Regulation of invasion and migration

by PKCK2 in *Helicobacter pylori*

infected gastric cancer cells

Yeo Song Lee

Department of Medical Science

The Graduate School, Yonsei University

Regulation of invasion and migration by PKCK2 in *Helicobacter pylori* infected gastric cancer cells

Directed by Professor Yong Chan Lee

The Doctoral Dissertation submitted to the Department of Medical Science, the Graduate School of Yonsei University in Partial Fulfillment of the Requirements for the Degree of Doctor of Philosophy

Yeo Song Lee

June 2014

This certifies that the Doctoral Dissertation

of Yeo Song Lee is approved.

from
Thesis Supervisor: Yong Chan Lee
A Fort
Thesis Committee Member#1: Kunhong Kim
Sun Y. Rha
Thesis Committee Member#2: Sun Young Rha
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Thesis Committee Member#3: Woo Jin Hyung
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Thesis Committee Member#4: Sang Kil Lee

The Graduate School Yonsei University

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June 2014

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ABSTRACT

Regulation of invasion and migration

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Yeo Song Lee

Department of Medical Science The Graduate School, Yonsei University

(Directed by Professor Yong Chan Lee)

Chronic infection with *Helicobacter pylori* (*H. pylori*) is causally linked with gastric inflammation and carcinogenesis. Virulent *H. pylori* strains harbor *cag* pathogenicity island (PAI) for delivery of the bacterial CagA into gastric epithelial cells. Induction of high motility and an elongated phenotype is considered to be CagA-dependent process. Epithelial-mesenchymal transition (EMT) is a complex cellular program involved in both development and cancer, and the induction of cell migration and invasion is the hallmark of the EMT. Protein kinase casein kinase 2 (PKCK2) plays a critical role in carcinogenesis through signaling pathways related to the epithelial mesenchymal transition (EMT). This study was aimed to investigate the effect of *H. pylori* infection on the PKCK2 mediated migration and invasion in gastric cancer cells. In *in-vivo* results, PKCK2 α immunostaining revealed strong expression in *H. pylori*-infected gastric cancer tissues and nuclear expression of PKCK2 β was decreased in *H.*

pylori-infected gastric cancer tissues. In *in vitro* data, *H. pylori* infection increases host cell PKCK2 activity and decreases PKCK2 β expression. Inhibition of PKCK2 with the chemical inhibitor 4,5,6,7-tetrabromo-2-azabenzimidazole (TBB), siRNA or shRNA significantly decreased both invasion / migration and dissociation of the membranous α/β -catenin complex in *H. pylori* infected gastric cancer cells. These results suggest that *H. pylori* induces PKCK2-mediated cell migration and invasion through α/β -catenin dissociation in gastric cancer cells.

Key words: *helicobacter pylori*, gastric cancer cells, PKCK2, α/β-catenin, migration, invasion

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I. INTRODUCTION

Helicobacter pylori (*H. pylori*) is a gram-negative, microaerophilic bacterium that selectively colonizes the human stomach. Virulent *H. pylori* strains harbor a *cag* pathogenicity island (PAI) for the translocation of the bacterial oncoprotein, CagA into gastric epithelial cells ¹⁻³. Induction of high motility and an elongated phenotype has been considered hallmark of *cag* PAI-dependent process ⁴. However, the pathogenic molecular mechanism which underlies the carcinogenesis and metastasis in gastric cancer by *H. pylori* remains largely unknown. The epithelial-mesenchymal transition (EMT) is a complex cellular program associated with induction of cell migration and invasion which plays critical roles in the development of *H. pylori* induced gastric carcinogenesis. Protein kinase casein kinase 2 (PKCK2) is a serine/threonine protein kinase ^{5, 6} that plays a key role in cell cycle control, cellular differentiation, transformation and tumorigenesis ⁷. PKCK2 can function as a monomer or a tetramer, composed of two catalytic subunits, CK2*a* and/or CK2*a*², and two CK2 β regulatory subunits ⁸. Although PKCK2 is traditionally considered a constitutively active kinase, studies

have shown that PKCK2 is activated in response to a diverse array of growth-factor stimuli, including EGF treatment ^{9, 10}. PKCK2 phosphorylates various intracellular molecules involved in tumor invasion, progression, or metastasis in multiple cancers, including cancers of the stomach, breast, lung, prostate, liver, and colon¹¹⁻¹³. Additionally, a selective inhibitor of PKCK2, 4,5,6,7-tetrabromo-2-azabenzimidazole (TBB), significantly inhibits invasion of human tongue cancer cells and lung adenocarcinoma cells ^{14, 15}. Although aberrant expression of PKCK2 is known to be involved in many cancers, the mechanism by which PKCK2 promotes tumorigenesis remains obscure.

Adherence junctions (AJs) are a network of membrane proteins and associated molecules, including E-cadherin and catenins. AJs are essential for establishing the polarity of epithelial cells and maintaining the integrity of the epithelial layer ^{16, 17}. The cytoplasmic domain of Ecadherin binds to the β -catenin, which in turn binds α -catenin ¹⁸⁻²⁰. Deleterious mutations in adhesion molecules correlate with tumor metastasis and invasion. Although it is obvious from several reports that cell elongation and migration are strongly enhanced by CagA²¹⁻²³, the precise role of CagA in the disruption of cellular AJs remains controversial and multiple mechanisms maybe involved in the disruption of cell adhesion during *H. pylori* infection. Phosphorylation of α -catenin at serine 641 by PKCK2 in response to EGF-ERK activation facilitates disruption of binding between α -catenin and β -catenin and intercellular adhesion, as well as promoting tumor cell migration 24 . In this study, we report that CagA⁺ H. pylori strongly activates PKCK2 without affecting the endogenous level of CK2 α protein. Furthermore, we show that CagA⁺ H. pylori induces PKCK2-mediated migration of gastric cancer cells by disrupting the membrane-bound α/β -catenin complex, resulting in loss of the E-cadherin complex and loss in cell to cell adhesive properties. Taken together, these data identify a new mechanistic link between CagA⁺ H. pylori and gastric cancer metastasis and invasion through activation of PKCK2 and subsequent induction of the EMT.

