# **PORCINE CIRCOVIRUS 3**



The mission of the Swine Health Information Center is to protect and enhance the health of the United States swine herd through coordinated global disease monitoring, targeted research investments that minimize the impact of future disease threats, and analysis of swine health data.

September 2016 | Updated July 2021

# **SUMMARY**

# **IMPORTANCE**

- Porcine circovirus 3 (PCV3) is a newly detected pathogen of swine. Although most cases are sublinical, PCV3 is also associated with clinical signs similar to those caused by porcine circovirus 2 (PCV2).
- Individual case definitions have been proposed for PCV3-associated reproductive disease and systemic disease to standardize diagnostic criteria (see *Infection in Swine*).

# **PUBLIC HEALTH**

Porcine circoviruses (PCVs), including PCV3, are not considered to be zoonotic.

# **INFECTION IN SWINE**

- PCV 3 has been associated with reproductive failure, porcine dermatitis and nephropathy syndrome (PDNS), and multi-systemic inflammation. However, only a few studies have demonstrated the presence of PCV3 in lesions. Anecdotally, PCV3 has been detected in neurological, respiratory, and enteric cases, but causation has not been established.
- In pathogenicity studies, PCV3 inoculation does not consistently lead to development of clinical disease.
- Individual diagnostic criteria proposed for PCV-3-reproductive disease (PCV-3-RD) include late reproductive problems and higher perinatal mortality, multi-systemic lymphoplasmacytic to lymphohistiocytic perivascular inflammation, and moderate to high amount of PCV-3 genome in damaged tissues.
- Individual diagnostic criteria proposed for PCV-3-systemic disease (PCV-3-SD) include weight loss, rough hair, neurological signs, multi-systemic lymphoplasmacytic to lymphohistiocytic perivascular inflammation, and moderate to high amount of PCV-3 genome in damaged tissues.

## TREATMENT

• There is no treatment for pigs infected with PCV3.

## **CLEANING AND DISINFECTION**

- Circoviruses are stable in the environment. In the laboratory, potassium perxoymonosulfate, sodium chloride, sodium hypochlorite, and sodium hydroxide are the most effective for PCV2.
- Specific information on the disinfection of PCV3 is not available.

# **PREVENTION AND CONTROL**

- Prevention for PCV2 is based on factors that can influence susceptibility. These include good nutrition, biosecurity, and vaccination.
- No specific control measures have been described for PCV3.

## TRANSMISSION

Both PCV2 and PCV3 spread through vertical and horizontal transmission. Virus is shed in most
secretions and excretions. Direct contact is the route of most significance but spread can also occur via
fomites and ingestion.

#### PATHOGENESIS

• Little is known about PCV3 pathogenesis. PCV3 utilizes a clathrin- and dynamin-2-mediated endocytic pathway. PCVs generally replicate in actively dividing cells.

## DIAGNOSIS

- PCV3 was first identified by metagenomic sequencing. Many polymerase chain reaction (PCR) assays have been developed for PCV3 in the research setting. Most are based on the ORF 2 gene (Cap protein). Virus can be demonstrated in lesions by immunohistochemistry (IHC) or in situ hybridization (ISH). PCV3 has only recently been isolated in cell culture (PK-15 cells).
- Although antibodies are not diagnostic for PCV3, a few enzyme-linked immunosorbent assays (ELISAs) have been described to detect recombinant Cap protein.
- PCV3 can be detected in many tissues and organs. Heart, lung, and lymphoid tissue are important for histology and IHC or ISH. When PDNS is suspected, skin lesions should be submitted, and in cases of reproductive failure, fetal tissues should be submitted. Oral fluids can be used with some PCR assays.
- Systemic vasculitis is an important feature associated with PCV3 cases.

# **EPIDEMIOLOGY**

- PCV3 is found in wild and domestic pigs. Antibodies to PCV3 have been detected in dogs, cattle, and mice. PCV3-positive mosquitoes have been found. Wild ruminants and ticks may also be reservoirs.
- PCV3 is found in many swine-producing regions of the world. Epidemiological studies have found that
  infection is widespread, with prevalence up to 100%. Little is known about the course of natural infection.
  A few studies have shown that prevalence is highest in piglets/weaners and decreases with age, but PCV3
  has been detected in pigs up to 23 weeks-of-age.

# **ETIOLOGY**

- PCV3 belongs to the family *Circoviridae*. Members are very small, non-enveloped viruses with a circular, covalently closed, single-stranded DNA genome.
- Currently, there are four recognized circoviruses of swine, designated PCV1–4. PCV3 is distinctly
  different from other PCVs and is closely related to canine and bat circoviruses.

## **HISTORY IN SWINE**

PCV1 was identified in 1974 as a contaminant in a pig kidney cell line. In 1997, PCV2 was recognized as
the cause of a novel wasting disease affecting Canadian pigs. In 2016, PCV3 was detected in tissues from
sows from North Carolina that died acutely with PDNS-like clinical signs and aborted fetuses. PCV4 was
discovered in samples from pigs with respiratory disease, enteric disease, and PDNS-like lesions in China
in 2019. All known PCVs have been circulating in swine long before they were first detected.

## IMMUNITY

- The PCV3 antibody response is mainly due to IgG, which can be detected experimentally at seven days post-infection (dpi). The duration of the humoral response during natural infection is unknown. However, experimental studies have demonstrated that specific PCV3 IgG antibodies remain detectable for up to 42 dpi. The protective role of the humoral response is unknown.
- While there are no commercially available vaccines for PCV3, herd-specific vaccines may be available through veterinary prescription. No cross-protection occurs between PCV2 and PCV3.

## **GAPS IN PREPAREDNESS**

- PCV3 has been associated with many PCVAD-like clinical signs. The strongest evidence of causality lies with PCV3-systemic and PCV3-reproductive disease.
- Like PCV2, many PCV3 infections are subclinical, and infection with other swine pathogens is common.
- In order to better understand the impact of PCV3 on the swine industry, more evidence is needed on pathogenesis, host genomics, viral genomics, and immunology.

# LITERATURE REVIEW: PORCINE CIRCOVIRUS 3

# IMPORTANCE

Porcine circovirus 3 (PCV3) is a newly detected pathogen of swine associated with clinical signs similar to those caused by porcine circovirus 2 (PCV2). However, many infections are subclinical, and pathogenicity studies have had mixed results. Individual case definitions for PCV3-associated reproductive disease and systemic disease have been proposed to standardize diagnostic criteria (see *Infection in Swine*).

# **PUBLIC HEALTH**

PCVs are not considered to be zoonotic. Anti-PCV antibodies have been detected in humans, mice, and cattle; however, this is most likely due to the presence of a similar virus not PCV.<sup>2</sup> Although PCV nucleic acids have been detected in vaccines produced for humans and pigs, this is thought to be due to poor quality control.<sup>3</sup>

# **INFECTION IN SWINE**

#### PCV 1 and PCV2

Porcine circovirus 1 (PCV1) is nonpathogenic in swine.<sup>3</sup> In the 1990s, PCV2 was associated with post-weaning multi-systemic wasting syndrome (PMWS) in pigs 2–4 months-of-age. Clinical signs of PMWS include enlarged subcutaneous lymph nodes, wasting, diarrhea, respiratory distress, pallor, and occasionally icterus.<sup>3</sup> Lymphoid depletion and lymphohistiocytic or granulomatous inflammation is a component of diagnosis.<sup>3</sup> Although PCV2 is a primary pathogen, most infections are subclinical.<sup>3,4</sup> Co-infection with other swine pathogens is common.<sup>5</sup> Collectively, PCV2 infections are now known as porcine circovirus-associated disease (PCVAD).<sup>4</sup> Clinical manifestations include systemic disease (PCV2-SD, formerly PMWS), porcine dermatitis and nephropathy syndrome (PDNS), lung disease (PCV2-LD), reproductive disease (PCV2-RD), enteric disease (PCV2-ED), and subclinical infection (PCV2-SI).<sup>4</sup>

## PCV3

PCV3 was first detected in sows with PDNS-like lesions (red-to-purple or dark crusted skin lesions, often on the hind limbs and perineum, along with enlarged and pale kidneys with petechiae) and aborted mummified fetuses.<sup>3</sup> The affected farm had sow mortality and chronic reproductive problems, including low-average conception rates.<sup>6</sup> Histological lesions included bronchointerstitial pneumonia, peribronchiolar and perivascular cuffing (lymphocytes and plasma cells), and intraluminal edema in the alveoli. In the skin, necrotizing vasculitis was seen, with fibroid changes and neutrophilic infiltration, hemorrhage, and fibrin exudation.<sup>6</sup> The lymph nodes showed diffuse granulomatous lymphadenitis (histiocytes and multinucleated giant cells) with moderate lymphoid depletion.<sup>6</sup> Diffuse membrane proliferative glomerulonephritis was seen in the kidneys.<sup>6</sup> Ten sows from the outbreak were also torque teno virus (TTV)-positive.<sup>6</sup> The role of TTV is unclear, but it has previously been linked to PCV2 infection.<sup>7</sup> Investigators also screened archived samples from PDNS cases that were PCV-2-negative by IHC; many were positive for PCV3 by quantitative polymerase chain reaction (qPCR). A subset also tested positive via IHC.<sup>6</sup>

