Research

The index herd with PMWS in Sweden: Presence of serum amyloid A, circovirus 2 viral load and antibody levels in healthy and PMWS-affected pigs

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Abstract

Background: Postweaning Multisystemic Wasting Syndrome (PMWS) is an emerging disease in pigs of multifactorial origin, but associated to porcine circovirus type 2 (PCV2) infection. PMWS was first diagnosed in Sweden at a progeny test station that received pigs aged five weeks from 19 different nucleus herds on the day after weaning. The objective of this study was to examine, for the first time in an index outbreak of PMWS, the relationship between PCV2 virus, antibodies to PCV2 and serum amyloid A (SAA) in sequentially collected serum samples from pigs with and without signs of PMWS.

Methods: Forty pigs of the last batch that entered the station at a mean age of 37.5 days were monitored for signs of PMWS during the first 55 days after arrival. Serum was collected on six occasions and analysed for presence of PCV2 DNA and antibodies to PCV2, as well as for levels of SAA.

Results: Four of the pigs (10%) were concluded to have developed PMWS, with necropsy confirmation in three of them. These pigs displayed low levels of maternal antibodies to PCV2, more than 10⁷ PCV2 viral DNA copies per ml serum and failed to mount a serological response to the virus. Starting between day 23 and 34 after arrival, an increase in PCV2 viral load was seen in all pigs, but PCV2 did not induce any SAA-response. Pigs that remained healthy seroconverted to PCV2 as the viral load was increased, regardless of initially having low or high levels of PCV2-antibodies.

Conclusion: In this index case of PMWS in Sweden, pigs affected by PMWS were not able to mount a relevant serum antibody response which contributed to the disease progression. The maximal PCV2 virus load was significantly higher and was also detected at an earlier stage in PMWS-affected pigs than in healthy pigs. However, a viral load above 10⁷ PCV2 DNA copies per ml serum was also recorded in 18 out of 34 pigs without any clinical signs of PMWS, suggesting that these pigs were able to initiate a protective immune response to PCV2.

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Introduction
Postweaning multisystemic wasting syndrome (PMWS) is a disease of pigs first recognised in Canada in 1991 that now is a global epizootic [1-3]. PMWS is regarded as a multifactorial disease although infection of pigs with porcine circovirus 2 (PCV2) is recognised as an essential component of the disease process. A difference in pathogenecity between various isolates of PCV2 has been suggested [4-8], but it is also generally accepted that the presence of other infectious or non-infectious factors is required for the development of the full clinical disease [9-12]. Experimental studies in colostrum deprived piglets have demonstrated that such factors can include co-infection with other microbes such as porcine parvovirus [9,10,13], porcine reproductive and respiratory syndrome virus [14,15] or Mycoplasma hyopneumoniae [16], but PMWS could also be induced by PCV2 in combinations with either immunostimulators [17] or immunosuppressors [18]. Experimental infections in both conventional and specific pathogen-free (SPF) pigs with tissue homogenates from PMWS-affected weaners have also induced mild PMWS [19,20]. In these experiments, all inoculated pigs seroconverted to PCV2, but not to any other known virus or bacteria. Transmission of PMWS has also been demonstrated by mixing healthy weaners with PMWS-affected pigs in previously emptied and cleaned facilities [21]. However, the reasons why some pigs develop PMWS while other pen mates remain healthy is still not clear [12,22].

PMWS was diagnosed for the first time in Sweden at a progeny test station in December 2003 [23]. As a consequence the station was closed down, but all pigs present at the station were reared to the weight of 100 kg before closure. To date there have been no reports in the literature on the investigation of the health status related to the load of PCV2 in blood, the level of antibodies to PCV2 virus and serum amyloid A (SAA) determined in sequentially collected serum samples from an on-going index case of PMWS. Within the last batch of pigs reared at the test station, this was determined in 40 pigs that also were monitored closely for clinical signs of PMWS.

Methods
Initial health status of the animals
Pigs in Sweden are free from all diseases listed by the Office International des Epizooties (OIE), including Aujeszky’s disease (AD), porcine reproductive and respiratory syndrome (PRRS), and also from porcine endemic diarrhoea (PED) and transmissible gastro-enteritis (TGE). The animals in this study emanated from purebred nucleus herds also declared free from atrophic rhinitis (toxin producing strains of Pasteurella multocida), Salmonella spp, swine dysentery (Brachyspira hyodysenteriae) and mange (Sarcoptes scabiei). Infections with Mycoplasma hyopneumoniae and Actinobacillus pleuropneumoniae are widespread in the conventional pig population in Sweden, but the influence of these diseases has decreased since the 1990s due to the commonly performed age segregated production from birth to slaughter [24].

