

An Acidic Extracellular pH Disrupts Adherens Junctions in HepG2 Cells by Src Kinases-Dependent Modification of E-Cadherin

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ABSTRACT

We have previously shown that culturing HepG2 cells in pH 6.6 culture medium increases the c-Src-dependent tyrosine phosphorylation of β -catenin and induces disassembly of adherens junctions (AJs). Here, we investigated the upstream mechanism leading to this pH 6.6-induced modification of E-cadherin. In control cells cultured at pH 7.4, E-cadherin staining was linear and continuous at cell-cell contact sites. Culturing cells at pH 6.6 was not cytotoxic, and resulted in weak and discontinuous junctional E-cadherin staining, consistent with the decreased levels of E-cadherin in membrane fractions. pH 6.6 treatment activated c-Src and Fyn kinase and induced tyrosine phosphorylation of p120 catenin (p120ctn) and E-cadherin. Inhibition of Src family kinases by PP2 attenuated the pH 6.6-induced tyrosine phosphorylation of E-cadherin and p120ctn, and prevented the loss of these proteins from AJs. In addition, E-cadherin was bound to Hakai and ubiquitinated. Furthermore, pH 6.6-induced detachment of E-cadherin from AJs was blocked by pretreatment with MG132 or NH₄Cl, indicating the involvement of ubiquitin-proteasomal/lysosomal degradation of E-cadherin. An early loss of p120ctn prior to E-cadherin detachment from AJs was noted, concomitant with a decreased association between p120ctn and E-cadherin at pH 6.6. PP2 pretreatment prevented the dissociation of these two proteins. In conclusion, pH 6.6 activated Src kinases, resulting in tyrosine phosphorylation of E-cadherin and p120ctn and a weakening of the association of E-cadherin with p120ctn and contributing to the instability of E-cadherin at AJs. *J. Cell. Biochem.* 108: 851–859, 2009. © 2009 Wiley-Liss, Inc.

KEY WORDS: EXTRACELLULAR ACIDIC PH; ADHERENS JUNCTION; E-CADHERIN; p120 CATENIN; c-Src; FYN KINASE

An acidic extracellular pH (pHe) is a feature of the tumor microenvironment [Tannock and Rotin, 1989; Webb et al., 1999] and has been associated with the invasive behavior of some types of tumor cells [Kato et al., 1992]. Several factors, including type IV gelatinase, metalloprotease-2, metalloprotease-9, cathepsins B and C, and angiogenic factors, have been reported to contribute to this acidic pHe-mediated cell invasion [Rofstad et al., 2006]. However, most studies have focused on the upregulation of metalloproteases under these circumstances.

Disruption of the adherens junction (AJ) is often associated with epithelial mesenchymal transition and cell invasiveness [Murai et al., 2004]. AJs, composed of E-cadherin and catenin (ctn), are responsible

for cell-cell contact. At the AJ, β -ctn binds to the C-terminus of E-cadherin and indirectly links to the actin cytoskeleton via α -catenin (α -ctn) [Takeichi, 1990; Takeichi et al., 1992]. The ctn family consists of α -, β -, γ -, and p120 ctn (p120ctn) [Anastasiadis and Reynolds, 2000]. The integrity of the AJ depends on the stable interactions between the molecules of the cadherin/ctn complex. Expression of cell adhesion molecules is regulated by transcriptional and posttranslational controls. The latter include cadherin endocytosis, cadherin/ctn phosphorylation and dephosphorylation by protein kinases and phosphatases, and the assembly of junctional actin filaments [Akhtar and Hotchin, 2001; Le et al., 2002]. Downregulation of E-cadherin is associated with increased migration and invasiveness of tumor cells

Abbreviations used: AJ, adherens junction; ctn, catenin; DAPI, 4',6-diamidino-2-phenylindole dilactate; IgG-HC, immunoglobulin heavy chain; IgG-LC, immunoglobulin light chain; MTT, 3-[4,5-dimethylthiazol-2-yl]-2, 5-diphenyltetrazolium bromide; PBS, phosphate-buffered saline; p120ctn, p120 catenin; pHe, extracellular pH; PI, propidium iodide; RIPA, radioimmunoprecipitation; siRNA, small interfering RNA; TBS, Tris-buffered saline.

Shih-Horng Huang and Seu-Mei Wang contributed equally to this work.

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[Lu et al., 2003; Guarino, 2007; Herzig et al., 2007]. Tyrosine phosphorylation of E-cadherin results in endocytosis of E-cadherin, disruption of cell-cell contacts, and cell dispersion [Takeichi et al., 1994; Daniel and Reynolds, 1997]. Fujita et al. [2002] reported that c-Src activation induces endocytosis of E-cadherin by ubiquitination of Hakai-bound E-cadherin [Fujita et al., 2002]. Whether this regulatory mechanism is involved in acidic pH-induced AJ disassembly deserves further study.

