elliott *et al*, 2001). Currently, the effects of EPEC infection on paracellular permeability are under investigation in a study introducing purified EspG and Orf3 into MDCK cells; EPEC effectors that regulate epithelial barrier function may be involved in this type of virulence. Indeed, a recent study demonstrated that both EspG and EspF are required for *C. rodentium* to achieve full virulence (Deng *et al*, 2004).

When EspG and Orf3 were translocated into host cells, both proteins appeared to interact with tubulins; this effect was found to promote the disruption of microtubule networks. A GST pull-down assay showed that EspG and Orf3

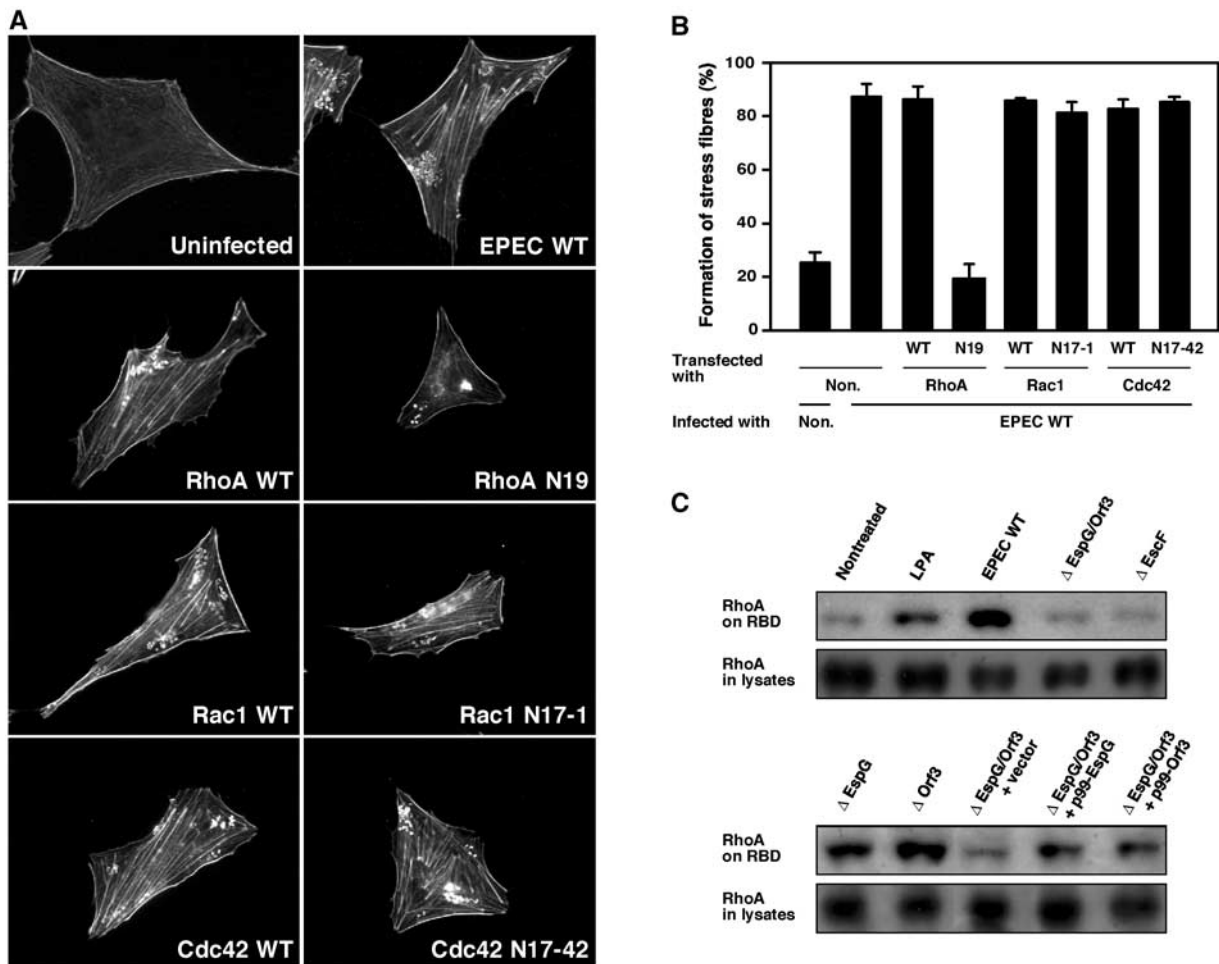


Figure 8 Activation of the RhoA signal pathway by EPEC infection. (A) Swiss 3T3 cells were transfected with plasmids encoding WT or dominant-negative forms (N19 for RhoA, N17-1 for Rac1, and N17-42 for Cdc42) of the Rho family GTPases, and the cells were infected with the EPEC WT for 3 h. The cells were then fixed and stained with rhodamine-labeled phalloidin and fluorescently labeled anti-FLAG antibodies in order to detect F-actin and transfected cells, respectively. FLAG-positive cells are shown in (A). (B) FLAG-positive cells with F-actin accumulation beneath the adherent bacteria were obtained, and the percentage of cells showing stress fiber formation is indicated. Percentages were based on counts of at least 150 cells, and the values given are the means \pm standard deviation from three independent experiments. (C) Cos-7 cells were transfected with FLAG-tagged WT RhoA, and then the cells were infected with the EPEC WT or mutants for 3 h, or they were treated with 1 μ g/ml of LPA for RhoA activation as a positive control for 3 min. Cells were lysed and the GTP-bound RhoA, the active form of RhoA, was co-precipitated with the GST-Rho binding domain (RBD) of rhotekin. The amount of RhoA bound to RBD and RhoA in the whole-cell lysates was analyzed by Western blotting with anti-FLAG antibodies.

