

Absence of a Human DnaJ Protein hTid-1_S Correlates with Aberrant Actin Cytoskeleton Organization in Lesional Psoriatic Skin*

Received for publication, October 14, 2011, and in revised form, May 17, 2012. Published, JBC Papers in Press, June 12, 2012, DOI 10.1074/jbc.M111.313809

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Background: HSP27 phosphorylation plays pivotal roles on F-actin polymerization and actin cytoskeleton organization.

Results: The loss of hTid-1_S expression was observed in lesional human psoriatic skin.

Conclusion: The binding of hTid-1_S with MK5 inhibits HSP27 phosphorylation and attenuates F-actin polymerization.

Significance: The lack of hTid-1_S expression correlates with the aberrant actin cytoskeleton organization in psoriatic human skin.

The biochemical mechanism by which the human tumorous imaginal disc_S (hTid-1_S) interferes with actin cytoskeleton organization in keratinocytes of human skin epidermis was investigated. We found that hTid-1, specifically hTid-1_S, interacts with MK5, a p38-regulated/activated protein kinase, and inhibits the protein kinase activity of MK5 that phosphorylates heat shock protein HSP27 in cultured HeLa cells. Thus, hTid-1_S expression inhibits the phosphorylation of HSP27 known to play important roles in F-actin polymerization and actin cytoskeleton organization. The interplay between MK5/HSP27 signaling and hTid-1_S expression was supported by the inhibition of HSP27 phosphorylation and MK5 activity in HeLa cells in response to hypoxia during which hTid-1_S expression was down-regulated. We also found that overexpression of hTid-1_S results in the inhibition of HSP27 phosphorylation, F-actin polymerization, and actin cytoskeleton organization in transduced HaCaT keratinocytes. This study further proposes that the loss of hTid-1_S expression in the basal layer of skin epidermis correlates with the enhanced HSP27 phosphorylation, keratinocyte hyperproliferation, and excess actin cytoskeleton organization in lesional psoriatic skin.

hTid-1² was first identified as a human homologue of *Drosophila* tumor suppressor protein Tid56 (1). hTid-1 belongs to the DnaJA3 family of proteins known to interact with HSP70 family proteins (2). At least two isoforms of hTid-1, namely, hTid-1_L and hTid-1_S, have been reported. Both isoforms of hTid-1 containing amino-terminal mitochondrial signal sequence reside mainly in the mitochondrial matrix. However,

these two isoforms of hTid-1 can interact with many cytoplasmic proteins when they are transiently retained in the cytoplasm before they transport to the mitochondria (3). The two alternative splice variants have opposing effects on the decision of cell fate. It has been reportedly known that proapoptotic hTid-1_L stimulates cytochrome *c* release from the mitochondria whereas antiapoptotic hTid-1_S inhibits cytochrome *c* release resulting in the inactivation of caspase-3 and subsequent inhibition of apoptosis in human osteosarcoma cells (4).

Using yeast two-hybrid screening (5), we have identified hTid-1_S as a new binding protein that interacts with mitogen-activated protein kinase (MAPK)-activated protein kinase 5 (MK5). MK5 and its human homologue p38-regulated/activated kinase (PRAK) belong to a Ca²⁺/calmodulin-dependent protein kinase and display approximately 40% amino acid sequence identity with other p38 MAPK-activated protein kinases MK2 and MK3 (6–8). In response to cellular stresses such as heat shock, UV light, oxidation, and proinflammatory cytokines, the activation phosphorylation site of MK5 at Thr-182 is phosphorylated by p38 MAPK and extracellular signal-regulated kinases (ERKs) (9–12). Studies have identified substrate proteins for MK5 including tumor suppressor p53 (13), human small heat shock protein HSP27, and its murine HSP25 homologue. As the phosphorylation of HSP27 is required for F-actin polymerization, MK5-induced HSP27 phosphorylation participates in actin filament dynamics and stabilization of actin cytoskeleton (14–16).

The epidermis of skin maintains its barrier function through continuous self-renewal and terminal differentiation processes. During these processes, keratinocytes leave the basal cell layer, proliferate, lose metabolic activities, and finally differentiate into dead cornified cells in outmost layer of the epidermis (17). To control the balance between self-renewal and terminal differentiation in the epidermis of skin, cytoskeleton remodeling should accompany the cellular events such as cell-cell and cell-

* This work was supported by National Research Foundation of Korea Grants 2010-0011268 and 2009-0085835 (to C. O. J.).

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² The abbreviations used are: hTid-1, human tumorous imaginal disc1; MK5, MAPK-activated kinase protein 5.

matrix adhesion, cell proliferation, and cell migration. Psoriasis is a common chronic inflammatory skin disease that is characterized by a marked hyperproliferation and altered differentiation of keratinocytes. This disease affecting ~1–3% of the US population (18) is commonly believed to have strong genetic components with altered expression of >1,300 genes in lesions of psoriatic skin (19, 20). This study delineates a biochemical mechanism by which the absence of hTid-1_s expression leads to aberrant actin filament dynamics in lesional psoriatic skin. Here, we propose that the absence of a hTid-1_s expression (3) stimulates the activity of MK5, increases the phosphorylation of HSP27, and enhances actin cytoskeleton organization in the hyperthickened epidermis of psoriatic skin.

EXPERIMENTAL PROCEDURES

Reagents and Antibodies—TNF- α and antibodies against tubulin and FLAG epitope were obtained from Sigma. Polyclonal rabbit antibody directed against MK5 (H-180) and monoclonal mouse antibody against hTid-1 (RS-11) or HSP27 (F-4) were supplied by Santa Cruz Biotechnology. Polyclonal rabbit antibody against phospho-HSP27 (Ser-82) was from Stressgen. [γ -³²P]ATP and protein G-Sepharose Fast Flow Resin were provided by Amersham Biosciences. Rhodamine-conjugated phalloidin was purchased from Invitrogen.

DNA Constructs—cDNA encoding MK5 was cloned into BamHI and EcoRV sites of pcDNA3-HA cloning vector (14) to generate HA-MK5 expression vector. A full-length cDNA encoding hTid-1_s or hTid-1_L was cloned into HindIII and BamHI sites of pFLAG-CMV2 cloning vector (Sigma) to generate expression vector for FLAG-hTid-1_s or FLAG-hTid-1_L. To generate recombinant adenoviral vector for FLAG-hTid-1_s, cDNA encoding FLAG-hTid-1_s was PCR-amplified and cloned into pAd/CMV/V5-DEST adenoviral vector (Invitrogen Gateway System). The resulting Ad FLAG-hTid-1_s expression vector was linearized by digestion with PacI, transfected, and amplified in 293A cells. The viral particles were purified by centrifugation in cesium chloride gradients.