P120ctn consists of multiple isoforms generated by alternative splicing [Anastasiadis and Reynolds, 2000]. By binding to the juxtamembrane domain of E-cadherin, p120ctn stabilizes E-cadherin at the cell membrane [Reynolds et al., 1994], and small interfering RNA (siRNA) knockdown of p120ctn accelerates cadherin degradation and reduces cell adhesion [Davis et al., 2003]. P120ctn is phosphorylated by Src kinase and receptor-tyrosine kinases [Downing and Reynolds, 1991; Reynolds et al., 1992, 1994] and by serine/threonine kinases, such as protein kinase C (PKC) [Mariner et al., 2001; Xia et al., 2003]. We have previously shown that an acidic pH activates c-Src [Chen et al., 2008], but it is not known whether it induces tyrosine phosphorylation of p120ctn.

In HepG2 cells, an acidic pH induces β -ctn tyrosine phosphorylation by c-Src kinase and reduces the binding of β -ctn to E-cadherin and subsequently disrupts AJs [Chen et al., 2008]. However, whether E-cadherin and p120ctn in HepG2 cells are affected by an acidic pH remains to be explored. We hypothesized that an acidic pH might disrupt AJs by modifying E-cadherin and p120ctn. In this study, we used acidic growth medium to mimic an acidic microenvironment to study the posttranslational modification of E-cadherin and p120ctn.

MATERIALS AND METHODS

CELL CULTURE

The human hepatocellular carcinoma cell line HepG2 (ATCC, HB8065) was purchased from the American Type Culture Collection (Rockville, MD) and was cultured in growth medium (DMEM containing non-essential amino acids, 1 mM sodium pyruvate, 100 IU/ml of penicillin, 100 μ g/ml of streptomycin, and 10% fetal calf serum; all from Gibco, Rockville, MD) in a 5% CO₂ humidified atmosphere at 37°C. Cells (2 \times 10⁵ cells/ml) were plated either in 35 mm culture dishes or on coverslips in culture dishes. For the preparation of acidic growth medium, growth medium was adjusted to pH 6.6 by addition of 1.2 N HCl.

INHIBITORS

The Src family kinase inhibitor PP2 and the ubiquitin-proteasome inhibitor MG132 were purchased from BioMol Lab (Plymouth Meeting, PA). NH₄Cl was from Sigma (St. Louis, MO).

CYTOTOXICITY ASSAY

HepG2 cells were plated in 24-well plate and incubated in growth medium for 24 h before cell viability assay. Cells were incubated in pH 7.4 or 6.6 for 3 and 5 h. After two washes with phosphate-buffered saline (PBS; 137 mM NaCl, 2.7 mM KCl, 1.5 mM KH₂PO₄, 8 mM Na₂HPO₄, pH 7.4), 500 μ l of growth medium containing

0.5 mg/ml of 3-[4,5-dimethylthiazol-2-yl]-2,5-diphenyltetrazolium bromide (MTT, Sigma) was added to each well for 4 h reaction. The reaction medium was then removed and the cells were lysed in 500 μ l DMSO per well, and the absorbance was measured at 590 nm on a spectrophotometer (Beckman Coulter Inc., Fullerton, CA).

IMMUNOFLUORESCENCE STAINING

HepG2 cells grown on coverslips were washed briefly in PBS. For E-cadherin staining, the cells were fixed in cold acetone for 10 min at -20°C. After a brief wash with PBS, they were incubated overnight at 4°C with mouse monoclonal anti-human E-cadherin antibodies (BD Biosciences, San Jose, CA) diluted 1:50 in PBS. After a PBS wash, bound antibodies were detected by incubation for 1 h at 37°C with FITC-conjugated goat anti-mouse IgG antibodies (Sigma) diluted 1:50 in PBS. The cells were counterstained for 15 min with 1 μ g/ml of 4',6-diamidino-2-phenylindole diacetate (DAPI; Sigma) in 0.9% NaCl. After washes in 0.9% NaCl, the coverslips were mounted in a mixture of 2% *n*-propyl gallate and 60% glycerol in 0.1 M phosphate-buffered saline, pH 8.0, and sealed with nail polish. Immunofluorescence images were examined using a Zeiss Axiophot microscope (Carl Zeiss, Oberkochen, Germany) equipped with epifluorescence, and the images were captured and digitized using a Nikon D1X digital camera (Nikon, Tokyo, Japan).

PREPARATION OF WHOLE CELL HOMOGENATES AND MEMBRANE FRACTIONS

All steps were at 4°C. HepG2 cells were suspended in 20 mM HEPES, 0.28 M sucrose, 50 mM NaCl, 20 mM sodium pyrophosphate, 2 mM EDTA, 2 mM PMSF, 1 mM NaF, 2 mM Na₃VO₄, 1 μ g/ml of aprotinin, 1 μ g/ml of leupeptin, and 1 μ g/ml of pepstatin, pH 7.4, and ultrasonicated for 2 \times 10 s. In some studies, the cell homogenate was used, while in others, membrane fractions were prepared by centrifuging the homogenate at 1,000g for 10 min to remove nuclei, then centrifuging the supernatant at 16,000g for 45 min. The membrane pellets were dissolved in radioimmunoprecipitation (RIPA) buffer (50 mM Tris-HCl, 0.25% Na deoxycholate, 150 mM NaCl, 1 mM PMSF, 1 mM NaF, 1 mM Na₃VO₄, 1 μ g/ml of aprotinin, leupeptin, and pepstatin, pH 7.4) containing 1% Nonidet-40.

