The cytotoxicity of infectious agents can be mediated by disruption of calcium signaling in target cells. Outer membrane proteins of the spirochete Treponema denticola, a periodontal pathogen, inhibit agonist-induced Ca\textsuperscript{2+} release from internal stores in gingival fibroblasts, but the mechanism is not defined. We determined here that the major surface protein (Msp) of T. denticola perturbs calcium signaling in human fibroblasts by uncoupling store-operated channels. Msp localized in complexes on the cell surface. Ratio fluorimetry showed that in cells loaded with fura-2 or fura-C18, Msp induced cytoplasmic and near-plasma membrane Ca\textsuperscript{2+} transients, respectively. Increased conductance was confirmed by fluorescence quenching of fura-2-loaded cells with Mn\textsuperscript{2+} after Msp treatment. Calcium entry was blocked with anti-Msp antibodies and inhibited by chelating external Ca\textsuperscript{2+} with EGTA. Msp pretreatment reduced the amplitude of [Ca\textsuperscript{2+}]\textsubscript{i} transients upon challenge with ATP or thapsigargin. In experiments using cells loaded with mag-fura-2 to report endoplasmic reticulum Ca\textsuperscript{2+}, Msp reduced Ca\textsuperscript{2+} efflux from endoplasmic reticulum stores when ATP was used as an agonist. Msp alone did not induce Ca\textsuperscript{2+} release from these stores. Msp inhibited store-operated influx of extracellular calcium following intracellular Ca\textsuperscript{2+} depletion by thapsigargin and also promoted the assembly of subcortical actin filaments. This actin assembly was blocked by chelating intracellular Ca\textsuperscript{2+} with 1,2-bis(2-aminophenoxy)ethane-N\textsubscript{2}N\textsubscript{3}N\textsubscript{4}N\textsubscript{1}-tetraacetic acid acetoxyethyl ester. The reduced amplitude of agonist-induced transients and inhibition of store-operated Ca\textsuperscript{2+} entry due to Msp were reversed by latrunculin B, an inhibitor of actin filament assembly. Thus, Msp retards Ca\textsuperscript{2+} release from endoplasmic reticulum stores, and it inhibits subsequent Ca\textsuperscript{2+} influx by uncoupling store-operated channels. Actin filament rearrangement coincident with conformational uncoupling of store-operated calcium fluxes is a novel mechanism by which surface proteins and toxins of pathogenic microorganisms may damage host cells.

What if pathogenic microorganisms were able to disrupt the regulation of intracellular calcium signals in host cells? They would disturb a critical signaling pathway by which eukaryotic cells respond to external stimuli (1) and would impair crucial downstream functions by which cells maintain homeostasis. For example, calcium ion concentration ([Ca\textsuperscript{2+}]\textsubscript{i}) regulates the assembly and turnover of actin filaments (2), which in turn control cell shape, volume regulation, locomotion, and phagocytosis. These are essential functions for maintaining tissue integrity, for promoting wound healing, and for mediating protection from infection.

Many human cells regulate calcium homeostasis using store-operated calcium fluxes, a system in which Ca\textsuperscript{2+} influx through store-operated channels (SOCs) in the plasma membrane is stimulated by the depletion of Ca\textsuperscript{2+} in internal stores in the lumen of the endoplasmic reticulum (ER) (3, 4). A variety of potential cytotoxins of insects, reptiles, and microorganisms affect ion channel function (5–7). These toxins may include the surface molecules of opportunistic or overtly pathogenic bacteria, including some that cause cytoskeletal perturbation or that form ion channels in the host cell plasma membrane. Several microbial products have been reported to alter ion transport, but none is known to uncouple Ca\textsuperscript{2+} ER stores and SOCs.

Although the mechanisms by which transient Ca\textsuperscript{2+} depletion in the ER triggers Ca\textsuperscript{2+} influx through SOCs are not well understood, one compelling model is based on the physical proximity between the plasma membrane and the ER (1). This system may involve the inositol triphosphate receptor on the ER as a mediator of the functional coupling of ER store depletion and Ca\textsuperscript{2+} entry through the plasma membrane (8, 9). Notably, the assembly and disassembly of actin filaments subjacent to the plasma membrane is a major determinant of reversible coupling of the ER and plasma membrane and thereby modulates the signals that regulate Ca\textsuperscript{2+} influx through SOCs (8, 10).