Cell Culture, Transfection, Cell Migration, and Cell Quantification—Human cervical cancer HeLa cells or immobilized human keratinocyte HaCaT cells were maintained in Dulbecco's modified Eagle's medium supplemented with 10% fetal bovine serum (Invitrogen) in a humidified atmosphere of 5% CO₂ at 37 °C. HeLa cells were transfected with indicated vector constructs using Lipofectamine reagent (Invitrogen). HaCaT keratinocytes was infected with adenoviruses harboring recombinant adenoviral vector encoding FLAG-hTid-1_s for 48 h. Cell migration assay was carried out employing a QCMTM Cell Migration Assay (Chemicon). The growth of HaCaT keratinocytes was measured by crystal violet staining according to Kueng *et al.* (21).

Immunoprecipitation—HeLa cells (~10⁷) were lysed in EBC lysis buffer (120 mM NaCl, 0.5% Nonidet P-40, 50 mM Tris-Cl, pH 8.0, 100 mM NaF, 200 mM sodium orthovanadate) containing 1 mM PMSF. Cell preparations were centrifuged at 4 °C for 10 min at 10,000 \times g, after which the supernatant constituted the cell lysate. The cell lysate was incubated with MK5 antibody for 4 h at 4 °C before the addition of 30 μ l of 50% protein G-Sepharose equilibrated with EBC lysis buffer. After additional incu-

bation for 12 h at 4 °C, beads were washed four times with the lysis buffer. Immunocomplexes were eluted from the resin by boiling in a sodium dodecyl sulfate (SDS) sample buffer (250 mM Tris-Cl, pH 6.8, 40% glycerol, 8% SDS, 4% 2-mercaptoethanol, 0.002% bromphenol blue) for 10 min. hTid-1 bound to MK5 was analyzed by 12% SDS-PAGE followed by immunoblot analysis using anti-hTid-1 antibody.

RNA Interference—To construct an expression vector for small hairpin RNA (hTid-1-shRNA), hTid-1-specific sequence containing a hairpin loop (5-GATCCCCAGCTACGGCTACGGAGACTTCAAGAGAGTCTCCGTAGCCGTAGCTGTT-TTTGGAAA-3) was cloned into the BglIII and HindIII sites of p3in1 vector (22). HeLa cells (2 \times 10⁶) were transfected with indicated amount of hTid-1-shRNA or p3in1 vector using Lipofectamine reagent. Experiments were carried out 24 h after hTid-1-shRNA transfection.

Immunoprecipitation MK5 Kinase Assay—Control or transfected HeLa cells overexpressing FLAG-hTid-1_s were grown for 24 h under normal or hypoxic conditions. Cells were washed twice in PBS and lysed in EBC lysis buffer. Lysates were centrifuged for 10 min at 10,000 \times g at 4 °C. MK5 protein kinase in the supernatant was immunoprecipitated using anti-MK5-protein G-Sepharose. Beads were mixed for 4 h at 4 °C and washed three times with the lysis buffer. Human recombinant HSP27 substrate protein (Abcam) (3 μ g) and beads containing immunoprecipitated MK5 were added to a total volume of 30 μ l of reaction mixture containing 50 mM Tris-HCl, pH 7.5, 10 mM MgCl₂, 0.1 mM ATP, 1 μ g/ μ l BSA, and 10 μ Ci of [γ -³²P]ATP. The reaction mixture was incubated for 20 min with shaking at 30 °C. The reaction was stopped by heating at 95 °C for 5 min after the addition of 10 μ l of 4 \times SDS sample buffer. The activity of MK5 as a protein kinase was evaluated by measuring the transfer of ³²P moiety from [γ -³²P]ATP to MK5 and HSP27 substrate protein. Autophosphorylation of MK5 and HSP27 phosphorylation was detected by autoradiography followed by 12% SDS-PAGE.

Staining of Actin Cytoskeleton—F-actin staining for immunofluorescence microscopy was performed as described previously (14). In brief, HeLa or HaCaT cells were fixed with 3.7% formaldehyde in PBS at room temperature, permeabilized with PBS containing 0.1% Triton X-100 at -20 °C, and blocked with 1% bovine serum albumin (BSA) in PBS. F-actin was stained with rhodamine-conjugated phalloidin (3 μ g/ml) in PBS containing 1% BSA for 20 min. F-actin polymerization was examined by monitoring stress fiber formation under a confocal scanning microscope.

Biopsies—Psoriasis patients and normal controls participated voluntarily and gave their written informed consents for this study. This study was approved by Institutional Review Board of each respective institution. Biopsies from the lesional plaque-type psoriatic skin were taken from the center of the plaque from psoriasis patients diagnosed and selected at the Department of Dermatology, Chungnam National University Hospital, South Korea, as described by Johansen *et al.* (23). Normal skin was obtained from skin transplants of patients. All epidermal tissue samples included the dermis and the epidermis. For immunofluorescence analysis, biopsies were fixed and

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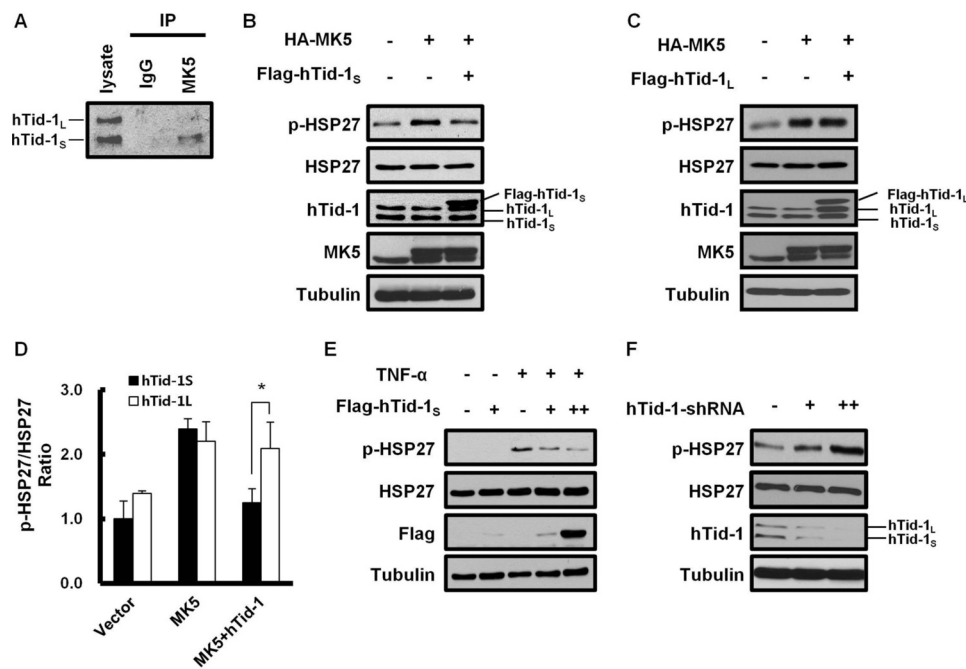