WESTERN BLOTTING

Gel electrophoresis and immunoblotting were performed as described previously [Towbin et al., 1979; Fritz et al., 1989] using a 10% resolving gel and a 3.5% stacking gel. After various treatments, HepG2 cells were homogenized or fractionated into membrane and cytosolic fractions. An equal volume of sample buffer was added and the mixture heated at 90°C for 3 min. Proteins (50 μ g per lane) were electrophoresed and transferred to a nitrocellulose membrane. Strips from the membrane were blocked with 5% non-fat milk in Tris-buffered saline (TBS: 150 mM NaCl, 50 mM Tris, pH 8.2) containing 0.1% Tween-20 and incubated overnight at 4°C with primary antibodies in TBS containing 0.1% Tween 20 and 5% non-fat milk. For detection of internal standards (GADPH or ATPase), rabbit anti-GADPH (Abcam, Cambridgeshire, UK) or rabbit anti-ATPase antibodies (Santa Cruz Biotechnology Inc., CA) was used. After washes with TBS-Tween, the strips were incubated with alkaline phosphatase-conjugated secondary

antibodies in TBS-Tween and positive bands visualized using nitro blue tetrazolium and 5-bromo-4-chloro-3-indolyl phosphate as chromogen. Densitometry was performed using Gel pro 3.1 (Media Cybernetics, Silver Spring, MD). The density of the band in the control sample was defined as 100% and the density of the band in the test sample expressed as a percentage of this value. All experiments were performed at least three times, and the values are expressed as the mean \pm SD.

IMMUNOPRECIPITATION

After treatment, HepG2 cells were lysed by ultrasonication in RIPA buffer containing 0.1% SDS and the lysate centrifuged at 16,000*g* for 10 min at 4°C. The supernatants were incubated for 24 h at 4°C on a shaker with mouse anti-human E-cadherin antibody (BD Biosciences) or anti-phosphotyrosine antibody (Zymed Lab, South San Francisco, CA) (1 μ g antibody for 400 μ g of protein). Prewashed aliquots of protein G-Sepharose beads (50 μ l of a 50% suspension in RIPA buffer) (GE Healthcare Bio-Sciences AB, Uppsala, Sweden) were added to the mixture for 1 h at 4°C on a shaker. After

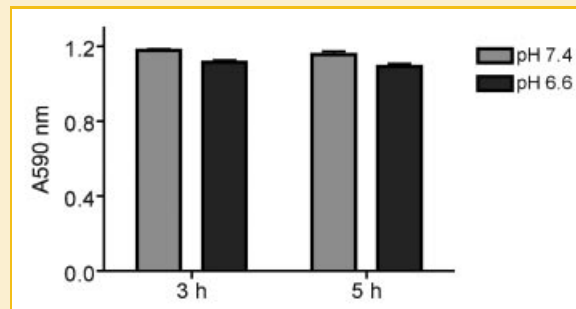


Fig. 1. Effect of acidic culture medium on cytotoxicity. HepG2 cells were incubated for 3 and 5 h in culture medium adjusted to pH 7.4 or 6.6, then assayed for viability by the MTT test. The results are expressed as the mean absorbance at 590 nm \pm SD, $n = 3$.

centrifugation at 16,000*g* for 1 min, the immunoprecipitates were washed, dissolved in 30 μ l of reducing SDS sample buffer, and analyzed by Western blotting with mouse monoclonal antibodies against phosphotyrosine (Zymed), human E-cadherin, ubiquitin,

