Infection with a wide variety of viruses often perturbs host cell signaling pathways including the Jun NH2-terminal kinase/stress-activated kinase (JNK/SAPK) and the p38 mitogen-activated protein kinase (p38/MAPK), which are important components of cellular signal transduction pathways. The present study demonstrated for the first time that porcine circovirus type 2 (PCV2), which is the primary causative agent of an emerging swine disease, postweaning multisystemic wasting syndrome, can activate JNK1/2 and p38 MAPK pathways in PCV2-infected PK15 cells. However, PCV2 at an early stage of infection, as well as UV-irradiated PCV2, failed to activate these two MAPK families, which demonstrated that PCV2 replication was necessary for their activation. We further found that PCV2 activated the phosphorylation of JNK1/2 and p38 MAPK downstream targets c-Jun and ATF-2 with virus replication in the cultured cells. The roles of these kinases in PCV2 infection were further evaluated using specific inhibitors: the JNK inhibitor 1 for JNK1/2 and SB202190 for p38. Inhibition of JNK1/2 and p38 kinases by these specific inhibitors did result in significant reduction of PCV2 viral mRNA transcription and protein synthesis, viral progeny release, and blockage of PCV2-induced apoptotic caspase-3 activation in the infected cells. Taken together, these data suggest that JNK/SAPK and p38 MAPK pathways play important roles in the PCV2 replication and contribute to virus-mediated changes in host cells.

Porcine circovirus (PCV) is classified in the genus Circovirus of the family Circoviridae (52). PCV was first discovered in 1974 as a contaminant of a continuous porcine kidney cell line (PK15) (49). Two genotypes of PCV have been identified. PCV type 1 (PCV1) is known to be nonpathogenic to pigs (1). A high prevalence of anti-PCV1 antibodies has been detected in the swine population, but no disease is correlated with PCV1 infection (33, 50). In contrast, infection with PCV2 occurs in all swine-producing areas of the world, and related PCV2-associated diseases are increasingly recognized as serious threats to global pig production (2, 7, 11, 13, 34, 39, 43). A primary manifestation of PCV2 infection is postweaning multisystemic wasting syndrome (PMWS), which appears in pigs aged 5 to 18 weeks and is clinically characterized by fever, wasting or unthriftiness, respiratory distress, enlarged lymph nodes and, occasionally, jaundice and diarrhea (8, 19, 43). Mortality rates may vary from 1 to 2% up to 30% in complicated cases when coinfections with porcine reproductive and respiratory syndrome virus, porcine parvovirus, or Mycoplasma hyoovipneumoniae (12). Microscopic lesions are characterized by lymphocyte depletion of follicular and interfollicular areas together with macrophage infiltration of lymphoid tissues in PMWS-affected pigs. Several lines of field and experimental evidence have suggested that severely PMWS affected pigs may develop immunosuppression (44).

PCV genome is a circular single-stranded DNA molecule of ~1.7 kb. Two major open reading frames (ORFs) have been recognized for PCV: ORF1, called the rep gene, which encodes a protein of 35.7 kDa involved in virus replication (35), and ORF2, called the cap gene, which encodes the major immunogenic capsid protein of 27.8 kDa (5, 38). In addition to the replicase ORF1 and the capsid protein ORF2, a novel protein, ORF3, has been detected in PCV2 productive infection and is not essential for PCV2 replication in the cultured cells but is involved in viral pathogenesis via an apoptotic function (30, 31). In a recent report, we demonstrated that PCV2 infection induces NF-kB activation in the cultured cells and further found the role of NF-kB activation in PCV2 replication and PCV2-induced apoptotic caspase activity (56). However, whether other signaling pathways may also contribute to PCV2 infection in the cultured cells remains unclear.