FIGURE 1. hTid-1_S interacts with MK5 to inhibit its kinase activity. *A*, coimmunoprecipitation was performed to prove intracellular interaction between MK5 and hTid-1_S in HeLa cells. Cell lysate proteins were mixed with anti-MK5 antibody. Immunoprecipitated proteins were separated in 12% SDS-PAGE and detected by immunoblot analysis using anti-hTid-1 antibody, which recognizes both hTid-1_S and hTid-1_L. *B* and *C*, HeLa cells (2×10^6) were transfected with expression vector for HA-MK5 (1 μ g) alone or together with expression vector for FLAG-hTid-1_S or FLAG-hTid-1_L (1 μ g each). After incubation for 24 h, cell lysates were resolved in 12% SDS-PAGE. Cellular level of phosphorylated HSP27 (pHSP27) or HSP27 was analyzed by immunoblot analysis using respective antibodies. Expression of FLAG-hTid-1_S, HA-MK5, or tubulin in cell lysates was also monitored by immunoblot analysis. *D*, relative band intensities in Western blot in *B* and *C* were quantified by using ImageJ software and the pHSP27/HSP27 ratio in transfected HeLa cells was determined. The band intensity of pHSP27 of untreated cells was set to 1.0. Values represent means \pm S.D. (*error bars*; $n \geq 3$ experiments). Statistical significance was determined by Student's *t* test. *, $p < 0.05$. *E*, HeLa cells (2×10^6) were transfected with expression vector for FLAG-hTid-1_S (1 or 2 μ g). After incubation for 24 h, transfected cells were further incubated in serum-free medium for 3 h and treated with 100 ng/ml TNF- α for 20 min. Cellular level of endogenous HSP27, pHSP27, tubulin, or ectopically expressed FLAG-hTid-1_S was examined by immunoblot analysis. *F*, HeLa cells (2×10^6) were treated with hTid-1-shRNA (1 or 2 μ g). After incubation for 24 h, cell lysate proteins were resolved in 12% SDS-PAGE. Cellular level of endogenous HSP27, pHSP27, hTid-1, or tubulin was examined by immunoblot analysis using respective antibodies.

embedded in paraffin. Alternatively, samples were frozen in liquid nitrogen and used for Western blot analysis.

Immunohistochemistry—Immunohistochemistry was performed using Chem-Mate EnVision Detection Kit (DAKO, Carpinteria, CA). Tissue sections were deparaffinized and rehydrated. After incubation with 0.3% H₂O₂ for 20 min, sections were placed in 10 mM EDTA, pH 8.0, and heated in a microwave for 5 min. Slides were stained with anti-hTid-1, anti-HSP27, or anti-phospho-HSP27 (Ser-82) antibody at room temperature for 1 h followed by the incubation with the secondary antibody for 30 min. F-actin was visualized by staining slides with rhodamine-conjugated phalloidin for 1 h. Tissue sections were developed with 3,3'-diaminobenzidine and counterstained with Mayer's hematoxylin.

RESULTS

hTid-1_S Interacts with MK5 to Inhibit HSP27 Phosphorylation—A potential interaction between MK5 and hTid-1_S by the yeast two-hybrid assay using MK5 as a bait prompted us to verify their interaction in cultured human cells. Intracellular interaction between the two endogenous proteins in HeLa cells was examined by coimmunoprecipitation analysis. A human DnaJ protein hTid-1_S was coimmunoprecipitated with MK5 (Fig. 1*A*). However, the hTid-1_L isoform, which differs from hTid-1_S only at carboxyl-terminal tail (4), fails to interact with MK5. We explored the possibility that the binding of hTid-1_S

with MK5 interferes with the catalytic activity of MK5. Because HSP27 is a prominent MK5 substrate protein, we attempted to explain the biological role of hTid-1_S interaction with MK5 by examining the effect of hTid-1_S on the phosphorylation of HSP27. Immunoblot analysis demonstrated that FLAG-hTid-1_S expression does not affect the intracellular level of HSP27 but inhibits the phosphorylation of HSP27 induced by the expression of HA-MK5 (Fig. 1, *B* and *D*). In addition, data in Fig. 1, *C* and *D*, indicate that the ability of MK5 to phosphorylate HSP27 was inhibited specifically by hTid-1_S isoform, which interacts with MK5, but not by hTid-1_L isoform. The phosphorylation of HSP27 induced by TNF- α , a proinflammatory cytokine known to activate MK5 (9), was also inhibited by the expression of hTid-1_S (Fig. 1*E*). Transfection of hTid-1-shRNA suppressed the expressions of both hTid-1_L and hTid-1_S and increased HSP27 phosphorylation in HeLa cells, supporting the negative role of hTid-1_S expression on the kinase activity of MK5 that phosphorylates HSP27 (Fig. 1*F*).

hTid-1_S Inhibits Actin Cytoskeleton Organization—The phosphorylation of HSP27 has been known to confer stabilization of actin cytoskeleton as well as F-actin polymerization in cultured mammalian cells under stress conditions induced by reactive oxygen metabolites, tumor necrosis factors, growth factors, or heat shock (24, 25). The expression of hTid-1_S, which interacts with MK5, is anticipated to interfere with HSP27