Whole cells and outer membrane proteins of the oral spirochete Treponema denticola exert profound effects on actin assembly in cultured human gingival fibroblasts and epithelial cells (11–14). These properties may contribute to bacterial pathogenicity by impeding actin-dependent functions that are crucial for homeostasis of gingival tissues (15, 16). Since actin assembly is highly sensitive to fluctuations in calcium concentration (2), it is notable that outer membrane fractions of T. denticola may interact with host cells to disrupt normal calcium signaling pathways.
*T. denticola* inhibit inositol phosphate and intracellular calcium responses of fibroblasts to chemical agonists (14, 17). Since cortical actin and the inositol trisphosphate receptor are evidently important in the regulation of store-operated calcium channels in a variety of cell types (8, 18–20), we hypothesized that the major surface protein of *T. denticola*, Msp,2 may be one of the outer membrane proteins that is responsible for bacterial perturbation of calcium flux in cells from the periodontium. We examined the effects of Msp on calcium homeostasis in gingival fibroblasts and report that Msp mediates calcium transients and inhibits SOC activation by promoting assembly of subcellular actin filaments. These filaments in turn uncouple the ER from SOCs. Our results provide a novel mechanism by which pathogenic bacteria may subvert host cell signaling systems and thereby contribute to loss of tissue homeostasis in chronic infections such as periodontitis.

**EXPERIMENTAL PROCEDURES**

**Reagents—**Bovine serum albumin, Texas Red-conjugated goat anti-rabbit antibodies, thapsigargin, ATP, ionomycin, latrunculin B, and valinomycin were purchased from Sigma. Fura-2/AM, mag-fura-2/AM, fura-C18, 1,2-bis(2-aminophenoxy)ethane-N,N'N'-tetraacetic acid acetoxyethyl ester, bis-1,3-dibutylbarbituric acid(trimethinoxenol, and rhodamine phalloidin were obtained from Molecular Probes, Inc. (Eugene, OR). 2-Aminoethoxydiphenyl borate (2-APB) was obtained from Calbiochem.

**Preparation of Msp—**Msp is an immunogenic surface protein of *T. denticola* that complexes with a serine protease, PrtP, in the outer sheath of the bacterium. Msp was prepared from *T. denticola* type strain ATCC 35405, as previously described (21, 22), with the following modifications. The starting material was ~8 g, wet weight, per 4-liter bacterial culture in late logarithmic growth phase. After the deoxycholate and the n-octylpolyoxylethylene extraction steps and ultracentrifugation, the enriched Msp solution was incubated for 7 days at 37 °C under the protection of the bacterium. Msp was prepared from Calbiochem.

**Gel Electrophoresis and Immunoblotting—**SDS-polyacrylamide gel electrophoresis was performed as previously described in 10% (w/v) acrylamide and N-benzoyl-DL-arginine-\(\text{NEt}_2\)-succinyl-L-alanyl-L-alanyl-L-prolyl-L-phenylalanine nitroanilide and N-benzoxy-\(\text{NEt}_2\)-arginine-p-nitroanilide (Sigma) (23). The enriched protein solution was concentrated ~30-fold by Amicon ultrafiltration (Amicon Concentric plus 80, Amicon Inc., Beverly, MA). After washing with 5 liters of 10 mM Tris (pH 8.0), five washings with double-distilled water, and ultracentrifugation for 2 h to remove trace amounts of detergent, the Msp was dissolved and dialyzed against double-distilled water. The protein content was determined by Bio-Rad assay using bovine serum albumin as a standard.

For all experiments (except dose-response experiments), a 1:100 dilution of Msp in calcium buffer was used. This preparation contained 0.41 mg of protein (dry weight) per mL. A final concentration of 30 μg of protein/mL corresponded to ~160 μM, was used for most challenge assays of cells. When pretreatment with Msp was specified, the Msp remained in the cell culture wells for the duration of the experiment unless mentioned. In some experiments, Msp was boiled for 10 min to denature proteins or heated at 60 °C for 30 min. In other experiments, Msp was pretreated with a mixture of Msp and anti-Msp antibodies at a 2.5:1 ratio (protein concentration) that had been preincubated for 60 min.

**Cell Culture—**Human gingival fibroblasts were derived from primary cultures as described (28). Cells from passages 6–15 were grown as monolayers in T-75 flasks in α-minimal essential medium with 10% fetal bovine serum and antibiotics. Cells from these cultures were harvested and replated onto glass coverslips 2 days before each experiment, as described previously (17). The cells were grown to confluence prior to all experiments except when sparse cultures were used.