Mitogen-activated protein kinases (MAPKs), including extracellular signal-regulated kinases (ERK1/2), c-Jun NH2-terminal kinase/stress-activated protein kinase (JNK/SAPK), and p38 MAPK, are central components of signal transduction pathways in the regulation of cell proliferation and differentiation, cytokine production, and apoptosis (17). ERK1/2 is primarily activated by growth factors, cytokines, and phagocytosis, whereas JNK and p38 are potently induced by proinflammatory cytokines, bacterial endotoxins, and environmental stresses (17, 26). Although there is coordinated regulation of JNK/SAPK and p38 MAPK, they have corresponding downstream targets. Activated JNK/SAPK and p38 MAPK can phosphorylate numerous substrates, including a variety
of transcription factors. The transcription factor c-Jun is specifically phosphorylated by JNK/SAPK, while ATF-2 is phosphorylated by both JNK/SAPK and p38 MAPK (9, 18, 29). Phosphorylation and activation of these downstream substrates ultimately alters gene expression, thereby manifesting the biological consequences of JNK/SAPK and p38 MAPK activation (57).

It has been shown that viruses are ultimately dependent on the host cell for their replication via altering cellular signal transduction pathways, including MAPK cascades. For instance, modulation of JNK and/or p38 MAPK pathways is required for infection and replication of human immunodeficiency virus type 1 (27), herpes simplex virus type 1 (HSV-1) (60), Epstein-Barr virus (16), Kaposi’s sarcoma-associated herpesvirus (41, 58), echovirus 1 (24), Sindbis virus (37), encephalomyocarditis virus (21), coxsackievirus B3 (46), and varicellazoster virus (42, 61). However, there is still no report on the activation of JNK/SAPK and p38 MAPK induced by PCV2 replication and the effects of their activation on PCV2 replication.

The present study was initiated to determine whether JNK/SAPK and p38 MAPK play a role in the replication of PCV2. Here, we report that phosphorylation of JNK/SAPK and p38 MAPK is increased during PCV2 replication and demonstrate that activation of these two MAPK pathways is essential for efficient PCV2 replication. Inhibition of these two MAPK activations reduces viral mRNA transcription and protein synthesis, viral progeny release, and PCV2-induced apoptotic caspase activity. These findings indicate that the JNK/SAPK and p38 MAPK pathways are involved in the PCV2 replication and contribute to virus-mediated changes in host cells.

MATERIALS AND METHODS

Virus and cells. The permanent PK15 cell line, which was free of PCV, was maintained in minimal essential medium supplemented with 5% heat-inactivated fetal bovine serum (FBS), 5% l-glutamine, 100 U of penicillin G/ml, and 100 µl of streptomycin/ml at 37°C in a humidified 5% CO₂ incubator. The PCV2 used in the study was originally isolated from a kidney tissue sample of a pig with naturally occurring PMWS (strain BJW) (31).

For PCV2 infection, PK15 cells seeded the day before were infected with PCV2 strain BJW. At 72 h, the cells were washed with PBS and fixed in 4% paraformaldehyde. After three washes, the cells were incubated with mouse anti-ORF2 antibody (56) diluted in 5% bovine serum albumin-PBS at room temperature for 1 h. After three further washes, the cells were incubated with HRP-conjugated anti-mouse immunoglobulin G (Sigma) at room temperature for 1 h and washed with PBS three times. Color development was carried out with 3,3’-diaminobenzidine tetrahydrochloride (Pierce) substrate and then stopped with distilled water. The cells were examined under a microscope, and cells positive for PCV2 viral antigens were counted in six fields of view.

Whole-cell lysates. Whole-cell lysate extracts from PK15 cells at various time points after infection were prepared with a nuclear extract kit (Active Motif) according to the manufacturer’s protocol. Protein concentration was determined by using a protein assay (Bio-Rad, Hercules, CA) with bovine serum albumin as a standard.