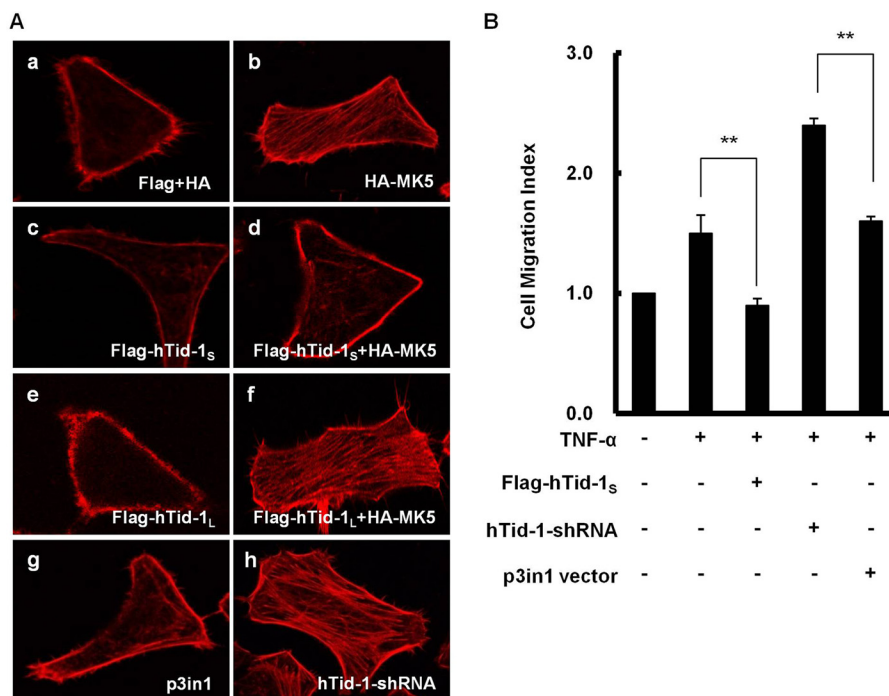


FIGURE 2. Effects of hTid-1_S expression on actin cytoskeleton organization and cell migration. *A*, disruption of MK5-mediated actin cytoskeleton organization by hTid-1_S expression. *a*, HeLa cells (5×10^5) were cotransfected with expression vectors for HA and FLAG (1 μ g each). *b* and *c*, cells were transfected with expression vector for HA-MK5 or FLAG-hTid-1_S (1 μ g each). *d*, cells were cotransfected with expression vectors for HA-MK5 and FLAG-hTid-1_S (1 μ g each). *e*, cells were transfected with expression vector for FLAG-hTid-1_L (1 μ g). *f*, cells were cotransfected with expression vectors for HA-MK5 and FLAG-hTid-1_L (1 μ g each). *g*, and *h*, cells were transfected with p3in1 vector or hTid-1-shRNA (1 μ g each). Transfected or cotransfected HeLa cells were incubated for 24 h, fixed, permeabilized, blocked, stained with rhodamine-conjugated phalloidin for F-actin, and imaged by confocal microscopy. *B*, inhibition of TNF- α -induced cell migration by hTid-1_S. HeLa cells (2×10^6) were transfected with expression vector for FLAG-hTid-1_S, hTid-1-shRNA, or p3in1 vector (1 μ g each). At 24 h after transfection, cells were starved for 24 h. Prior to cell migration experiment, cells were left untreated or treated with 100 ng/ml TNF- α for 20 min. After incubation for 12 h in 6.5-mm Transwell plates, cell migration was measured colorimetrically on a standard microplate reader. Cell migration was expressed as a Migration Index (Chemicon). Cell migration of untreated HeLa cells was set to 1.0. Data represent results from three independent experiments means \pm S.D. (error bars; $n \geq 3$ experiments). **, $p < 0.01$.

phosphorylation by MK5. We examined whether the expression of hTid-1_S plays negative roles on MK5-mediated HSP27 phosphorylation and subsequent actin cytoskeleton organization in HeLa cells coexpressing HA-MK5 and FLAG-hTid-1_S. Actin cytoskeleton organization leading to stress fiber formation was increased by MK5 expression. Although the expression of hTid-1_S decreased MK5-induced stress fiber formation, the expression of hTid-1_L did not influence actin cytoskeleton organization in HeLa cells coexpressing MK5 and hTid-1_L. The ability of hTid-1_S to inactivate MK5/HSP27 signaling was further supported by showing the increased stress fiber formation in HeLa cells transfected with hTid-1-shRNA. The silencing of hTid-1_S by hTid-1-shRNA treatment led to the increase in actin cytoskeleton organization in HeLa cells (Fig. 2*A*). Cell migration is a cellular process that requires F-actin polymerization and actin cytoskeleton stabilization (26). Therefore, we examined whether the expression of hTid-1_S that disrupts actin cytoskeleton organization can affect cell migration. Overexpression of hTid-1_S resulted in the inhibition of TNF- α -induced cell migration, whereas the silencing of hTid-1_S by hTid-1-shRNA stimulated cell migration in HeLa cells treated with TNF- α (Fig. 2*B*).

Activation of MK5/HSP27 Signaling under Hypoxic Conditions—Transcriptional and translational down-regulation of hTid-1_L in cultured human cells under hypoxic conditions has been described previously (27). In this study, we examined

mRNA and protein levels of hTid-1_S in HeLa cells grown under hypoxic conditions. CoCl₂ was chosen as a hypoxia-mimicking agent to maintain hypoxic signal throughout the experiments (28, 29). Intracellular levels of hTid-1_S mRNA and protein were reduced in HeLa cells treated with 200 μ M CoCl₂ for 24 h (Fig. 3, *A* and *B*). Our data also illustrate that phosphorylation of HSP27 was elevated by hypoxia. However, the phosphorylation of HSP27 was severely inhibited by the expression of hTid-1_S in transfected HeLa cells grown under hypoxic conditions (Fig. 3, *C* and *D*). Moreover, immunoprecipitation kinase assay in Fig. 3*E* strongly supports that the increased HSP27 phosphorylation under hypoxic conditions is attributable to the activation phosphorylation of MK5. MK5 phosphorylation as well as HSP27 phosphorylation was elevated in HeLa cells grown under hypoxic conditions. However, the expression of hTid-1_S completely inhibited the phosphorylation of HSP27 and autophosphorylation of MK5 in transfected HeLa cells under hypoxic conditions. Increased stress fiber formation in HeLa cells under hypoxic conditions also implicates that down-regulation of hTid-1_S during hypoxia increases MK5-dependent HSP27 phosphorylation and facilitates actin cytoskeleton organization (Fig. 3*F*). In addition to the stimulation of actin cytoskeleton organization, cell migration was enhanced in HeLa cells by down-regulating hTid-1_S expression during hypoxia (Fig. 3*G*).