II. MATERIALS AND METHODS

1. Bacteria, cell culture, and H. pylori infection

H. pylori strains Hp60190 (CagA⁺) and Hp8822 (CagA⁻) were used. *H. pylori* strains were cultured on agar plates containing 10% horse serum at 37°C in a microaerobic atmosphere using the Campy Container System (BBL, Sparks, MD, USA). Human gastric cancer cells (AGS and MKN28) were cultured in RPMI-1640 medium (Gibco, Grand Island, NY, USA) containing 10% fetal bovine serum (Gibco). The cells were incubated at 37°C in a humidified atmosphere with 5% CO₂. Confluent cells were incubated overnight in serum-free and antibiotic-free media before experiments. Human gastric cancer cells with/without pretreatment with PKCK2 inhibitor (4,5,6,7-tetrabromo-2-azabenzimidazole [TBB]), PI3K inhibitor (LY294002), Akt inhibitor (Akt1/2 kinase inhibitor) were infected with *H. pylori* at multiplicity of infection (MOI) of 100:1 for various times.

2. Western blot analysis

Whole cell lysates were prepared with lysis buffer containing 50 mM Tris (pH 7.5), 5 mM EDTA, 100 μ M NaCl, 1% Triton X-100, 1 mM Na₃VO₄, and protease inhibitors (Roche Molecular Biochemical, Indianapolis IN, USA). Lysates were separated by 10% SDS-PAGE and transferred to polyvinylidene fluoride (PVDF) membranes. Immunodetection was performed using an enhanced chemiluminescence (ECL) reagent (Intron, Seoul, Korea), according to the manufacturer's instructions. In experiments using the kinase inhibitor, cells were incubated with TBB for 1 hour before co-incubation with *H. pylori*. Anti- α -catenin, anti- β -catenin, and anti-CK2 α antibodies were from Santa Cruz Biotechnology (Santa Cruz, CA, USA), anti-p- α -catenin antibody was from Signalway Antibody (Pearland, TX, USA) and anti-MMP-7 antibody was from Abcam (Cambridge, UK).

3. siRNA and plasmid DNA transfection

CK2 α -specific siRNAs and control siRNA were purchased from Bioneer (Daejeon, Korea). In brief, cells were seeded in a 6-well plate at a density of 10⁶ cells/well one day before transfection, with a target of 40-50% confluency at the time of transfection. Cells were transfected with 50 nM of siRNA using Lipofectamine 2000 (Invitrogen, Carlsbad, CA, USA) according to the manufacturer's protocol. Adequate inhibition of the siRNA-mediated knockdown was confirmed by western blot analysis. The CK2 α expression vector pRC/CMV-CK2 α -HA or control pRC/CMV-HA plasmid vectors were transfected into AGS cells using Lipofectamine 2000 (Invitrogen, Carlsbad, CA, USA), according to the manufacturers protocol. Cells were harvested for western blot analysis or used in invasion and migration assays at the indicated time intervals.

4. Immunoprecipitation

Cells grown in a 100-mm dish were infected with *H. pylori* at an MOI of 100. Cells were washed twice with PBS and incubated in lysis buffer (50 mM Tris-Hcl pH 7.5, 5 mM EDTA, 100µM NaCl, 1% Triton X-100, 1 mM Na₃VO₄, and protease inhibitors (Complete, Roche Molecular Biochemical) for 30 minutes at 4°C. Lysates were incubated with the appropriate antibodies for 4 hours at 4°C and immune complexes were trapped on protein A/G-agarose beads (Santa Cruz Biotechnology) overnight. Beads were washed 3-5 times with cold lysis buffer (50 mM Tris-HCl pH 7.5, 150 mM NaCl, protease inhibitors). The total cell lysates and immunoprecipitated materials were subjected to SDS-PAGE and the proteins were transferred to PVDF membranes, which were incubated with solutions containing the appropriate antibodies and then visualized using ECL reagent (Intron).

5. In vitro PKCK2 kinase assay

PKCK2 activity in cell lysates was determined using a PKCK2 assay kit (MBL International, Woburn, MA, USA) according to the manufacturer's instructions. Each treatment was repeated in triplicate and statistical significance was determined as p<0.05.

6. Quantitative reverse transcription-polymerase chain reaction (qRT-PCR)

Total RNA was extracted with TRIzol (Invitrogen) according to the manufacturer's instructions. First-strand complementary DNA was synthesized from 1µg total cellular RNA with random primers using a RNA PCR kit (Intron). Quantitative RT-PCR was performed on a 7500 fast realtime PCR system (Applied Biosystems, Foster City, CA, USA) using Power SYBR Green PCR Master Mix (Applied Biosystems). The oligonucleotide primers used for qRT-PCR were as follows: $CK2\alpha$, 5'-CTTCTCAGGGGAGGCAGGA-3' and 5'-CACACTTCCACAAGAGCCACT-3'; MMP7, 5'-GGAGATGCTCACTTCGATGA -3' and 5'-ATACCCAAAGAATGGCCAAG -3'; β -actin, 5'-TTGCCGACAGGATGCAGAAGA-3' and 5'-AGGTGGACAGCGAGGCCAGGAT-3'.

7. CK2a stable knockdown using lentiviral short hairpin RNA

Four premade lentiviral CK2α short hairpin RNA (shRNA) constructs and a negative control construct in the same vector system (pLKO.1) were purchased from Open Biosystems (Huntsville, AL, USA). Lentiviral helper plasmids (pCMV-dR8.2 dvpr and pCMV-VSV-G) were obtained from Addgene (Cambridge, MA, USA). Transient lentivirus stocks were prepared in 293T cells according to the manufacturer's protocol. AGS cells that stably expressed shRNA constructs were selected with 0.5-2µg/ml puromycin 48 hours after lentivirus infection. After 2 weeks of selection, monolayers of stably infected clones were harvested for use and cryopreservation.