Western blotting. The whole-cell lysate extracts prepared as described above were diluted in 2× sample buffer and boiled for 5 min. Portion (20 µg) of each extract were resolved by sodium dodecyl sulfate–10% polyacrylamide gel electrophoresis (SDS–10% PAGE) and blotted onto nitrocellulose (NC) membranes (Stratagene) with a semidyel transfer cell (Bio-Rad). The membranes were blocked for 2 h at room temperature in blocking buffer TBST (20 mM Tris-HCl [pH 7.4], 150 mM NaCl, 0.1% Tween 20) containing 5% skim milk powder to prevent nonspecific binding and then incubated with specific primary antibodies raised against ORF2, JNK1/2, p38, c-Jun, ATF-2, p-JNK1/2, p-p38, p-c-Jun, and p-ATF-2, as well as β-actin, at room temperature for 2 h. The membranes were washed three times with TBST buffer and then incubated for 2 h at room temperature with HRP-conjugated secondary antibodies diluted in blocking buffer (1:2,000). Immunoreactive bands were visualized by using an enhanced chemiluminescence system (Amersham Biosciences).

Fluorometric assay of caspase-3 activity. Spectrofluorometric assays of proteolytic activity were carried out using the synthetic fluorogenic substrate 7-amino-4-trifluoromethyl coumarin (AFC) to measure caspase-3 activity. A BD ApoAlert caspase fluorescent assay kit (Clontech) was used to determine caspase-3 activity. In brief, 80% confluent monolayers of PK15 cells grown on 12-well plates were infected with PCV2 at an MOI of 1 TCD₅₀ with various amounts of JNK1/2 or p38 MAPK inhibitor treatment. At 72 h postinfection, cells were harvested at 400 × g for 5 min. Cells (2 × 10⁶ to 3 × 10⁶) were lysed in 50 µl of lysis buffer on ice for 10 min and centrifuged at 16,000 × g for 10 min, and the supernatant was collected. A 50-µl supernatant was added to an equal volume of 2× reaction-dithiothreitol buffer supplemented with caspase-3 substrate DEVD-AFC and then incubated at 37°C for 2 h. The optical densities at 405 nm for caspase-3 were determined. The nanomoles of AFC (released) expressed per hour were calculated from the standard curve.

Statistical analysis. Results are presented as averages ± the standard deviation (SD) or standard errors of the means, as indicated. Statistical comparisons are made by using a Student t test, and differences between groups were considered significant if the P value was <0.05.
RESULTS

Activation of JNK/SAPK and p38 MAPK during PCV2 infection. It was reported that JNK/SAPK and p38 MAPK are phosphorylated during various virus infection. In order to assess whether activation of these two MAPK signaling pathways occurred in the cultured cells during the course of PCV2 infection, the degrees of JNK1/2 and p38 phosphorylation in the PCV2-infected cells were examined by Western blotting. PK15 cells were infected with PCV2 strain BJW at an MOI of 1 TCID_{50}, and whole-cell lysates were prepared and resolved by SDS-PAGE, transferred to nitrocellulose membranes, and immunoblotted. The protein levels of JNK1/2 and p38 and their phosphorylated forms, as well as PCV2 viral capsid protein, were analyzed. The amounts of β-actin were also assessed to monitor the equal loadings of protein extracts. As shown in Fig. 1A, infection with PCV2 led to progressive accumulation of p-JNK1/2 signals over time, and the maximal induction was seen at 72 h postinfection. The kinetics of p-p38 accumulation in the PCV2-infected cells was similar to that of the accumulation of phosphorylated JNK (Fig. 1A). The increased levels of JNK and p38 MAPK phosphorylation were concurrent with the expression of viral capsid protein ORF2 in the infected cells (Fig. 1A). In contrast, the protein levels of total amounts of JNK1/2 and p38 remained unchanged in the PCV2-infected cells at various time points after infection compared to that in the mock-infected cells. A loading control, β-actin in each sample, was comparable.

To further determine activated JNK and p38 MAPK quantitatively in the infected cells, we used FACE assays to investigate the levels of ability of JNK and p38 MAPK phosphorylation at the indicated time points after PCV2 infection. As shown by Fig. 1B and C, there was a time-dependent increase in the JNK1/2 and p38 phosphorylation in the PCV2-infected cells at 72 h postinfection which decreased thereafter. At 72 h after infection, the activations of phosphorylated JNK and p38 showed approximately 4.3- and 3.1-fold increases, respectively,
versus those in the mock-infected cells. In addition, the levels of total JNK and p38 remained unchanged in the PCV2-infected cells at various time points after infection compared to that in the mock-infected cells. These data indicate that PCV2 infection induces the activation of JNK and p38 MAPK signaling pathways.