Loss of hTid-1_s Expression in Psoriatic Skin

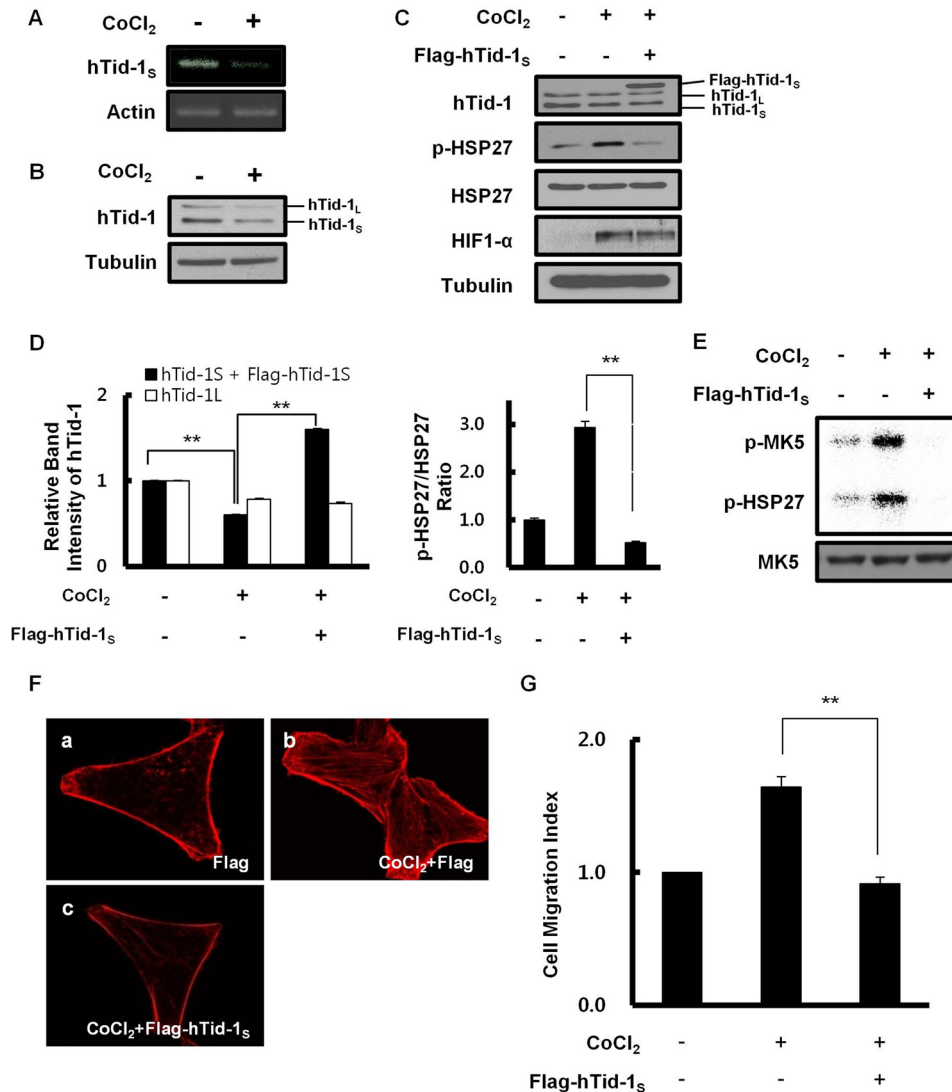


FIGURE 3. Down-regulation of hTid-1_s expression under hypoxic conditions enhances actin cytoskeleton organization. *A*, HeLa cells (2×10^6) were treated with $200 \mu\text{M}$ CoCl_2 for 24 h. Reverse transcription-PCR analysis was carried out using specific primer for hTid-1_s (forward, 5'-TCAGGGTGCAGAAAAGC-CCT-3'; reverse, 5'-CTAGTTCCAGTGGATCTTTTC-3'). β -Actin served as a loading control. *B*, intracellular level of hTid-1_s or hTid-1_L was examined by immunoblotting using anti-hTid-1 antibody. Tubulin was used as a loading control. *C*, HeLa cells (2×10^6) were transiently transfected with expression vector for FLAG-hTid-1_s ($2 \mu\text{g}$). After incubation for 24 h, transfected cells were treated with $200 \mu\text{M}$ CoCl_2 for 24 h. Cellular level of endogenous HSP27, phosphorylated HSP27, hTid-1, HIF1- α , or tubulin was examined by immunoblot analysis. *D*, relative band intensities in Western blot were quantified, and the pHSP27/HSP27 ratio in HeLa cells is presented. The band intensity of hTid-1_s plus FLAG-hTid-1_s or hTid-1_L in untreated cells was, respectively, set to 1.0 (left). The pHSP27/HSP27 ratio of untreated cells was set to 1.0 (right). Data represent mean \pm S.D. (error bars; $n \geq 3$ experiments). **, $p < 0.01$. *E*, HeLa cells ($\sim 10^7$) were left untreated or treated with $200 \mu\text{M}$ CoCl_2 for 24 h. Cell lysate proteins were immunoprecipitated using anti-MK5 antibody. MK5 kinase activity was determined by measuring the transfer of ^{32}P -labeled phosphate moiety from [γ - ^{32}P]ATP to MK5 and HSP27. Activation phosphorylation of MK5 or HSP27 phosphorylation was visualized by autoradiography (upper). Cellular level of MK5 present in total cell lysates was examined by immunoblot analysis using anti-MK5 antibody (lower). *F*, effect of hTid-1_s expression on hypoxia-induced actin cytoskeleton organization was evaluated. *a* and *b*, HeLa cells (5×10^5) were left untreated (*a*) or treated with $200 \mu\text{M}$ CoCl_2 for 24 h (*b*). *c*, HeLa cells (5×10^5) were transfected with expression vector for FLAG-hTid-1_s ($1 \mu\text{g}$). After incubation for 24 h, cells were treated with $200 \mu\text{M}$ CoCl_2 for 24 h. F-actin was visualized after staining cells with rhodamine-conjugated phalloidin. *G*, effect of hTid-1_s expression on hypoxia-induced cell migration is shown. HeLa cells were transfected with expression vector for hTid-1_s. After incubation for 24 h, control or transfected HeLa cells were treated with $200 \mu\text{M}$ CoCl_2 for 24 h. Cell migration was expressed as a Migration Index as in Fig. 2*B*. Data represent mean \pm S.D. (error bars; $n \geq 3$ experiments). **, $p < 0.01$.

Absence of hTid-1_s Expression and Aberrant Actin Cytoskeleton Organization in Psoriatic Human Skin—There have been studies showing the involvement of activated MAPK p38 (30) and MAPK-activated protein kinase that phosphorylates HSP27 (31) in hyperproliferative epidermis of psoriatic skin. Our experimental results in Fig. 4 imply that hTid-1_s expression negatively regulates HSP27 phosphorylation and actin cytoskeleton organization in human skin epidermis. A comparative analysis of hTid-1 expression between lesional psoriatic

and normal skin was carried out. Monoclonal mouse antibody employed in this study recognizes both hTid-1_s and hTid-1_L. In normal skin, hTid-1 expression was focal in the basal layer of epidermis (Fig. 4*Aa*). On the contrary, hTid-1 expression was not detected in the epidermis of psoriatic skin (Fig. 4*Ab*). HSP27 was expressed in both epidermis of normal and hyperthickened psoriatic skin (Fig. 4*A, e* and *f*). The localization of phosphorylated HSP27 that stabilizes actin cytoskeleton (24, 25) markedly differs in normal and lesional psoriatic skin. Phos-