Herd, animals and experimental design
The present study that was conducted at a progeny test station was approved by the ethical committee in Uppsala, Sweden (C38/4). The test station was established in March 2002, and introduced intensified rearing strategies previously not used in the country with the aim of improving genetic selection. Briefly, boars from 19 nucleus herds (pure bred Landrace, Yorkshire or Hampshire) were allocated to the test station on the day after weaning at the age of approximately five weeks. On arrival they were mixed with boars of the same age from other herds, and the animals were remixed according to weight every fortnight four times before entering the pen for individual testing. During the individual test period the boars were still group housed, but individually fed via transponders.

In December 2003 PMWS was diagnosed in this herd as the index case of Sweden [23] by employing the internationally accepted criterias for diagnosing PMWS at individual and herd levels [25,26]. As a consequence, the station was closed down, but animals already at the station were reared to echo-sounding at market weight before being slaughtered.

The 40 pigs selected for this study belonged to the last group that entered the test station before closing. The pigs that were mixed with each other on arrival came from 10 nucleus herds (Table 1). The health status of the animals was recorded during the first 55 days after arrival. Pigs attended for clinical signs resembling PMWS (i.e. underweight or obvious loss of weight) during this time were denoted as “thin”. Pigs that died or were euthanized during the observation period were sent for necropsies whenever suitable. The necropsies were carried out at Analyce AB (Lidköping, Sweden), and formalin fixed samples were sent to the National Veterinary Institute SVA for histological and PCV 2 immunohistochemical analyses.

Collection of blood and analyses performed
Blood samples without additives were collected by jugular vein punctures on days 9, 17, 23, 34, 43 and 55 after arrival. The sera were separated and stored at -20°C until analysed.

Presence of PCV2 in individual serum samples was measured using a quantitative real time PCR assay previously described [27], with a detection limit of 1,100 DNA copies per ml (Log 3.04). In brief, nucleic acids were extracted
from 200 μl serum using an EasyMag nucleic acid extractor (Biomerieux, Durha, USA) and eluted in 55 μl elution buffer. For the quantitative PCR, 2.5 μl of each elute was run in a 25 μl reaction with primers and probe previously described [27] on an MxPro 3005 PCR machine (Stratagene, La Jolla, USA). The detection limit of the PCR was 1.1 × 10^3 (10 Log 3.04) genome copies per ml serum, and results are presented as 10-logarithms.

Antibodies specific to PCV2 in serum were measured using an immuno-peroxidase-monolayer-assay (IPMA) method previously described. [28]. In brief, freshly trypsinized cells of the PCV-free continuous cell line PK15 A were inoculated with PCV 2 (Stoon-1010) [29]. The inoculated cell-suspension was seeded in 96-well cell culture plates and incubated for 5 days at 37°C (5% CO₂). The culture medium was removed and the cells were washed with physiological saline. The plates were then fixed in 99.5% ethanol for one hour. The ethanol was removed and glycerol (87%) diluted 1:1 in PBS was added and the plates were kept at -20°C until further use. The serum samples were diluted in twofold dilutions from 1:10 to 1:20,480 (Log 1.0 to Log 4.3), The results are presented as Log 10 levels of the antibody titres and seroconversion between two consecutive samplings was defined as an increase with at least two titre steps, corresponding to an increase with at least Log 0.6.

The serum levels of the acute phase protein Serum Amyloid A (SAA) were analysed using a commercial kit (Serum Amyloid A Assay TP-802, Tridelta, Maynooth, Ireland) according to the instructions of the manufacturer. The results are presented as mg SAA per L serum.

**Statistics**
All results in the text are given as mean values ± standard deviations. Groups of pigs were compared using Student’s t-tests in pair wise comparisons between groups. For comparisons within groups over time, consecutive recordings were compared with each other using paired t-tests.

**Results**
The pigs studied were mixed with each other on arrival to the test station at a mean age of 37.5 ± 4.0 days (Table 1). Six of the 40 pigs used in this study were denoted as "thin" during the observation period (Table 2). One of these pigs came from an SPF-herd and was denoted as "thin" on day 20.