**PCV2 replication is required for JNK/SAPK and p38 MAPK phosphorylation.** It has been shown that replicating virus can be detected in PCV2-infected PK15 cells as early as 14 h postinfection (59). Therefore, we used the FACE assay to determine whether PCV2 replication was required for JNK and p38 MAPK phosphorylation during the early phase of PCV2 infection. As shown in Fig. 2, no significant increase in activation of phosphorylated JNK or p38 MAPK was observed from 0.25 h through 5 h postinfection. In contrast, the levels of JNK and p38 MAPK phosphorylation increased by 1.8- and 2.3-fold, respectively, at 24 h postinfection. The levels of total JNK and p38 remained unchanged in the PCV2-infected cells at various time points after infection compared to that in the mock-infected cells. In addition, no replicating virus was detected in the PCV2-inoculated cells from 0.25 h to 5 h postinoculation, whereas the accumulation of PCV2 cap spliced transcript was observed at 24 h postinoculation (data not shown).

We further used an UV light-irradiated PCV2 to determine whether phosphorylation of JNK1/2 and p38 MAPK was associated with PCV2 replication. Culture fluid from both PCV2-infected PK15 cells and mock-infected PK15 cells was collected at 72 h postinfection and exposed to UV light (wavelength, 253 nm) for 20 min. In the UV-irradiated PCV2-infected cells at 72 h postinfection, the level of JNK or p38 phosphorylation fell to its basal level as seen in the mock-infected cells (Fig. 2). These results demonstrated that PCV2 replication was required for JNK and p38 MAPK phosphorylation and eliminated the possibility that attachment of PCV2 to cell surface receptors alone was responsible for JNK and p38 MAPK phosphorylation in the infected cells.

**Phosphorylated JNK/SAPK and p38 MAPK in infected cells activate downstream targets.** Activated JNK and p38 perform Ser/Thr phosphorylation of c-Jun inducing transcription factor and ATF-2, respectively. To confirm that phosphorylation of JNK and p38 MAPK after PCV2 infection truly activated their downstream substrates c-Jun and ATF-2, respectively, the phosphorylation of these two downstream substrates was measured in the infected cells by Western blotting. As shown in Fig. 3A, no activation of c-Jun and ATF-2 was detected in the mock-infected PK15 cells. Consistent with the activation of JNK and p38, a concomitant increase in c-Jun and ATF-2 phosphorylation was observed, which peaked at 72 h postinfection and then declined (Fig. 3A). In contrast, the protein levels of total amounts of c-Jun and ATF-2 remained unchanged in the PCV2-infected cells at various time points after infection compared to that in the mock-infected cells. Cells exposed to the UV-irradiated virus sample also failed to phosphorylate c-Jun or ATF-2 (Fig. 3A). The level of β-actin, used as a loading control, remained constant.

We then determined the effects of JNK inhibitor (the JNK inhibitor 1) or p38 MAPK inhibitor (SB202190) on the activation of JNK or p38 MAPK downstream substrates. As determined by testing for cell viability by trypan blue exclusion staining, the concentrations (5 to 20 μM for the JNK inhibitor 1 and SB202190) had no toxic effects on the treated cells (data not shown). In addition, the selected concentrations of these two inhibitors were tested to show their effectiveness in inhibiting virus-induced JNK or p38 MAPK activity by the FACE assay, which demonstrated that activation of JNK and p38 MAPK decreased dose dependently after treatment with their corresponding inhibitors (data not shown).