**Intracellular Calcium—**For measurement of whole cell intracellular calcium ion concentration ([Ca\(^{2+}\)]_i), cells on coverslips were loaded with 3 μM fura-2/AM (a Ca\(^{2+}\)-sensitive fluorescent dye) for 20 min at 37 °C (17). For measurement of near-plasma membrane Ca\(^{2+}\) concentration, cells attached to coverslips were briefly permeabilized with saponin and incubated with 2 μM fura-2 AM. For determination of extracellular calcium, cells on coverslips were incubated with mag-fura-2/AM (4 μM) for 150 min at 37 °C, in α-minimal essential medium containing fetal bovine serum (10%) (31). The calcium-free buffer consisted of a bicarbonate-free medium containing 150 mM NaCl, 5 mM KCl, 10 mM glucose, 1 mM MgSO\(_4\), 1 mM NaH\(_2\)PO\(_4\), and 20 mM HEPES at pH 7.4 with an osmolarity of 291 mosM. For experiments requiring external calcium, 1 mM Ca\(_{\text{Cl}}\), was added to the buffer; for experiments requiring chelation of external Ca\(^{2+}\), 2 mM EGTA was added. F\(_{\text{max}}\) and F\(_{\text{min}}\) were determined using ionomycin and EGTA as described previously (17).

After incubation with fura-2/AM, inspection of cells by fluorescence microscopy demonstrated no vesicular compartmentalization of fura-2, suggesting that the dynamic compartmentalization of fura-2 does not occur with the cytosolic fraction of *T. denticola*. Visual inspection of fura-18-loaded cells and mag-fura-2-loaded cells showed fluorescent labeling of the plasma membrane and intracellular organelles, respectively. Estimates of whole cell [Ca\(^{2+}\)]_i, and near plasma membrane [Ca\(^{2+}\)]_i, were calculated according to the equation of Grynkiewicz (32) from the emitted fluorescence measurements, using a Nikon Diaphot II inverted microscope optically interfaced to a Deltascan 4000, dual beam, epifluorescence spectrophotometer and analysis system (Photon Technology Int., London, Ontario, Canada), as described previously (17).

**Msp Perturbation of Calcium Fluxes—**To examine the effect of Msp on internal release of Ca\(^{2+}\), ATP (100 μM), and thapsigargin (1 μM; Sigma) were used as agonists and added to resting cells in calcium-free buffer with EGTA (17). Cells that were pretreated with Msp for 40 min at 25 °C were compared with untreated cells. To compare the function of SOCs in these cells, the influx of extracellular calcium following depletion of intracellular stores was also studied. Fibroblasts were exposed to 1 μM thapsigargin in 1 mM EGTA buffer for 30 min to deplete internal Ca\(^{2+}\) and to chelate residual extracellular Ca\(^{2+}\). CaCl\(_2\) (2 mM) was added, and [Ca\(^{2+}\)]_i increases due to calcium influx were measured. These experiments were repeated in cells pretreated with 1 μM latrunculin B (30 min, 25 °C), an inhibitor of actin polymerization.

In some experiments, the influx pathway was examined after the addition of Msp or water vehicle in cells loaded with fura-2. We measured the fluorescence of single cells at the isosbestic point (356 nm) in the presence of MnCl\(_2\) (1 mM) in the standard Ca\(^{2+}\)-containing buffer (33). For indirect assessment of general cation permeability of the plasma membrane to monovalent cations, we measured membrane potential using flow cytometry and the bisoxonol dye bis-[1,3-dibutylbarbituric acid]trimethinoxenol (3) as described (34).

**Immunocytochemistry and Confocal Microscopy—**To examine the location of Msp binding to cells, indirect immunofluorescence microscopy was performed on Msp-pretreated cells using polyclonal anti-Msp antibodies as the primary antibody. Cells grown on coverslips were fixed and permeabilized with methanol at ~20 °C for 10 min, blocked with 1:1000 mouse serum in PBS for 10 min, incubated with primary antibody (1:100 dilution) for 1 h at room temperature, washed three times with PBS containing 0.2% bovine serum albumin, and incubated with Texas Red-conjugated goat-anti-rabbit antibodies (1:100). The coverslips were washed with PBS and mounted with an antifade mounting medium (ICN, Montreal, Canada). The fibroblasts were examined using a × 40, 1.3 NA oil immersion objective under epifluorescence optics (Leica CLSM, Heidelberg, Germany).

Laser-scanning confocal microscopy (Leica CLSM) was used to examine actin filament assembly induced by Msp. The cells were fixed in 3% paraformaldehyde for 15 min, permeabilized in 0.2% Triton X-100 for 5 min, and stained with rhodamine phalloidin (35). The excitation wavelength was 530/20 nm, and the emitted fluorescence was detected through an emission filter (620/40 nm). Cells were imaged with a × 63 oil immersion lens, 1.4 NA, and transverse optical sections were obtained in 1-μm steps from the level of cell attachment at the substratum to the dorsal surface of the cell (33). In some experiments, the fluorescence intensity of rhodamine phalloidin staining was measured with a fluorescence spectrophotometer (Leitz MWP, Wetzlar, Germany) (28) in

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2 The amino acid sequence of Msp protein can be accessed through NCBI protein data base under NCBI accession number AAB47939 (51).
square sampling zones (25 μm²) immediately beneath the cell membrane.