### Table 1: Herd of origin, breed, mean weight and age of the 40 pigs examined.

| Herd of origin | Breed | Age Days | Weight Kg | PCV2 antibodies Range | Denoted “thin” |
|---------------|-------|----------|-----------|-----------------------|----------------|
| Y1-SPF        | 3     | -        | 41.3 ± 1.2| 11.9 ± 10.6           | 2.2 – 2.8      |
| Y2-conv       | 4     | -        | 36.5 ± 1.3| 10.8 ± 0.9            | 2.2 – 3.1      |
| Y3-conv       | 4     | -        | 39.0 ± 0.8| 10.1 ± 1.6            | 2.2 – 3.1      |
| Y4-conv       | 4     | -        | 39.5 ± 1.3| 13.3 ± 2.0            | 2.2 – 2.8      |
| L1-conv       | -     | 4        | 36.8 ± 2.2| 10.8 ± 0.7            | 2.5 – 3.4      |
| L2-conv       | -     | 4        | 33.3 ± 3.3| 11.5 ± 1.0            | 2.2 – 3.1      |
| L3-conv       | -     | 4        | 33.5 ± 5.7| 10.7 ± 1.9            | 2.2 – 2.8      |
| L&H-conv      | -     | 3 2      | 39.9 ± 4.9| 11.3 ± 2.1            | 2.2 – 3.1      |
| H1-conv       | -     | 4        | 40.1 ± 2.9| 12.1 ± 2.0            | 2.2 – 3.7      |
| H2-conv       | -     | 4        | 35.3 ± 3.7| 11.9 ± 1.4            | 2.5 – 3.1      |
| Overall       | 15    | 15 10    | 37.5 ± 4.0| 11.4 ± 1.6            | 2.2 – 3.7      |

Y = Yorkshire, L = Landrace, H = Hampshire; SPF = SPF-herd

The table also shows the Log-range of serum antibody titres to PCV2 when determined at nine days after arrival, the number of pigs denoted as thin or diagnosed with PMWS before day 55 after arrival.
12 after arrival at 49 days of age. The other five pigs were recorded as "thin" between day 20 and day 46 after arrival (ranging from 59 to 86 days of age). Two of these six pigs (581-SPF and 1008) were alive at the end of the observation period on day 55. One of the pigs (666) died from wasting on day 39 after arrival, and the remaining three pigs (1037, 842 and 418) were euthanised due to wasting on day 46. PMWS was confirmed as the cause of death in all these three pigs (Table 2) according to international standards [25,26], which apart from wasting included enlarged lymph nodes macroscopically with typical histological lesions and the presence of an abundance of PCV2 antigen in these lesions.

The serum levels of PCV2 virus DNA, antibodies to PCV2 and SAA of the six pigs denoted as "thin" are shown individually in Figure 1. No increased amounts of PCV2 virus DNA in serum was recorded in SPF-pig 581 or pig 1008 when they were denoted as "thin" on day 12 and day 20 after arrival, respectively. Both these pigs expressed a PCV2 virus DNA load above 10^7 at a later stage during the observation period (day 43 and 34 after arrival), and at that time also responded by seroconversion to PCV2 (Figure 1). Thus, these two pigs were not diagnosed as having PMWS, and it is noteworthy that they were the only two pigs denoted as "thin" that were still alive at day 55 after arrival.

In four of the six pigs diagnosed as having PMWS (Table 2), the wasting coincided in time with serum levels of PCV2 exceeding 10^7 per ml. None of these pigs showed a clear seroconversion to PCV2 in relation to this increased serum load of PCV2 (Figure 1). As outlined above the remaining two "thin" pigs in this group showed an active seroconversion to PCV2. A serum antibody titer of Log 2.2 and Log 2.5, respectively. Pig 1037 was diagnosed with PMWS by necropsy on day 46, but the load of PCV2 had decreased to below 10^7 per ml serum at day 43. Thus it cannot be excluded that this pig was in an early phase of recovery from PMWS at the time for necropsy.

As the first sampling occasion occurred nine days after weaning at a mean age of 46.5 ± 4.0 days, the antibody status at that time is referred to as remaining maternal immunity. At that time, the mean serum antibody titre of the six pigs denoted as "thin" ranged from Log 2.2 to 2.5 with a mean value of Log 2.36 ± 0.18 for the four pigs diagnosed with PMWS (Figure 1) and of 2.2 for each of the two other pigs. The 34 pigs that remained free from signs of PMWS during this period were divided into two groups according to the level of maternal antibodies to PCV2. One group had a similar range (log 2.2 to 2.5) as the pigs later diagnosed with PMWS with a mean titre of 2.29 ± 0.15 (n = 17), while the other group had higher amounts of maternal antibodies (log 2.8 to 3.7) with a mean titre of 3.00 ± 0.26 (n = 17). Despite decreasing (p < 0.01) from arrival to day 34, the amounts of serum antibodies to PCV2 in the latter group was significantly (p < 0.001 to p < 0.02) higher than in the other two groups until day 23 after arrival (Figure 2).