8. Cell migration assay

Cell migration assays were performed using μ -Dish 35-mm Culture Inserts (Ibidi, Martinsried, Germany) according to the manufacturer's protocols. In brief, cells were seeded into each well of culture inserts and incubated at 37°C in a humidified atmosphere with 5% CO₂. On the day before experiment, confluent cells were incubated overnight with serum-free RPMI 1640. After cell attachment, the culture inserts were gently removed using sterile tweezers and cells were infected with *H. pylori* for 24 hours at an MOI of 100.

9. Matrigel invasion assay

Invasion of cells was measured using Matrigel (BD Biosciences, San Jose, CA, USA)-coated Transwell inserts (6.5 mm, Costar, Cambridge, MA, USA) containing polycarbonate filters with 8-µm pores. After exposure to *H. pylori* for 24 hours, cells $(2 \times 10^5$ cells in 200 µl of serum-free medium) were plated in the upper chamber, and 600 µl of medium containing 10% FBS was added to the lower well. After incubation for 24 hours at 37°C, non-invaded cells on top of the transwell were scraped off with a cotton swab. The filters (with attached invaded cells on the lower side) were washed with PBS, fixed with Diff Quik fixative (Sysmex corporation, Kobe, Japan), and stained with Diff Quik solutions I, II (Sysmex corporation). The invaded cells were counted under a light microscope (Olympus BX40) in 10 randomly selected fields at ×200 magnification. Each experiment was performed in triplicate. Invasion of cells under different conditions was normalized to the control and expressed as fold change.

10. Immunofluorescence microscopy

After co-incubation with *H. pylori*, cells were washed twice with cold PBS, fixed in methanol for 10 minutes at 4°C, washed with PBS, and permeabilized with 0.1% Triton X-100 in PBS for 10 minutes. Non-specific binding was blocked with 3% BSA in 0.1% Triton X-100/PBS for 30

minutes, followed by incubation with primary antibody against α -catenin, E-cadherin or MMP-7 in 1% BSA/0.1% Triton X-100/PBS at 4°C overnight. After washing with PBS, immunolabeled proteins were visualized by treatment with fluorescence-conjugated secondary antibodies for 60 minutes at room temperature. Cells were washed with PBS, mounted with DAPI mounting medium, sealed with cover slips, and examined using a confocal laser scanning microscope (LSM 510; Carl Zeiss, Thornwood NY, USA).

11. Enzyme-linked immunosorbent assay (ELISA)

On the day before experimentation, confluent cells were incubated overnight with serum-free RPMI 1640 and then infected with Hp60190 or Hp8822 for 8 hours with or without pretreatment with TBB for 1 hour. MMP-7 concentrations in culture supernatant were determined using commercially available ELISA kit (Abcam, Cambridge, UK). Method was as described in the manufacturer's protocol. The optical density at 450 nm was read using an automated microplate photometer, and concentrations of MMP-7 were determined by comparison with the MMP-7 standard curve.

12. Immunohistochemical staining

Immunohistochemistry (IHC) for α -catenin, β -catenin, and E-cadherin was performed on 4-mm sections of frozen clinically obtained gastric cancer tissues from our hospital tissue bank. The study protocol and expression of inferred consents were approved by the Institutional Review Board of the Severance Hospital (4-2010-0265). Tissue sections were deparaffinized with xylene, hydrated in serial dilutions of alcohol, and immersed in 3% H₂O₂. Following antigen retrieval in citrate buffer (pH 6.0), the tissue sections were incubated with protein blocking agent (Immunotech, Marseille, France) to block non-specific antibody binding for 30 minutes at room temperature and then incubated overnight at 4°C with primary antibody against CK2 α ,

CK2 β , α -catenin, β -catenin, or E-cadherin (Santa Cruz Biotechnology; 1:200), in a humidified chamber. After washing with PBS three times, the sections were incubated with a biotinylated secondary antibody and streptavidin conjugated to horseradish peroxidase (Immunotech) for 60 minutes at room temperature, followed by a PBS wash. The chromogen was developed for five minutes with liquid 3, 3'-diaminobenzidine (Immunotech) followed by counterstaining with Meyer's hematoxylin. Slides were examined under a light microscope.

13. Statistical analysis

The resulting data from invasion, migration, *In vitro* PKCK2 kinase assay, reporter assay, quantitative RT-PCR and ELISA were analyzed using GraphPad Prism 5.0 software (GraphPad Software, Inc., La Jolla, CA, USA), applying one-way ANOVA with post hoc analysis using Bonferroni post hoc test. To evaluate the immunohistochemistry results, statistical analyses were carried out using SPSS 20.0 software (SPSS, Chicago, IL, USA), applying fisher's exact test.

III. RESULTS

1. *H. pylori* induces the invasion and migration in gastric cancer cells through PKCK2 activation

H. pylori induces an elongated 'hummingbird' morphologic phenotype in epithelial cells that is associated with the loss of cell polarity and enhanced cell migration $^{22, 25}$. To determine whether *H. pylori* activates PKCK2 to induce cancer cell invasion and migration, we treated MKN28 (Fig. 1A, C) and AGS (Fig. 1B, D) cells with the PKCK2 inhibitor TBB or CK2a-specific siRNA before infection with CagA⁺ (Hp60190) and CagA⁻ (Hp8822) *H. pylori*. The invasiveness (Fig. 1A, B) and migration (Fig. 1C, D) of cells were more pronounced after infection with CagA⁺ than CagA⁻ *H. pylori*. Inhibition of PKCK2 by TBB in *H. pylori* infected MKN28 cells profoundly suppressed cell invasiveness (Fig. 1A) and motility (Fig. 1C). Similarly, down regulation of CK2a using siRNA resulted in a significant decrease in the invasiveness (Fig. 1B) and migratory activities in AGS cells (Fig. 1D). MKN28 and AGS cells with down-regulated PKCK2 showed significantly impaired invasion and migration compared with control cells (P<0.05). These results indicate that increased cellular invasion and migratory activities induced by *H. pylori* are mediated through PKCK2.