Cell Viability—Propidium iodide staining was used to determine if Msp was toxic to cells during the duration of the experiments. Flow cytometry analyses were performed as described (36). Briefly, cells were trypsinized and suspended in PBS, pelleted, resuspended in PBS to a cell concentration of 1 × 10⁶/ml, and stained with propidium iodide (10 μl/ml; Calbiochem). After a 5-min incubation, cells were analyzed by flow cytometry.

Data Analysis—Means and S.E. values were calculated for [Ca²⁺], measurements. Calcium measurements were restricted to (i) base-line [Ca²⁺]; (ii) percentage change of the transient [Ca²⁺], above base-line; (iii) net change in [Ca²⁺], above base line; and (iv) time to peak [Ca²⁺]. For continuous variable data, means and S.E. of [Ca²⁺], were computed, and, when appropriate, comparisons between two groups were made with the unpaired Student’s t test with statistical significance set at p < 0.05.

RESULTS

Preparation of Msp—The Msp preparation was highly enriched by sequential detergent extraction and autoproteolysis of T. denticola extracts. We isolated a fraction that contained primarily Msp oligomers, which migrated as a single ∼190-kDa band in SDS-polyacrylamide gel electrophoresis (21) (Fig. 1A). Boiled samples demonstrated a 53-kDa polypeptide that is known to be the monomer (27, 37). The 95-kDa chymotrypsin-like protease that usually associates with Msp in the outer membrane was not detected in the gels; nor was its peptidase activity detected. Immunoblots of the preparation using anti-Msp antibodies showed almost exclusive staining of the prominent 190-kDa oligomer and the 53-kDa monomer in native and heat-denatured preparations, respectively (Fig. 1B), confirming that these bands contained the authentic Msp from T. denticola.

Previous studies demonstrated that Msp binds to fibronectin, fibrinogen, and laminin and associates with epithelial cell surfaces and cytoplasmic membrane proteins (22, 37). When the Msp preparation was incubated with fibroblasts, immunostaining showed that Msp associated with the fibroblast surface (Fig. 1C). After a 30-min incubation, intense Msp staining was clustered in one area of the cell, indicating a capping phenomenon and possible aggregation of ligand-bound receptors.

Msp Induces Ca²⁺ Transients—Single cell ratio fluorimetry was used to test whether the association of Msp with the fibroblast plasma membrane caused perturbation of calcium homeostasis. For all experiments, [Ca²⁺], was stable in untreated cells, even over prolonged time course experiments (>120 min). In calcium buffer, exposure of cells to Msp induced a robust Ca²⁺ transient that often returned to a level ~15–30% higher than base-line within 90 s of the peak (Fig. 2, A–C); subsequent smaller transients could also be detected over a period of 30 min in some cells (Fig. 2B, inset). Treatment of cells with increasing concentrations of Msp (1, 10, 30 μg/ml) showed a dose-response relationship with the amplitude of the calcium peak (Fig. 2H). In control cells treated with the vehicle (distilled H₂O added to cell culture buffer), there was no Ca²⁺ response (Fig. 2D), indicating that the Msp-induced Ca²⁺ responses were not simply due to physical disturbance or an anisomotic change. Further, only cells with bright fura-2 staining in the cytoplasm were included in analyses to overcome the possibility of measuring dead or dying cells (see data on cell death below). In cells challenged with Msp that was preincubated with anti-Msp antibodies, there was no Ca²⁺ response detected upon the addition of Msp (Fig. 2E), indicating that the Msp-induced transient was due to a specific interaction between Msp and the cells. Boiling the Msp eliminated the Ca²⁺ response (Fig. 2F); the amplitude of the Ca²⁺ transient was much lower than controls when the Msp was heated to 60 °C for 30 min (Fig. 2G). Thus, the ability of Msp to interact with cells and to generate Ca²⁺ responses evidently depends on the unadorned protein complex. Finally, cells that were incubated repeatedly with fresh doses of Msp showed progressively diminished calcium responses (Fig. 2J).

To determine if Msp could induce Ca²⁺ transients subjacent to the cell membrane, cells were loaded with fura-C18. In these experiments Msp evoked large amplitude calcium transients, demonstrating a sharp elevation of [Ca²⁺], near the plasma membrane (Fig. 2J). These transients occurred more rapidly than the cytoplasmic calcium signals following Msp treatment (n = 10 experiments; p < 0.05), indicating that Msp may stimulate Ca²⁺ influx through plasma membrane-permeable channels. We determined the source of Ca²⁺ for the observed rise in [Ca²⁺], For the EGTA buffer, Fura-2-loaded cells were incubated in Ca²⁺-free buffer containing EGTA. The cells were rapidly switched to the EGTA buffer prior to the addition of Msp to avoid depletion of intracellular stores. Under these conditions, Msp treatment failed to evoke a defined Ca²⁺ transient (Fig. 2K), indicating that most of the Ca²⁺ for the robust intracellular response was probably derived from influx through plasma membrane-per-
measurable channels. We examined the influx pathway further by performing quenching experiments in which the extracellular bathing buffer was supplemented with 1 mM MnCl₂ and the fluorescence of fura-2 was excited at 356 nm (the isosbestic or calcium-insensitive point) (33). When single emission photon counts of single cells showing a sharp decline in the fluorescence of fura-2 excited at 356 nm ~2 min after the addition of Msp in buffer containing manganese chloride (1 mM). The loss of fluorescence after the addition of Msp was contemporaneous with Msp-induced calcium transients (~90 s; see Fig. 1C). The data are representative of five independent experiments. The inset in L shows Msp-treated cells in medium without manganese chloride.