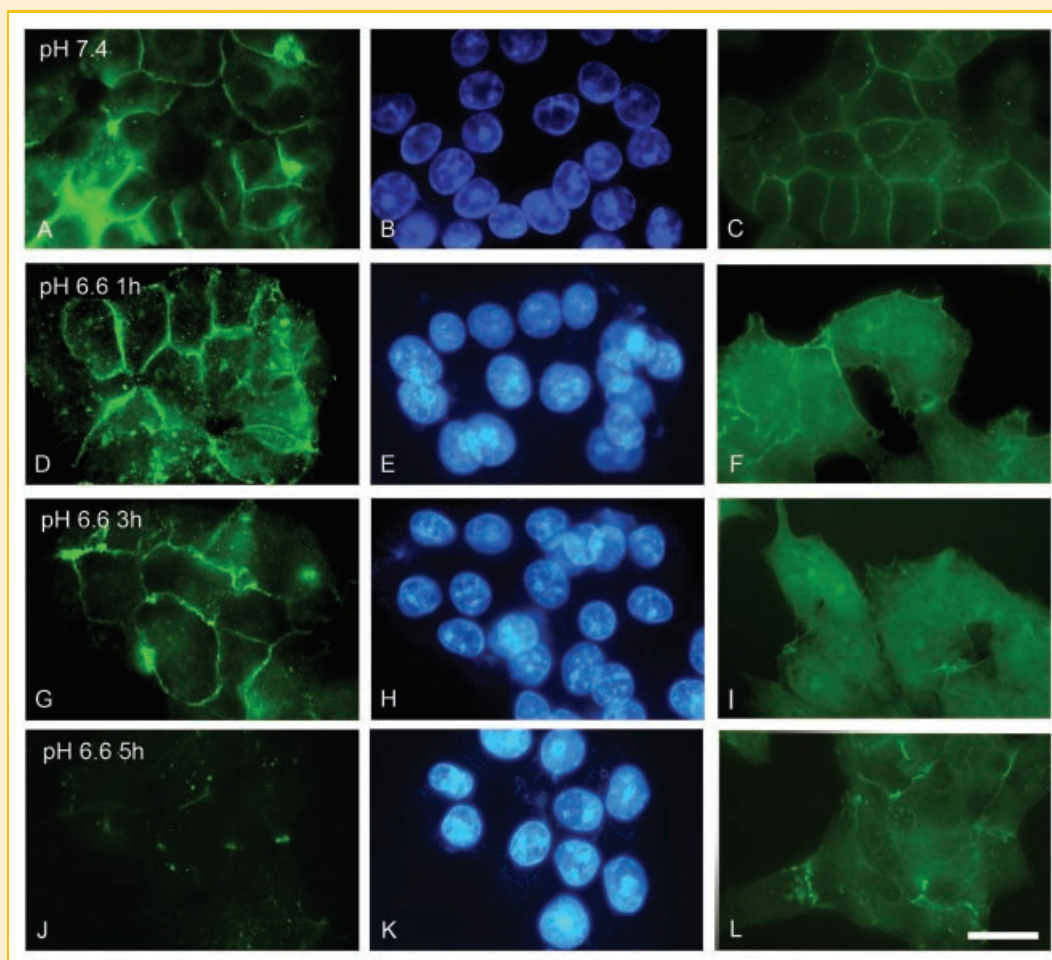


Fig. 2. Distributional change of E-cadherin and p120ctn in HepG2 cells at pH 6.6. HepG2 cells were incubated in culture media at pH 7.4 (A–C) for 5 h or at pH 6.6 for 1 h (D–F), 3 h (G–I), and 5 h (J–L). All were immunostained for E-cadherin (A, D, G, J) or p120ctn (C, F, I, L) and counterstained with DAPI (B, E, H, K). Bar = 20 μ m. [Color figure can be viewed in the online issue, which is available at www.interscience.wiley.com.]

Hakai or human p120ctn (BD Biosciences, San Jose, CA), followed by alkaline phosphatase-conjugated anti-mouse immunoglobulin G antibodies (Promega Corporation, Madison, WI).

STATISTICAL ANALYSIS

All results are expressed as the mean \pm SD. Statistical differences between means were evaluated using Student's *t*-test. Differences were considered significant if $P < 0.05$.

RESULTS

ACIDIC CULTURE MEDIUM DECREASES E-CADHERIN STAINING AT CELL-CELL CONTACT SITES

We first examined whether pH 6.6 caused cell cytotoxicity. MTT assay showed that cells incubated in pH 6.6 culture medium for 3 and 5 h did not affect cell viability (Fig. 1). As previously stated by Chen et al., [2008], proidium iodide vital staining and 4',6-diamidino-2-phenylindole dilactate staining showed very few apoptotic or necrotic cells after 24 h incubation at pH 6.6, the percentages being comparable to those in the pH 7.4 group [Chen et al., 2008].

A time course study on the distributional change of E-cadherin was carried out. The cells were counterstained with DAPI to show the location of the nuclei in pH 7.4-treated cells (Fig. 2B) and pH 6.6-treated cells (Fig. 2E,H,K). HepG2 cells grown in pH 7.4 culture medium showed linear, continuous staining of E-cadherin at cell junctions (Fig. 2A,B). Intracellular punctates positive for E-cadherin were increased after 1 h of pH 6.6 treatment (Fig. 2D,E). A noticeable decrease in E-cadherin immunoreactivity at AJ was detected after 3 h of pH 6.6 treatment (Fig. 2G,H). E-cadherin staining became faint after 5 h of pH 6.6 treatment (Fig. 2J,K).

ROLE OF c-Src IN THE PH 6.6-INDUCED DOWNREGULATION OF E-CADHERIN

We next examined the role of Src family kinases in the pH 6.6-induced downregulation of E-cadherin at AJs. Phosphorylation of Src (Tyr416) family kinases was increased at 30 min and remained high up to at least 90 min of pH 6.6 treatment (Fig. 3A). Immunoprecipitation and Western blotting showed that both c-Src kinase and Fyn kinase were activated by tyrosine phosphorylation in response to pH 6.6, and pharmacological inhibition of Src kinases by PP2 prevented pH 6.6-induced activation of both kinases (Fig. 3B). Furthermore, PP2 pretreatment prevented the pH 6.6-induced loss of E-cadherin from AJs, as the intensity of E-cadherin immunostaining in the PP2 + pH 6.6 group was comparable to that in the pH 7.4 group (Fig. 4A). In agreement with the immunofluorescence data, E-cadherin levels in the membrane fraction were reduced at pH 6.6 compared to pH 7.4, and this effect was prevented by PP2 pretreatment (Fig. 4B). To examine whether pH 6.6 induced tyrosine phosphorylation of E-cadherin, cell lysates were immunoprecipitated with anti-phosphotyrosine antibodies, and the amounts of E-cadherin in the immunoprecipitates were analyzed. Figure 5A shows a significant increase in tyrosine phosphorylation of E-cadherin after 1 h of pH 6.6 treatment, and this pH 6.6-induced