Myocarditis and multi-systemic inflammation have also been linked to PCV3 in post-weaning pigs.<sup>8</sup> In a 2–3 week-old pig with weight loss and swollen joints, lesions included fibrinous arthritis/synovitis with necrotizing arteriolitis of the synovial capsule, necrotizing arteriolitis in the esophagus, diffuse lymphohistiocytic interstitial pneumonia, and multifocal lymphoplasmacytic and histiocytic myocarditis and arteriolitis.<sup>8</sup> A second pig, 9–10-

weeks-old from another state, had a history of respiratory disease and rectal prolapse. Several swine pathogens were identified; unexplained perivascular lymphocytic encephalitis and meningitis in the cerebellum was later linked to PCV3.<sup>8</sup> The third pig, 19-days old and from another state, presented with severe dyspnea and neurologic disease. Similar lesions were seen as those previously described. In this study, all three cases were co-infected with porcine astrovirus, which may or may not have contributed to pathogenesis.<sup>8</sup> Individuals were also infected with rotavirus A, porcine cytomegalovirus, and porcine hemagglutinating encephalomyelitis<sup>8</sup>

Following these early reports, PCV3 nucleic acids have been detected in pigs with neurological disease,<sup>8-10</sup> reproductive failure,<sup>10-20</sup> respiratory disease,<sup>6, 8, 17, 21-25</sup> enteric disease,<sup>23-26</sup> and PDNS.<sup>6, 10, 12</sup> PCV3 has also been detected in healthy pigs.<sup>24, 27-34</sup> Relatively few studies have demonstrated the presence of PCV3 within lesions.<sup>8, 10, 17, 20, 35, 36</sup> Based on the lesions most commonly associated with PCV3, heart, lung, and lymphoid tissue appear to be most important for diagnosis.<sup>36</sup>

Published studies on experimental PCV3 infection have had mixed results.

- In 4- and 8-week-old SPF pigs, intranasal inoculation with an infectious PCV3 clone led to fever, anorexia, diarrhea, and respiratory distress, skin lesions (described as PDNS-like) and sudden death. PCV3 was demonstrated in lungs, heart, lymph nodes, liver, kidneys, and small intestines using IHC.<sup>37</sup> However, results of this study have been questioned by other researchers due to the atypical appearance of the skin lesions and time frame for reproduction of the glomerular lesions, as well as inability to reproduce the infectious PCV3 clone used to inoculate pigs.
- In 6-week-old CD/CD pigs, PCV3 intranasal and intramuscular inoculation (plus subcutaneous administration of an immunostimulant two days before and after inoculation) did not lead to development of clinical disease.<sup>38</sup> However, multi-systemic inflammation and perivasculitis were observed via histology, and PCV3 nucleic acid was detected in lesions by ISH.<sup>38</sup>
- In 5-week-old CD/CD pigs, PCV3-positive tissue homogenate was used for intramuscular and intranasal inoculation. Pigs were re-inoculated after seven days. No clinical signs were seen during the study. Viremia occurred in PCV3-inoculated pigs at three days post-infection (dpi) and continued until the end of the study. Histology demonstrated multi-systemic inflammation and perivasculitis. PCV3 was confirmed in tissues via qPCR and ISH.<sup>39</sup>
- In 4-week-old pigs, PCV3 intranasal inoculation led to moderate clinical signs, including anorexia, emaciation, and coughing at 12 dpi. At >12 dpi, shivering and tachypnea were seen, plus development of multifocal papules on the skin. Histology revealed mucosal epithelial cell necrosis and lymphocyte necrosis in tissue from the small intestine. Presence of PCV3 antigen was confirmed by IHC.<sup>40</sup>
- In 3-week-old CD/CD pigs, PCV3 intranasal and intramuscular inoculation did not lead to development
  of clinical disease. Despite this, PCV3 nucleic acids were detected in many organs/tissues via qPCR,
  with the highest amounts in lung and inguinal lymph node. PCV3 replication was detected in these tissues
  by ISH. Histology revealed lymphocyte reduction and inflammatory cell infiltration in the lymph nodes
  and epithelial cell proliferation, inflammatory cells infiltration, and thickened alveolar septum in the
  lungs. Passage in PK-15 cells failed.<sup>41</sup>

To standardize diagnostic criteria and improve case finding, Saporiti and colleagues<sup>1</sup> have proposed case definitions for reproductive and systemic disease caused by PCV3 (see *Table 1*). There is no proposed case definition for PCV3-PDNS as of yet since the authors do not believe an etiological association has been fully established. For PCV2-PDNS, detection of PCV2 is not a diagnostic requirement, with criteria relying on clinical signs and histology, including:

- Presence of hemorrhagic and necrotizing skin lesions, primarily on the hind limbs and perineal area, and/or swollen and pale kidneys with generalized cortical petechiae
- Presence of systemic necrotizing vasculitis and necrotizing fibrinous glomerulonephritis

Table 1. Proposed Diagnostic Criteria for PCV3-Associated Diseases*		
Proposed Name	Main Clinical Signs	Individual Diagnostic Criteria
PCV-3-reproductive disease (PCV-3-RD)	Late abortion, malformations, mummified fetuses, stillborn fetuses, weak-born piglets	<ol> <li>Late reproductive problems and higher perinatal mortality</li> <li>Multi-systemic lymphoplasmacytic to lymphohistiocytic perivascular inflammation</li> <li>Moderate to high amount of PCV-3 genome in damaged tissues</li> </ol>
PCV-3-systemic disease (PCV-3-SD)	Wasting, weight loss, ill thrift or poor-doers, neurological signs	<ol> <li>Weight loss, rough hair, neurological signs</li> <li>Multi-systemic lymphoplasmacytic to lymphohistiocytic perivascular inflammation</li> <li>Moderate to high amount of PCV-3 genome in damaged tissues</li> </ol>

\*Diagnostic criteria proposed by Saporiti et al. Porcine circovirus 3 (PCV-3) as a causal agent of disease in swine and a proposal of PCV-3 associated disease case definition. *Transbound Emerge Dis.* 2021. doi: 10.1111/tbed.14204

#### PCV4

PCV4 has been recently described, and more information is needed to definitively know whether it causes clinical illness in pigs (see *History in Swine*).

#### TREATMENT

There is no treatment for pigs infected with PCVs. Antibiotics may be used to treat secondary bacterial infections.

## **CLEANING AND DISINFECTION**

#### SURVIVAL

Circoviruses are very stable in the environment.<sup>42</sup> PCV1 survives at 70°C (158°F) for 15 minutes, and PCV2 survives at 75°C (167°F) for 15 minutes or 56°C (133°F) for one hour.<sup>3</sup> PCV2 has been found in different water sources, including those treated by chlorination for human or swine consumption.<sup>43</sup>

In fresh pork, PCV2b can survive for two dpi at room temperature, six dpi at refrigeration temperature  $(4^{\circ}C/39.2^{\circ}F)$ , and 30 dpi at freezer temperature  $(-20^{\circ}C/-4^{\circ}F)$ .<sup>44</sup> PCVs are also resistant to extreme pH. PCV2 retains some infectivity at pH of 2 and pH of 11–12.<sup>3</sup> No information was found on the survival of PCV3.

#### DISINFECTION

In the laboratory, potassium peroxymonosulfate and sodium chloride, sodium hypochlorite (bleach), and sodium hydroxide appear to be the most effective virucidal agents for PCV2. Other potentially effective products include quaternary ammonium compounds and phenolics Chlorhexidine, ethanol, aldehydes, and iodine products are generally not effective disinfectants for PCVs.<sup>45-47</sup> No information was found on the disinfection of PCV3.

## **PREVENTION AND CONTROL**

#### **DISEASE REPORTING**

PCVs are not OIE-listed. There are no restrictions for the importation of animals from countries or zones affected by PCVs. Any suspicious clinical or necropsy findings should always be reported to the USDA and your State Animal Health Official.