We treated the infected cells with the JNK inhibitor 1 or SB202190 at a concentration of 20 μM and determined the
levels of phosphorylated c-Jun and ATF-2 by Western blotting analysis. As shown in Fig. 3B, the phosphorylation of ATF-2, which is phosphorylated by the p38 MAPK pathway, was blocked by the p38 MAPK-specific inhibitor SB202190, while the activation of ATF-2 was substantially increased after PCV2 infection. In addition, the phosphorylation of c-Jun, which is phosphorylated by the JNK pathway, was also inhibited by treatment with SB202190 (Fig. 3B). While we examined the phosphorylation of c-Jun and ATF-2 after treatment with the JNK inhibitor 1, we also found that the inhibitor was able to reduce PCV2-induced c-Jun and ATF-2 phosphorylation (Fig. 3B).

Inhibition of ATF-2 activation by treatment with JNK inhibitor 1 and p38 inhibitor SB202190 was due to both JNK and p38 phosphorylated ATF-2 in the PCV2-infected cells. In addition, we also observed that p38 inhibitor SB202190 reduced c-Jun activation, which is only phosphorylated by JNK. This suggests that an indirect inhibition of c-Jun activation might be related to reduction of PCV2 replication when the PCV2-infected cells were treated with p38 inhibition. Therefore, these data suggested that phosphorylation of these two MAPKs as a consequence of PCV2 infection led to the activation of downstream targets.

**JNK/SAPK and p38 MAPK regulate PCV2 replication.** To determine whether activated JNK/SAPK and p38 MAPK play any role in the replication of PCV2, we examined the effect of these kinases on progeny virus production in the PCV2-infected PK15 cells by treatment with these kinase inhibitors. We infected PK15 cells with PCV2 in the presence of JNK inhibitor 1 (5 to 20 μM) or the p38 inhibitor SB202190 (5 to 20 μM) and determined the virus titers in the cell culture supernatant at 72 h postinfection by using an ICS assay. PCV2 replication in PK15 cells was not significantly altered by the presence of JNK or p38 inhibitors at 5 μM (Fig. 4). At 10 and 20 μM, the JNK inhibitor reduced PCV2 growth in the cultured cells by 54 and 83%, respectively. Treatment with the p38 inhibitor SB202190 at 10 and 20 μM reduced PCV2 growth by 59 and 87%, respectively. As expected, the UV-irradiated PCV2 failed to grow compared to the mock-infected cells (Fig. 4). These results suggested that the activation of MAPK pathways involving JNK and p38 regulate the efficient replication of PCV2 in the PK15 cells.

**Effect of JNK/SAPK and p38 MAPK inhibition on PCV2 viral transcription and protein synthesis.** Because the JNK inhibitor and SB202190 inhibited PCV2 replication in PK15
we determined which phase of the replication cycle, viral transcription or protein synthesis, is affected. The PCV2 viral transcription was examined by RT-PCR analysis with RNA extracted from the PCV2-infected PK15 cells 24 h after treatment with the JNK inhibitor 1 or p38 inhibitor SB202190 at various concentrations. The abundance of the target fragment corresponding to the spliced cap mRNA was dose dependently decreased in the PCV2-infected cells with these two inhibitors tested (Fig. 5A). In contrast, the fragment was detected in the DMSO-treated infected cells as seen in the DMSO-untreated infected cells (Fig. 5A and data not shown). No product was amplified from RNA extracted from cells infected with the UV-irradiated PCV2, as well as from the mock-infected PK15 cells (Fig. 5A).

To help delineate the effect of JNK and p38 inhibition on virus translation, the expression of PCV2 capsid protein ORF2 in the PK15 cells infected with PCV2 in the presence or absence of the inhibitors was analyzed by ICS. The ORF2 protein expressions were significantly reduced when cells were treated with these two inhibitors, as demonstrated by the increased number of PCV2-positive cells observed in the DMSO-treated infected cells (Fig. 5B). No significant differences were seen in the ORF2 protein expression between DMSO-treated infected cells and untreated infected cells (data not shown). No ORF2 protein expression was detected in the UV-irradiated PCV2-infected cells, as well as in the mock-infected cells (Fig. 5B). These results suggested that inhibitors of these two MAPK pathways decreased PCV2 viral RNA transcription and protein synthesis.