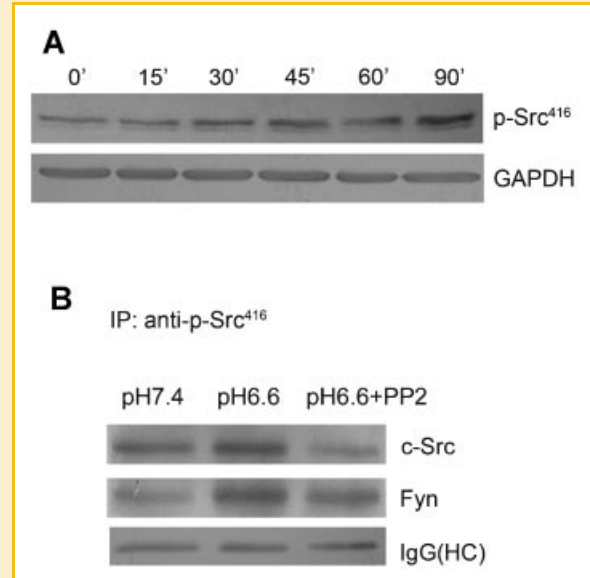


Fig. 3. pH 6.6 induces c-Src activation. A: HepG2 cells were cultured at pH 6.6 for 0, 15, 30, 45, 60, or 90 min, then the cell homogenates were analyzed for phosphorylated Src kinase (Tyr416) and GAPDH (loading control) on Western blots. B: Cells were cultured at pH 7.4 or pH 6.6 for 1 h or were pretreated for 30 min with 10 μ M PP2 at pH 7.4 for 30 min, then at pH 6.6 for 1 h, then the cell lysates were immunoprecipitated with anti-phosphorylated Src kinase (Tyr416), and the resulting immunoprecipitates analyzed for c-Src kinase, Fyn kinase and mouse IgG heavy chain (IgG-HC) (loading control). The results shown in (A) and (B) are representative of those obtained in two separate experiments.

tyrosine phosphorylation of E-cadherin was Src family kinase-dependent, as it was greatly attenuated by PP2 pretreatment (Fig. 5A). Tyrosine phosphorylation of E-cadherin by v-Src promotes the interaction with Hakai, a c-Cbl-like ubiquitin ligase, which has a high affinity for tyrosine phosphorylated E-cadherin, and the binding of Hakai with E-cadherin triggers the ubiquitination of E-cadherin [Fujita et al., 2002]. We then examined the binding of Hakai with these E-cadherin molecules. As expected, the amounts of Hakai bound to the E-cadherin increased by 50% 1 h after pH 6.6 treatment (Fig. 5B). We further evaluated the levels of E-cadherin ubiquitination. The E-cadherin-bound ubiquitin analyzed by co-immunoprecipitation was increased 1 h after pH 6.6 treatment (Fig. 5C).

We had used MG132, a cell-permeable proteasome inhibitor, which reduces the degradation of ubiquitin-conjugated proteins [Holecek et al., 2006], to examine its effect in preventing the loss of E-cadherin from cell junctions induced by pH 6.6. Figure 6A shows the continuous and linear staining of E-cadherin at cell contacts in pH 7.4-treated group, in contrast to the weak and diffuse E-cadherin staining in pH 6.6-treated groups (Fig. 6B). Pretreatment with MG132 could block pH 6.6-induced detachment of E-cadherin from cell junctions (Fig. 6C). In addition, pretreatment with NH₄Cl, a lysosome inhibitor, abrogated pH 6.6-induced loss of E-cadherin from cell-cell junctions (Fig. 6D). These results indicated that pH 6.6 reduced the levels of junctional E-cadherin through ubiquitin-proteasome/lysosome pathway.