had the ability to bind to the soluble form of tubulins (Figure 4). In order to investigate in more detail whether or not EspG and Orf3 could bind to microtubules, taxol-stabilized tubulins, prepared as previously described (Yoshida *et al*, 2002), were used for the pull-down assay. However, we were unable to detect the binding of EspG or Orf3 to taxol-stabilized tubulins (data not shown). It should be noted that similar results have been reported in the case of the *Shigella* VirA effector (Yoshida *et al*, 2002). The oncoprotein Op18/stathmin is known to sequester tubulins, and the presence of this protein resulted in a reduction in the tubulin concentration required for microtubule assembly, followed by a promotion of microtubule destabilization (Cassimeris, 2002). EspG and Orf3 may exert an effect on microtubules much like that of Op18/stathmin (Cassimeris, 2002). Although EspG and Orf3 in EPEC, as well as VirA in *Shigella*, can exert effects on microtubule networks, EspG and Orf3 induce the formation of actin stress fibers and VirA induces membrane ruffling (Yoshida *et al*, 2002). The different mechan-

isms underlying the cytoskeletal rearrangement of host cells, induced by these factors, remain to be elucidated.

Figure 10 depicts the action of EPEC type III effectors (e.g., Tir, EspF, and EspG/Orf3) in host cells. In EPEC-infected cells, the rearrangement of actin organization is promoted by two different pathways, that is, by the formation of a pedestal structure induced by Tir, and by actin stress fiber assembly induced by EspG and Orf3. The *tir* and *eae* mutants still possessed the ability to induce stress fiber formation, but they could not form the pedestal structure (Figure 1B). In contrast, the formation of a pedestal structure was still induced by infection with the *espG/orf3* double mutant (Figure 2A). Although actin stress fiber formation was inhibited by the introduction of dominant-negative RhoA, and also by treatment with Y27632, Tir-induced pedestal formation was not affected (Figures 8 and 9). These results indicate that Tir and EspG/Orf3 independently contribute to the induction of host cytoskeletal rearrangement using different host signal pathways. Like Tir, EspG and its homolog Orf3 are