Inhibition of JNK/SAPK and p38 MAPK activation prevents PCV2-mediated caspase activation. PCV2 has been shown to induce apoptosis in cultured cells through the activation of caspase-8, followed by activation of the caspase-3 pathway (31). As demonstrated above, PCV2 replication was strongly prevented after treatment with these kinase inhibitors. Therefore, to determine whether JNK1/2 and p38 MAPK activation is required for PCV2-induced apoptosis, caspase-3 activity was assayed in the PCV2-infected PK15 cells at 72 h postinfection after treatment with the indicated amounts of these inhibitors. As expected, PCV2 alone induced activation of caspase-3 in the infected cells, whereas its activity was significantly reduced when the infected cells were treated with Ac-DEVD-CHO, a peptide inhibitor of caspase-3 activity (Fig. 6). A basal caspase-3 activity was detected in the UV-irradiated PCV2-

FIG. 5. Inhibition of JNK1/2 and p38 MAPK phosphorylation decreases viral transcription and protein synthesis in the PCV2-infected PK15 cells. (A) Effects of JNK and p38 inhibitors on PCV2 spliced cap mRNA synthesis in infected cells. Total RNAs (1 μg) isolated from PCV2-infected cells at 24 h after treatment with JNK inhibitor 1 or SB202190 at the indicated concentrations were subjected to RT-PCR analysis of PCV2 spliced cap mRNA synthesis. The products were separated on a 1% agarose gel. β-Actin was used as an internal control. The results are representative of at least three independent experiments. (B) JNK inhibitor 1 and SB202190 reduce PCV2 protein expression. PCV2-infected PK15 cells at 72 h in the presence of the JNK inhibitor 1 (20 μM) or SB202190 (20 μM), as well as UV-irradiated PCV2-infected cells at 72 h, were assayed for the amount of PCV2 viral antigen by ICS. The amounts of PCV2 ORF2 protein expression are shown as percentages of PCV2-positive signals in cells infected with PCV2 alone. The data are mean values ± the SD based on three independent experiments; error bars show the SD.
infected cells, as well as in the mock-infected cells (Fig. 6). Furthermore, activation of caspase-3 was dose dependently decreased in the infected cells when treated with increasing amounts of the JNK or p38 inhibitor (Fig. 6). Therefore, we believe that the activation of MAPK pathways involving JNK and p38 is required for PCV2-induced apoptosis, but this can be an indirect effect due to a decrease in virus replication when PCV2-infected cells were treated with JNK or p38 inhibition (Fig. 4).

**DISCUSSION**

Viral infections are known to activate various cellular signaling pathways, which can affect cellular function and virus replication. In this report, we show that JNK1/2 and p38 MAPK pathways were activated during the course of PCV2 infection in PK15 cells and that activation of these two MAPK pathways was required for active replication of PCV2. We further show that inhibitors of these two MAPK pathways lead to a reduction of viral activity, as determined by decreases in viral transcription, virus protein synthesis and progeny production, and decreased cell death. This study has demonstrated for the first time that the MAPK signaling pathways involving JNK1/2 and p38 are manipulated by PCV2 and that these two pathways play beneficial roles in the PCV2 replication.

The effect of JNK and p38 MAPK on virus replication has been studied in a number of viruses. We determined whether PCV2-induced JNK and p38 MAPK are involved in viral replication by examining the effects of inhibitors of the MAPK pathways on PCV2 production and found that JNK inhibitor 1 and SB202190 significantly reduced the production of progeny virus (Fig. 4). This shows that optimal PCV2 replication requires these two MAPK activities and raises the possibility that PCV2 has acquired the ability to activate these two MAPK kinases to aid its replication. MAPK activation has been found to be required for the optimal replication of many other viruses. For instance, rotavirus replication was significantly reduced after JNK and p38 inhibition (22). JNK and p38 activation by HSV requires the expression of particular viral proteins, and blockage of JNK translocation to the nucleus resulted in decreased virus production (10, 20, 36). Coxsackievirus B3 activates JNK and p38 MAPK, which appears to facilitate virus replication and virion release (46). Treatment of HSV-1-infected cells with p38 inhibitor resulted in a decrease in virus titer (60). Human immunodeficiency virus type 1 production was increased in p38 MAPK-activated cells (45). Like the above-mentioned viruses, PCV2-induced activation of JNK and p38 MAPK are truly involved in the replication of PCV2 in PK15 cells.