In the six pigs denoted as "thin" within the first 55 days after arrival to the test station.

| Pig ID | Breed | Arrival Age (days) | Weight (kg) | DWG (from birth) | Day after arrival | Age (days) | Status | Day 55 after arrival | Diagnose at necropsy |
|-------|-------|-------------------|-------------|------------------|-----------------|------------|--------|---------------------|---------------------|
| 581   | Y-SPF | 40                | 11.8        | 258              | 12              | 52         | Alive  | -                   | -                   |
| 1008  | Y     | 39                | 9.6         | 208              | 20              | 59         | Alive  | -                   | -                   |
| 666   | L     | 42                | 9.0         | 179              | 20              | 62         | Dead   | Not done            | PMWS                |
| 1037  | Y     | 36                | 11.0        | 264              | 32              | 68         | Euthanized | PMWS            | PMWS                |
| 842   | Y     | 38                | 11.4        | 261              | 41              | 79         | Euthanized | PMWS            | PMWS                |
| 418   | H     | 40                | 13.7        | 305              | 46              | 86         | Euthanized | PMWS            | PMWS                |

Y = Yorkshire, L = Landrace, H = Hampshire; SPF = of SPF-origin

The birth weight was standardised to 1.5 kg when the daily weight gains were calculated.
Figure 1
Individual recordings for pigs attended as "thin". The black arrows indicate date when the pig was recorded as "thin" for the first time. The Y-axis shows the PCV2 genome copies per ml serum (grey circles) and serum antibody titres to PCV2 (black squares) as Log 10 values. The serum level of SAA (white diamonds) are presented as mg per L serum x 10. Enlarged white diamonds indicate that the level is above 100 mg SAA per L serum (equal to 10 on that scale). The retarded growth recorded for pigs 581 (a) and 1008 (b) was not accompanied with development of other symptoms of PMWS and these pigs were still alive on day 55. The other four thin pigs got the diagnosis PMWS and were dead on day 55. PMWS was confirmed by necropsy in three euthanized pigs (d-e), whereas necropsy not was performed on pig c that died of wasting.
The amount of PCV2 in serum, measured by PCR detection of nucleic acid, increased to Log 6.53 ± 2.77 at day 23 after arrival in the four pigs later to be diagnosed with PMWS (Figure 3). On day 34, both groups with low levels of maternal antibodies had higher (p < 0.005) load of PCV2 in serum than the pigs with high levels of maternal antibodies.

At an individual level, the pigs peaked in PCV2-load in serum on day 23, 34 or 43 after arrival (Table 3). When comparing the peak load of virus regardless of when it took place in time, pigs with the diagnosis PMWS expressed a higher peak viral load than both the other groups (p < 0.005). Furthermore, the healthy pigs with low levels of maternal antibodies to PCV2 peaked with a higher (p < 0.05) viral load than pigs with high levels of maternal antibodies. Pigs with the diagnosis PMWS peaked at 33.5 days after arrival, whereas healthy pigs with high levels of maternal antibodies peaked at day 40 (Table 3).

As shown in figure 4, the serum antibody levels of the four pigs with the diagnosis PMWS was similar when the viral load exceeded $10^7$ as it was the week before (Log 2.20 ± 0.25 vs Log 2.28 ± 0.29), and it still remained at that level one week later (Log 2.51 ± 0.31). In contrast, pigs with a viral load of PCV2 exceeding $10^7$ that remained healthy increased (p < 0.05 to 0.001) their antibody levels between the corresponding sampling occasions, regardless of having low (n = 11) or high (n = 7) levels of maternal antibody levels. One week after that the viral load of PCV2 exceeded $10^7$, both these groups differed (p < 0.01) significantly (p < 0.01) to that of the pigs that developed PMWS with respect to level of serum antibodies to PCV2 (Figure 4).