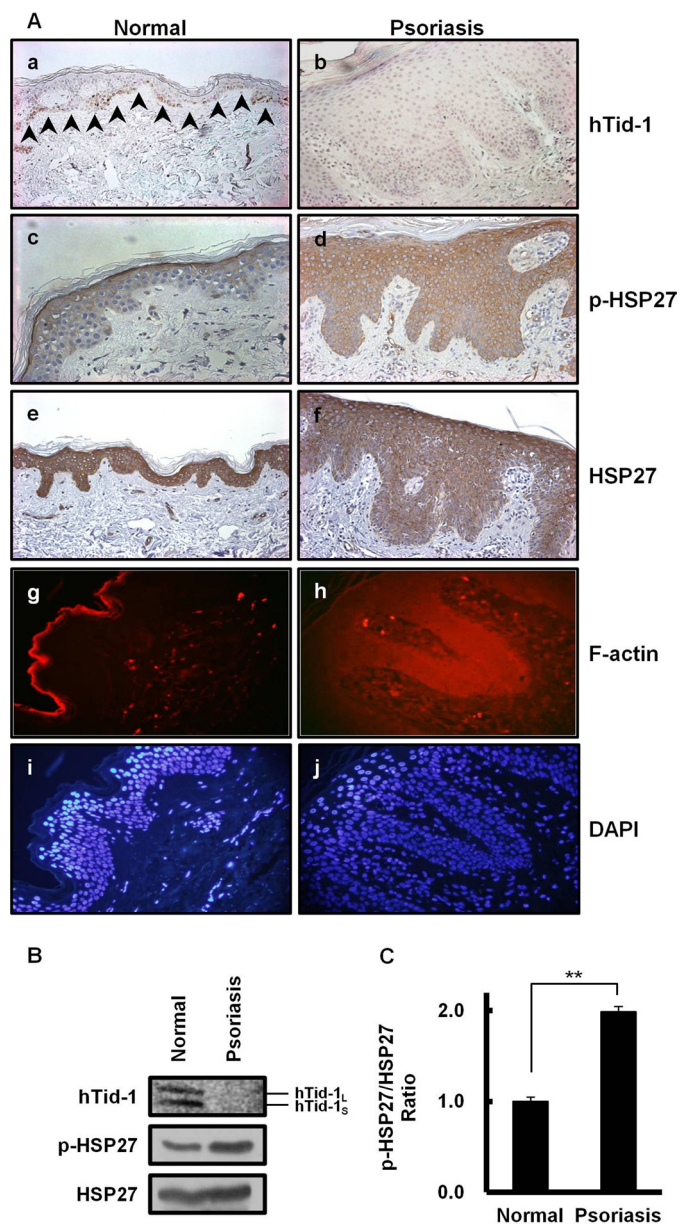


FIGURE 4. Absence of hTid-1 expression and enhanced actin cytoskeleton organization in lesional psoriatic human skin. *A*, sections of skin epidermis from normal controls (*a*, *c*, *e*, and *g*) and lesional psoriatic skin (*b*, *d*, *f*, and *h*) were immunostained with anti-hTid-1, anti-HSP27, or anti-phospho-HSP27 antibody. Arrowheads indicate positive staining of cytosolic hTid-1 in the basal layer of the normal epidermis (*a*). Sections of normal (*g*) and lesional psoriatic skin (*h*) were immunostained with rhodamine-conjugated phalloidin for F-actin. The cyan color corresponds to DAPI staining of nuclei of cell in *g* (*i*) and *h* (*j*) (original magnification, $\times 400$). *B*, HSP27 phosphorylation is significantly enhanced in lesional psoriatic skin lacking hTid-1_s expression. Immunoblotting was performed on cell extracts prepared from the epidermis of keratome biopsies from normal (three different subjects) or lesional psoriatic skin (three different psoriatic patients). Data are the representative Western blots of hTid-1, HSP27, and phosphorylated HSP27 in human keratinocytes. *C*, band intensities in Western blots were quantified. The pHSP27/HSP27 ratio in the epidermis from psoriatic skin was compared with that from normal controls. The pHSP27/HSP27 band intensity ratio in the epidermis from normal skin was set to 1.0. Values represent means \pm S.D. (error bars; $n \geq 3$ samples). **, $p < 0.01$.

phorylated HSP27 was restricted to the granular layer of normal skin, whereas phosphorylated HSP27 was localized in the entire hyperthickened epidermis of lesional psoriatic skin (Fig. 4*A*, *c* and *d*). In normal skin, actin filament formation was observed

extensively in the granular layer of epidermis where phosphorylated HSP27 is present (Fig. 4*Ag*). In psoriatic skin, F-actin polymerization and stress fiber formation were intensively stimulated throughout the whole hyperthickened epidermis including basal layer, stratum spinosum, and granular layer (Fig. 4*Ah*). To further support the reciprocal relationship between hTid-1_s expression and HSP27 phosphorylation, the levels of HSP27 and its phosphorylation form in the epidermal tissue from lesional psoriatic skin were compared with those from normal controls. To evaluate the important role of hTid-1_s expression on HSP27 phosphorylation, the pHSP27/HSP27 ratio in the epidermis from psoriatic human skin was compared with that from normal control by immunoblotting using HSP27 as a loading control. Western blot analysis ascertained the loss of hTid-1_s expression together with the enhanced HSP27 phosphorylation in the epidermis from lesional psoriatic skin (Fig. 4*B*).

Inhibition of F-actin Polymerization by hTid-1_s Expression in Human HaCaT Keratinocytes—To examine whether hTid-1_s expression also attenuates actin cytoskeleton organization in the epidermis of human skin, we evaluated the effect of hTid-1_s expression on MK5/HSP27 signaling in cultured human HaCaT keratinocytes. Ectopic expression of hTid-1_s in HaCaT cells infected with recombinant adenoviruses was confirmed by immunoblotting (Fig. 5*A*). The expression of hTid-1_s in HaCaT keratinocytes interfered with the phosphorylation of HSP27 induced by TNF- α (Fig. 5*B*). Overexpression of hTid-1_s inhibited F-actin polymerization, resulting in reduced stress fiber formation in transduced HaCaT cells treated with TNF- α (Fig. 5*C*). Cell migration induced by TNF- α was also inhibited by hTid-1_s expression in HaCaT cells (Fig. 5*D*). As actin cytoskeleton rearrangement is required for cell proliferation as well as cell motility (32), we examined whether hTid-1_s expression affects proliferation of HaCaT keratinocytes. The retarded keratinocyte proliferation was accompanied by the reduced stress fiber formation in transduced HaCaT keratinocytes overexpressing hTid-1_s (Fig. 5*E*).