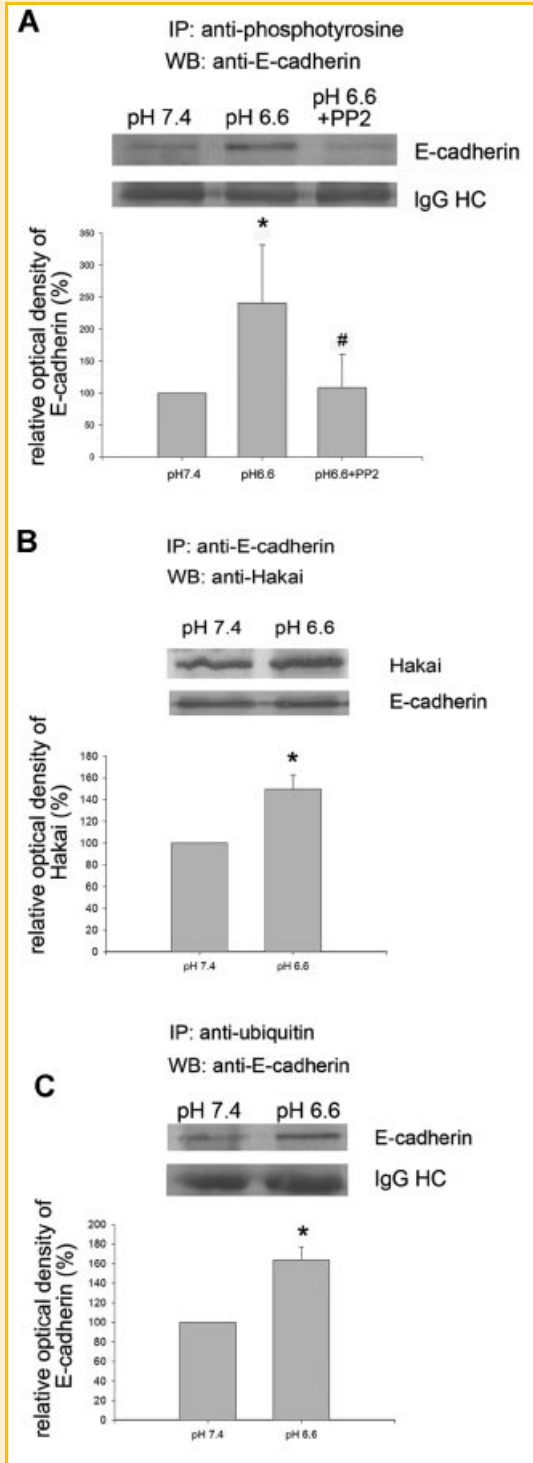


Fig. 5. pH 6.6 induces tyrosine phosphorylation and ubiquitination of E-cadherin. The cells were cultured for 1 h at pH 7.4 or 6.6. Whole cell lysates were immunoprecipitated with anti-phosphotyrosine antibodies (A), anti-E-cadherin antibodies (B) or anti-ubiquitin (C), and the immunoprecipitates analyzed for E-cadherin (A,C), Hakai (B) or IgG-HC (loading control). Pretreatment with 10 μ M PP2 (pH 6.6 + PP2) prevents pH 6.6-induced tyrosine phosphorylation of E-cadherin. A representative gel is shown and the results for three experiments are shown in the lower panel expressed as a percentage of the pH 7.4 result. * $P < 0.05$, compared to pH 7.4. # $P < 0.05$, compared to pH 6.6.

growth factor and hepatocyte growth factor [Cozzolino et al., 2003]. To study the impact of tyrosine phosphorylation of p120ctn on the association with E-cadherin, we performed immunoprecipitation with anti-p120ctn antibodies and analyzed the immunoprecipitates for E-cadherin by blotting. The amount of E-cadherin associated with p120ctn was drastically reduced after 1 h of pH 6.6 treatment, and PP2 pretreatment prevented this effect (Fig. 7B). Thus, the weakening of the binding of p120ctn to E-cadherin at pH 6.6 might contribute to the early loss of p120ctn from AJs and accelerate the destabilization of E-cadherin.

DISCUSSION

Loss of E-cadherin function can result from transcriptional or post-translational processes. Previous study has shown that tyrosine phosphorylation of E-cadherin by v-Src promotes the interaction with Hakai [Fujita et al., 2002]. In this study, we showed that pH 6.6 induced the activation of c-Src kinase and Fyn kinase and tyrosine phosphorylation of E-cadherin, which was subsequently bound to Hakai and ubiquitin. The modified E-cadherin has been proposed to enter endosomal compartments, targeted to lysosomes or recycling [Pece and Gutkind, 2002]. In our results, degradation of E-cadherin from cell junctions induced by pH 6.6 condition was both proteasome- and lysosome-dependent (Fig. 6). In mouse enterocytes, the degradation of E-cadherin from cell-cell contacts upon anoikis is mediated by both proteasome- and lysosome-dependent pathways [Fouquet et al., 2004].

The increased numbers of punctuate structures positive for E-cadherin in the cytoplasm might represent the processing of E-cadherin, which might account for the decrease in the junctional E-cadherin at pH 6.6. Thus, modifications of E-cadherin by c-Src kinase contribute to the disruption of AJs in HepG2 cells in an acidic environment.

The Src family kinase members, c-Src and Fyn, regulate the interaction between cadherin and ctns (α -ctn, β -ctn, and p120-ctn) by tyrosine phosphorylation of both cadherin and catenin. c-Src kinase and Fyn kinase can phosphorylate E-cadherin, β -ctn, and p120-ctn [Roura et al., 1999; Lilien et al., 2002; Piedra et al., 2003]. Different sites of tyrosine phosphorylation on p120ctn have opposite effects on RhoA binding. c-Src phosphorylates p120ctn at Tyr217 or Tyr228 and stimulates p120ctn and RhoA binding, whereas Fyn kinase phosphorylates p120ctn at Tyr112 and inhibits the interaction between p120ctn and RhoA [Castano et al., 2007]. Fyn kinase exerts a dominant effect, since simultaneous phosphorylation of p120ctn by Src and Fyn diminished the amount of RhoA associated to p120ctn [Castano et al., 2007]. Tyrosine phosphorylation of p120ctn by either Fyn kinase or c-Src promotes an increased association between p120ctn and E-cadherin [Castano et al., 2007]. In this study, however, we observed a decreased association between E-cadherin and p120ctn when both c-Src kinase and Fyn kinase were activated by pH 6.6. It is possible that pH 6.6 treatment triggers other signaling pathways which regulate the interaction between p120ctn and E-cadherin.