To assess whether Msp causes a general increase in cation permeability, we measured the membrane potential with the bisoxonol dye bis-[1,3-dibutylbarbituric acid]trimethineoxonol and flow cytometry. Measurement of bis-[1,3-dibutylbarbituric acid]trimethineoxonol fluorescence after vehicle or Msp treatment showed only a 2% reduction of fluorescence, while KCl depolarization in the presence of valinomycin (3 μM) produced large reductions (~25% reductions after each of the sequential additions of 50 mM KCl; 5000 cells/run). Experiments conducted with calcium buffer and ionomycin (2 μM) treatments indicated that cells preincubated with Msp or vehicle responded with an equivalent increase in [Ca²⁺] (p > 0.2; n = 5 cells each). These data indicated that the cell membranes of Msp-treated cells were intact. Consistent with these observations, the basal [Ca²⁺], was unchanged in Msp-treated and vehicle-treated cells (~150 nM).

We determined if Msp is cytotoxic to fibroblasts in the short term (i.e. under the conditions used for these experiments) because of a rise in intracellular Ca²⁺ (38). Propidium iodide staining experiments were done as described under “Experimental Procedures.” Flow cytometry analysis showed that there were ~15% propidium iodide-permeable cells after treatment with Msp (30 μg/ml, 25°C) for 40 min compared with ~9% for control cells (n = 5; p < 0.05). Therefore, most of the cells that were pretreated with Msp remained viable under the conditions of the experiments; those that were permeable to propidium iodide also exhibited less fura-2 staining and were not included in the measurements of [Ca²⁺]. In contrast, after 24 h of exposure to Msp (10 μg/ml, 37°C), there were ~66% propidium iodide-permeable cells after treatment (n = 5; p < 0.05), which is consistent with both the concentration and time course for the cytotoxicity of Msp complex reported previously (39). An alternative viability assay based on a replating and cell attachment method (40) showed no difference in viability...
between control and Msp-challenged cells over a 60-min incubation (data not shown).

Msp Inhibits Ca\(^{2+}\) Release from Internal Stores—Outer membrane extracts of \textit{T. denticola} inhibit agonist-induced increases of [Ca\(^{2+}\)]\(_i\), in human gingival fibroblasts (17). We used two agonists capable of generating internal calcium release to study the effect of Msp pretreatment on calcium release from intracellular stores. ATP (200 \(\mu\)M) or thapsigargin (1 \(\mu\)M) was incubated with cells in calcium-free buffer containing 1 mM EGTA, a protocol that ensures that the induced calcium transients were progressively reversed during increased time periods following wash-out of free Msp from the assay buffer. 

Summary data showing basal [Ca\(^{2+}\)]\(_i\) changes caused by Msp, control cells, and ATP are mean ± S.E. from \(n = 8\) experiments showing maximum [Ca\(^{2+}\)]\(_i\) change (in nM) induced by ATP or thapsigargin treatment. 