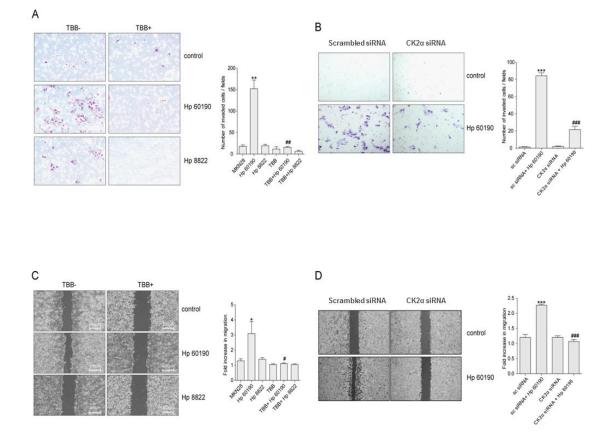
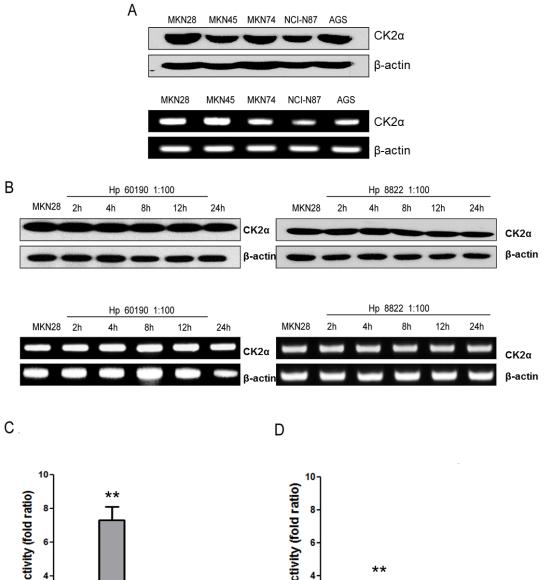


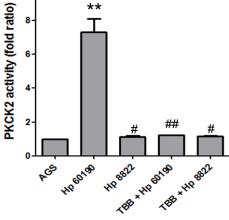
Figure 1. *H. pylori* induces the invasion and migration in gastric cancer cells through PKCK2 activation. (A) MKN28 cells were pretreated with TBB (100 μ M) for 1 hour followed by infection with CagA⁺ (Hp60190) and CagA⁻ (Hp8822) *H. pylori* for 24 hours. (B) The repeated experiments in AGS cells infected only with Hp60190 were performed following transfection with scrambled siRNA and CK2a siRNA. After invasion, the membranes were fixed and stained, and the invaded cells were counted. (C) MKN28 cells were pretreated with TBB (100 μ M) for 1 hour followed by infection with Hp60190 and Hp8822 for 24 hours. (D) The repeated experiments except for infection with only Hp60190 were performed following transfection of

AGS cells with scrambled siRNA and CK2a siRNA. Quantitative analysis of the invasive and migratory cells was performed (right panels). Data was assessed by one-way ANOVA followed by Bonferroni method (* p < 0.05, ** p < 0.01, *** p < 0.001 vs control, [#] p < 0.05, ^{##} p < 0.01, ^{###} p < 0.001 vs Hp60190-infected cells without TBB).

2. *H. pylori* activates PKCK2 without affecting CK2α protein expression levels in gastric cancer cells

PKCK2 is a ubiquitous, highly conserved serine, threonine kinase that is up-regulated in rapidly dividing cells including most human tumors. However, the mechanism by which PKCK2 is activated in gastric cancer remains unclear. CK2 α was expressed in all five gastric cancer cell lines tested with variation in the expression level among these cell lines (Fig. 2A). Most importantly, CK2 α protein and mRNA expression levels were not altered in *H. pylori* infected MKN28 cell lines (Fig. 2B). To examine whether CagA translocation by *H. pylori* infection increases PKCK2 activity, we infected AGS cells (Fig. 2C) and MKN28 cells (Fig. 2D) with CagA⁺ (Hp60190), CagA⁻ (Hp8822) *H. pylori*. Although there were no significant differences in CK2 α protein and mRNA expression levels in *H. pylori* infected AGS and MKN28 cells, infection with CagA⁺ (Hp60190) *H. pylori* significantly increased PKCK2 activities while CagA⁻ (Hp8822) *H. pylori* infected gastric cancer cells profoundly suppressed the activity of PKCK2. These data indicate that CagA affects host cell PKCK2 activity without affecting the transcriptional or translational level of CK2 α .





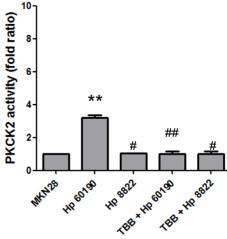
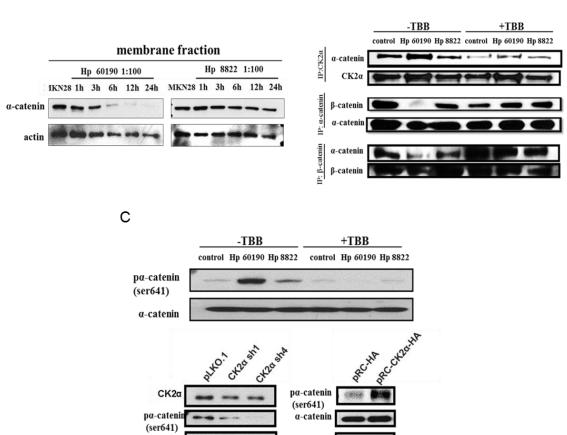


Figure 2. *H. pylori* activates PKCK2 without affecting CK2 α expression levels in gastric cancer cells. (A) Western blot and RT-PCR analysis of CK2 α expression in five gastric cancer cell lines. (B) MKN28 cells were infected with *H. pylori* for the indicated times. Actin was used as a loading control. AGS (C) and MKN28 (D) cells were pretreated with 100 μ M TBB for 1 hour followed by infection with *H. pylori* for 6 hours. The PKCK2 activity was determined using a Cyclex PKCK2 activity kit. Data was assessed by one-way ANOVA followed by Bonferroni method (** *p* < 0.01 vs control, ^{##} *p* < 0.01 vs Hp60190-infected cells without TBB).