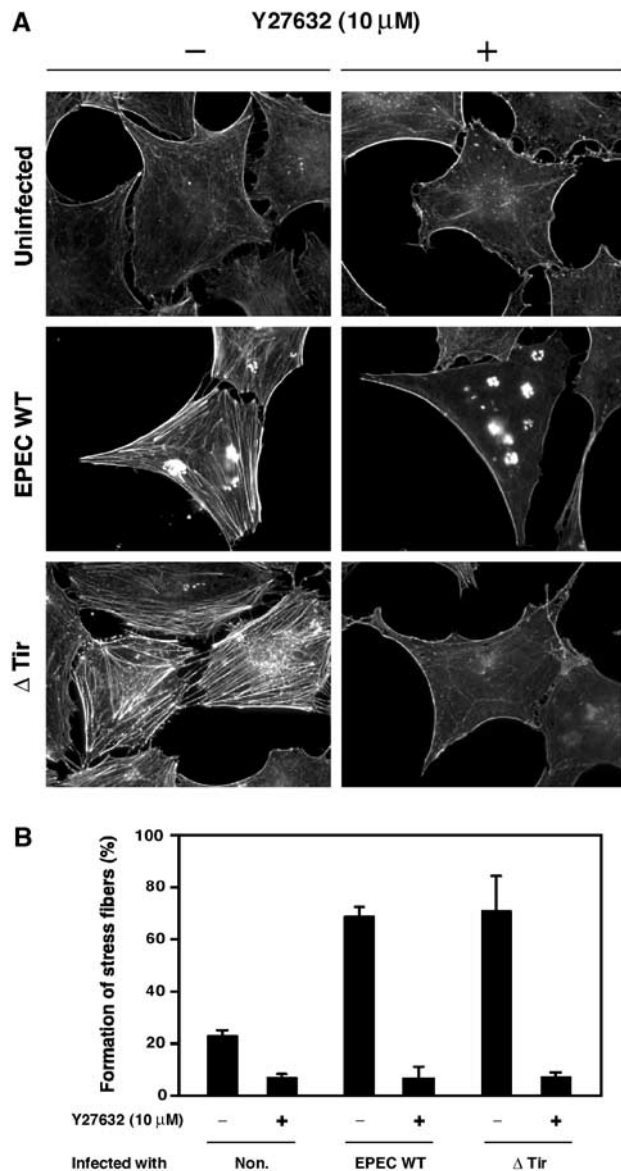


Figure 9 Effect of Y27632 on EPEC-induced actin stress fiber formation. (A) Swiss 3T3 cells were maintained in the presence or absence of 10 μ M Y27632 for 1 h, and were infected with EPEC WT and the *tir* mutant for 3 h in the presence or absence of 10 μ M Y27632. The cells were fixed and stained with rhodamine-labeled phalloidin in order to detect F-actin. All fluorescence images were taken with the same exposure time. (B) The percentage of Swiss 3T3 cells showing marked formation of actin stress fibers is indicated. Swiss 3T3 cells were infected with EPEC WT and the *tir* mutant for 3 h in the presence or absence of 10 μ M Y27632 as in (A). Percentages are based on counts of at least 150 cells, and the values are the means \pm standard deviation from three independent experiments.

secreted and injected into host cells via type III machinery. EspG/Orf3 can in turn bind directly to tubulins, and thereby induce the destabilization of microtubules (Figures 3 and 4). These effects of EspG/Orf3 on tubulins cause the release of GEF-H1 from the cytoskeleton to the cytosol, and may activate a host cell signaling pathway involving GEF-H1, RhoA, and ROCK (Figures 5–9).

We demonstrated here that EspG and Orf3 effectors regulate the crosstalk between microtubules and actin stress