#### **DISEASE PREVENTION**

PCV2 prevention is based on factors that can influence susceptibility. These include good nutrition, biosecurity, and vaccination (see *Immunity*). Since females have a lower risk of developing PMWS, sorting nursery pigs by sex can be beneficial.<sup>3</sup> Increased risk in males may be due to infection following castration and genetic or hormonal influence.<sup>48</sup> Poor management, including overcrowding, lack of ventilation, and frequent commingling, may contribute to disease severity. Management factors linked to lower disease risk include:

- Housing pregnant sows in groups
- Weaning at higher weights
- Vaccinating sows for atrophic rhinitis
- Treating for ectoparasites
- Adding spray-dried plasma to nursery rations<sup>3</sup>

No specific preventive measures have been described for PCV3. Vaccines are discussed under Immunity.

#### **DISEASE CONTROL**

There are no specific control measures for PCV3. However, standard biosecurity practices should be in place on all swine premises.

#### TRANSMISSION

As described by Opriessnig et al.,<sup>5</sup> both PCV2 and PCV3 are spread by vertical and horizontal transmission. PCVs are transmitted mainly through direct contact. Virus can be detected in nasal, ocular, tonsillar, and bronchial secretions, as well as saliva, urine, feces, milk, colostrum, and semen.<sup>3, 22</sup> Fomites, contaminated feed, and biologics, hypodermic needles, and biting insects may play a role in transmission. Pigs might become infected by eating the raw tissues of viremic animals.<sup>3</sup> For PCV2, fetal infection can occur if the dam is exposed during pregnancy or inseminated with virus-containing semen. The virus can also be spread fetus-to-fetus, and the timing of in utero infection determines the clinical outcome.<sup>49</sup>

## PATHOGENESIS

Circoviruses replicate in actively dividing cells of young animals. PCV infection is enhanced when the immune system is stimulated, and more lymphocytes are available for replication.<sup>50</sup> Likewise, virus replication can occur in other cells with a high mitotic index, such as endothelial cells, epithelial cells, and macrophages.<sup>3</sup> One recent study found that PCV3 utilizes a clathrin- and dynamin-2-mediated endocytic pathway, entering both early and late endosomes, the latter of which requires an acidic environment.<sup>51</sup>

In 6-week-old CD/CD experimentally inoculated pigs, ISH confirmed PCV3 replication in the myocardiocytes, tunica media, and endothelial cells of the arteries in addition to the tubular renal epithelium and tunica media arteries in the kidney.<sup>38</sup> Moreover, PCV3 replication in 5-week-old CD/CD pigs was observed in the interstitial myocarditis, inflammatory infiltrate of the cortical renal insterstitium, white pulp of the spleen, endothelial cells of the hepatic sinusoids, Peyer's patches of the small intestine, and inflammatory foci of the intestinal serosa.<sup>39</sup>

Experimentally, PCV3 infection leads to disruption of gut microbiota.<sup>40</sup>

## DIAGNOSIS

PCVs are widely distributed in the global swine population. Diagnosis of PCVAD is based on a combination of clinical signs, characteristic gross and microscopic lesions, and detection of the virus in lesions.<sup>3</sup>

## TESTS TO DETECT NUCLEIC ACIDS, VIRUS, OR ANTIGENS

The ORF 2 gene, encoding for the highly conserved Cap protein, is often the basis for PCR. PCR assays that have been described for PCV3 include:

- qPCR targeting the *cap* gene<sup>11, 52</sup>
- TaqMan qPCR detecting the *cap* gene<sup>6,9,10,13,15,53,54</sup>
- SYBR green-based qPCR detecting the *rep* gene<sup>55</sup> and *cap* gene<sup>15</sup>
- Direct (no DNA required) and qPCR targeting the rep gene<sup>56, 57</sup>
- Loop-mediated isothermal amplification (LAMP) assay detecting the cap gene<sup>58, 59</sup>
- Real-time recombinase polymerase amplification (rt-RPA) assay detecting the *cap* gene <sup>60</sup>
- Colorimetric isothermal multiple-self-matching-initiated amplification (IMSA) detecting the cap gene<sup>61</sup>
- Polymerase spiral reaction (PSR) assay detecting the *cap* gene<sup>62</sup>
- Multiplex qPCR to detect and differentiate PCV3 and PCV2<sup>63</sup>
- Multiplex PCR to detect and differentiate PCV1, PCV2, and PCV3<sup>64</sup>
- Duplex qPCR for simultaneous detection of PCV2 and PCV3<sup>65</sup>
- Multiplex qPCR to detect and differentiate PCV3, PCV2a, PCV2b, and PCV2d<sup>66</sup>
- Duplex qPCR to differentiate PCV2 and PCV3<sup>67, 68</sup>
- Duplex qPCR to detect PCV2 and PCV3<sup>39</sup>
- Duplex qPCR to detect PCV3 and PCV4<sup>69</sup>
- Multiplex qPCR to detect and differentiate PCV1, PCV2, PCV3, and PCV4<sup>70</sup>
- Duplex qPCR assay to simultaneously detect PCV3 and porcine epidemic diarrhea virus<sup>71</sup>
- Duplex qPCR to simultaneously detect PCV3 and pseudorabies virus<sup>72</sup>
- Duplex qPCR to simultaneously detect PCV3 and classical swine fever virus<sup>73</sup>

In addition to PCR, demonstration of virus within lesions is important for PCV diagnosis. Methods include IHC and ISH.<sup>74</sup> PCV3 was first identified by metagenomic sequencing<sup>6, 8</sup> and confirmed in tissues from sows with PDNS-like lesions (by PCR and IHC)<sup>6</sup> and post-weaning pigs with weight loss and swollen joints (by ISH).<sup>8</sup> Next-generation sequencing has been described in combination with ISH to detect challenging or emerging pathogens like PCV3.<sup>75</sup>

Porcine kidney cells (PK-15),<sup>76</sup> Vero cells, and other porcine-derived cell lines can be used for isolation of PCV2.<sup>3</sup> Until recently, PCV3 had not been isolated in cell culture. One successful study found that PCV3 did not induce a cytopathic effect in primary porcine kidney cells, but infection was confirmed with ISH.<sup>77</sup> PCV3 has also been isolated from diagnostic case samples (from weak-born, stillborn, or mummified pigs)<sup>10</sup> in PK-15 cells and confirmed by qPCR and next-generation sequencing.<sup>38</sup>

#### **TESTS TO DETECT ANTIBODY**

Antibodies are not diagnostic for PCV infection, but they may be useful for confirmation at the herd level. Tests described for PCV3 include several enzyme-linked immunosorbent assays (ELISAs) to detect recombinant Cap protein.<sup>6, 38, 78-80</sup> One assay utilized both a *cap*-based peptide and a *rep*-based peptide.<sup>81</sup>

#### SAMPLES

PCV3 can be detected in many sample types, including oral fluids.<sup>82</sup> Heart, lung, and lymphoid tissue should always be submitted for histology.<sup>36</sup> In cases of reproductive failure, fetal tissues should be submitted, including the myocardium. For suspected cases of PDNS, diagnostic submissions should include skin lesions.<sup>3</sup>

# **EPIDEMIOLOGY**

#### **SPECIES AFFECTED**

As described by Opriessnig et al.,<sup>5</sup> circoviruses are found in mammals, fish, birds, and insects. PCVs are found in both wild<sup>83-90</sup> and domestic pigs, but clinical disease seems to occur only in the latter (see *Infection in Swine*). PCV2 has been detected in farmed minks with diarrhea,<sup>91</sup> buffaloes,<sup>92</sup> beef products,<sup>93</sup> dogs,<sup>94</sup> farmed shellfish,<sup>95</sup> and flies.<sup>96</sup> While PCV2 antibodies have been found in rodents,<sup>2</sup> few studies have confirmed PCV in field samples.<sup>97</sup> Species susceptible to PCV3 include dogs,<sup>98, 99</sup> cattle,<sup>100</sup> mice,<sup>101</sup> and mosquitoes.<sup>102</sup> One study found evidence of PCV3 infection in chamois, roe deer, and ticks (*Ixodes ricinus*).<sup>83</sup> However, others have found that wildlife (including rodents and *Ixodes* spp. ticks,<sup>103</sup> and free-ranging ruminants and lagomorphs)<sup>104</sup> play little to no role in PCV3 epidemiology.