As seen in Figure 1, individual pigs expressed high levels of SAA in serum at different time points, and pigs could have increased levels of serum-PCV2 without a contemporaneous SAA-response and vice versa. With one exception, no significant differences in SAA levels in serum were

Figure 2
Mean log titres of antibodies to PCV2 in serum. Black circles represent pigs diagnosed with PMWS (n = 4). All these pigs were dead at day 55. Pigs without clinical signs of PMWS were grouped as having low (squares, n = 17) or high (triangles, n = 17) maternal immunity to PCV2. All these pigs were alive at day 55. Significant differences between the group with high maternal immunity and the other groups are indicated in the figure (p < 0.05 = *, p < 0.01 = **, p < 0.001 = ***).
obtained between the three groups at any occasion (Table 4). The highest level of SAA in serum was obtained at the first sampling occasion at day 9 after arrival. At that time more than 100 mg SAA per L serum was obtained in 10 out of the 40 pigs. However, high levels of SAA could be seen occasionally in individual pigs during the entire period studied.

**Discussion**

The close examination of 40 randomly selected pigs suggests that four pigs denoted as “thin” actually developed PMWS. This was confirmed by necropsies in three of them, and necropsy still is the golden standard for diagnosing PMWS in individual pigs [1-3,26]. It is notable that all PMWS-affected pigs had low levels of maternal antibodies to PCV2, and that none out of the 17 pigs with high levels of maternal antibodies to PCV2 developed clinical signs resembling PMWS. This concurs well with suggestions that antibodies to PCV2 can hinder the development of PMWS [30-33]. The IPMA-method used in this study does not measure truly neutralising antibodies, but a positive correlation between neutralising antibodies and total amount of antibodies has previously been reported [34,35].

**Figure 3**

Mean log levels of PCV2 DNA copy number in serum. Black circles represent pigs diagnosed with PMWS (n = 4). All these pigs were dead at day 55. Pigs without clinical signs of PMWS were grouped as having low (squares, n = 17) or high (triangles, n = 17) maternal immunity to PCV2. All these pigs were alive at day 55. ** illustrates that the indicated groups differ (p < 0.01) from the group with high maternal immunity to PCV2 at that day.

| Category                                | N   | Peak Day after arrival | Peak Log PCV2 | t-test versus |
|-----------------------------------------|-----|------------------------|---------------|---------------|
| PMWS                                    | 4   | 33.5 ± 8.2             | 8.9 ± 0.4     | P < 0.001     | p < 0.001     |
| Healthy, low level of maternal antibodies | 17  | 36.7 ± 5.7             | 7.3 ± 1.0     | -             | p < 0.05     |
| Healthy, high level of maternal antibodies | 17  | 39.8 ± 9.1             | 6.5 ± 0.9     | p < 0.05      | -             |
Results from the present study support the important role of the maternal immunity in preventing development of PMWS, as also suggested by others [31,36]. However, Table 1 shows that every nucleus herd sending pigs to the station had delivered individual pigs with low levels of maternally derived antibodies to PCV2, i.e. pigs that potentially could develop PMWS but did not. This is consistent with an earlier report showing that some farm pigs with low levels of PCV2 antibodies in serum did not develop PMWS whereas some pigs with higher levels did [34]. The present study confirms this finding and suggests that low levels of maternal antibodies to PCV2 in piglets do not necessarily lead to development of PMWS. Indeed, 17 pigs with low levels of antibodies to PCV2 on arrival remained free from PMWS. These pigs responded better to the PCV2 exposure than pigs developing PMWS in terms of a rapid development of antibodies to PCV2. Pigs that developed PMWS basically did not seroconvert to PCV2 as they became diseased. The absence of a proper immune response to PCV2 in these pigs undoubtedly contributed to the excessive proliferation of PCV2 which is commonly seen in pigs affected by PMWS [25,1-3].

As stated above, every nucleus herd had sent pigs that potentially could develop PMWS to the test station. Accordingly PMWS had been diagnosed by necropsies in pigs that developed PMWS were dead at day 55. Pigs without clinical signs of PMWS were grouped as having low (squares, n = 11) or high (triangles, n = 7) maternal immunity to PCV2. Significant differences to other groups are indicated in the figure (p < 0.05 = *, p < 0.01 = **, p < 0.001 = ***).