The stability of E-cadherin depends on its association with p120ctn, since p120ctn knockdown by siRNA results in complete

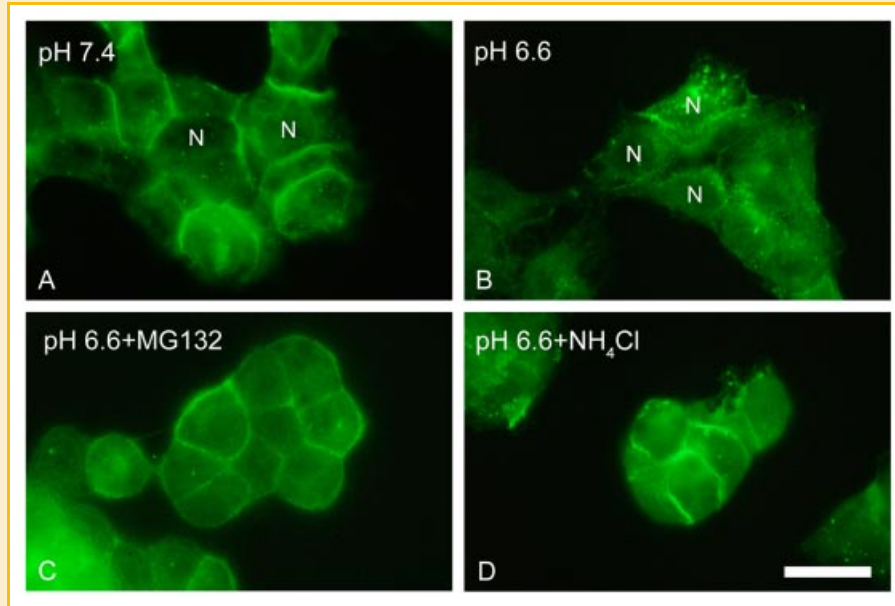


Fig. 6. pH 6.6 induces the detachment of E-cadherin from junctional complex by ubiquitin-proteosomal or lysosomal degradation. HepG2 cells were incubated in culture media for 5 h at pH 7.4 (A) or 6.6 (B) or pretreated for 30 min with 25 μ M MG132 or 20 mM NH₄Cl at pH 7.4, then at pH 6.6 for 5 h in the presence of MG132 (C, pH 6.6 + MG132) or 20 mM NH₄Cl (D, pH 6.6 + NH₄Cl), then immunostained for E-cadherin. N, nuclei. Bar = 20 μ m. [Color figure can be viewed in the online issue, which is available at www.interscience.wiley.com.]

loss of cell-cell adhesion [Davis et al., 2003; Xiao et al., 2003]. By binding to the cytoplasmic tail of cadherin, p120ctn prevents the interaction of cadherin with the endocytic membrane trafficking machinery [Davis et al., 2003; Hoshino et al., 2005; Xiao et al., 2007]. Consistent with the E-cadherin-protective role of p120ctn, E-cadherin levels are dramatically decreased in p120ctn-null SW48 colon carcinoma cells [Ireton et al., 2002]. Here, we detected tyrosine phosphorylation of p120ctn concomitant with the loss of p120ctn from AJs after 1 h of pH 6.6 treatment, suggesting that tyrosine phosphorylation of p120ctn caused this distributional change of p120ctn. Detachment of p120ctn occurred prior to E-cadherin detachment, and this observation correlated with the decreased binding of p120ctn to E-cadherin after 1 h of pH 6.6 treatment. Failure of p120ctn to protect E-cadherin might contribute to the degradation of E-cadherin from AJs by internalization. Our results provide support for a role of p120ctn in stabilizing junctional E-cadherin.

The role of tyrosine phosphorylation of p120ctn in maintaining the integrity of AJs remains obscure. P120ctn tyrosine phosphorylation is absent until transiently induced by various stimuli [Mariner et al., 2001, 2004]. Epidermal growth factor (EGF) receptor-induced tyrosine phosphorylation of p120ctn in BEL-704 hepatoma cells results in translocation of p120ctn to the nucleus and decreased cell adhesion [Huang et al., 2003]. In vitro tyrosine phosphorylation of p120ctn increases the binding of the Fer/Fyn-p120ctn complex to E-cadherin [Piedra et al., 2003]. Tyrosine phosphorylation of p120ctn correlates with an increased affinity for cadherins, both in vivo [Roura et al., 1999] and in vitro [Calautti et al., 1998; Kinch et al., 1995; Piedra et al., 2003]. However, other investigators reported that no significant change in the p120ctn/E-cadherin

association was found, even when p120ctn was heavily tyrosine-phosphorylated [Reynolds et al., 1994; Shibamoto et al., 1994; Papkoff, 1997]. Moreover, tyrosine-phosphorylated p120ctn in v-Src-transfected L cells has been shown to dissociate from the cadherin complex, probably due to a conformational change in tyrosine residues in p120ctn [Ozawa and Ohkubo, 2001]. On the other hand, modification of cadherin and the catenin complex affects the localization of p120ctn at AJs. Tyrosine phosphorylation of VE-cadherin can lead to dissociation of p120ctn from AJs [Fujita et al., 2002; Potter et al., 2005; Tricaud et al., 2005], while tyrosine phosphorylation of β -ctn or p120ctn is known to disrupt the p120ctn/E-cadherin interaction [Ezaki et al., 2007]. pH 6.6 treatment was shown to increase tyrosine phosphorylation of β -ctn in our previous study [Chen et al., 2008] and to increase tyrosine phosphorylation of E-cadherin in this study. The amount of β -ctn-associated E-cadherin is decreased by \sim 20%, as reported in our previous study [Chen et al., 2008]. In this study, we observed that, when both E-cadherin and p120ctn were tyrosine-phosphorylated, the association between these two molecules became weak. P120ctn at AJ is properly phosphorylated at Ser/Thr [Fukumoto et al., 2008]. Serine/threonine phosphorylation of p120ctn could occur in response to PKC activation and vascular endothelial growth factor [Xia et al., 2003]. PKC activation by PMA dephosphorylates p120ctn at Ser/Thr residues and redistributes p120ctn to the cytoplasm [Xia et al., 2004]. Whether acidic pH affects p120ctn Ser/Thr phosphorylation to cause the detachment of p120ctn from AJs awaits for elucidation.