#### **GEOGRAPHIC DISTRIBUTION**

PCV1 may have been widespread at one time. PCV2 and PCV3 are currently found in many swine producing regions of the world, including North America,<sup>6,8</sup> South America,<sup>16,105-108</sup> Europe,<sup>12-14, 32, 33, 57, 109-112</sup> and Asia.<sup>11, 21, 22, 29, 52, 113-117</sup> To date, PCV4 has been described only in Asia.<sup>69, 118-121</sup>

#### **MORBIDITY AND MORTALITY**

The current prevalence of PCV1 is thought to be low, but this is uncertain since PCV1 primers are not included in assays that detect PCV2 and PCV3.<sup>5</sup> Both PCV2 and PCV3 are common in pigs. For PCV3 specifically, up to 100% seroprevalence occurs as described by Ouyang et al.<sup>82</sup> Prevalence of the recently emerged PCV4 is unclear. However, estimates from China show that 5–45% of swine samples are PCV4-positive.<sup>69, 119, 120</sup> A study from Guangxi Province found that nearly 70% of samples testing positive for PCV4 also contained PCV2 or PCV3.<sup>120</sup>

PCV3 seems to affect pigs of all ages.<sup>116</sup> A few studies have shown that prevalence is highest in piglets and/or weaners then decreases with age.<sup>10, 115, 122</sup> However, a longitudinal study found that PCV3 infection spanned from 4–23 weeks.<sup>123</sup> Evidence also shows that primiparous sows shed PCV3 at higher levels in colostrum<sup>30, 31</sup> and have more PCV3-infected fetuses compared to multiparous sows. Like PCV2, cases of PCV3 often involve co-infection with other swine pathogens.<sup>1, 82</sup>

Morbidity for PCVAD is variable but generally low to moderate. However, mortality can be quite high, especially for pigs with PDNS vs. pigs with wasting syndrome.<sup>3</sup> Since their introduction in 2006, commercial vaccines for sows, gilts, and piglets have significantly reduced PCV2-associated mortality infection in the United States. There is also some indication that disease susceptibility<sup>124</sup> and vaccination effectiveness is influenced by genetic differences (e.g., breed).

#### **ETIOLOGY**

#### **CHARACTERISTICS OF CIRCOVIRUSES**

PCVs are members of the family *Circoviridae*. Circoviruses are very small (15–25 nm), non-enveloped viruses that contain a circular, covalently closed, single-stranded DNA genome.<sup>42</sup> As of 2020, the family includes two genera, *Circovirus* and *Cyclovirus*, which contain 29 and 49 species, respectively.<sup>125</sup>

#### **CHARACTERISTICS OF PORCINE CIRCOVIRUSES**

Like other circoviruses, PCVs use an ambisense transcription strategy. They have at least two major open reading frames (ORFs). ORF1 encodes the replication-associated protein (Rep, encoded on the virion sense strand), and ORF2 encodes the capsid protein (Cap, encoded on the complementary sense strand).<sup>42</sup>

Currently, there are four recognized circoviruses of swine, designated PCV1–4.<sup>125</sup> PCV2 is highly diverse compared to PCV1, particularly within ORF2.<sup>126</sup> PCV2 is divided into five genotypes (PCV2a-2e).<sup>3</sup> PCV3 is distinct from PCV1 and PCV2, with only 31% and 48% homology, respectively.<sup>111</sup> However, a high degree of

homology occurs between PCV3 viruses. Several classification schemes have been suggested based on *cap* features, including two (a, b) or three (a, b, c) clades; and, clade a has been divided into either two (a1, a2) or three (a1, a2, a3) subclades.<sup>5</sup> Importantly, no differences in pathogenicity between PCV3 clades have been observed.

Phylogenetic analyses show that PCV3 is related to canine and bat circoviruses.<sup>6,8</sup> Additionally, the PCV3 *cap* gene has similarities to avian ones, suggesting a recombinant origin.<sup>127</sup> It is likely that new PCVs will continue to emerge. Additionally, classification reshuffling will likely continue, in part due to the increasing recognition of recombinant PCVs.<sup>105</sup>

# **HISTORY IN SWINE**

In 1974, a small, spherical virus-like contaminant was detected in a pig kidney cell line (PK-15).<sup>128</sup> The virus, later identified as PCV1, was found to be widespread in pigs but has never been associated with disease.<sup>129</sup> As described by Harding and Clark,<sup>130</sup> a novel wasting disease affecting Canadian pigs emerged in the mid-1990s, reaching epidemic proportions in countries across the world. In 1997, the causative agent was determined to be PCV2, which is antigenically and genetically distinct from PCV1.<sup>131</sup>

In 2016, PCV3 was detected in tissues from sows that died acutely with PDNS-like clinical signs and aborted fetuses using metagenomic sequencing.<sup>35</sup> Since that time PCV3 has been widely found in swine (see *Epidemiology*). PCV4 was discovered in samples from pigs with respiratory disease, enteric disease, and PDNS-like lesions in China in 2019.<sup>118</sup> Additional PCV4s have since been detected in Asia, with high homology compared to the original reported sequence.<sup>69, 119, 120, 132</sup> Current evidence suggests that all known PCVs were circulating in swine long before they were first detected.<sup>5, 82</sup>

# IMMUNITY

#### **POST-EXPOSURE**

Many swine herds have anti-PCV2 antibodies indicating previous exposure; however, these are not necessarily protective.<sup>3</sup> In piglets, seroconversion occurs as colostral antibody wanes around seven weeks-of-age.<sup>48</sup> PCV2 seroconversion occurs in both subclinical and clinical cases, although some studies have shown decreased humoral immunity, and specifically fewer neutralizing antibodies, in clinically affected pigs. This corresponds with a higher concentration of virus in the serum and increased viral shedding among this cohort. Cell-mediated immunity may also be a factor in viral clearance.<sup>3</sup>

A few experimental studies have examined the immune response to PCV3 infection.

- In 4–6-week-old CD/CD pigs, PCV3 antibody response (primarily IgM) was detected 7–10 days following intranasal and intramuscular inoculation.<sup>38</sup>
- In 5-week-old CD/CD pigs, an IgG response was dominant, appearing at 7 dpi and persisting at 42 dpi following intramuscular and intranasal inoculation with PCV3.<sup>39</sup>
- In 3-week-old CD/CD pigs, IgG was variable following intranasal and intramuscular inoculation with PCV3. Of the nine piglets tested, four showed an obvious IgG response at 7–10 dpi, one showed an obvious IgG response at 14 dpi, and the remaining four piglets either had a mild IgG response or did not seroconvert by the end of the study (28 dpi).<sup>41</sup>

#### VACCINES

Several vaccines are commercially available for PCV2 in swine. They confer humoral and cellular immunity,

reducing mortality while improving average daily weight gain, feed conversion, and uniformity at slaughter. In the United States, PCV2 prevalence has decreased due to widespread vaccination.<sup>133</sup> While there are no commercially available vaccines for PCV3, herd-specific vaccines may be available through veterinary prescription.

# **CROSS-PROTECTION**

Commercial vaccines confer cross-protection between PCV2 subtypes a–d.<sup>5</sup> However, PCV2 vaccines are not expected to protect against other PCVs due to a lower degree of homology. A recent study found that PCV3 viremia was not related to PCV2 vaccine efficacy.<sup>134</sup> Similarly, a field study found that vaccinating pigs for PCV2 did not affect PCV3 presence.<sup>57</sup>

# **GAPS IN PREPAREDNESS**

PCV3 has been associated with many PCVAD- like clinical signs. According to Saporiti and colleagues, the strongest evidence of causality lies with PCV3-systemic and PCV3-reproductive disease. Like PCV2, many PCV3 infections are subclinical, and infection with other swine pathogens is common. In order to better understand the impact of PCV3 on the swine industry, more evidence is needed on pathogenesis, host genomics, viral genomics, and immunology.

