

Molecular mechanism for PEDV entry

Cell entry of porcine epidemic diarrhea coronavirus is activated by lysosomal proteases

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Running title: Molecular mechanism for PEDV entry

Keywords: cysteine protease; membrane fusion; protease inhibitor; proteolysis; virus entry

ABSTRACT

Porcine epidemic diarrhea coronavirus (PEDV) is currently devastating the US pork industry by causing 80-100% fatality rate in infected piglets. Coronavirus spike proteins mediate virus entry into cells, a process that requires the spike proteins to be proteolytically activated. It has been a conundrum what proteases activate PEDV entry. Here we systematically investigated the roles of different proteases in PEDV entry using pseudovirus entry, biochemical, and live virus infection assays. We found that PEDV spike is activated by lysosomal cysteine proteases, but not proprotein convertases or cell-surface serine proteases. Extracellular trypsin activates PEDV entry when lysosomal cysteine proteases are inhibited. We further

pinpointed cathepsin L and cathepsin B as the lysosomal cysteine proteases that activate PEDV spike. These results advance our understanding of the molecular mechanism for PEDV entry and identify potential antiviral targets for curbing the spread of PEDV.

Since 2013, porcine epidemic diarrhea coronavirus (PEDV) has swept throughout the United States, causing 80-100% fatality rate in piglets and wiping out more than 10% of America's pig population in less than a year (1-3). Currently there is no effective strategy available to keep the spread of PEDV in check. PEDV belongs to the α -genus of the coronavirus family. An envelope-anchored spike protein guides coronavirus entry into

host cells (4-7). Its ectodomain contains a receptor-binding subunit S1 and a membrane-fusion subunit S2. The spike exists in two different conformations: the pre-fusion conformation is a clove-shaped trimer with three individual S1 heads and a trimeric S2 stalk; the post-fusion conformation is a trimeric S2 that has been structurally rearranged to fuse the viral and host membranes (8-12). During virus entry, S1 first binds to a receptor on host cell surface for viral attachment, and then S2 transitions to the post-fusion conformation for membrane fusion. For the spike to undergo conformational transitions, it needs to be proteotically activated by one or more host proteases.

The host proteases mainly come from four different stages of the virus infection cycle: (i) proprotein convertases during virus packaging in virus-producing cells; (ii) extracellular proteases after virus release from virus-producing cells and before virus entry into virus-targeted cells; (iii) cell-surface proteases after virus attachment to virus-targeted cells; (iv) lysosomal proteases after virus endocytosis in virus-targeted cells (9,10). In addition, proprotein convertases were shown to activate MERS-CoV spike after virus endocytosis in virus-targeted cells (13). PEDV is unique among coronaviruses in that its propagation in cell culture requires exogenous trypsin, and thus it is commonly believed that extracellular trypsin-like proteases in pig intestines are essential for cell entry of PEDV (14-16). However, these cell culture studies used live PEDV particles for cell entry, and did not differentiate PEDV entry from other steps of the PEDV infection cycle such as virus replication or release. Therefore, it remains to be a conundrum what proteases activate cell entry of PEDV.

We recently characterized the receptor usage and cell entry of PEDV (17). We found that PEDV S1 uses human and porcine aminopeptidase N (APN) as its main receptor and sugar as its co-receptor. In addition, PEDV infects both human and porcine cells. Importantly, we established a PEDV-spike-mediated pseudovirus entry assay. In this assay, a replication-deficient retrovirus

becomes pseudotyped with PEDV spike and is used to enter PEDV-susceptible host cells. The pseudovirus assay only concerns virus entry, but not virus replication or release. Hence the pseudovirus entry assay has advantages over the live PEDV infection assay in virus entry studies. Here using this PEDV pseudovirus entry assay along with biochemical and live PEDV infection assays, we systematically investigated what proteases process and activate PEDV spike, revealing the molecular mechanism for PEDV entry. Identification of the PEDV-spike-processing proteases provided potential targets for development of antiviral drugs to block PEDV entry.

RESULTS

Role of proprotein convertases in PEDV pseudovirus entry- To identify the proteases that activate PEDV entry, we examined potential spike-processing proteases from different stages of the virus infection cycle. We first analyzed whether proprotein convertases cleave PEDV spike during virus packaging. To this end, we packaged retrovirus particles pseudotyped with PEDV spike (i.e., PEDV pseudoviruses) in HEK293T cells (human embryonic kidney cells), and performed western blot analysis to detect the cleavage state of PEDV spike. Here the PEDV spike contained a C-terminal C9 tag, and hence could be detected using anti-C9 tag monoclonal antibody. Our result showed that PEDV spike remained intact on the pseudovirus surface (Fig. 1A). As a positive control, MERS-CoV spike, which also contained a C-terminal C9 tag, had been cleaved on the pseudovirus surface (Fig. 1A), consistent with previous observations that MERS-CoV entry could be activated by proprotein convertases during virus packaging in virus-producing cells (18,19). Thus, proprotein convertases from virus-producing cells do not proteolytically activate PEDV spike or PEDV entry.

We also examined whether proprotein convertases from virus-targeted cells cleave PEDV spike during virus endocytosis. Our result showed that a proprotein convertase inhibitor, Dec-RVCR-CMK, did not affect PEDV pseudovirus entry into Huh-7 cells

(human liver) or PK-15 cells (porcine kidney) (Fig. 1B, 1C). As a positive control, MERS-CoV pseudoviruses demonstrated decreased entry into Huh-7 cells in the presence of the proprotein convertase inhibitor (Fig. 1D), consistent with previous observations that MERS-CoV entry could be activated by proprotein convertases after virus endocytosis in virus-targeted cells (13). Thus, proprotein convertases from virus-targeted cells do not proteolytically activate PEDV spike or PEDV entry either.