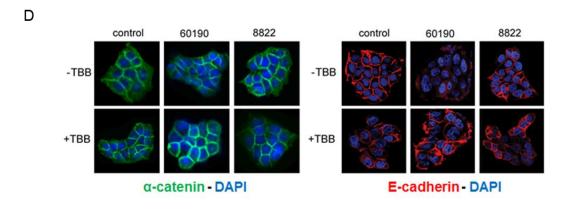
3. Phosphorylation of α -catenin by PKCK2 results in dissociation of the α/β -catenin complex in *H. pylori* infected gastric cancer cells

It was previously reported that, EGF associated ERK activates PKCK2 to phosphorylates α catenin at serine 641. Phosphorylation of α -catenin promotes disruption of α/β -catenin complex, resulting in EMT²⁴. Like EGF, *H. pylori* is known to induce ERK activation, which may be dependent or independent of CagA in gastric epithelial cells. Therefore, we examined whether H. *pylori* disrupts the α/β -catenin complex by selective measurement of membranous α -catenin expression. As expected, infection of MKN28 cells with CagA⁺ (Hp60190), but not CagA⁻ (Hp8822), significantly depleted the membranous α -catenin levels in a time-dependent manner (Fig. 3A). The E-cadherin complex is composed of E-cadherin, β -catenin, α -catenin and other compounds. Dissociation of α/β -catenin enhances disruption of the E-cadherin complex and induces cellular invasion and migratory activities of epithelial cells 26 . To investigate whether H. pylori influences a/β-catenin disruption through activation of PKCK2, MKN28 cells were pretreated with TBB (100 µM) for 1 hour before infection with H. pylori for 24 hours. Cell lysates were immunoprecipitated with antibody against α -catenin, β -catenin or CK2 α and then immunoblotted with antibody against β -catenin or α -catenin as indicated. Through immunoprecipitation analyses, we found that Hp60190 increased the interaction between $CK2\alpha$ and α -catenin and induced disruption of the α/β -catenin interaction (Fig. 3B). Next, to determine whether *H. pylori* increases α -catenin phosphorylation at serine 641 through activation of PKCK2, we performed Western blot analysis in gastric cancer cells that were pretreated with TBB prior to H. pylori infection. Infection with Hp60190 resulted in an increased level of p-acatenin at serine 641 compared with Hp8822 infection. TBB pretreatment decreased the upregulation of p-α-catenin in cells infected with Hp60190 (Fig. 3C). Considering that H. pylori infected gastric cancer cells expressed the phosphorylated form of α -catenin through the activity of PKCK2, we next determined whether inhibition of CK2a expression directly affects α - catenin phosphorylation. We used CK2 α -specific shRNA lentivirus and puromycin to select for stable pools of AGS cells with CK2 α down-regulation and control lines transfected with pLKO.1. Analysis of CK2 α protein expression by immunoblotting confirmed that CK2 α protein expression was significantly suppressed in the AGS cells expressing CK2 α shRNA (sh1 and sh4), but not in the cells transfected with control pLKO.1. Although CK2 α knockdown had no effect on total α -catenin expression in AGS cell lines, the level of p- α -catenin was prominently decreased in CK2 α knockdown cell lines (Fig. 3C, lower left panel). Moreover, overexpression of CK2 α protein resulted in significant up-regulation of α -catenin phosphorylation (Fig. 3C, lower right panel). Immunofluorescence studies showed that *H. pylori* infection for 24 hours resulted in disruption of membranous α -catenin and E-cadherin. In addition, more pronounced effects were observed for infection with Hp60190 compared with Hp8822 (Fig. 3D). These results show that PKCK2 mediated phosphorylation of α -catenin leads to membranous α -catenin depletion through dissociation of the α/β -catenin complex in *H. pylori* infected gastric cancer cells in CagA dependent manner.



HA

tubulin



a-cateni

tubulin

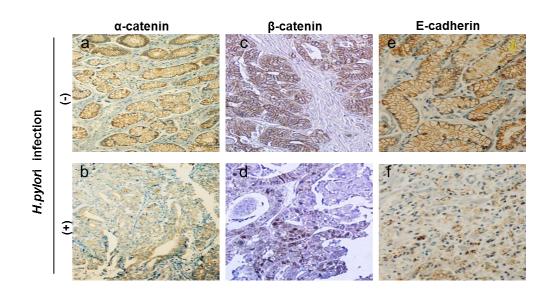
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Figure 3. PKCK2-mediated phosphorylation of α -catenin results in dissociation of the α/β catenin complex in H. pylori-infected gastric cancer cells. (A) MKN28 cells were infected with Hp60190 or Hp8822 for the indicated times. The change in membranous α -catenin level was analyzed by immunoblotting. (B) MKN28 cells were pretreated with TBB (100 μ M) for 1 hour followed by infection with Hp60190 or Hp8822 for 24 hours. Endogenous CK2a, α-catenin or β -catenin was immunoprecipitated, and binding of CK2 α to α -catenin (first panel), β -catenin to α -catenin (second panel) and α -catenin to β -catenin (third panel) was analyzed by immunoblotting. (C) MKN28 cells were pretreated with TBB for 1 hour followed by infection with Hp60190 or Hp8822 for 3 hours (upper panel). Western blot analysis was performed to determine the a-catenin and phospho-a-catenin levels in shRNA-mediated CK2a knockdown AGS cells (lower left panel) and the expression of α -catenin and phospho- α -catenin in AGS cells transfected with pRC-CK2a-HA (lower right panel). (D) MKN28 cells were pretreated with TBB (100 µM) for 1 hour followed by infection with Hp60190 and Hp8822 for 24 hours at a MOI of 1:100. The samples were subjected to immunofluorescent staining with antibody against a-catenin or E-cadherin and mounted in DAPI-containing mounting solution; 400× magnification.