# REFERENCES

- 1. Saporiti V, Franzo G, Sibila M, Segalés J. Porcine circovirus 3 (PCV-3) as a causal agent of disease in swine and a proposal of PCV-3 associated disease case definition. *Transbound Emerg Dis.* Jun 2021. doi:10.1111/tbed.14204
- 2. Tischer I, Bode L, Apodaca J, et al. Presence of antibodies reacting with porcine circovirus in sera of humans, mice, and cattle. *Arch Virol*. 1995;140(8):1427-1439. doi:10.1007/BF01322669
- 3. Segalés J, Allan G, Domingo M. Circoviruses. In: Zimmerman J, Karriker L, Ramirez A, Schwartz K, Stevenson G, Zhang J, eds. *Diseases of Swine*. 11th ed. John Wiley & Sons, Inc.; 2019:473-488:Chap 26.
- 4. Segalés J. Porcine circovirus type 2 (PCV2) infections: clinical signs, pathology and laboratory diagnosis. *Virus Res.* Mar 2012;164(1-2):10-9. doi:10.1016/j.virusres.2011.10.007
- 5. Opriessnig T, Karuppannan AK, Castro AMMG, Xiao CT. Porcine circoviruses: current status, knowledge gaps and challenges. *Virus Res.* 09 2020;286:198044. doi:10.1016/j.virusres.2020.198044
- 6. Palinski R, Pineyro P, Shang P, et al. A novel porcine circovirus distantly related to known circoviruses is associated with porcine dermatitis and nephropathy syndrome and reproductive failure. *J Virol.* 2017;91(1). doi:10.1128/jvi.01879-16
- 7. Krakowka S, Hartunian C, Hamberg A, et al. Evaluation of induction of porcine dermatitis and nephropathy syndrome in gnotobiotic pigs with negative results for porcine circovirus type 2. *Am J Vet Res.* Dec 2008;69(12):1615-22. doi:10.2460/ajvr.69.12.1615
- 8. Phan TG, Giannitti F, Rossow S, et al. Detection of a novel circovirus PCV3 in pigs with cardiac and multi-systemic inflammation. *Virol J.* 2016;13(1):184. doi:10.1186/s12985-016-0642-z
- 9. Chen GH, Mai KJ, Zhou L, et al. Detection and genome sequencing of porcine circovirus 3 in neonatal pigs with congenital tremors in South China. *Transbound Emerg Dis.* Dec 2017;64(6):1650-1654. doi:10.1111/tbed.12702
- 10. Arruda B, Piñeyro P, Derscheid R, et al. PCV3-associated disease in the United States swine herd. *Emerg Microbes Infect*. 2019;8(1):684-698. doi:10.1080/22221751.2019.1613176
- 11. Ku X, Chen F, Li P, et al. Identification and genetic characterization of porcine circovirus type 3 in China. *Transbound Emerg Dis.* Jun 2017;64(3):703-708. doi:10.1111/tbed.12638
- 12. Yuzhakov AG, Raev SA, Alekseev KP, et al. First detection and full genome sequence of porcine circovirus type 3 in Russia. *Virus Genes*. Aug 2018;54(4):608-611. doi:10.1007/s11262-018-1582-z
- 13. Deim Z, Dencső L, Erdélyi I, et al. Porcine circovirus type 3 detection in a Hungarian pig farm experiencing reproductive failures. *Vet Rec.* 07 2019;185(3):84. doi:10.1136/vr.104784

- 14. Faccini S, Barbieri I, Gilioli A, et al. Detection and genetic characterization of porcine circovirus type 3 in Italy. *Transbound Emerg Dis.* Dec 2017;64(6):1661-1664. doi:10.1111/tbed.12714
- 15. Zou Y, Zhang N, Zhang J, et al. Molecular detection and sequence analysis of porcine circovirus type 3 in sow sera from farms with prolonged histories of reproductive problems in Hunan, China. *Arch Virol*. Oct 2018;163(10):2841-2847. doi:10.1007/s00705-018-3914-7
- 16. Tochetto C, Lima DA, Varela APM, et al. Full-genome sequence of porcine circovirus type 3 recovered from serum of sows with stillbirths in Brazil. *Transbound Emerg Dis.* Feb 2018;65(1):5-9. doi:10.1111/tbed.12735
- 17. Kim SH, Park JY, Jung JY, et al. Detection and genetic characterization of porcine circovirus 3 from aborted fetuses and pigs with respiratory disease in Korea. *J Vet Sci*. Sep 2018;19(5):721-724. doi:10.4142/jvs.2018.19.5.721
- 18. Dal Santo AC, Cezario KC, Bennemann PE, Machado SA, Martins M. Full-genome sequences of porcine circovirus 3 (PCV3) and high prevalence in mummified fetuses from commercial farms in Brazil. *Microb Pathog.* Apr 2020;141:104027. doi:10.1016/j.micpath.2020.104027
- Tochetto C, de Lima DA, Varela APM, et al. Investigation on porcine circovirus type 3 in serum of farrowing sows with stillbirths. *Microb Pathog*. Dec 2020;149:104316. doi:10.1016/j.micpath.2020.104316
- 20. Saporiti V, Valls L, Maldonado J, et al. Porcine circovirus 3 detection in aborted fetuses and stillborn piglets from swine reproductive failure cases. *Viruses*. 02 2021;13(2). doi:10.3390/v13020264
- 21. Shen H, Liu X, Zhang P, et al. Genome characterization of a porcine circovirus type 3 in South China. *Transbound Emerg Dis.* Feb 2018;65(1):264-266. doi:10.1111/tbed.12639
- 22. Kedkovid R, Woonwong Y, Arunorat J, et al. Porcine circovirus type 3 (PCV3) infection in grower pigs from a Thai farm suffering from porcine respiratory disease complex (PRDC). *Vet Microbiol*. Feb 2018;215:71-76. doi:10.1016/j.vetmic.2018.01.004
- 23. Qi S, Su M, Guo D, et al. Molecular detection and phylogenetic analysis of porcine circovirus type 3 in 21 Provinces of China during 2015-2017. *Transbound Emerg Dis.* Mar 2019;66(2):1004-1015. doi:10.1111/tbed.13125
- 24. Zhai SL, Zhou X, Zhang H, et al. Comparative epidemiology of porcine circovirus type 3 in pigs with different clinical presentations. *Virol J*. Nov 13 2017;14(1):222. doi:10.1186/s12985-017-0892-4
- 25. Saporiti V, Cruz TF, Correa-Fiz F, Núñez JI, Sibila M, Segalés J. Similar frequency of porcine circovirus 3 (PCV-3) detection in serum samples of pigs affected by digestive or respiratory disorders and agematched clinically healthy pigs. *Transbound Emerg Dis.* Jan 2020;67(1):199-205. doi:10.1111/tbed.13341
- 26. Zhang F, Yuan W, Li Z, et al. Porcine circovirus type 3 in pig farms experiencing diarrhea in Jiangxi, China: prevalence, genome sequence and pathogenicity. *Animals (Basel)*. Dec 2020;10(12). doi:10.3390/ani10122324
- 27. Wen S, Sun W, Li Z, et al. The detection of porcine circovirus 3 in Guangxi, China. *Transbound Emerg Dis.* Feb 2018;65(1):27-31. doi:10.1111/tbed.12754
- 28. Zheng S, Wu X, Zhang L, et al. The occurrence of porcine circovirus 3 without clinical infection signs in Shandong Province. *Transbound Emerg Dis.* Oct 2017;64(5):1337-1341. doi:10.1111/tbed.12667
- 29. Kwon T, Yoo SJ, Park CK, Lyoo YS. Prevalence of novel porcine circovirus 3 in Korean pig populations. *Vet Microbiol.* Aug 2017;207:178-180. doi:10.1016/j.vetmic.2017.06.013
- 30. Kedkovid R, Woonwong Y, Arunorat J, et al. Porcine circovirus type 3 (PCV3) shedding in sow colostrum. *Vet Microbiol.* Jul 2018;220:12-17. doi:10.1016/j.vetmic.2018.04.032
- 31. Saporiti V, Martorell S, Cruz TF, et al. Frequency of detection and phylogenetic analysis of porcine circovirus 3 (PCV3) in healthy primiparous and multiparous sows and their mummified fetuses and stillborn. *Pathogens*. Jul 2020;9(7). doi:10.3390/pathogens9070533
- 32. Franzo G, Legnardi M, Hjulsager CK, et al. Full-genome sequencing of porcine circovirus 3 field strains from Denmark, Italy and Spain demonstrates a high within-Europe genetic heterogeneity. *Transbound Emerg Dis.* Feb 16 2018. doi:10.1111/tbed.12836
- 33. Stadejek T, Wozniak A, Milek D, Biernacka K. First detection of porcine circovirus type 3 on commercial pig farms in Poland. *Transbound Emerg Dis.* Oct 2017;64(5):1350-1353. doi:10.1111/tbed.12672