Role of cell-surface proteases in PEDV pseudovirus entry- Next we investigated whether cell-surface proteases activate PEDV entry. Previous studies demonstrated that Huh-7 cells do not express cell-surface serine protease TMPRSS2 (19,20). Here we showed that PEDV pseudoviruses entered Huh-7 cells efficiently, indicating that PEDV entry does not require activation by TMPRSS2 (Fig. 2A). Furthermore, exogenously expressing TMPRSS2 in Huh-7 cells did not enhance PEDV pseudovirus entry (Fig. 2A). Additionally, a TMPRSS2 inhibitor, camostat, had no impact on PEDV pseudovirus entry into Huh-7 cells exogenously expressing or not expressing TMPRSS2 (Fig. 2A). These results all suggest that TMPRSS2 does not activate PEDV entry into host cells. As a positive control, MERS-CoV pseudovirus entry was enhanced in Huh-7 cells exogenously expressing TMPRSS2 (Fig. 2A, 2B). Moreover, the enhanced MERS-CoV pseudovirus entry in Huh-7 cells exogenously expressing TMPRSS2 could be reversed by camostat (Fig. 2B). These results were consistent with previous observations that MERS-CoV entry could be activated by TMPRSS2 (21,22). As a negative control, camostat did not affect MERS-CoV pseudovirus entry into Huh-7 cells not expressing TMPRSS2 (Fig. 2B). Therefore, we can rule out the role of cell-surface serine proteases in processing PEDV spike and activating PEDV entry.

Role of lysosomal cysteine proteases in PEDV pseudovirus entry- Then we examined whether lysosomal cysteine proteases activate PEDV entry. To this end,

we carried out PEDV pseudovirus entry into Huh-7 or PK-15 cells in the presence of lysosomal acidification inhibitor, bafilomycin A1, or lysosomal cysteine protease inhibitor, E-64d. We found that both inhibitors significantly reduced PEDV pseudovirus entry into host cells in a dose-dependent manner (Fig. 3A and 3B). As a control, only bafilomycin A1, but not E-64d, significantly reduced VSV pseudovirus entry into Huh-7 and PK-15 cells (Fig. 3C and 3D). The result from the control experiment is consistent with previous reports that VSV entry into host cells depends on endocytosis, but not lysosomal cysteine proteases (23,24). The control experiment also showed that the inhibitors did not have non-specific cytotoxic effects on target cells. Thus, lysosomal cysteine proteases play a critical role in PEDV entry.

We went further to pinpoint the specific lysosomal cysteine proteases that cleave PEDV spike and activate PEDV entry. We focused on cathepsin L and cathepsin B because both of these cathepsins have been previously identified to process the spike proteins from other coronaviruses including SARS and MERS coronaviruses (19,24-27). To identify the role of cathepsin L and cathepsin B in PEDV entry, we carried out PEDV pseudovirus entry in the presence of inhibitors that are specific for cathepsin L (i.e., inhibitor Z-FY-CHO) or cathepsin B (i.e., CA-074 Me), respectively. The result showed that both inhibitors dramatically reduced PEDV pseudovirus entry into Huh-7 and PK-15 cells (Fig. 4A, 4B).

To provide direct biochemical evidence that cathepsin L and cathepsin B cleave PEDV spike, we expressed PEDV spike in HEK293T cells, lysed the cells, and treated the cell-expressed PEDV spike with recombinant cathepsin L and cathepsin B, respectively, at pH 5.6 (i.e., the working pH for cathepsins). We then detected the cleavage state of the cell-expressed PEDV spike using western blot analysis. Our result showed that at relatively low concentrations (e.g., 1 μ g/ml), cathepsin L cleaved PEDV spike to S2 (Fig. 4C). At higher concentrations (e.g. 4 μ g/ml), cathepsin L further cleaved PEDV S2. On the other hand, at relative low

concentrations (e.g., 1 $\mu\text{g/ml}$), cathepsin B did not cleave PEDV spike efficiently. At higher concentrations (e.g. 10 $\mu\text{g/ml}$), cathepsin B cleaved PEDV spike to S2, but failed to further cleave PEDV S2 (Fig. 4C). Hence PEDV spike is more sensitive to the cleavage of cathepsin L than to the cleavage of cathepsin B. In sum, host lysosomal cysteine proteases, particularly cathepsin L and cathepsin B, process PEDV spike and activate PEDV entry.

Recognizing that lysosomal cysteine proteases differ in their expression levels among different tissues (28), we selected IPI-21 cells (porcine small intestines) for repeating PEDV pseudovirus entry in the presence of lysosomal cysteine proteases (Fig. 5). Porcine small intestines are the major target organ for PEDV infections (29-31). Our results showed that PEDV pseudovirus entry into IPI-21 cells could be inhibited by lysosomal acidification inhibitor bafilomycin A1, lysosomal cysteine protease inhibitor E-64d, and cathepsin-L- and cathepsin-B-specific inhibitors. Therefore, lysosomal cysteine proteases activate PEDV entry into cells from porcine small intestines.

Role of extracellular proteases in PEDV pseudovirus entry- We also tackled the confounding role of extracellular protease trypsin in PEDV entry. Previous studies showed that exogenous trypsin could activate the entry of SARS and MERS coronaviruses into host cells after the viruses had already been attached to host cells (19,26,32). Hence we added trypsin after PEDV pseudoviruses had been attached to Huh-7 or PK-15 cells. Our result revealed that trypsin slightly reduced PEDV pseudovirus entry into Huh7 and PK-15 cells (Fig. 6A, 6B). On the other hand, in the presence of cathepsin L or cathepsin B inhibitor, the dramatically reduced PEDV pseudovirus entry into host cells could be partially rescued by extracellular trypsin (Fig. 4A, 4B). Taken together, extracellular trypsin has the potential to process PEDV spike when lysosomal cysteine proteases are inhibited or unavailable; however, when available, lysosomal cysteine proteases play the major role in PEDV entry into host cells.