DISCUSSION

The p38 MAPK signal pathway has been shown to participate in F-actin polymerization, actin cytoskeleton organization, and actin filament dynamics in various types of cells (33, 34). F-actin organization by activated p38 MAPK has been known to be largely mediated by MK2, which phosphorylates HSP27 (35, 36). Recently, we and others reported that MK5 is also involved in F-actin polymerization and actin cytoskeleton organization (14, 15). Both MK5 and MK2 phosphorylate their target protein HSP27, which plays important roles in the regulation of F-actin polymerization. Here, we propose that hTid-1_s is a new regulator that controls HSP27 phosphorylation. A coimmunoprecipitation assay provides evidence for the intracellular binding of hTid-1_s with MK5. We found that hTid-1_s interacts with MK5 and inhibits HSP27 phosphorylation in HeLa cells (Fig. 1). Despite functional and structural similarities between MK5 and MK2, hTid-1_s failed to interact with MK2 (data not shown). The binding ability of hTid-1_s with MK5 seems to be important for the modulation of MK5 kinase activity because the phosphorylation of HSP27 in transfected HeLa cells was not affected

Loss of hTid-1_S Expression in Psoriatic Skin

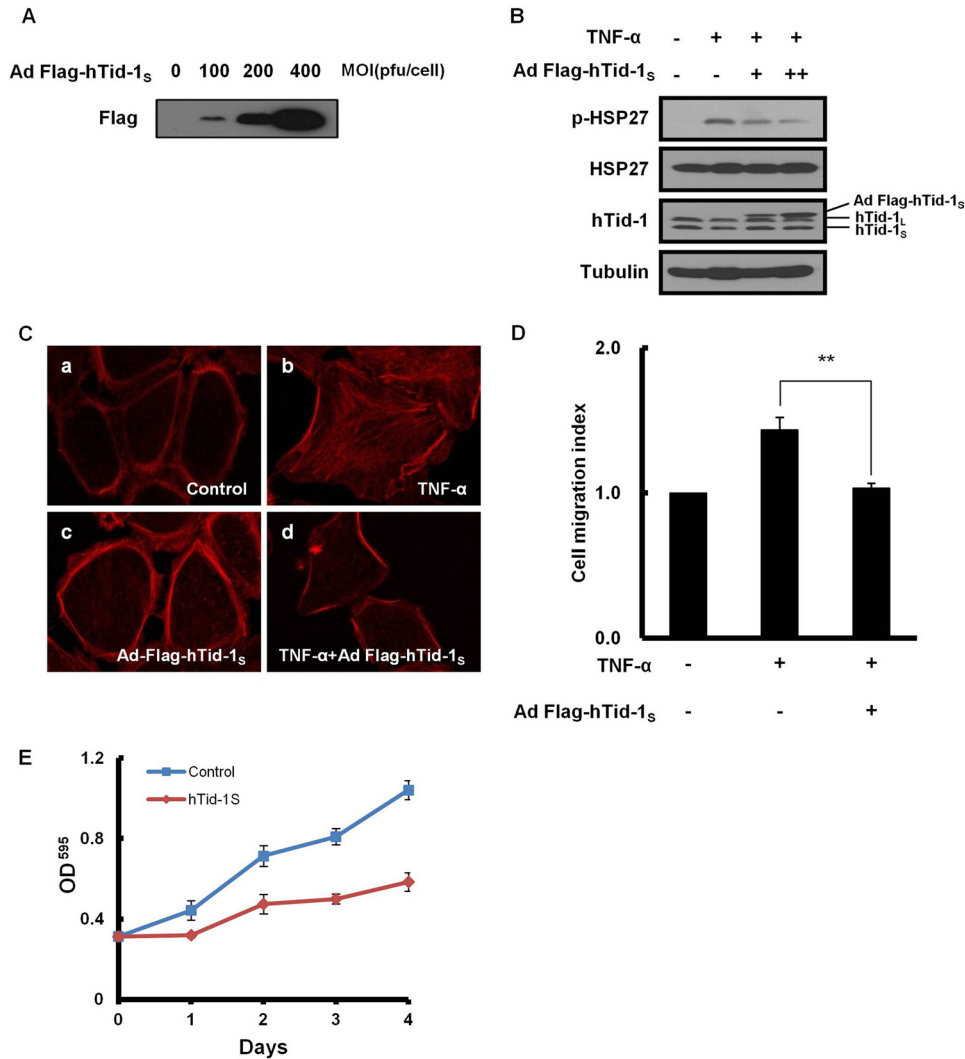


FIGURE 5. Inactivation of MK5/HSP27 signaling by hTid-1_S expression in human HaCaT keratinocytes. *A*, ectopic expression of hTid-1_S in HaCaT keratinocytes infected with recombinant adenoviruses expressing FLAG-hTid-1_S (100, 200, or 400 pfu/cell) examined by immunoblotting using antibody for FLAG epitope. *B*, inhibition of HSP27 phosphorylation by hTid-1_S overexpression in HaCaT keratinocytes treated with TNF-α (100 ng/ml). Cell extracts prepared 48 h after viral infection (100 or 200 pfu/cell) were subjected to immunoblot analysis for the level of HSP27 or pHSP27. *C*, transduced HaCaT keratinocytes examined for the effect of hTid-1_S overexpression on TNF-α-induced actin cytoskeleton organization. *D*, inhibition of TNF-α-induced cell migration in HaCaT keratinocytes. HaCaT keratinocytes (2×10^6) were infected with either recombinant adenoviral vector for FLAG-hTid-1_S (100 pfu/cell each). At 24 h after infection, cells were starved for 24 h. Cell migration was examined as in Fig. 2*B*. *E*, retarded HaCaT keratinocyte proliferation by hTid-1_S expression. HaCaT cells seeded on 9-well plates (5×10^5) were infected with recombinant adenoviral vector for FLAG-hTid-1_S (100 pfu/cell). After incubation for the stated amount of time, cell number was determined colorimetrically using the crystal violet viability assay. Data represent mean \pm S.D. (*error bars*; $n \geq 3$ experiments).

by the expression of hTid-1_L, which lacks the binding ability with MK5 (Fig. 1*C*).

In addition to ATP-dependent chaperonin activity (37, 38), HSP27 has been known as an actin filament-capping protein that modulates actin polymerization by its state of phosphorylation. Unphosphorylated form of HSP27 inhibits F-actin polymerization. During cellular stress and growth, HSP27 undergoes rapid phosphorylation and promotes F-actin polymerization leading to actin cytoskeleton organization and stress fiber formation (35, 39–41). Immunofluorescence analysis of actin stress fiber formation in HeLa cells coexpressing hTid-1_S and MK5 illustrates the inhibition of MK5-induced F-actin polymerization by hTid-1_S. Enhanced actin cytoskeleton organization by silencing of hTid-1_S using hTid-1-shRNA supports the negative role of hTid-1_S on F-actin polymerization (Fig. 2*A*). Because cell migration is a multistep cellular process

initiated by forming protrusive structures like filopodia, lamellipodia, and invadopodia/podosomes that require F-actin polymerization (42), it is reasonable to anticipate that the inhibition of HSP27 phosphorylation by hTid-1_S expression may well interfere with cell migration. The expression of hTid-1_S in HeLa cells inhibited TNF-α-induced cell migration, but the silencing of hTid-1_S by hTid-1-shRNA treatment enhanced cell migration (Fig. 2*B*), supporting the negative role of hTid-1_S on cell migration.