| Table 4: Mean levels of Serum amyloid A (SAA) in serum (mg per L). |
| Day after arrival | PMWS pigs (n = 4) | Healthy pigs, low level of maternal antibodies (n = 17) | Healthy pigs, high level of maternal antibodies (n = 17) |
|-------------------|------------------|-------------------------------------------------|-------------------------------------------------|
| Day 9             | 424 ± 475        | 148 ± 306                                       | 115 ± 250                                       |
| Day 17            | 24 ± 17          | 27 ± 33                                         | 57 ± 76                                         |
| Day 23            | 111 ± 114        | 134 ± 304                                       | 94 ± 134                                        |
| Day 34            | 39 ± 26          | 57 ± 62 *                                       | 17 ± 8 *                                        |
| Day 43            | 92 ± 120         | 71 ± 109                                        | 31 ± 41                                         |
| Day 55            | -                | 38 ± 75                                         | 42 ± 87                                         |

Mean values within day indicated with *differ (p < 0.05) from each other
All pigs that developed PMWS were dead at day 55. Pigs without clinical signs of PMWS were grouped as having low or high maternal immunity to PCV2. All these pigs were alive at day 55.
pigs from every nucleus herd that had delivered pigs to the test station as previously reported [37]. As clinical signs resembling PMWS significantly less often had been attended in pure bred conventional Hampshire boars (2.8%; n = 497) than in pure bred conventional Yorkshire (8.8%; n = 509) or Landrace pigs (11.3%; n = 655) [37], a genetic difference in resistance to development of the disease between breeds may be indicated. This has also been indicated by others, suggesting a lower resistance towards development PCV2-associated lesions of Landrace pigs [38,39]. However, the station mixed pigs from different sources and also the effect of stressors and pathogen load at the herds of origin should be taken into account. Indeed, there was a variation in the incidence of pigs with clinical signs resembling PMWS within breeds depending on the herd of origin [37].

A higher level of PCV2 genome copies in serum was recorded in pigs that developed PMWS than in pigs that remained healthy. All four PMWS-affected pigs had expressed levels well above log 7 of PCV2 per ml serum. However, serum concentrations above log 7 of PCV2 per ml were also recorded in several pigs that were not denoted as "thin" (11 out of 17 pigs with log 7 of PCV2 per ml) [41], in which makes detection of PCV2 virus in serum unsuitable as a single diagnostic tool to diagnose PMWS. However, as significantly lower peak levels of PCV2 were recorded in pigs with high levels of maternal antibodies, an important role of antibodies to PCV2 in preventing an excessive proliferation of the virus was again indicated [30-33,35].

It has been reported that an unrestrained growth of PCV2 in pigs with low levels of serum antibodies with concurrent infections and/or another stressor are required for development of PMWS [1-3,34,40]. Production of high levels of SAA in pigs can be indicative of acute bacterial infections [41], and SAA has also been reported to be increased in pigs diseased with PCV2 [42]. However, these authors compared the serum levels of several acute phase proteins in pigs of different sources and ages affected by different diseases with that of SPF pigs aged ten weeks, and it cannot be ruled out that the levels of acute phase proteins they reported could have been partly age and herd dependent [42]. Such an effect of age has previously been shown with respect to the acute phase protein haptoglobin [43]. Furthermore, individual serum levels of both pig-MAP and haptoglobin in PCV2 negative pigs could exceed that of equally aged PCV2-positive pigs in the same herd [44]. In the present study, no association between SAA levels and PCV2 viral load was detected. Instead, the concentrations of SAA peaked on day nine after arrival, mirroring the effect of mixing pigs of different origin and thereby exposing them to an unfamiliar flora of microorganisms [45,46]. Accordingly, this peak in SAA concentrations is likely to decay over time due to an adaptation of the immune system to the new environmental flora [46], and acute phase proteins appears to be less valuable as indicators for PMWS.

In conclusion, the higher PVC2 viral load observed in pigs that developed PMWS agrees with suggestions of the importance of a rapid and relevant immune response in preventing PMWS [30-33]. The peak viral load was also seen earlier in pigs that developed PMWS, possibly indicating an impaired immune function in pigs developing PMWS. However, it is also of interest that a majority of the pigs with low maternally derived antibodies to PCV2 did not develop PMWS. This study was carried out in a progeny test station allocating and mixing recently weaned piglets at an early age. Thus, both the age of the pigs in relation to stressors, as well as their age at weaning, may be of importance for the development of PMWS.

Competing interests
The authors declare that they have no competing interests.

Authors' contributions
PW, GBe initiated in the study and designed it in cooperation with CF, FW and GA. IMB and CMJ was responsible for the PCR-analyses and GBII for the iPMI-analysis. PW was head writer of the manuscript with help from the other authors. All authors read and approved the final manuscript.

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