Role of lysosomal cysteine proteases in live PEDV entry- Last we investigated the role of lysosomal cysteine proteases in live PEDV infection in cell culture (Fig. 7). Without trypsin, PEDV replicated inefficiently in Vero CCL81 cells (monkey kidney), but still at a detectable level. PEDV replication in Vero CCL81 cells was reduced to nearly undetected levels by lysosomal cysteine protease E-64d, cathepsin L inhibitor, or cathepsin B inhibitor. These results are consistent with pseudovirus entry assay, confirming that lysosomal cysteine proteases play critical roles in PEDV entry into host cells.

DISCUSSION

Our study has elucidated a long-standing puzzle regarding what proteases activate PEDV entry into host cells. Previous studies identified extracellular protease trypsin as required for PEDV infection in cell culture, which led to the conclusion that intestinal proteases are essential for PEDV entry (14-16). However, these previous studies all used PEDV live virus particles, and thereby were unable to differentiate between PEDV entry and other steps in the PEDV infection cycle such as virus replication or release. Indeed, an electron microscopic study showed that PEDV release is a limiting step in the PEDV infection cycle and that trypsin is required for PEDV release (33). To separate PEDV entry from other steps of the PEDV infection cycle, we performed a PEDV pseudovirus entry assay in which PEDV-spike-packaged pseudovirus particles can only enter host cells, but cannot replicate or be released. Thus the PEDV pseudovirus entry assay provides a simplified system for studying PEDV entry (17). Using this assay, we showed that PEDV entry does not depend on proprotein convertases or cell-surface proteases. Instead, PEDV entry is activated by lysosomal cysteine proteases. Using both pseudovirus entry and direct biochemical assays, we further identified cathepsin L and cathepsin B as the specific lysosomal cysteine proteases that can process PEDV spike. We obtained the result using several cell lines including IPI-21 cells from porcine small intestines, the major target organ for PEDV infections. Hence our finding

is relevant for pig infection and development of antivirals. Further, we confirmed this result using live PEDV infection assay in cell culture. Our study also elucidated the puzzling role of extracellular trypsin in PEDV entry. When cathepsins are available, trypsin is not as efficient as cathepsins in activating PEDV entry. However, when cathepsins are inhibited or unavailable, trypsin can fill in the role of cathepsins by activating PEDV entry. Because the current study focuses on the PEDV entry, the role of trypsin in other steps of the PEDV infection cycle such as PEDV release remains to be investigated using other research approaches. Nevertheless, our study has laid out a blueprint for systematically examining the roles of proteases in virus entry, and provided insight into the puzzling molecular mechanism for PEDV entry.

PEDV is currently sweeping through America's pig populations with little hindrance, as neither vaccines nor antiviral drugs are available to curb its spread. Our study suggests that cysteine protease inhibitors, such as MDL 28170, can serve as a class of antiviral agents that potentially block PEDV infections (34). Moreover, our finding suggests that cysteine protease inhibitors alone may not be sufficient to block PEDV entry because trypsin can serve a backup role in activating PEDV entry. Instead, a combinational use of cysteine protease and trypsin inhibitors may be more effective to block PEDV entry and treat infected pigs.

MATERIALS AND METHODS

Cell lines and plasmids- HEK293T (human embryonic kidney), PK-15 (porcine kidney), and Vero CCL81 (monkey kidney) cells were obtained from American Type Culture Collection. IPI-21 (porcine small intestine) cells were purchased from Sigma-Aldrich. Huh-7 (human hepatoma) cells were kindly provided by Dr. Charles M. Rice at Rockefeller University. These cell lines were cultured in Dulbecco's modified Eagle medium (DMEM) supplemented with 10% fetal bovine serum (FBS), 2mM L-glutamine, 100 units/mL penicillin, and 100 µg/mL streptomycin (Life Technologies).

The genes of MERS-CoV spike (GenBank accession number AFS88936.1) and PEDV spike (GenBank accession number AGO58924.1) were each cloned into pcDNA3.1(+) vector (Life Technologies) with a C-terminal C9 tag. The genes of cathepsin L (GenBank accession number CAA30981.1) and cathepsin B (GenBank accession number AAA52129.1) were synthesized from GenScript, and were each cloned into pFastBac1 vector (Life Technologies) with a C-terminal His₆ tag. The plasmid of human TMPRSS2 was kindly provided by Dr. Tom Gallagher at Loyola University Medical Center.

Protein expression and purification- Cathepsin L and cathepsin B were expressed as inactive proform in insect cells and purified as previously described (19). Briefly, the full length proteases containing a C-terminal His₆ tag were expressed in Sf9 insect cells using the Bac-to-Bac expression system (Life Technologies), secreted to cell culture medium, and purified using HiTrap Chelating HP column and Superdex 200 gel filtration column (GE Healthcare), sequentially. Purified pro-cathepsin L and pro-cathepsin B were auto-activated as previously described (35). Briefly, purified pro-cathepsin L or pro-cathepsin B were diluted 10 fold using 100 mM sodium acetate, pH 4.0, and incubated at 37°C for 1 hour. The activated mature proteases were further purified on a Superdex 200 gel filtration column.

Pseudovirus entry into human and pig cells- Retroviruses pseudotyped with PEDV spike, MERS-CoV spike, or VSV envelope glycoprotein were generated as previously described (17,19). Briefly, HEK293T cells were co-transfected with a plasmid carrying Env-defective, luciferase-expressing HIV-1 genome (pNL4-3.luc.R-E-) and a plasmid encoding PEDV spike or MERS-CoV spike using Lipofectamine 3000 reagent (Life Technologies) according to the manufacturer's instructions. Supernatants containing pseudoviruses were harvested 72 hours after transfection, and centrifuged at 1200 × g for 10 min to remove cell debris. Pseudoviruses were concentrated using Amicon Ultra centrifugal filter units with a 100 kDa

molecular weight cut-off, and were then aliquoted and frozen at -80°C for later use.