4. Expression of CK2 subunits and junctional proteins in an *in vivo* gastric cancer tissue model

To examine the relationship between *H. pylori* infection and expression of junctional proteins and cellular CK2 α , CK2 β , gastric cancer tissues of patients with *H. pylori* infection (n=17) or without *H. pylori* infection (n=34) were subjected to immunohistochemical analysis using antibodies to α -catenin, β -catenin, E-cadherin (Fig. 4A) and CK2 α , CK2 β (Fig. 4B). Membranous α -catenin immunostaining revealed strong expression in non-infected gastric cancer tissues (Fig. 4A,a) and low expression in H. pylori-infected gastric cancer tissues (Fig. 4A,b). Although membranous β -catenin expression was dominant (Fig. 4A,c), nuclear β -catenin immunostaining was frequently seen in H. pylori-infected gastric cancer tissues (Fig. 4A,d). Membranous expression of E-cadherin (Fig. 4A,e) was also decreased in H. pylori-infected gastric cancer tissue (Fig. 4A,f). CK2 α immunostaining revealed more strong expression in H. pylori-infected gastric cancer tissues than non-infected gastric cancer tissues and nuclear expression of CK2 β in *H. pylori* -infected gastric cancer tissues (Fig. 4d) were lower than noninfected gastric cancer tissues (Fig. 4B). We performed statistical analysis by fisher's exact test to examine the relation between H. pylori-infection and CK2a expression according to the staining grade (0<10%, 1=10-49%, 2=50-100% in CK2a scoring and 0= no, 1=weak, 2=modest in CK2 β scoring). Although CK2 α expression had no statistically significant correlation with H. *pylori* infection (p=0.2127), CK2 α expression tended to be higher in *H. pylori* infected gastric cancer tissues. In contrast, there was an significant inverse relationship between CK2^β expression and *H. pylori* infection (*p*=0.0002).



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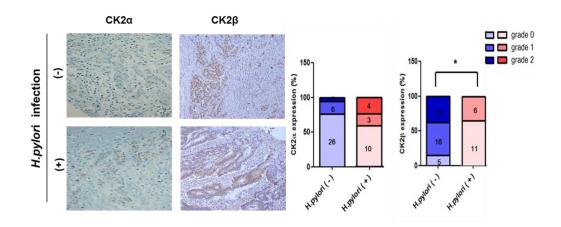


Figure 4. Immunohistochemical detection of α -catenin, β -catenin, E-cadherin, CK2 α and CK2 β expression in *H. pylori* non-infected and infected gastric cancer tissues.

Gastric cancer tissues of patients with or without *H. pylori* infection were subjected to immunohistochemical analysis using antibodies to (A) α -catenin, β -catenin and E-cadherin and (B) CK2 α , CK2 β . Magnification, ×200. Bar graphs represented percent expression of CK2 α (left panel) and CK2 β (right panel) according to the staining grade. Numbers of patients are defined in the graphs (*p*<0.05).

5. *H. pylori* induced α/β -catenin dissociation resulted in β -catenin nuclear translocation and increased MMP7 expressions

To examine the effect of *H. pylori*-induced dissociation of the α/β -catenin complex on β -catenin transactivation, we next examined T-cell factor-1 (TCF1) and lymphoid enhancing factor-1 (LEF1) transcriptional activity profiles. TCF/LEF-1 luciferase reporter analysis showed that Hp60190 induced greater β -catenin transactivation than Hp8822 in gastric cancer cells. Furthermore, Hp60190-enhanced TCF/LEF-1 transcriptional activity was blocked by treatment with the PKCK2 inhibitor TBB, showing that activation of PKCK2 by H. pylori plays an important role in β -catenin transactivation (Fig. 5A). Recent studies have shown that the nuclear translocation of β -catenin increases expression of the MMP7 gene expressions ²⁷ in lung and gastric epithelial cell lines and that MMP7 is strongly involved in EMT^{28, 29}. Therefore, we examined expression of the β -catenin/Tcf downstream target gene MMP7 and production of MMP7 by real-time qRT-PCR and ELISA, respectively in MKN28 cells that were pretreated with TBB (100 µM) before H. pylori infection. As expected, the level of MMP7 mRNA expression and MMP7 production were increased only in CagA⁺ H. pylori infected MKN28 cells. Moreover, H. pylori-induced MMP7 expression and production of MMP7 were blocked by TBB pretreatment (Fig. 5B, C). Next, to determine whether H. pylori increases MMP7 through PKCK2 activation, we performed immunofluorescence studies in MKN28 cells that were pretreated with TBB prior to H. pylori infection. Infection with Hp60190 resulted in an increased level of MMP7 compared with Hp8822 infection. TBB pretreatment decreased the upregulation of MMP7 in cells infected with Hp60190 (Fig. 5D). Taken together, these findings indicate that up-regulation of PKCK2 by CagA⁺ H. pylori plays an important role in β -catenin nuclear translocation and EMT-related MMP7 expression in gastric cancer cells.

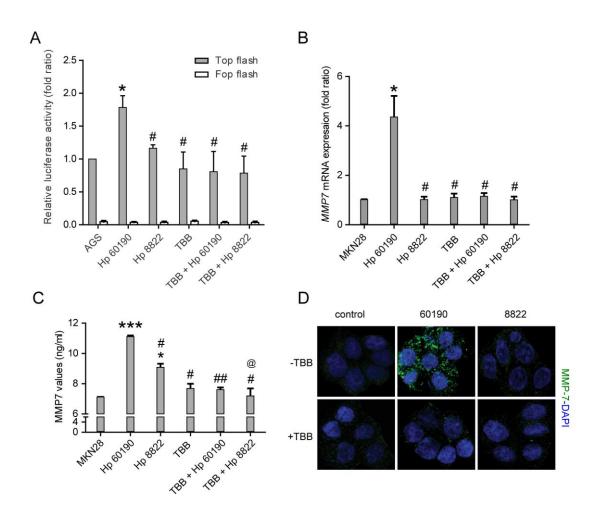


Figure 5. *H. pylori* promotes β -catenin/TCF promoter activity and downstream MMP7 expression by increasing PKCK2 activity. (A) AGS cells were co-transfected with TopFlash (multiple copies of an optimal TCF-binding site) and FopFlash (mutant Tcf binding sites) vectors together with TK-*Renilla* (control) vector and then infected with Hp60190 or Hp8822 for 24 hours with or without pretreatment with 100 μ M TBB for 1 hour. (B) MKN28 cells were infected with Hp60190 or Hp8822 for 6 hours at a MOI of 100. *MMP7* expression was measured by qRT-PCR. (C) MKN28 cells were pretreated with TBB (100 μ M) for 1 hour