- 34. Saporiti V, Huerta E, Correa-Fiz F, et al. Detection and genotyping of porcine circovirus 2 (PCV-2) and detection of porcine circovirus 3 (PCV-3) in sera from fattening pigs of different European countries. *Transbound Emerg Dis.* Nov 2020;67(6):2521-2531. doi:10.1111/tbed.13596
- 35. Palinski R, Piñeyro P, Shang P, et al. A novel porcine circovirus distantly related to known circoviruses is associated with porcine dermatitis and nephropathy syndrome and reproductive failure. *J Virol.* Jan 2017;91(1). doi:10.1128/JVI.01879-16
- 36. De Conti ER, Resende TP, Marshall-Lund L, Rovira A, Vannucci FA. Histological lesions and replication sites of PCV3 in naturally infected pigs. *Animals (Basel)*. May 2021;11(6). doi:10.3390/ani11061520
- 37. Jiang H, Wang D, Wang J, et al. Induction of porcine dermatitis and nephropathy syndrome in piglets by infection with porcine circovirus type 3. *J Virol*. 02 2019;93(4). doi:10.1128/JVI.02045-18
- 38. Mora-Díaz J, Piñeyro P, Shen H, et al. Isolation of PCV3 from perinatal and reproductive cases of PCV3associated disease and in vivo characterization of PCV3 replication in CD/CD growing pigs. *Viruses*. 2020;12(2). doi:10.3390/v12020219
- 39. Temeeyasen G, Lierman S, Arruda BL, et al. Pathogenicity and immune response against porcine circovirus type 3 infection in caesarean-derived, colostrum-deprived pigs. *J Gen Virol*. 2021;102(1). doi:10.1099/jgv.0.001502
- 40. Hou L, Wang J, Zhang W, et al. Dynamic alterations of gut microbiota in porcine circovirus type 3infected piglets. *Front Microbiol*. 2020;11:1360. doi:10.3389/fmicb.2020.01360
- 41. Wang M, Yu Y, Wu J, et al. Pathogenic characterization of a porcine circovirus type 3 isolate from Heilongjiang, China. *Dis Markers*. 2021;9434944. doi:10.1155/2021/9434944
- 42. Breitbart M, Delwart E, Rosario K, Segalés J, Varsani A, ICTV Report Consortium. ICTV Virus Taxonomy Profile: *Circoviridae. J Gen Virol.* Aug 2017;98(8). doi:10.1099/jgv.0.000871
- 43. Garcia LA, Viancelli A, Rigotto C, et al. Surveillance of human and swine adenovirus, human norovirus and swine circovirus in water samples in Santa Catarina, Brazil. *J Water Health*. Sep 2012;10(3):445-52. doi:10.2166/wh.2012.190
- 44. Abin M, Mor S, Popowski J, Cox R, Goyal S, Patnayak D. Effect of temperature on the survival of porcine circovirus type 2b in fresh pork. *Vet J*. 2013;197(3):898-899. doi:10.1016/j.tvjl.2013.03.015
- 45. Royer R, Nawagitgul P, Halbur P, Paul P. Susceptibility of porcine circovirus type 2 to commercial and laboratory disinfectants. *J Swine Health Prod*. 2001;9(6):281-4.
- 46. Kim HB, Lyoo KS, Joo HS. Efficacy of different disinfectants in vitro against porcine circovirus type 2. *Vet Rec.* May 9 2009;164(19):599-600.
- 47. Martin H, Le Potier MF, Maris P. Virucidal efficacy of nine commercial disinfectants against porcine circovirus type 2. *Vet J.* Sep 2008;177(3):388-93. doi:10.1016/j.tvj1.2007.06.016
- 48. Rodríguez-Arrioja G, Segalés J, Calsamiglia M, et al. Dynamics of porcine circovirus type 2 infection in a herd of pigs with post-weaning multi-systemic wasting syndrome. *Am J Vet Res.* 2002;63(3):354.
- 49. Yaeger MJ. Disorders of Pigs. In: Njaa BL, ed. *Kirkbride's Diagnosis of Abortion and Neonatal Loss in Animals*. John Wiley & Sons; 2011. Vol. Ed.4.
- 50. *Circoviridae*. In: MacLachlan NJ, Dubovi EJ, eds. *Fenner's Veterinary Virology*. 4th ed. Academic Press; 2011:237-242:Chap 13.
- 51. Shi R, Hou L, Wei L, et al. Porcine circovirus type 3 enters into PK15 cells through clathrin- and dynamin-2-mediated endocytosis in a Rab5/Rab7 and pH-dependent fashion. *Front Microbiol*. 2021;12:636307. doi:10.3389/fmicb.2021.636307
- 52. Tan CY, Opaskornkul K, Thanawongnuwech R, Arshad SS, Hassan L, Ooi PT. First molecular detection and complete sequence analysis of porcine circovirus type 3 (PCV3) in Peninsular Malaysia. *PLoS One*. 2020;15(7):e0235832. doi:10.1371/journal.pone.0235832
- 53. Wang J, Zhang Y, Liu L, Pang X, Yuan W. Development of a TaqMan-based real-time PCR assay for the specific detection of porcine circovirus 3. *J Virol Methods*. Oct 2017;248:177-180. doi:10.1016/j.jviromet.2017.07.007
- 54. Yuan L, Liu Y, Chen Y, et al. Optimized real-time fluorescence PCR assay for the detection of porcine circovirus type 3 (PCV3). *BMC Vet Res.* Jul 2020;16(1):249. doi:10.1186/s12917-020-02435-y
- 55. Chen GH, Tang XY, Sun Y, et al. Development of a SYBR green-based real-time quantitative PCR assay to detect PCV3 in pigs. *J Virol Methods*. Jan 2018;251:129-132. doi:10.1016/j.jviromet.2017.10.012

- 56. Franzo G, Legnardi M, Centelleghe C, et al. Development and validation of direct PCR and quantitative PCR assays for the rapid, sensitive, and economical detection of porcine circovirus 3. *J Vet Diagn Invest*. Jul 2018;30(4):538-544. doi:10.1177/1040638718770495
- 57. Plut J, Jamnikar-Ciglenecki U, Golinar-Oven I, Knific T, Stukelj M. A molecular survey and phylogenetic analysis of porcine circovirus type 3 using oral fluid, faeces and serum. *BMC Vet Res*. Aug 2020;16(1):281. doi:10.1186/s12917-020-02489-y
- 58. Park YR, Kim HR, Kim SH, et al. Loop-mediated isothermal amplification assay for the rapid and visual detection of novel porcine circovirus 3. *J Virol Methods*. Mar 2018;253:26-30. doi:10.1016/j.jviromet.2017.12.006
- 59. Zheng S, Wu X, Shi J, et al. Rapid specific and visible detection of porcine circovirus type 3 using loopmediated isothermal amplification (LAMP). *Transbound Emerg Dis.* Mar 4 2018. doi:10.1111/tbed.12835
- 60. Wang J, Zhang Y, Zhang R, et al. Recombinase polymerase amplification assay for rapid detection of porcine circovirus 3. *Mol Cell Probes*. Dec 2017;36:58-61. doi:10.1016/j.mcp.2017.09.001
- 61. Gou H, Bian Z, Cai R, et al. The colorimetric isothermal multiple-self-matching-initiated amplification using cresol red for rapid and sensitive detection of porcine circovirus 3. *Front Vet Sci.* 2020;7:407. doi:10.3389/fvets.2020.00407
- 62. Ji J, Xu X, Wang X, et al. Novel polymerase spiral reaction assay for the visible molecular detection of porcine circovirus type 3. *BMC Vet Res.* Sep 2019;15(1):322. doi:10.1186/s12917-019-2072-9
- 63. Wang Y, Feng Y, Zheng W, et al. A multiplex real-time PCR assay for the detection and differentiation of the newly emerged porcine circovirus type 3 and continuously evolving type 2 strains in the United States. *J Virol Methods*. 07 2019;269:7-12. doi:10.1016/j.jviromet.2019.03.011
- 64. Yang K, Jiao Z, Zhou D, Guo R, Duan Z, Tian Y. Development of a multiplex PCR to detect and discriminate porcine circoviruses in clinical specimens. *BMC Infect Dis.* Sep 2019;19(1):778. doi:10.1186/s12879-019-4398-0
- 65. Li X, Qiao M, Sun M, Tian K. A duplex real-time PCR assay for the simultaneous detection of porcine circovirus 2 and circovirus 3. *Virol Sin.* Apr 2018;33(2):181-186. doi:10.1007/s12250-018-0025-2
- 66. Bai J. Detection and Differentiation of PCV3 from PCV2a, PCV2b, and the Highly Prevalent PCV2d Mutant Strains. *Diagnostic Assay Catalog*. 2016. https://www.swinehealth.org/wpcontent/uploads/2020/09/SHIC-diagnostic-assay-catalog-21Sept2020.pdf
- 67. Kim HR, Park YR, Lim DR, et al. Multiplex real-time polymerase chain reaction for the differential detection of porcine circovirus 2 and 3. *J Virol Methods*. Dec 2017;250:11-16. doi:10.1016/j.jviromet.2017.09.021
- 68. Zhao Y, Han HY, Fan L, et al. Development of a TB green II-based duplex real-time fluorescence quantitative PCR assay for the simultaneous detection of porcine circovirus 2 and 3. *Mol Cell Probes*. 06 2019;45:31-36. doi:10.1016/j.mcp.2019.04.001
- 69. Hou CY, Zhang LH, Zhang YH, et al. Phylogenetic analysis of porcine circovirus 4 in Henan Province of China: A retrospective study from 2011 to 2021. *Transbound Emerg Dis.* Jun 2021. doi:10.1111/tbed.14172
- 70. Chen N, Xiao Y, Li X, et al. Development and application of a quadruplex real-time PCR assay for differential detection of porcine circoviruses (PCV1 to PCV4) in Jiangsu province of China from 2016 to 2020. *Transbound Emerg Dis.* May 2021;68(3):1615-1624. doi:10.1111/tbed.13833
- 71. Han HY, Zheng HH, Zhao Y, et al. Development of a SYBR green I-based duplex real-time fluorescence quantitative PCR assay for the simultaneous detection of porcine epidemic diarrhea virus and porcine circovirus 3. *Mol Cell Probes*. 04 2019;44:44-50. doi:10.1016/j.mcp.2019.02.002
- 72. Tian RB, Jin Y, Xu T, Zhao Y, Wang ZY, Chen HY. Development of a SYBR green I-based duplex realtime PCR assay for detection of pseudorabies virus and porcine circovirus 3. *Mol Cell Probes*. 10 2020;53:101593. doi:10.1016/j.mcp.2020.101593
- 73. Zheng HH, Zhang SJ, Cui JT, et al. Simultaneous detection of classical swine fever virus and porcine circovirus 3 by SYBR green I-based duplex real-time fluorescence quantitative PCR. *Mol Cell Probes*. 04 2020;50:101524. doi:10.1016/j.mcp.2020.101524
- 74. Tan CY, Lin CN, Ooi PT. What do we know about porcine circovirus 3 (PCV3) diagnosis so far?: A review. *Transbound Emerg Dis.* Jun 2021. doi:10.1111/tbed.14185