Retroviruses pseudotyped with PEDV spike, MERS-CoV spike, VSV envelope glycoprotein, or empty vector (mock) were used to transduce Huh-7 cells, Huh-7 cells transiently expressing human TMPRSS2, PK-15 cells, or IPI-21 cells in 96-well plates. For trypsin processing of pseudoviruses, the initial DMEM medium was removed after pseudovirus attachment for 2 hours, and was subsequently replaced with serum-free DMEM with 0 µg/ml, 10 µg/ml, or 40 µg/mL TPCK-treated trypsin (Sigma-Aldrich). After incubating with trypsin at 37°C for 10 min, DMEM supplemented with 75 µg/mL soybean trypsin inhibitor (Sigma-Aldrich) was added to neutralize trypsin. Cells were incubated at 37°C for another 12 hours, and medium was replaced with fresh DMEM. 48 hours later, cells were washed with phosphate-buffered saline (PBS) and lysed. Aliquots of cell lysates were transferred to Optiplate-96 plate (PerkinElmer), and luciferase substrate (Promega) was added. Relative luciferase units (RLU) were measured using EnSpire plate reader (PerkinElmer).

Inhibition of pseudovirus entry into human and pig cells using inhibitors- Inhibition of pseudovirus entry using various protease inhibitors was carried out as previously described (18). Briefly, target cells were pre-incubated with 10 µM or 50 µM proprotein convertases inhibitor Dec-RVKR-CMK (Enzo Life Sciences), 20 µM or 100 µM camostat mesylate (Sigma-Aldrich), 20 nM or 100 nM bafilomycin A1 (Sigma-Aldrich), 10 µM or 50 µM E-64d (Sigma-Aldrich), 50 µM cathepsin L inhibitor Z-FY-CHO (Santa Cruz Biotechnology), or 50 µM cathepsin B inhibitor CA-074 Me (Santa Cruz Biotechnology) at 37°C for 1 hour. The cells were subsequently transduced by retroviruses pseudotyped with PEDV spike, MERS-CoV spike, or VSV envelope glycoprotein. The cells were incubated at 37°C for 6-8 hours, and then medium was replaced with fresh DMEM. 48 hours later, the cells were lysed and measured for luciferase activity.

Western blot analysis of spike cleavage by proprotein convertases- PEDV and MERS-CoV pseudoviruses were packaged in HEK293T cells, lysed and then subjected to western blot analysis. The C9-tagged spikes were detected using anti-C9 tag monoclonal antibody (Santa Cruz Biotechnology).

Western blot analysis of spike cleavage by lysosomal cysteine proteases- HEK293T cells were transfected with plasmids encoding PEDV spike or MERS-CoV spike. 48 hours after transfection, the cells were harvested, washed with PBS, and lysed by sonication. Cell lysates were then incubated with activated cathepsin L or cathepsin B at gradient concentrations at pH 5.6 and 37°C for 30 minutes, and subjected to western blot analysis. The C9-tagged spikes were detected using anti-C9 tag monoclonal antibody (Santa Cruz Biotechnology).

Inhibition of live PEDV entry into host cells using inhibitors- Vero CCL81 cells were washed 3 times with DMEM, and were pre-treated with 50 µM E-64d, 50 µM Cathepsin L inhibitor, or 50 µM Cathepsin B inhibitor in DMEM. After 1 hour, cells were infected with PEDV strain Ohio VBS2 at a multiplicity of infection (MOI) of 0.5 as previously described (17). 2 hours after infection, cells were washed with DMEM 3 times to remove unbound PEDV particles. DMEM supplemented with 0.018% (w/v) Tryptose Phosphate Broth (TPB) (Sigma), 0.02% yeast extract (Sigma), and the respective inhibitor at the above concentration was then added. 24 hours after infection, cells were washed twice with PBS and fixed with 4.0% (v/v) paraformaldehyde and 0.2% (v/v) glutaraldehyde in 0.1 M potassium phosphate buffer (PPB), pH 7.4, at 22°C for 15 minutes. They were then washed 3 times with PBS. After permeabilization with 0.1% Triton X-100 in PBS for 15 minutes, the cells were washed with PBS, blocked with PBS containing 2% bovine serum albumin at room temperature for 1 hour. Cells were then incubated with fluorescein isothiocyanate (FITC)-labeled mouse anti-PEDV N protein antibody (Medgene labs, Brookings, SD) in 0.2% BSA in PBS at dilution of 1:100 at 4°C overnight. Cells were examined under an Olympus fluorescent microscope system.

Conflict of interest: The authors declare that they have no conflicts of interest with the contents of this article.

Author contributions: CL, YZ, SJ, LD, JL, FL designed experiments and wrote the manuscript; CL, YM, YY, YZ, JS performed the experiments; CL, YM, YY, YZ, JS, YZ, SJ, LD, JL, FL analyzed the data.

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FOOTNOTES

This work was supported by NIH grants R01AI089728 (to FL).

FIGURE LEGENDS

FIGURE 1. Proprotein convertases do not activate PEDV pseudovirus entry. (A) Western blot analysis of PEDV spike in pseudovirus particles. Retroviruses pseudotyped with PEDV spike (i.e., PEDV pseudoviruses) were produced in HEK293T cells and then subjected to western blot analysis using antibody against its C-terminal C9 tag. (B) Huh-7 cells or (C) PK-15 cells were pre-incubated with PPCi (proprotein convertases inhibitor, Dec-RVCR-CMK) at indicated concentrations, and then transduced by PEDV pseudoviruses. Empty vector-packaged pseudoviruses (mock) were used as a negative control. (D) As a positive control, Huh-7 cells were transduced by retroviruses pseudotyped with MERS-CoV spike (i.e. MERS-CoV pseudoviruses). The pseudovirus entry efficiency was characterized as luciferase activity accompanying the entry. The pseudovirus entry in target cells without any inhibitor treatment was taken as 100%. Error bars indicate SEM (n = 5).