Mps Inhibits Ca\(^{2+}\) Release from Internal Stores—Outer membrane extracts of \textit{T. denticola} inhibit agonist-induced increases of [Ca\(^{2+}\)]\(_i\), in human gingival fibroblasts (17). We used two agonists capable of generating internal calcium release to study the effect of Msp pretreatment on calcium release from intracellular stores. ATP (200 \(\mu\)M) or thapsigargin (1 \(\mu\)M) was incubated with cells in calcium-free buffer containing 1 mM EGTA, a protocol that ensures that the induced calcium transients were attributable solely to release from internal calcium stores. Prior to the addition of agonists, the calcium-free buffer was switched immediately before experiments to reduce intracellular calcium depletion. The calcium-deficient media did not significantly reduce basal [Ca\(^{2+}\)]\(_i\) over the time course of these experiments. Data are mean ± S.E. from \(n = 8\) experiments showing maximum [Ca\(^{2+}\)]\(_i\) change (in nM) induced by ATP or thapsigargin treatment. Pretreatment of cells with latrunculin B reverses the decrease in [Ca\(^{2+}\)]\(_i\), change caused by Msp, and summarize data (mean ± S.E.) from \(n = 4–8\) experiments per group showing that the effects of Msp on ATP-induced Ca\(^{2+}\) transients were progressively reversed during increasing time periods following wash-out of free Msp from the assay buffer. D-G, Mag-fura-2 ratio data for estimation of [Ca\(^{2+}\)]\(_i\). Msp has no acute effect on baseline mag-fura-2 ratio (D). After the addition of ATP (200 \(\mu\)M), Msp pretreatment (30 \(\mu\)M) blocks ATP-induced reduction of mag-fura-2 ratio (E). In control cells with vehicle, the mag-fura-2 ratio is transiently reduced and then returns to baseline after refilling of ER stores (F). Summary data showing basal [Ca\(^{2+}\)]\(_i\), and peak ATP-induced change of mag-fura-2 ratio (G; means ± S.E.; data are from \(n = 9\) experiments). Note that pretreatment with Msp reduces the basal mag-fura-2 ratio by −25%.

For estimation of Ca\(^{2+}\) concentration in the ER stores, cells were loaded with mag-fura-2 according to the methods of Hofer et al. (30), and the mag-fura-2 emission ratio was calculated. Fluorescence microscopic examination of these cells showed that the dye was compartmentalized in discrete vesicular zones that closely resembled previously published images (30). Digitonin experiments as well as stimulation with ATP showed that the mag-fura-2 loading protocol was indeed reporting on calcium stores with properties similar to that of the endoplasmic reticulum. Msp produced no abrupt change in the mag-fura-2 ratio (Fig. 3D), but it did cause a slight decline in base-line measurements over the time course of the experiment. Pretreatment of cells with Msp (25 °C, 40 min) followed by incubation with ATP (200 \(\mu\)M) blocked the ATP-induced reduction of
Indeed, both the amplitude (Fig. 4, A and B) and the kinetics of the transients were not different in control and Msp-pretreated cells. Shown are typical calcium signals evoked after thapsigargin (1 µM)-EGTA, showing different kinetics for [Ca\(^{2+}\)] transient integration in control (C) and Msp-pretreated cells (D). The amplitude of the [Ca\(^{2+}\)] transient is diminished, and the Ca\(^{2+}\) release is prolonged in the Msp-pretreated cells. E, summary data (mean ± S.E.) from n = 4–8 experiments per group showing that internal stores of both control and Msp-pretreated cells are comparably depleted of Ca\(^{2+}\) after thapsigargin (1 µM)-EGTA treatment, by measuring subsequent ionomycin (2 µM)-releasable Ca\(^{2+}\) transients.

The mag-fura-2 ratio (Fig. 3E). In contrast, treatment with ATP alone induced a sharp reduction of the mag-fura-2 ratio followed by a rapid recovery to baseline (Fig. 3F), indicating that ATP induced calcium efflux from ER stores. Traces of the mag-fura-2 ratio of single cells (Fig. 3, E and F) and summary analyses (Fig. 3G) showed that pretreatment with Msp lowered the basal mag-fura-2 ratio and the rate or amount of Ca\(^{2+}\) release from the ER stores upon ATP challenge. Collectively, these data seemed to suggest that Msp may block calcium release or affect the kinetics of depletion and replenishment cycles of internal Ca\(^{2+}\) stores.

Alternatively, we considered that the combined effects of Msp and ER agonists may have so completely depleted intracellular stores that there was no more calcium to be released after ATP or thapsigargin treatment. Accordingly, we performed experiments in which cells were rapidly switched over to an EGTA (1 mM), calcium-free buffer prior to Msp or vehicle treatment and then treated with ionomycin (2 µM). In contrast to the ATP- or thapsigargin-induced response (Fig. 3, A and B; also see Fig. 4, C and D), the ionomycin-triggered Ca\(^{2+}\) transients were not different in control and Msp-pretreated cells. Indeed, both the amplitude (Fig. 4, A and B; summary analyses: vehicle = 263 ± 22 nm; Msp = 250 ± 25 nm; p > 0.2; n = 4 per treatment) and the kinetics of the transients were similar in vehicle- and Msp-pretreated fibroblasts. These results indicate that the inhibitory effect of Msp on the release of Ca\(^{2+}\) from the stores was not due to depletion of the stores prior to adding the agonists. Rather, Msp appears to specifically reduce the rate and/or the amount of agonist-induced Ca\(^{2+}\) release, per se. To determine whether Msp affected the total amount of the released Ca\(^{2+}\) or the kinetics of the release, we further analyzed the thapsigargin-induced [Ca\(^{2+}\)] transient. To this end, the cells were incubated in the presence of EGTA (1 mM) and then challenged by thapsigargin. Finally, to assess the amount of releasable Ca\(^{2+}\) after thapsigargin treatment, ionomycin (2 mM) was added. The total Ca\(^{2+}\) released by thapsigargin, determined by measuring the area under the curve of [Ca\(^{2+}\)] traces, was only marginally lower for the Msp-pretreated cells (vehicle = 9.13 ± 0.25 × 10\(^{-9}\) nM·s; Msp = 7.75 ± 0.19 × 10\(^{-6}\) nM·s; p < 0.005). Yet, Msp pretreatment clearly reduced the amplitude of the Ca\(^{2+}\) transient and slowed down the rate of Ca\(^{2+}\) release and return of the transient to baseline levels (Fig. 4, C and D). Moreover, the thapsigargin-sensitive Ca\(^{2+}\) stores of the Msp-pretreated and control cells were comparably depleted, as shown by the similar residual Ca\(^{2+}\) release upon the addition of ionomycin (Fig. 4E). Taken together, this series of experiments indicates that Msp interferes with the process of Ca\(^{2+}\) release from ER stores, seen as the lower amplitude and altered kinetics of [Ca\(^{2+}\)] transients, despite its minimal effect on the amount of Ca\(^{2+}\) released over time.