followed by infection with Hp60190 and Hp8822 for 8 hours at a MOI of 1:100. MMP-7 secretions were measured by ELISA kit. (D) The repeated experiments in MKN28 cells were subjected to immunofluorescent staining with antibody against MMP-7 (green) and mounted in DAPI-containing mounting solution. Data was assessed by one-way ANOVA followed by Bonferroni method (* p < 0.05, *** p < 0.001 vs control, * p < 0.05, *# p < 0.01 vs Hp60190-infected cells without TBB, [@] p < 0.05 vs Hp8822-infected cells); 400× magnification.

6. *H. pylori* induces CK2β degradation through PI3K/Akt pathway

Recent studies have shown that low CK2 β expression correlated with characteristic EMT markers, including Snail1, Zeb2 or Twist1³⁰ and leads to morphological transformations associated with activation of EMT pathways. Therefore, we examined whether *H. pylori* increases CK2 β degradation by measurement of CK2 β protein expression. As expected, infection of AGS and MKN28 cells with CagA⁺ (Hp60190), significantly depleted the CK2 β levels in a time-dependent manner while CK2 α level did not change (Fig. 6A, B). However, both CK2 α and CK2 β mRNA expression levels were not altered in *H. pylori* infected AGS and MKN28 cell lines (Fig. 6C, D). To investigate which signaling pathway involved in CK2 β degradation, we used PI3K-specific inhibitor, LY294002 and Akt specific inhibitor, Akt1/2 kinase inhibitor. As expected, *H. pylori*-induced CK2 β degradation were blocked by LY294002 (Fig. 6E) and Akt1/2 kinase inhibitor (Fig. 6F) in AGS cells. Taken together, these results indicate that *H. pylori* infection increases EMT-related CK2 β degradation through PI3K-Akt pathway.

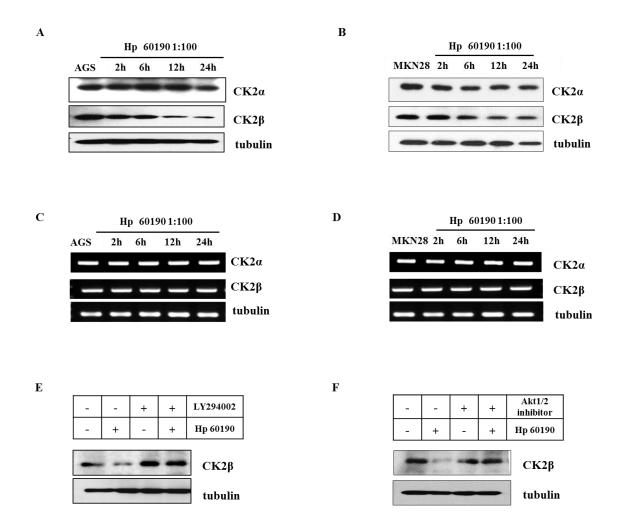


Figure 6. *H. pylori* induced CK2β degradation through PI3K/Akt pathway. AGS cells (A, C) were infected with CagA⁺ (Hp60190) at multiplicity of infection (MOI) of 100:1 for various times. (B, D) The repeated experiments in MKN28 cells. (E) AGS cells were pretreated with LY294002 (10µM) for 1 hour followed by infection with CagA⁺ (Hp60190) *H pylori* for 4 hours.
(F) The repeated experiments except for pretreatment with Akt1/2 kinase inhibitor were performed.

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IV. DISCUSSION

The Helicobacter pylori (H. pylori) cag pathogenicity island (cag PAI) is a 35-40-kb genetic element that encodes a type IV secretion system and is strongly associated with gastric malignant progression ³¹⁻³³. Virulent *H. pylori* strains harbor a PAI for the translocation of the bacterial oncoprotein, CagA into gastric epithelial cells ¹⁻³. Differences have been observed in the abilities of CagA⁺ and CagA⁻ H. pylori strains to activate various signaling cascades in infected gastric epithelial cells ^{34, 35}. Using CagA-inducible gastric epithelial model, Murata-Kamiya et al. demonstrated that intracellular CagA interacts with E-cadherin and disrupts the formation of E-cadherin– β -catenin complexes, as well as inducing nuclear accumulation of β catenin³⁶, suggesting that inhibition of cell-cell adhesion may be an intracellular target of CagA. In our study, *H. pylori* infection induced β -catenin transactivation and down-regulation of Ecadherin expressions which induced cell invasiveness and motility in gastric epithelial cells in CagA-dependent manner. These findings are consistent with previous reports that investigated the mechanisms regulating invasiveness and motility in *H. pylori*-infected gastric epithelial cells. Protein kinase casein kinase 2 (PKCK2) is a serine/threonine protein kinase ^{5, 6} that plays a key role in cell cycle control, cellular differentiation and transformation⁷. Many studies have shown that dysregulation of PKCK2 is associated with tumorigenesis ^{12, 37, 38}. Moreover, Zou et al. reported that $CK2\alpha$ modulates the process of epithelial mesenchymal transition (EMT), thereby affecting regulation of cell migration and invasion in colorectal cancer cells and PKCK2specific inhibitors significantly inhibit membrane invasion, adhesion, and migration of ovarian carcinoma cells³⁹. In agreement with these results, our study demonstrated that activity of PKCK2 in gastric epithelial cells increased in a CagA-dependent manner during H. pylori infection.