FIGURE 2. Cell-surface serine proteases do not activate PEDV pseudovirus entry. Huh-7 cells transiently transfected with empty pCAGGS vector or TMPRSS2 in pCAGGS vector were pre-incubated with camostat (cell-surface serine proteases inhibitor) at indicated concentrations, and were transduced by PEDV pseudoviruses (A) or MERS-CoV pseudoviruses (B). The pseudovirus entry in empty pCAGGS vector-transfected Huh-7 cells without any inhibitor treatment was taken as 100%. Error bars indicate SEM (n = 5).

FIGURE 3. Lysosomal cysteine proteases activate PEDV pseudovirus entry. (A) Huh-7 cells or (B) PK-15 cells were pre-incubated with Baf-A1 (Bafilomycin A1) (lysosomal acidification inhibitor) or E-64d (lysosomal cysteine protease inhibitor) at indicated concentrations, and then transduced by PEDV pseudoviruses. (C) and (D) Retroviruses pseudotyped with VSV envelop glycoprotein (i.e. VSV pseudoviruses) were used as a control. The pseudovirus entry in target cells without any inhibitor treatment was taken as 100%. Error bars indicate SEM (n = 5).

FIGURE 4. Cathepsin L and cathepsin B activate PEDV pseudovirus entry. Before being infected by PEDV pseudoviruses, Huh-7 cells (A) or PK-15 cells (B) were pre-incubated with 50 μ M cathepsin L inhibitor (i.e., Z-FY-CHO) or 50 μ M cathepsin B inhibitor (i.e., CA-074 Me). After pseudovirus attachment to target cells, unbound pseudovirus particles were removed and bound pseudovirus particles were either treated or not treated with 40 μ g/ml exogenous trypsin. PEDV pseudovirus entry in the absence of inhibitor or exogenous trypsin was taken as 100% in each cell line. PEDV pseudovirus entry in the absence of inhibitor and in the present of trypsin was shown separately in Fig. 6. Error bars indicated SEM (n = 4). (C) PEDV spike was transiently expressed in HEK293T cells, the cells were lysed through sonication, and the expressed spike protein was subsequently subjected to cathepsin L or cathepsin B cleavage at gradient concentrations at pH 5.6. The spike was detected using an antibody against its C-terminal C9 tag.

FIGURE 5. Lysosomal cysteine proteases activate PEDV pseudovirus entry into porcine small intestine cells. IPI-21 cells from porcine small intestines, which are the primary target organ for PEDV, were pre-incubated with Baf-A1, E-64d, cathepsin L (CTSL) inhibitors (i.e., Z-FY-CHO) and cathepsin B (CTSL) inhibitors (i.e., CA-074 Me) at indicated concentrations, and then transduced by PEDV pseudoviruses. Empty pcDNA vector-packaged pseudoviruses (mock) were used as a negative control. The pseudovirus entry mediated by PEDV spike in the absence of any inhibitor was taken as 100%. Error bars indicate SEM (n = 4).

FIGURE 6. Extracellular protease trypsin serves a backup role in activating PEDV pseudovirus entry. After PEDV pseudoviruses were incubated with Huh-7 cells (A) or PK-15 cells (B), unbound PEDV pseudovirus particles were removed and bound PEDV pseudovirus particles were either treated or not treated with exogenous trypsin at indicated concentrations. Empty pcDNA vector-packaged pseudoviruses (mock) were used as a negative control. The pseudovirus entry mediated by PEDV spike in the absence of exogenous trypsin was taken as 100%. Error bars indicate SEM (n = 4).

FIGURE 7. Lysosomal cysteine proteases activate live PEDV entry into cells. Vero CCL81 cells were pre-treated with one of lysosomal cysteine protease inhibitors (E64d, cathepsin L inhibitor, or cathepsin B inhibitor), before they were infected by live PEDV. 24 hours post-infection, cells were chemically fixed and incubated with fluorescein isothiocyanate (FITC)-labeled mouse anti-PEDV N protein antibody. PEDV-positive cells were observed using a fluorescence microscope.