**Msp Inhibits Store-operated Calcium Influx**—We determined if Msp may inhibit the refilling of internal Ca\(^{2+}\) stores by blocking the influx of extracellular calcium following depletion of these stores. Cells were pretreated with thapsigargin, and the medium was augmented with EGTA (1 mM) to deplete the stores and prevent refilling. As shown above, after sufficient
time, the ER stores were similarly depleted in control and Msp-pretreated cells. After the addition of CaCl_2 (2 mM), [Ca^{2+}]), rose to a peak and then returned to a somewhat higher base-line (Fig. 5A). The maximum influx of Ca^{2+} was diminished >2-fold (p < 0.05) in Msp-treated cells (Fig. 5, B and C). This finding suggests that Msp may have inhibited the influx of extracellular calcium through SOCs and uncoupled the feedback connection between internal Ca^{2+} release and store-operated calcium entry.

**Msp Induces Actin Reorganization in Human Gingival Fibroblasts**—Since *T. denticola* causes actin rearrangement in fibroblasts (11), we determined if Msp may inhibit activation of SOCs by promoting actin assembly (5, 7). Cells were pretreated with latrunculin B (1 μM) to prevent subcortical actin assembly and incubated with vehicle or Msp, and then intracellular calcium was depleted with thapsigargin and EGTA as described for cells in Fig. 5A. Under these conditions, internal Ca^{2+} stores of Msp-pretreated and control cells were comparably depleted. The amplitude of acute Ca^{2+} transients in response to Msp (30 μg/ml) was unaffected by latrunculin B (maximum [Ca^{2+}]), change (in nM): control cells, 258.8 ± 50.0; latrunculin B-pretreated cells, 221.7 ± 15.9; n = 4 per treatment). Yet, under the same conditions, Msp barely inhibited Ca^{2+} depletion-induced Ca^{2+} influx (Fig. 5, D–F) when compared with its strong inhibition in cells that were not treated with latrunculin B (Fig. 5C). Since these data indicated that the mechanism by which Msp uncouples SOCs is related to its effects on actin assembly, we used confocal microscopy to examine the effects of Msp on the distribution of actin filaments (Fig. 5, G–I). After a 5-min treatment with Msp (30 μg/ml), there was an almost complete loss of rhodamine-phalloidin-stained actin filaments in stress fibers and a greatly increased staining in the subcortical region. After treatment of cells with Msp for 45 min induces almost complete depolymerization of stress fibers and the formation of a dense subcortical actin filament network (I).
T. denticola Msp Uncouples Store-operated Ca\textsuperscript{2+} Channels

The central finding in this investigation is that the major surface protein of the spirochete T. denticola promotes subcortical actin assembly that subsequently perturbs calcium signaling in human fibroblasts. Since calcium signals are tightly and reciprocally linked to regulation of the actin cytoskeleton (2), microbial perturbation of this system has profound effects on cell regulation. Not only is the activity of actin-binding and -severing proteins determined by local concentrations of calcium, but the assembly of subcortical actin filaments evidently affects the coupling of two key components of store-operated calcium flux: depletion/refill cycles of ER store Ca\textsuperscript{2+} and Ca\textsuperscript{2+} entry through SOCs (5–7). Since store-operated flux is a major pathway by which intracellular Ca\textsuperscript{2+} concentration is regulated in response to extracellular agonists, it occurred to us that pathogenic microorganisms may perturb host cells by affecting their store-operated calcium pathways. In contrast to the exotoxins or surface proteins of bacteria that stimulate host cells as a target cell and Msp as a biologically relevant antagonist, we provide corroborating evidence for actin assembly-dependent conformational coupling of SOCs in an additional cell type, perturbs store-operated calcium flux by uncoupling SOCs, and may be due to increased conductance across the plasma membrane, indicating that there was no general increase of cation conductance (e.g. sodium). Thus, it appears that Msp represents a novel type of bacterial cytotoxin whose immediate pathogenic effect is the specific promotion of increased calcium conductance.