- 75. Resende TP, Marshall Lund L, Rossow S, Vannucci FA. Next-generation sequencing coupled with in situ hybridization: a novel diagnostic platform to investigate swine emerging pathogens and new variants of endemic viruses. *Front Vet Sci.* 2019;6:403. doi:10.3389/fvets.2019.00403
- 76. Gagnon C, Music N, Fontaine G, Tremblay D, Harel J. Emergence of a new type of porcine circovirus in swine (PCV): a type 1 and type 2 PCV recombinant. *Vet Microbiol*. 7/29/ 2010;144(1–2):18-23. doi:10.1016/j.vetmic.2009.09.072
- 77. Oh T, Chae C. First isolation and genetic characterization of porcine circovirus type 3 using primary porcine kidney cells. *Vet Microbiol.* Feb 2020;241:108576. doi:10.1016/j.vetmic.2020.108576
- 78. Deng J, Li X, Zheng D, et al. Establishment and application of an indirect ELISA for porcine circovirus 3. *Arch Virol*. Feb 2018;163(2):479-482. doi:10.1007/s00705-017-3607-7
- 79. Zhang S, Wang D, Jiang Y, et al. Development and application of a baculovirus-expressed capsid proteinbased indirect ELISA for detection of porcine circovirus 3 IgG antibodies. *BMC Vet Res*. Mar 2019;15(1):79. doi:10.1186/s12917-019-1810-3
- 80. Wang Y, Wang G, Duan WT, et al. Self-assembly into virus-like particles of the recombinant capsid protein of porcine circovirus type 3 and its application on antibodies detection. *AMB Express*. Jan 2020;10(1):3. doi:10.1186/s13568-019-0940-0
- 81. Bai J. Development and Evaluation of Antibody Detection Assay for PCV3. *Diagnostic Assay Catalog*. Swine Health Information Center; 2017. https://www.swinehealth.org/wpcontent/uploads/2020/09/SHIC-diagnostic-assay-catalog-21Sept2020.pdf
- 82. Ouyang T, Niu G, Liu X, Zhang X, Zhang Y, Ren L. Recent progress on porcine circovirus type 3. *Infect Genet Evol.* 09 2019;73:227-233. doi:10.1016/j.meegid.2019.05.009
- 83. Franzo G, Grassi L, Tucciarone CM, et al. A wild circulation: high presence of porcine circovirus 3 in different mammalian wild hosts and ticks. *Transbound Emerg Dis.* Jul 2019;66(4):1548-1557. doi:10.1111/tbed.13180
- 84. Klaumann F, Dias-Alves A, Cabezón O, et al. Porcine circovirus 3 is highly prevalent in serum and tissues and may persistently infect wild boar (*Sus scrofa scrofa*). *Transbound Emerg Dis*. Jan 2019;66(1):91-101. doi:10.1111/tbed.12988
- 85. Franzo G, Tucciarone CM, Drigo M, et al. First report of wild boar susceptibility to porcine circovirus type 3: high prevalence in the Colli Euganei Regional Park (Italy) in the absence of clinical signs. *Transbound Emerg Dis.* Aug 2018;65(4):957-962. doi:10.1111/tbed.12905
- 86. Prinz C, Stillfried M, Neubert LK, Denner J. Detection of PCV3 in German wild boars. *Virol J*. 02 2019;16(1):25. doi:10.1186/s12985-019-1133-9
- 87. Dei Giudici S, Franzoni G, Bonelli P, et al. Genetic characterization of porcine circovirus 3 strains circulating in Sardinian pigs and wild boars. *Pathogens*. May 2020;9(5). doi:10.3390/pathogens9050344
- 88. Song S, Park KN, Choe S, et al. Complete genome sequences of two type 3 porcine circoviruses, WB17KW and WB20GG, isolated from Korean wild boar. *Microbiol Resour Announc*. Feb 2021;10(6). doi:10.1128/MRA.01386-20
- 89. Dhandapani G, Yoon SW, Noh JY, et al. Detection of porcine circovirus 3 from captured wild boars in Korea. *Vet Med Sci.* May 2021. doi:10.1002/vms3.518
- 90. Varela APM, Loiko MR, Andrade JDS, et al. Complete genome characterization of porcine circovirus 3 recovered from wild boars in Southern Brazil. *Transbound Emerg Dis.* Mar 2021;68(2):240-247. doi:10.1111/tbed.13679
- 91. Wang GS, Sun N, Tian FL, et al. Genetic analysis of porcine circovirus type 2 from dead minks. *J Gen Virol*. Jun 20 2016. doi:10.1099/jgv.0.000529
- 92. Zhai S-L, Chen R-A, Chen S-N, et al. First molecular detection of porcine circovirus type 2 in bovids in China. *Virus Genes*. 2014// 2014;49(3):507-511. doi:10.1007/s11262-014-1117-1
- 93. Li L, Shan T, Soji OB, et al. Possible cross-species transmission of circoviruses and cycloviruses among farm animals. *J Gen Virol*. Apr 2011;92(Pt 4):768-72. doi:10.1099/vir.0.028704-0
- 94. Herbst W, Willems H. Detection of virus particles resembling circovirus and porcine circovirus 2a (PCV2a) sequences in feces of dogs. *Res Vet Sci.* Dec 2017;115:51-53. doi:10.1016/j.rvsc.2017.01.014
- 95. Krog JS, Larsen LE, Schultz AC. Enteric porcine viruses in farmed shellfish in Denmark. *In J Food Microbiol*. 9/1/2014;186:105-109. doi:10.1016/j.ijfoodmicro.2014.06.012