Previous studies have shown that the organization of actin cytoskeleton and the formation of stress fiber are activated (43), but the expression of hTid-1_L is down-regulated in response to hypoxia in cultured human cells (27). However, our data demonstrate that hTid-1_S is involved in the modulation of actin cytoskeleton organization during hypoxia. Cellular levels of hTid-1_S mRNA and protein were decreased after the treatment

of HeLa cells with hypoxia-mimicking agent CoCl₂ (Fig. 3, A and B). Our results imply that the decreased hTid-1_S expression under hypoxic conditions enhances HSP27 phosphorylation and mediates the activation of MK5/HSP27 signal pathway (Fig. 3, C and D) leading to the enhanced actin cytoskeleton organization. MK5 is a protein kinase that requires autophosphorylation at Thr-182 in the catalytic domain for its activation (9). We found that the activity of MK5 as a protein kinase becomes activated in response to hypoxia, during which hTid-1_S expression is repressed. The activation phosphorylation of MK5 at Thr-182 was proven to be completely inhibited by hTid-1_S expression in transfected HeLa cells treated with CoCl₂ (Fig. 3E). Inhibition of MK5 protein kinase activity by hTid-1_S expression in cells under hypoxic conditions strongly supports our hypothesis that hTid-1_S binding to MK5 inhibits MK5 activity and subsequent HSP27 phosphorylation resulting in the down-regulation of F-actin polymerization. As F-actin polymerization is inhibited by ectopic expression of hTid-1_S during hypoxia (Fig. 3F), cell migration under hypoxic conditions was also affected by hTid-1_S expression as shown in Fig. 3G.

In this study, we attempted to interpret the biological roles of hTid-1_S expression on MK5/HSP27 signaling in association with the etiology of psoriasis skin disease. Rationales for the involvement of hTid-1 signaling in the etiology of psoriasis disease are as follows. (i) The activity of transcription factor NF- κ B is up-regulated in the epidermis of psoriatic lesions (44), but hTid-1 expression represses NF- κ B activity (45, 46). (ii) The expression of HIF1- α , which mediates oxygen homeostasis and angiogenesis under hypoxic conditions, is up-regulated in psoriatic skin (47); however, the expression of hTid-1 inhibits angiogenesis by destabilizing HIF1- α (27). (iii) Th1-derived IFN γ is increased greatly in the epidermis of psoriatic lesions (48), where hTid-1_L and hTid-1_S interact with IFN γ to inhibit its transcriptional competence (49).

The direct bearing of actin cytoskeleton organization on the induction of psoriasis was first raised by Wanger and Sundqvist (50), who showed excess F-actin polymerization in the epidermis of lesional skin from patients with psoriasis. More recently, proteomic analysis has demonstrated that concentrations of cytoskeleton proteins including actin are increased in the plasma samples from patients with psoriasis compared with normal controls, emphasizing the relevance of actin filament dynamics to the etiology of psoriasis (51). The important role of hTid-1_S expression on actin filament dynamics through F-actin polymerization in the epidermis of human skin is exemplified in Fig. 4. Immunohistochemistry data disclose the absence of hTid-1 expression, enhanced HSP27 phosphorylation, and the stimulation of actin cytoskeleton organization in hyperthickened psoriatic epidermis (Fig. 4A). Western blot analysis showing the loss of hTid-1_S expression together with the increased pHSP27/HSP27 ratio in lesional epidermis from patients with psoriasis disease strongly suggests that the aberrant actin cytoskeleton organization correlates with the absence of hTid-1_S expression in the epidermis of psoriatic skin (Fig. 4, B and C). These demonstrations make a fine agreement with the experimental results showing the inhibition of HSP27 phosphorylation and reduced actin cytoskeleton organization by hTid-1_S

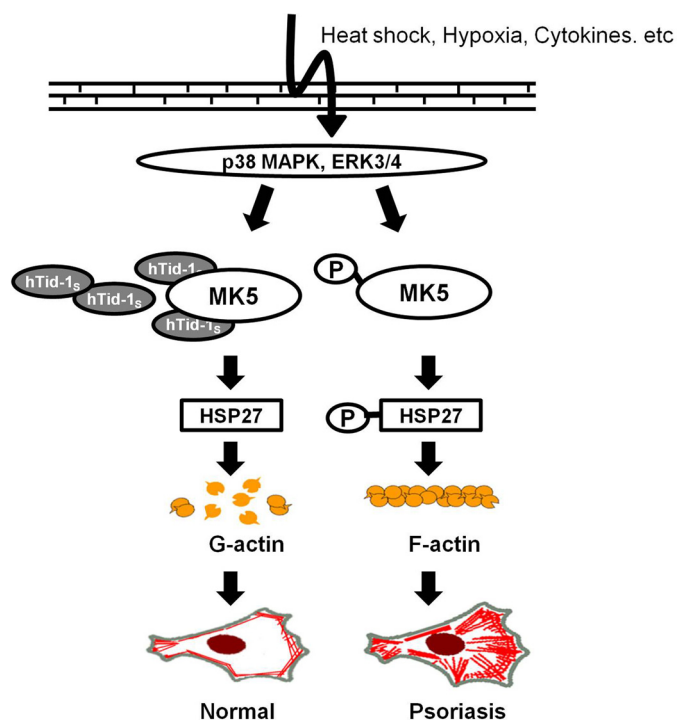


FIGURE 6. Proposed model for the aberrant actin cytoskeleton organization in the epidermis of psoriatic skin. p38 MAPK and ERK activate MK5. The expression of hTid-1_S inhibits the kinase activity of MK5 which phosphorylates HSP27 and attenuates F-actin polymerization in keratinocytes of normal skin. However, the loss of hTid-1 expression in the hyperthickened epidermis of psoriatic skin fails to inhibit MK5 activity, leading to the increased HSP27 phosphorylation, aberrant actin cytoskeleton organization, and hyperproliferation of keratinocytes.

expression in HeLa cells (Fig. 3). From the *in vitro* study using fibroblastic HeLa cells, can it be concluded that the loss of hTid-1_S expression stimulates HSP27 phosphorylation and F-actin polymerization in the epidermis? The correlation between the loss of hTid-1_S expression and the enhancement of HSP27 phosphorylation was also investigated in cultured human keratinocyte HaCaT cells. In accordance with the inactivation of MK5/HSP27 signaling by hTid-1_S expression, actin cytoskeleton organization was inhibited in HaCaT keratinocytes expressing FLAG-hTid-1_S (Fig. 5, A–C). Inhibition of cell migration (Fig. 5D) and antiproliferative potential by hTid-1_S in HaCaT keratinocytes (Fig. 5E) also support that the loss of hTid-1_S expression is involved in the aberrant actin cytoskeleton organization and keratinocyte hyperproliferation in lesional psoriatic skin. Inordinate actin cytoskeleton organization may disrupt cytoskeletal plasticity in the hyperthickened epidermis of psoriatic skin (Fig. 6). Here, we propose hTid-1_S as one of potential targets in understanding the etiology of psoriatic skin disease.

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