In the current study we show for the first time that $CagA^+$ (Hp60190) *H. pylori* up-regulates cellular invasiveness and migration activity through PKCK2 activation whereas $CagA^-$

(Hp8822) *H. pylori* does not induce cell invasiveness or migration. Furthermore, *H. pylori* affects PKCK2 activity in gastric cancer cells without affecting the transcriptional or translational expression of CK2 α . Although EMT is typically observed in the invasive progression of cancer cells, *in vivo* evidence suggests that loss of epithelial adhesion is sufficient to initiate invasive phenotype in the pancreas and stomach. In adherens junctions, α -catenin links the cadherin-beta-catenin complex to the actin-based cytoskeleton and stabilizes cell polarity²⁰. Our results showed that CagA-mediated PKCK2 activation disrupts these adherens junction complexes through α/β -catenin dissociation in gastric epithelial cells. Because it has been suggested that CK2 α disrupts α -catenin binding to β -catenin through CK2 α/α -catenin interaction ²⁴, it is tempting to speculate that CagA is a mediator of this interaction. However, many questions remain unanswered, mainly concerning the association between CagA and CK2 α . In any case, the results presented here have revealed new perspectives on the role of *H. pylori* CagA and epithelial PKCK2 activation in the regulation of cellular migration and invasion in gastric carcinogenesis.

Immunohistochemical analysis revealed that membranous α -catenin, β -catenin and E-cadherin expression was lowered in *H. pylori*-infected gastric cancer tissues. Although CK2 α expression in gastric cancer tissues had no statistically significant correlation with *H. pylori* infection, CK2 α expression tended to be higher in *H. pylori* infected gastric cancer tissues. The lack of significance in CK2 α expression and *H. pylori* infection may be due to the small number of patients in the *H. pylori* infected group. However, the expression of CK2 β significantly lowered in *H. pylori* infected gastric cancer tissues and these results matched with our in-vitro results.

The MMP-7 gene is one of target gene of β -catenin transactivation signals, which is critical for remodeling of the extracellular matrix for a variety of physiological and pathological processes²⁷. It has been suggested that *H. pylori*-induced MMP expression plays a role in inflammation, carcinogenesis and cancer metastasis^{28, 29}, and our findings clearly showed that *H.*

pylori being able to induce MMP-7 protein expression inducing epithelial cell invasiveness and motility *in vitro*. In this study, we observed that *H. pylori*-induced α/β -catenin dissociation resulted in β -catenin nuclear translocation and increased MMP7 expression. Furthermore, we found that up-regulation of PKCK2 by CagA⁺ *H. pylori* played an important role in β -catenin nuclear translocation and EMT-related MMP7 expression in gastric cancer cells.

Evidence of the mechanistic interplay between CagA and PKCK2-induced α/β -catenin dissociation along with β -catenin transactivation provides important insight into cell invasion and migration in gastric cancer. Taken together, these data suggest that targeting oncogenic PKCK2 activity may be a potential therapeutic strategy for treating human gastric cancer associated with *H. pylori* infection.

Recent studies have shown that unbalanced expression of PKCK2 subunits is sufficient to drive epithelial-to-mesenchymal transition by Snail1 induction⁴⁰. We show for the first time that CagA⁺ (Hp60190) *H. pylori* infection increases CK2 β degradation through PI3K-Akt pathway. Reduced CK2 β may represent a novel molecular alteration during malignant tumor progression. Interactions between *H. pylori* and cancer cells play an important role in the development of epithelial mesenchymal transition. Our finding will not only improve our understanding of *H. pylori*-induced epithelial mesenchymal transition but also provide mechanistic insight into other malignancies.

V. CONCLUSION

In this study, we showed for the first time that CagA⁺ (Hp60190) *H. pylori* up-regulates cancer cell invasiveness and migration activity through PKCK2 activation, whereas CagA⁻ (Hp8822) *H. pylori* does not induce cell invasiveness or migration. Evidence of the mechanistic interplay between CagA and PKCK2-induced α/β -catenin dissociation along with β -catenin transactivation provides important insight into cell invasion and migration in gastric cancer. Taken together, these data suggest that targeting oncogenic PKCK2 activity may be a potential therapeutic strategy for treating human gastric cancer associated with *H. pylori* infection.

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헬리코박터 파일로리 감염 위암세포에서의 CK2 매개

침윤과 이동 조절

<지도교수 이 용 찬>

연세대학교 대학원 의과학과

이여송

만성 헬리코박터 파일로리 감염은 위 점막의 염증과 위암 발생의 주요 인자가 된다. 병독 성 헬리코박터 파일로리 균 주는 *cag* pathogenicity island (PAI) 유전자를 가지고 있어, 암 단백인 CagA 를 숙주 위 상피세포에 침투시킨다. 세포의 이동성 증가와 표현형의 변화는 이러한 CagA 의존적으로 유발되는 과정이다. 상피-중간엽세포 이행은 발생과정과 암세포에서 유발되는 복잡한 세포현상이며, 세포이동과 침윤성의 증가는 이러한 상피-중간엽세포 이행의 대표적인 특징이다. Casein kinase 2 (CK2)는 상피-중간엽세포 이행 과정과 관련한 여러 신호전달체계에 중요한 역할을 한다. 본 연구의 목적은 헬리코박터 파일로리 감염이 유발하는 CK2 매개 위암세포의 침윤과 이동의 증가 기전을 이해하는 것이다. 더 높게 나타났으며, CK2β의 발현은 헬리코박터 파일로리 감염 위암 환자의 조직에서 감소함을 확인하였다. *In-vitro* 결과에서 헬리코박터 파일로리 감염은 CK2의 활성을 높이고, CK2β 의 발현을 감소시킴을 확인하였다. CK2억제제 (TBB), siRNA 또는 shRNA를 통한 CK2 억제는 헬리코박터 파일로리 감염된 위암세포의 이동과 침윤능력을 효과적으로 차단하였고, 세포막에 존재하는 α/β-catenin 복합체의 해체를 억제하였다.

이번 연구를 통해 헬리코박터 파일로리가 CK2활성화를 유도함으로써 α/β-catenin 해체를 통해 세포의 이동과 침윤을 조절함을 확인 할 수 있었다.

핵심 되는 말: 헬리코박터 파일로리, 위암세포, PKCK2, α/β-catenin, 세포이동, 침윤