Longer term experiments (~30 min) showed that Msp reduced ATP- and thapsigargin- induced [Ca\textsuperscript{2+}], transients and blocked subsequent Ca\textsuperscript{2+} entry following thapsigargin treatment, indicating that Msp uncouples physiologic store-operated calcium flux. These findings were not due to prior store depletion, since ionomycin induced immediate and equivalent calcium release in control and Msp-pretreated cells in calcium-free buffer. Further experiments with ionomycin showed that thapsigargin-sensitive Ca\textsuperscript{2+} stores were comparably depleted in control and Msp-pretreated cells. Yet, the amplitude of thapsigargin-induced calcium transients was clearly decreased, and the kinetics of Ca\textsuperscript{2+} release were clearly altered in the Msp-pretreated cells. The mechanisms accounting for this unprecedented effect of a bacterial toxin are not entirely clear. They may involve perturbation of Ca\textsuperscript{2+} release and/or reuptake into ER or other internal Ca\textsuperscript{2+} stores and may be affected by de novo actin filament assembly near the ER and other calcium-containing organelles. Notably, the reduction of thapsigargin- and ATP- induced Ca\textsuperscript{2+} transients by Msp was reversed in latrunculin B-pretreated cells. Thus, the altered kinetics of Ca\textsuperscript{2+} release and/or replenishment of ER stores following incubation of Msp-treated cells with agonists may be caused, in part, by actin filament reorganization. Evidently, the key element in the perturbation of store-operated calcium flux by Msp is that it induces a calcium-dependent assembly of subcortical actin filaments in gingival fibroblasts, suggesting that Msp may uncouple the ER and Ca\textsuperscript{2+} channels in the plasma membrane.

There is growing experimental support for conformational coupling as a principal regulatory mechanism that links Ca\textsuperscript{2+} store depletion and Ca\textsuperscript{2+} influx through SOCs (1). Experimental strategies using drugs that affect actin assembly, translocation of actin to the cell periphery, and the association of actin and actin-binding proteins with the plasma membrane support the hypothesis that de novo assembly of an actin filament network subjacent to the plasma membrane impairs conformational coupling (5, 7). We have shown previously that T. denticola induces rearrangement of actin (11, 12, 15), with a proportional shift in actin filaments from the stress fiber-rich ventral interface to the dorsal area of the cell (14). Here, we observed that the Msp-induced calcium transient is required for subcortical actin filament assembly, and this actin network is formed contemporaneously with the block of store-operated calcium flux. However, in cells pretreated with latrunculin B to inhibit actin filament assembly, both Ca\textsuperscript{2+} release from thapsigargin-sensitive stores and Ca\textsuperscript{2+} entry following Ca\textsuperscript{2+} store depletion were hardly affected upon challenge with Msp. We interpret these findings to indicate that the Msp of T. denticola perturbs store-operated calcium flux by uncoupling SOCs, and we provide corroborating evidence for actin assembly-depend- ent conformational coupling of SOCs in an additional cell type, the human gingival fibroblast.

The binding of many pathogenic bacterial species or their toxins to host cells can elicit intracellular calcium signals that are often associated with substantial cytoskeletal rearrangement (44–48). Some bacterial outer membrane porins can even translocate to the plasma membrane of target cells and induce apoptosis secondary to the rapid influx of Ca\textsuperscript{2+} (49, 50). For another example, cholera toxin increases Ca\textsuperscript{2+} current through calcium release-activated calcium channels in mast cells (11). Therefore, pathogens may induce cytotoxicity by causing ab-
normally elevated cytoplasmic Ca\textsuperscript{2+} levels in target cells, which is one of the acute effects of \textit{T. denticola} outer membrane proteins (17), specifically Msp. Although presently unique, our central finding is that the major surface protein of the spirochete \textit{T. denticola} inhibits calcium flux by uncoupling store-operated calcium channels. Conceivably, this may well become a more general theme of pathogenic host-parasite interactions. Indeed, our data suggest a general mechanism by which pathogens can block one of the key cellular signaling pathways for responses to natural agonists such as growth factors, extracellular matrix proteins, cytokines, and a variety of other mediators of intercellular communication networks. Pharmacological intervention to block this pathway of bacterial pathogenicity may provide a novel approach to inhibit the cytotoxic effects of some species of spirochetes and other pathogens that cause chronic disease.

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