MAPK regulation of viral replication can act at specific steps of the replication cycle, including entry, gene transcription, protein expression, and assembly. In some cases, inhibition of MAPK activation did not affect viral transcription and/or protein synthesis. For instance, inhibition of p38 MAPK by treatment of HSV-1-infected cells with inhibitor SB203580 did not affect the viral transcription program (25). Coxsackievirus B3 viral protein synthesis was reported to be unaffected by treatment with JNK or p38 inhibitors (46). On the other hand, inhibition of p38 MAPK reduced EMCV replication via inhibition of EMC viral protein synthesis, as well as a secondary effect on viral RNA synthesis (21). Reduced murine coronavirus production in p38 inhibitor SB203580-treated cells was in part due to a decrease in virus-specific protein synthesis and virus-specific mRNA accumulation (3). Efficient translation in HSV-1-infected cells is regulated by p38-dependent eukaryotic translation initiation factor 4E phosphorylation (54). PCV rep-
licates via rolling-circle replication involving an intermediate double-stranded replicative form of DNA (6). In the present study, we further determine which step of PCV2 replication is affected by inhibition of JNK or p38 MAPK activation. Our data showed that treatment of cells with JNK inhibitor 1 or p38 inhibitor SB202190 suppressed PCV2 viral RNA transcription and protein synthesis, which is due to that viral transcription and protein synthesis are closely related in the replication of PCV2. Thus, the results suggest that JNK or p38 MAPK inhibition reduced the accumulation of both viral mRNA and structural protein, followed by causing a lowered production of infectious PCV2 particles in cells. However, our study does not exclude the possibility that these specific inhibitors have a preferential effect on PCV2 viral protein synthesis and/or RNA transcription.

Apoptosis has been shown to be regulated by the JNK and p38 MAPK cellular stress pathways (28, 48, 53). Virus infection can trigger an apoptogenic kinase cascade, leading to the phosphorylation and/or activation of JNK and p38 MAPK, which are responsible for inducing apoptotic cell death (47). Hepatitis B virus X protein-mediated apoptosis was reported to involve sustained activation of JNK and p38 pathways (55). Severe acute respiratory syndrome coronavirus 3a protein-induced apoptosis was dependent on the activation of p38 MAPK (40). In the present study, using the specific inhibitor of JNK or p38 MAPK activation, we demonstrated that interference with the JNK and p38 MAPK pathways leads to reduction of PCV2-induced apoptotic caspase activity (Fig. 6). This result suggests that JNK and p38 MAPK signaling pathways play a role in the mechanism by which PCV2 triggers an apoptotic program in infected cells. In nonstressed cells, nonphosphorylated JNK is complexed to p53, causing its ubiquitination, followed by proteasomal degradation (14, 15). Upon stimulation, the phosphorylated JNK mediates dissociation of p53, followed by its stabilization (14, 15). The role of p38 MAPK kinase in transcription of the p53-regulated Bax and Noxa genes also involves p53 phosphorylation (4, 23). Binding of PCV2 ORF3 protein to porcine p53-induced RING-H2 was reported to interfere with its function of ubiquitination of p53, followed by causing increased p53 level and thus suggesting that the PCV2-induced apoptosis in the cultured cells may be p53 mediated (32). Therefore, activation of JNK and p38 after PCV2 infection may participate in p53 stabilization and phosphorylation, ultimately leading to apoptosis.

In conclusion, the results reported here establish that PCV2 infection induces the activation of JNK and p38 and their involvement in c-Jun and ATF-2 activation in the cultured cells and demonstrate that the activation of JNK or p38 is required for efficient PCV2 replication, as well as PCV2-induced apoptotic cell death. The role of JNK and p38 activation in PCV2 infection and replication will contribute important information about the molecular mechanism of PCV2 pathogenesis.

REFERENCES