P120ctn can act upstream of the Rho family GTPase and plays a regulatory role in cell motility [Grosheva et al., 2001]. Mislocalization of p120ctn to the cytoplasm promotes cell motility, as

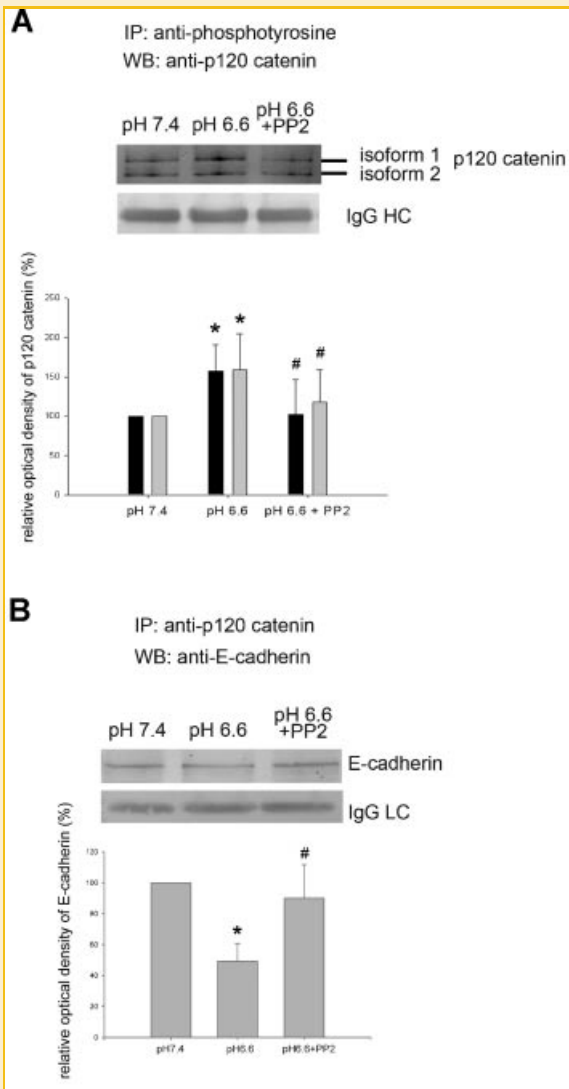


Fig. 7. pH 6.6 induces tyrosine phosphorylation of p120catenin and decreases its association with E-cadherin. A: Cells were cultured at pH 7.4 or pH 6.6 for 1 h or pretreated for 30 min with 10 μ M PP2 at pH 7.4, then at pH 6.6 in the presence of PP2 for 1 h (pH 6.6 + PP2), then whole cell lysates were incubated with anti-phosphotyrosine antibody, followed by immobilized protein G-agarose, and the immunoprecipitates analyzed for p120catenin or IgG-HC. The upper panel shows a typical result and the lower panel the mean \pm SD for the band density in three separate experiments expressed as a percentage of the pH 7.4 value. Black column, isoform 1. Gray column, isoform 2. * $P < 0.05$, compared to pH 7.4. # $P < 0.05$, compared to pH 6.6. B: Cells were cultured at pH 7.4 or pH 6.6 for 1 h or pretreated for 30 min with 10 μ M PP2 at pH 7.4, then at pH 6.6 in the presence of PP2 for 1 h (pH 6.6 + PP2), then whole cell lysates were incubated with anti-p120catenin antibody, followed by immobilized protein G-agarose, and the immunoprecipitates analyzed for E-cadherin or IgG-LC. The results show are typical of those obtained in four separate experiments. * $P < 0.05$, compared to pH 7.4. # $P < 0.01$, compared to pH 6.6.

cytoplasmic p120catenin inhibits RhoA activity [Anastasiadis and Reynolds, 2000; Anastasiadis, 2007] and activates Rac [Noren et al., 2000; Grosheva et al., 2001], which favors cell migration. Overexpression of p120catenin in fibroblasts and epithelial cells induces pronounced changes in cell motility, a result of a significant increase

in Cdc42 and Rac activity [Grosheva et al., 2001]. All four p120catenin isoforms are able to activate Rac [Yanagisawa et al., 2008]. In this study, we observed loss of p120catenin from AJs and increased diffuse cytoplasmic p120catenin staining at pH 6.6. A high migratory and invasive behavior of HepG2 cells is seen at pH 6.6 [Chen et al., 2008] and the interaction of cytoplasmic p120catenin with Rac might be involved in this invasive property of HepG2 cells.

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