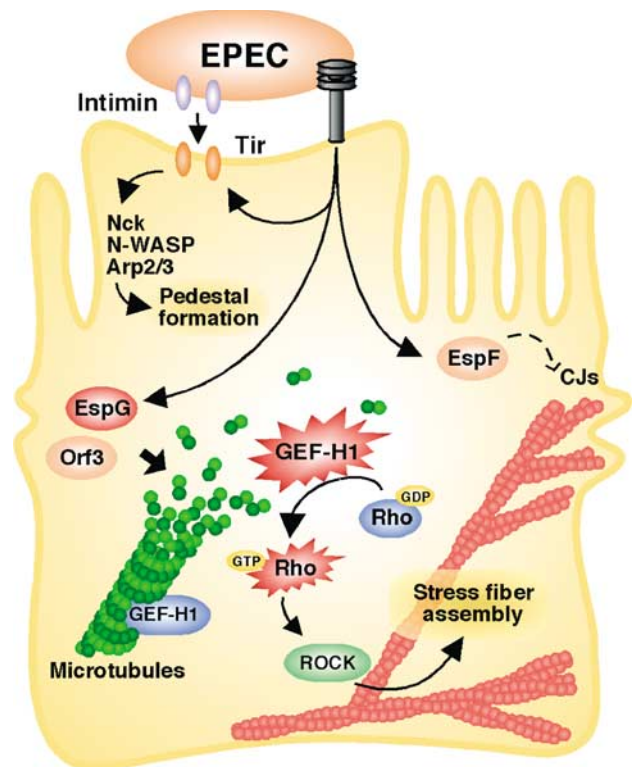


Figure 10 A presumed model for the action of EPEC type III effectors in host cells. EPEC injects the effectors (e.g., Tir, EspF, EspG, and the EspG homolog Orf3) via a type III secretion machinery into the host cells. The interaction of a bacterial membrane protein, intimin, and translocated Tir triggers the host signal transduction pathway, which includes the recruitment of Nck, N-WASP, and the Arp2/3 complex, resulting in the formation of a pedestal structure beneath the adherent bacteria. On the other hand, it is possible that EspF and EspG/Orf3 influence the function of cell junctions (CJs). Although the precise function of EspF is not clear at present, EspF is known to exert an effect on the morphology of tight junction strands. EspG and Orf3 associate with tubulins, resulting in the disruption of microtubule networks. GEF-H1 switches to the active form as a result of its dissociation from microtubule networks. Activated GEF-H1 promotes the binding of GTP to RhoA, resulting in the activation of RhoA. Finally, the activation of ROCK, which is located downstream of the RhoA signal, induces actin stress fiber assembly and probably exerts influence on paracellular permeability in EPEC-infected cells.

fibers by exploiting GEF-H1. These events are likely to be involved in an increase in paracellular permeability. A/E lesion formation due to Tir-intimin interactions, and the disruption of tight junctions by EspF have an effect on the overall morphology and physiology of host cells. The synergistic effects caused by multiple effectors are expected to contribute to triggering EPEC-mediated diarrhea.

Materials and methods

Bacterial strains, plasmids, transfection, infection assay, in vitro tubulin binding assay, and ligand overlay assay
See Supplementary Materials and methods available at *The EMBO Journal* Online.

In vitro microtubule-destabilization assay

Polymerization of twice-cycled microtubule proteins was monitored spectrophotometrically (absorbance at 340 nm) at 37°C as described previously (Gaskin, 1982; Horwitz *et al*, 1997; Yoshida *et al*, 2002). To assess the effect of EspG or Orf3 on microtubule stabilization,

equal volumes of PM-1.3 M buffer (PM buffer containing 1.3 M glycerol) alone or PM-1.3 M buffer containing GST alone, GST-EspG, or GST-Orf3 were added to the cuvettes either before or after inducing the polymerization of the microtubules.

Small-interference RNA assay

Duplex siRNAs were constructed against sequences coding for different GEF-H1 regions (GEF-H1-siRNA-I, 5'-AACAAGAGCATCA CAGCCAAGdTdT-3'; GEF-H1-siRNA-II, 5'-AATGTGACTATCCACAA CCGCdTdT-3'), as described elsewhere (Benais-Pont *et al*, 2003), and nonsilencing control siRNA with a random sequence was purchased from QIAGEN. These siRNA oligonucleotides were introduced into cultured cells with RNAiFect transfection reagent (QIAGEN) and the transfection efficiency (approximately 90%) was confirmed by using a fluorescein-labeled siRNA. The total RNA was purified from siRNA-transfected cells using Concert Micro-to-Midi Total RNA Purification System (Invitrogen, Carlsbad, CA), and then reverse transcription was performed using a Superscript III First Strand Synthesis system (Invitrogen). Transcription of GEF-H1, p115-RhoGEF, and β -actin mRNA was analyzed by agarose gel electrophoresis after RT-PCR using the respective primers.

Differential cytoskeletal extraction

Differential cytoskeletal extraction was performed as described previously (van Bergen en Henegouwen *et al*, 1992; Ding *et al*, 1996). HeLa cells stably expressing GFP-tagged GEF-H1 were infected with EPEC WT or its mutants for 3 h, or the cells were treated with 10 μ M nocodazole for 1 h as a positive control. After infection, the cells were washed four times in PBS and lysed in cytoskeleton-stabilizing buffer (10 mM PIPES-NaOH pH 6.8,

250 mM sucrose, 3 mM MgCl₂, 150 mM KCl, 1 mM EGTA, and 1 mM PMSF) containing 0.15% Triton X-100 for 5 min at 37°C. Cell lysates were then centrifuged at 14 000 g at room temperature for 10 min, and the supernatant and the pellet were considered as the cytosolic and cytoskeletal fractions, respectively. The relative amounts of actin, tubulin, and GEF-H1 were analyzed by Western blotting using anti-actin antibodies (CHEMICON, Temecula, CA), anti-tubulin antibodies (Sigma, St Louis, MO), or anti-GFP antibodies (Molecular Probes, Eugene, OR).

GST-RBD pull-down assay

See Supplementary Materials and methods available at *The EMBO Journal Online*.

Supplementary data

Supplementary data are available at *The EMBO Journal Online*.

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