- 96. Blunt R, McOrist S, McKillen J, McNair I, Jiang T, Mellits K. House fly vector for porcine circovirus 2b on commercial pig farms. *Vet Microbiol.* May 2011;149(3-4):452-5. doi:10.1016/j.vetmic.2010.11.019
- 97. Zhai SL, Lu SS, Wei WK, et al. Reservoirs of porcine circoviruses: a mini review. *Front Vet Sci.* 2019;6:319. doi:10.3389/fvets.2019.00319
- 98. Zhang J, Liu Z, Zou Y, et al. First molecular detection of porcine circovirus type 3 in dogs in China. *Virus Genes*. Feb 2018;54(1):140-144. doi:10.1007/s11262-017-1509-0
- 99. Sun W, Wang W, Xin J, et al. An epidemiological investigation of porcine circovirus 3 infection in dogs in the Guangxi Province from 2015 to 2017, China. *Virus Res.* 09 2019;270:197663. doi:10.1016/j.virusres.2019.197663
- 100. Wang W, Sun W, Cao L, et al. An epidemiological investigation of porcine circovirus 3 infection in cattle in Shandong province, China. *BMC Vet Res.* Feb 2019;15(1):60. doi:10.1186/s12917-019-1793-0
- 101. Jiang S, Zhou N, Li Y, An J, Chang T. Detection and sequencing of porcine circovirus 3 in commercially sourced laboratory mice. *Vet Med Sci.* 05 2019;5(2):176-181. doi:10.1002/vms3.144
- 102. Ha Z, Li JF, Xie CZ, et al. First detection and genomic characterization of porcine circovirus 3 in mosquitoes from pig farms in China. *Vet Microbiol*. Jan 2020;240:108522. doi:10.1016/j.vetmic.2019.108522
- 103. Grassi L, Tagliapietra V, Rizzoli A, et al. Lack of evidence on the susceptibility of ticks and wild rodent species to PCV3 infection. *Pathogens*. Aug 2020;9(9). doi:10.3390/pathogens9090682
- 104. Czyżewska-Dors E, Núñez JI, Saporiti V, et al. Detection of porcine circovirus 3 in wildlife species in Spain. *Pathogens*. May 2020;9(5). doi:10.3390/pathogens9050341
- 105. Franzo G, Cortey M, Olvera A, et al. Revisiting the taxonomical classification of porcine circovirus type 2 (PCV2): still a real challenge. *Virol J*. 2015;12:131. doi:10.1186/s12985-015-0361-x
- 106. Saraiva GL, Vidigal PMP, Fietto JLR, Bressan GC, Silva Junior A, de Almeida MR. Evolutionary analysis of porcine circovirus 3 (PCV3) indicates an ancient origin for its current strains and a worldwide dispersion. *Virus Genes*. Mar 3 2018. doi:10.1007/s11262-018-1545-4
- 107. Souza TCGD, Gava D, Schaefer R, Leme RA, Porto GDS, Alfieri AA. Porcine circovirus 3a field strains in free-living wild boars in Paraná state, Brazil. *Animals (Basel)*. May 2021;11(6). doi:10.3390/ani11061634
- 108. Vargas-Bermudez DS, Campos FS, Bonil L, Mogollon D, Jaime J. First detection of porcine circovirus type 3 in Colombia and the complete genome sequence demonstrates the circulation of PCV3a1 and PCV3a2. *Vet Med Sci.* 05 2019;5(2):182-188. doi:10.1002/vms3.155
- 109. Allan GM, McNeilly F, Kennedy S, et al. Isolation of porcine circovirus-like viruses from pigs with a wasting disease in the USA and Europe. *J Vet Diagn Invest*. Jan 1998;10(1):3-10. doi:10.1177/104063879801000102
- 110. Collins PJ, McKillen J, Allan G. Porcine circovirus type 3 in the UK. Vet Rec. 2017:181(22):599.
- 111. Ye X, Berg M, Fossum C, Wallgren P, Blomström AL. Detection and genetic characterisation of porcine circovirus 3 from pigs in Sweden. *Virus Genes.* Jun 2018;54(3):466-469. doi:10.1007/s11262-018-1553-4
- 112. Woźniak A, Miłek D, Stadejek T. Wide range of the prevalence and viral loads of porcine circovirus type
   3 (PCV3) in different clinical materials from 21 Polish pig farms. *Pathogens*. May 2020;9(5).
   doi:10.3390/pathogens9050411
- 113. Liu Y, Zhang S, Song X, et al. The prevalence of novel porcine circovirus type 3 isolates in pig farms in China. *Transbound Emerg Dis.* Sep 2019;66(5):2143-2151. doi:10.1111/tbed.13266
- 114. Cai L, Ni J, Xia Y, et al. Identification of an emerging recombinant cluster in porcine circovirus type 2. *Virus Res.* Apr 2012;165(1):95-102. doi:10.1016/j.virusres.2012.01.008
- 115. Hayashi S, Ohshima Y, Furuya Y, et al. First detection of porcine circovirus type 3 in Japan. *J Vet Med Sci*. Sep 2018;80(9):1468-1472. doi:10.1292/jvms.18-0079
- 116. Sukmak M, Thanantong N, Poolperm P, et al. The retrospective identification and molecular epidemiology of porcine circovirus type 3 (PCV3) in swine in Thailand from 2006 to 2017. *Transbound Emerg Dis.* Jan 2019;66(1):611-616. doi:10.1111/tbed.13057
- 117. Bera BC, Choudhary M, Anand T, et al. Detection and genetic characterization of porcine circovirus 3 (PCV3) in pigs in India. *Transbound Emerg Dis*. May 2020;67(3):1062-1067. doi:10.1111/tbed.13463

- 118. Zhang HH, Hu WQ, Li JY, et al. Novel circovirus species identified in farmed pigs designated as porcine circovirus 4, Hunan province, China. *Transbound Emerg Dis.* May 2020;67(3):1057-1061. doi:10.1111/tbed.13446
- 119. Tian RB, Zhao Y, Cui JT, et al. Molecular detection and phylogenetic analysis of porcine circovirus 4 in Henan and Shanxi Provinces of China. *Transbound Emerg Dis.* Mar 2021;68(2):276-282. doi:10.1111/tbed.13714
- 120. Sun W, Du Q, Han Z, et al. Detection and genetic characterization of porcine circovirus 4 (PCV4) in Guangxi, China. *Gene*. Mar 2021;773:145384. doi:10.1016/j.gene.2020.145384
- 121. Li G, He W, Zhu H, et al. Origin, genetic diversity, and evolutionary dynamics of novel porcine circovirus 3. *Adv Sci (Weinh)*. Sep 2018;5(9):1800275. doi:10.1002/advs.201800275
- 122. Franzo G, Legnardi M, Tucciarone CM, et al. Porcine circovirus type 3: a threat to the pig industry? *Vet Rec.* 01 2018;182(3):83. doi:10.1136/vr.k91
- 123. Klaumann F, Correa-Fiz F, Sibila M, Núñez JI, Segalés J. Infection dynamics of porcine circovirus type 3 in longitudinally sampled pigs from four Spanish farms. *Vet Rec.* 05 2019;184(20):619. doi:10.1136/vr.105219
- 124. Opriessnig T, Fenaux M, Thomas P, et al. Evidence of breed-dependent differences in susceptibility to porcine circovirus type-2-associated disease and lesions. *Vet Pathol*. May 2006;43(3):281-93. doi:10.1354/vp.43-3-281
- 125. International Committee on the Taxonomy of Viruses (ICTV). Virus Taxonomy Update: Circovirus. Accessed July 14, 2021. https://talk.ictvonline.org/ictv-reports/ictv\_online\_report/ssdnaviruses/w/circoviridae/659/genus-circovirus
- 126. Cortey M, Segalés J. Low levels of diversity among genomes of porcine circovirus type 1 (PCV1) points to differential adaptive selection between porcine circoviruses. *Virology*. 1/20/ 2012;422(2):161-164. doi:10.1016/j.virol.2011.10.014
- 127. Franzo G, Segales J, Tucciarone CM, Cecchinato M, Drigo M. The analysis of genome composition and codon bias reveals distinctive patterns between avian and mammalian circoviruses which suggest a potential recombinant origin for Porcine circovirus 3. *PLoS One*. 2018;13(6):e0199950. doi:10.1371/journal.pone.0199950
- 128. Tischer I, Rasch R, Tochtermann G. Characterization of papovavirus-and picornavirus-like particles in permanent pig kidney cell lines. *Zentralbl Bakteriol Orig A*. Feb 1974;226(2):153-67.
- 129. Tischer I, Gelderblom H, Vettermann W, Koch MA. A very small porcine virus with circular singlestranded DNA. *Nature*. Jan 1982;295(5844):64-6. doi:10.1038/295064a0
- 130. Harding J, Clark E. Recognizing and diagnosing post-weaning multi-systemic wasting syndrome (PWS). *J Swine Health Prod.* 1997;5(5):201-203.
- 131. Nayar GP, Hamel A, Lin L. Detection and characterization of porcine circovirus associated with postweaning multi-systemic wasting syndrome in pigs. *Can Vet J.* Jun 1997;38(6):385-6.
- 132. Ha Z, Yu C, Xie C, et al. Retrospective surveillance of porcine circovirus 4 in pigs in Inner Mongolia, China, from 2016 to 2018. *Arch Virol*. Jul 2021;166(7):1951-1959. doi:10.1007/s00705-021-05088-w
- Dvorak C, Yang Y, Haley C, Sharma N, Murtaugh M. National reduction in porcine circovirus type 2 prevalence following introduction of vaccination. *Vet Microbiol*. 2016;189:86-90. doi:10.1016/j.vetmic.2016.05.002
- 134. Woźniak A, Miłek D, Bąska P, Stadejek T. Does porcine circovirus type 3 (PCV3) interfere with porcine circovirus type 2 (PCV2) vaccine efficacy? *Transbound Emerg Dis.* Jul 2019;66(4):1454-1461. doi:10.1111/tbed.13221