Figure 1

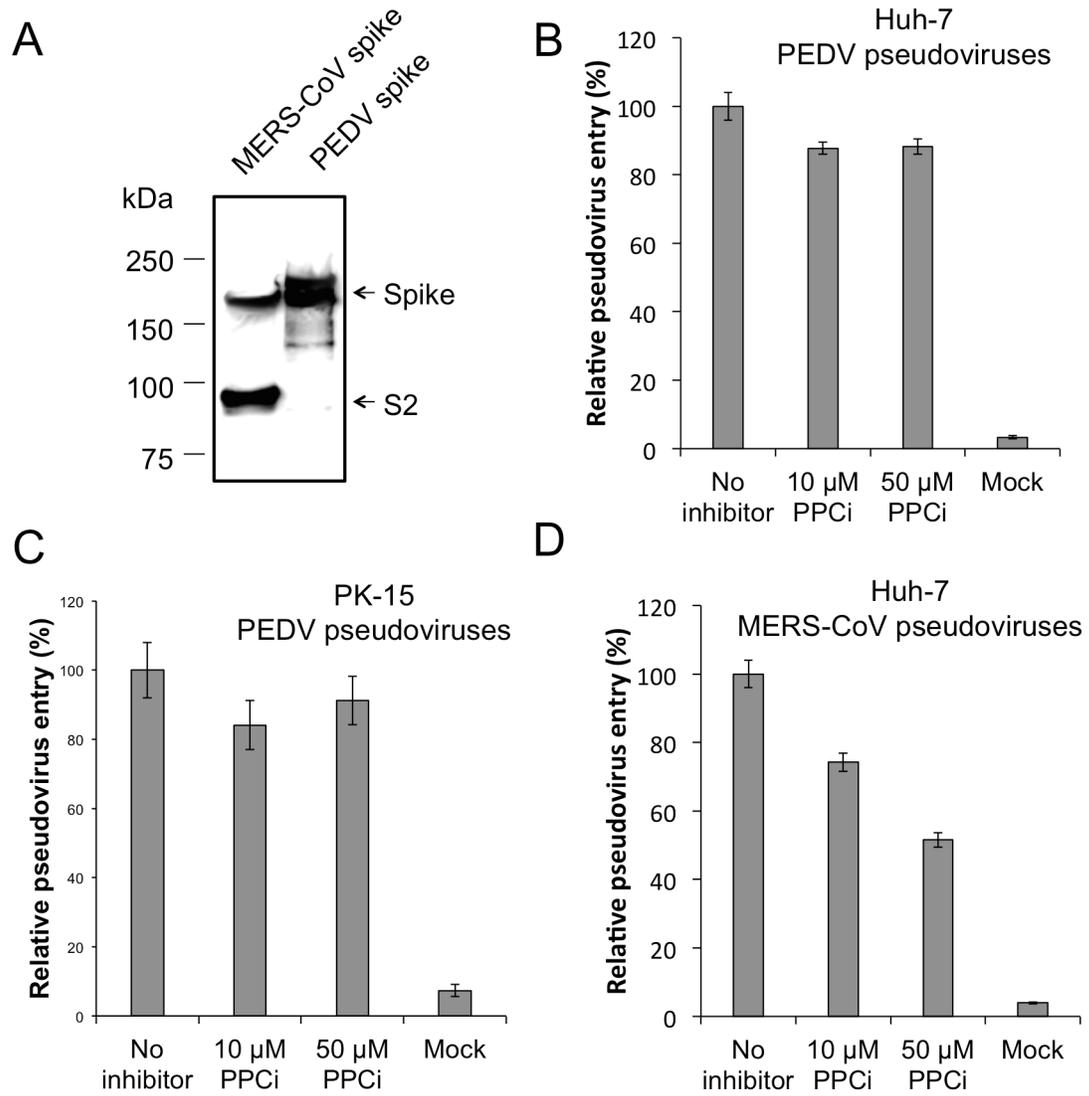


Figure 2

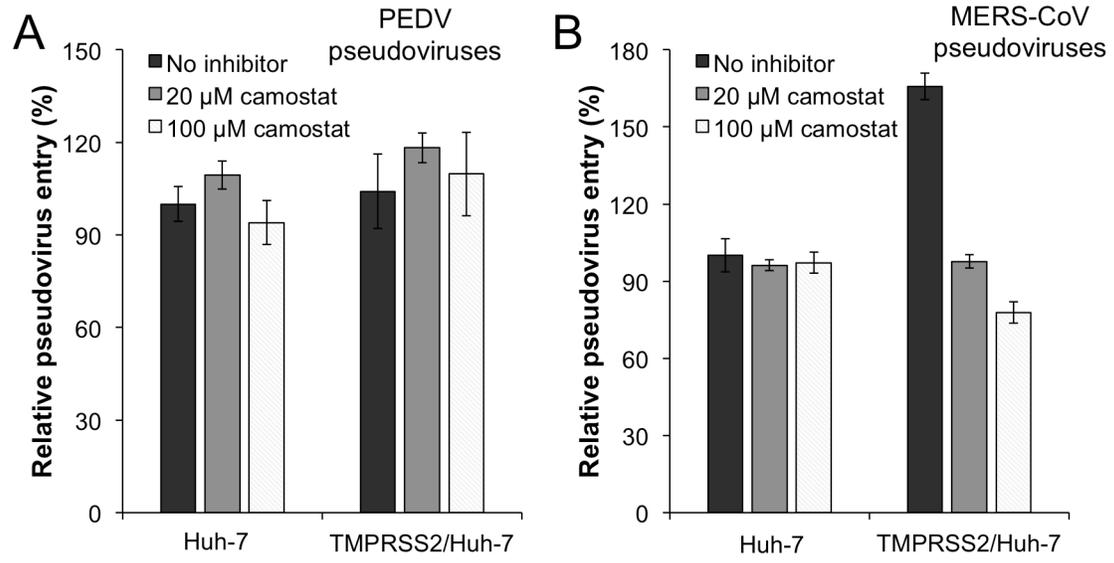


Figure 3

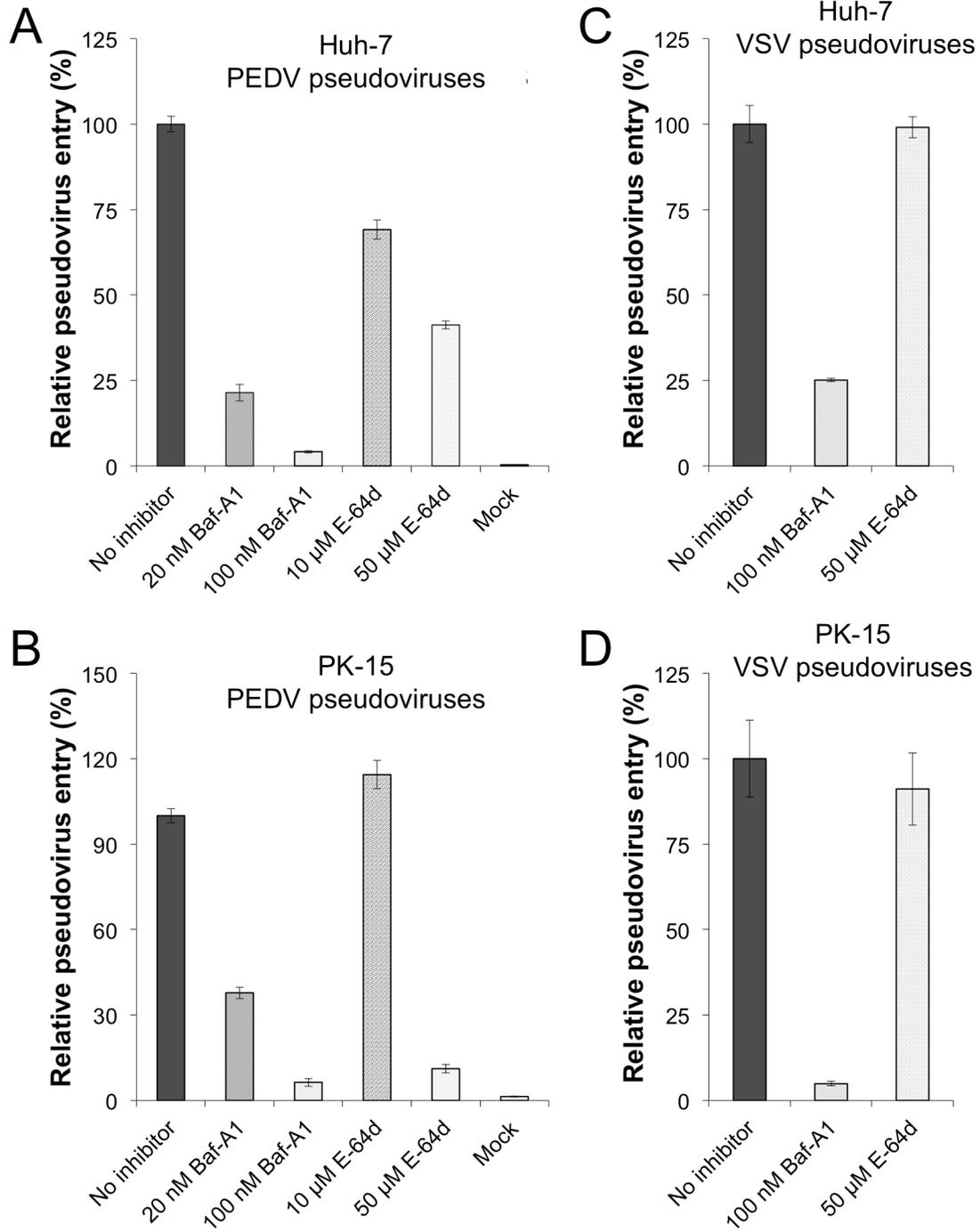


Figure 4

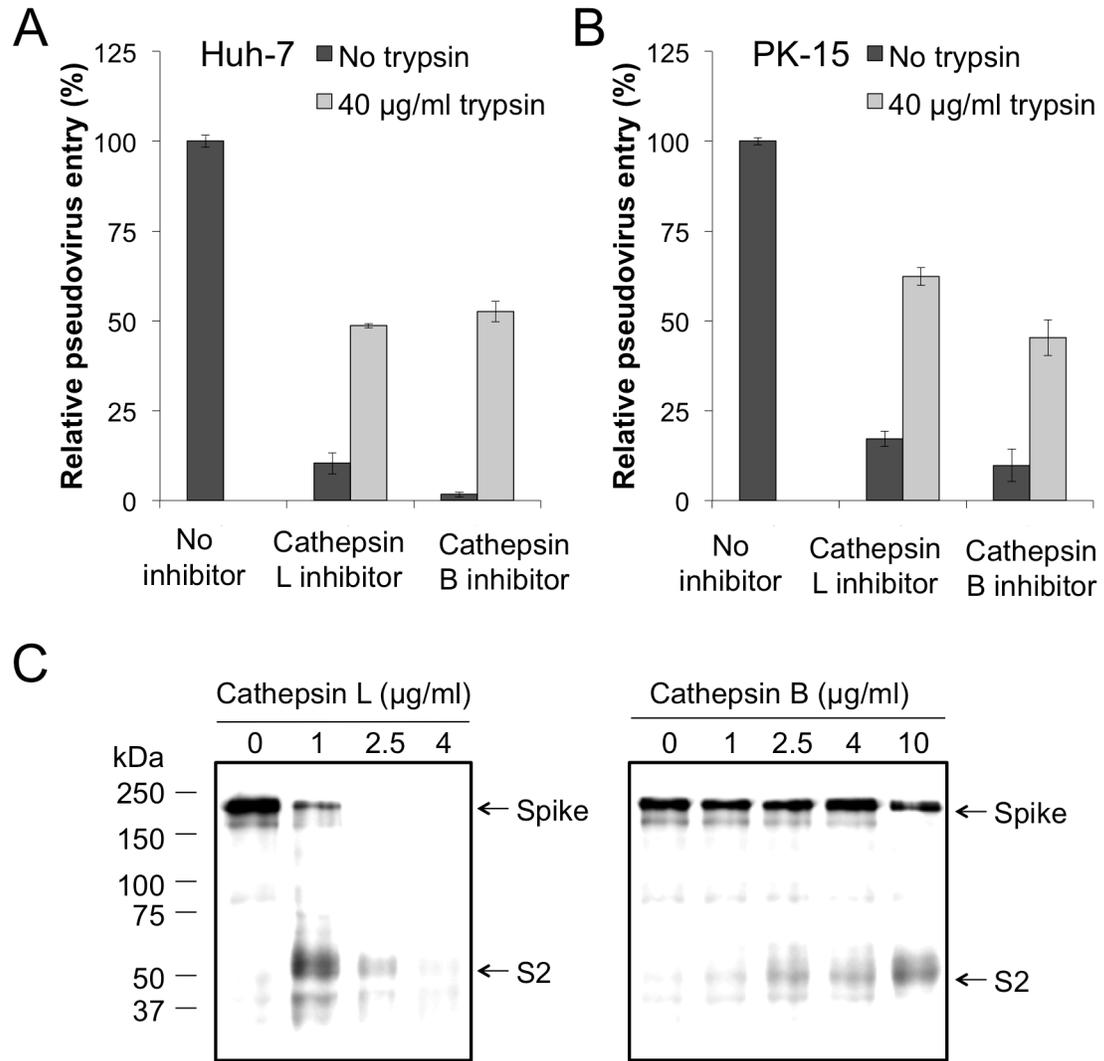


Figure 5

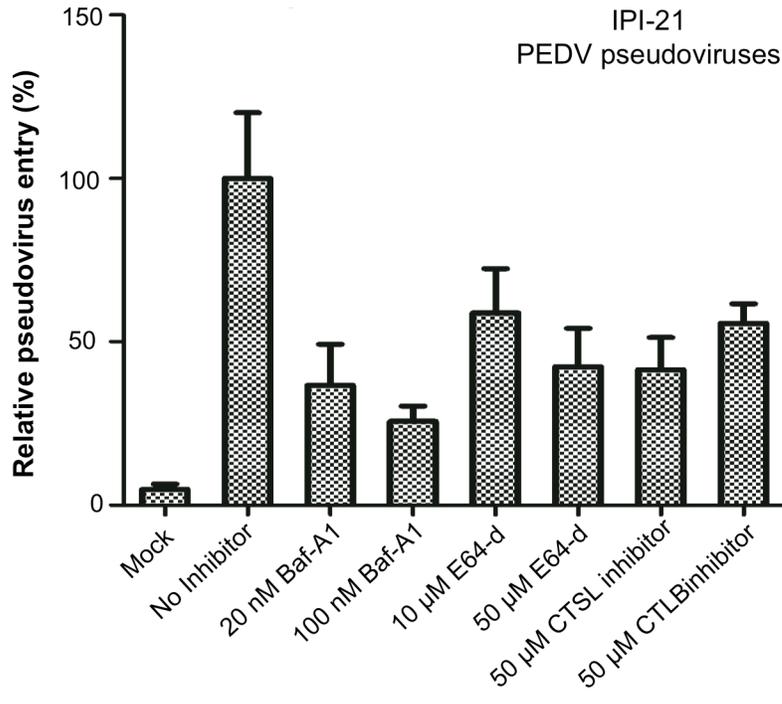


Figure 6

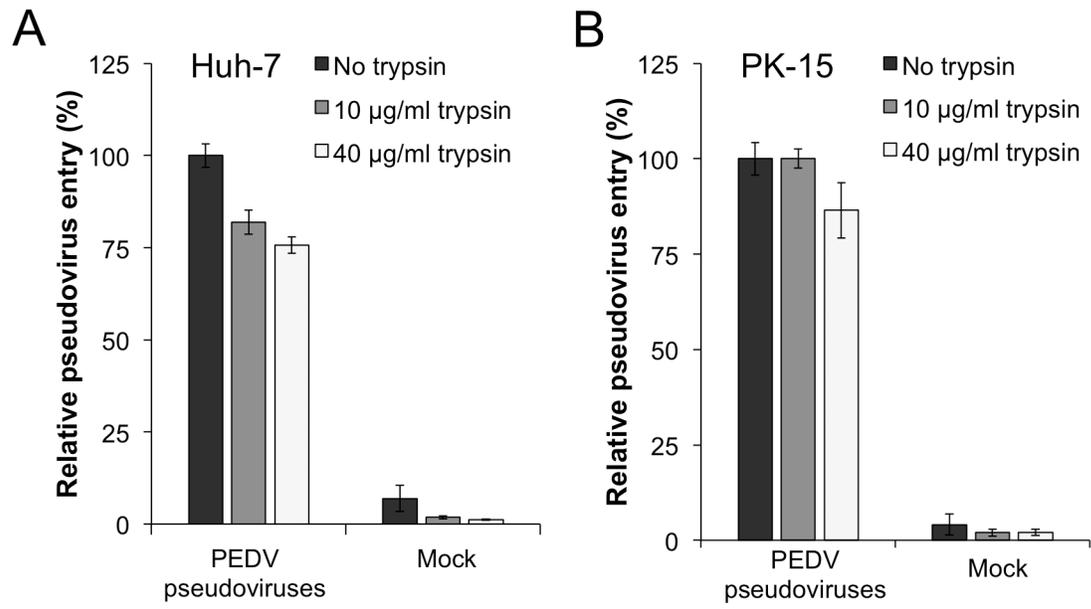
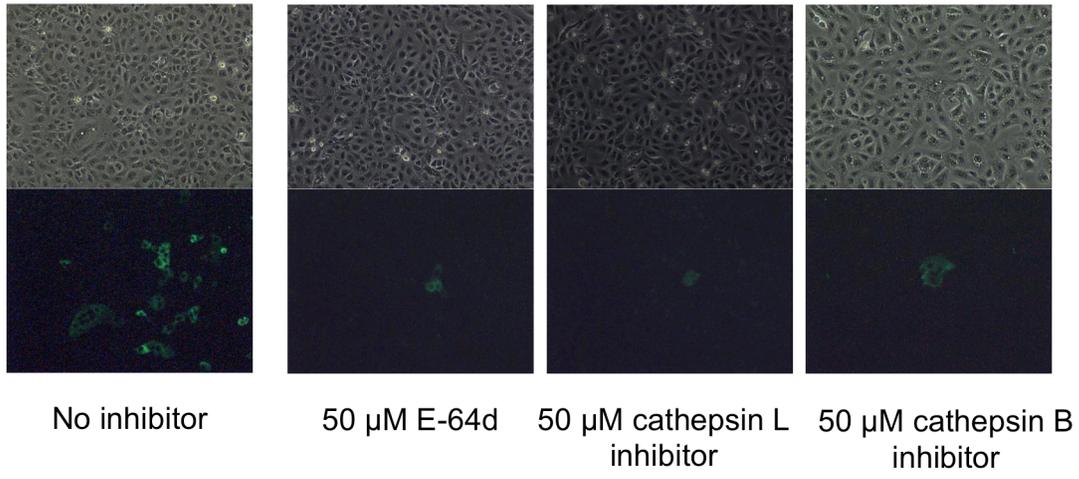


Figure 7



Cell entry of porcine epidemic diarrhea coronavirus is activated by lysosomal proteases

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J. Biol. Chem. published online October 11, 2016

Access the most updated version of this article at doi: [10.1074/jbc.M116.740746](https://doi.org/10.1074/jbc.M116.740746)

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