

1 **Evaluation of porcine epidemic diarrhea virus RNA contamination on swine industry**  
2 **transportation vehicles**

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13 **Abstract**

14 Porcine epidemic diarrhea virus (PEDV) is one of the most devastating pathogens of global swine  
15 health and welfare. It is well known that contaminated fomites and vehicle movements play an  
16 important role in farm-to-farm PEDV spread, but the efficacy of cleaning and disinfection (C&D)  
17 protocols on the reduction in dissemination risk via vehicles and trailers remains unclear. This  
18 study used swine industry data to determine how frequently vehicles and trailers were  
19 contaminated with PEDV RNA before and after C&D. Environmental RNA samples were  
20 collected at three eastern North Carolina C&D sites from four different vehicle types: crew trucks,  
21 feed trucks, pigs-to-farm trucks and trailers, and pigs-to-market trucks and trailers. A total of 2,004  
22 samples were collected from truck cabins, trailers, and tires before and after C&D with two  
23 commercial disinfectants at two different concentrations. An in-house RT-qPCR assay was used

24 to detect the presence of PEDV RNA only (not infectivity status). Results suggest that pigs-to-  
25 market trucks hauling live pigs were the most likely to be contaminated with PEDV (82.06% of  
26 trucks tested positive before C&D and 89.55% tested positive after C&D), while feed trucks were  
27 the least likely contaminated (11.73% of trucks testing positive before C&D and 11.25% testing  
28 positive after C&D). Based on PEDV RNA detection, we demonstrated that quaternary ammonium  
29 and glutaraldehyde is a more effective disinfectant compared to advanced hydrogen peroxide in  
30 eliminating detectable PEDV RNA. Results also show that truck cabins are just as contaminated  
31 as the exterior of their vehicles. Based on these results, vehicle biosecurity measures should be  
32 evaluated and modified to prevent the spread of PEDV.

33

34 **Keywords:** swine, PEDV, RNA detection, between farm virus transmission, RT-qPCR, vehicle  
35 cleaning and disinfection, glutaraldehyde, quaternary ammonium

36

## 37 1. Introduction

38 Porcine epidemic diarrhea virus (PEDV) is one of the most devastating pathogens of global swine  
39 health and welfare. This highly contagious enteric pathogen of pigs causes vomiting, diarrhea, and  
40 dehydration in swine of all ages, as well as high mortality in neonatal piglets ([Stevenson et al.,](#)  
41 [2013](#); [Lei et al., 2024](#)). PEDV was introduced into the U.S. swine industry in 2013 when 16 states  
42 reported a total of 218 cases within the first nine weeks ([Stevenson et al., 2013](#); [Mole, 2013](#); [Lowe](#)  
43 [et al., 2014](#); [Lee, 2015](#); [Machado et al., 2019](#)). It is now estimated that roughly 55 - 60% of U.S.  
44 commercial breeding herds have been exposed to the virus at one time or another (AASV, 2013;  
45 [Goede et al., 2015](#); [Kikuti et al., 2022](#)). Moreover, evidence from field veterinarians shows that

46 PEDV remains ranked among the top concerns from veterinary and production standpoints  
47 ([VanderWall and Deen, 2018](#)). Over a decade since its introduction into the U.S., PEDV continues  
48 to be an important issue for the industry due to the economic burden from both direct and indirect  
49 costs associated with the occurrence of an outbreak, which can reach up to USD \$300,000 (based  
50 on the closure of a 700-head breeding herd) ([Weng et al., 2016](#)). As of April 2024, the Morrison  
51 Swine Health Monitoring Project (MSHMP) reported an incidence of 3.8% of PEDV cases in U.S.  
52 sow herds ([MSHMP Chart 1](#)). This has been an improvement since 2015, but the virus still costs  
53 the U.S. swine industry over USD \$50 million per year ([National Hog Farmer, 2024](#)).

54 PEDV can be transmitted directly through the fecal-oral route, as well as via contaminated  
55 fomites ([Stevenson et al., 2013](#); [Bowman et al., 2015](#); [Lei et al., 2024](#); [Houston et al., 2024](#)). It is  
56 also known that vehicle movements play a role in disease spread ([Lee et al., 2019](#); [Lowe et al.,](#)  
57 [2014](#); [Boniotto et al., 2018](#)). However, the importance of pathogen transmission via truck cabins,  
58 live-haul trailers, feed trucks, and crew trucks in the swine industry is under-researched ([Stevenson](#)  
59 [et al., 2013](#); [Houston et al., 2024](#)). Biosecurity measures including decontamination via cleaning  
60 and disinfecting (C&D) protocols are in place to help reduce virus transmission, but studies have  
61 shown that these methods are not always effective ([Boniotto et al., 2018](#); [Houston et al., 2024](#); [Li](#)  
62 [et al., 2020](#)). Therefore, understanding how vehicle movement and decontamination affect disease  
63 outbreaks is crucial to decreasing the transmission of PEDV ([Galvis and Machado, 2024](#)).

64 The objective of this study was to use swine industry data to determine how frequently  
65 swine production vehicles were contaminated with PEDV RNA before and after C&D following  
66 the industry SOP which included washing (including soaking and removal of organic material)  
67 and disinfection. While addressing this question, the effectiveness of selected commercial  
68 disinfectants at eliminating detectable PEDV RNA, as well as how PEDV RNA contamination

69 varied across truck classifications and sampling locations (e.g., interior of truck cabin vs. trailer or  
70 tire), were evaluated. Notably, this current study did not assess whether PEDV RNA detected on  
71 swine vehicles was capable of causing infection.

72

## 73 **2. Materials and Methods**

74

75 Figure 1 provides a schematic of the key steps in this study's experimental design, which are  
76 described in detail in the following subsections.

### 77 *2.1 Selection of truck types and disinfectants*

78 A total of four different truck types used by the participating commercial swine company were  
79 sampled in this study: 1) pigs-to-farm trucks, which transport live pigs between farms (trucks are  
80 classified depending on whether they are coming from farms that were experiencing a PEDV  
81 outbreak [referred to as “PED+ trucks”] and trucks coming from farms that were not experiencing  
82 a PEDV outbreak [referred to as “PED- trucks”]), 2) pigs-to-market trucks, which transport pigs  
83 to markets (i.e. slaughterhouse, packing plants; referred to as “market trucks”), 3) feed trucks,  
84 which carry feed from the feed mill to farms, and 4) crew trucks, which are used to move personnel  
85 performing a wide range of farm tasks (e.g. vaccination, power washing at closeouts, pig loading,  
86 and unloading). Pigs-to-farm and market trailers were power washed by the swine company  
87 personnel to remove debris and organic material, then disinfectant was sprayed onto to the inside  
88 and outside of the trailers. The lower portion of feed trucks (e.g. tires and surfaces close to the  
89 ground) are sprayed with disinfectant in between loads and crew trucks are washed using a  
90 company car wash after daily shifts. The sample size for each truck type was determined using a  
91 power analysis with a 95% confidence interval assuming 30% of the selected 398 trucks were

92 positive for PEDV (*pers comm*). Trucks were sampled at three specific C&D stations located in  
93 eastern North Carolina, U.S. selected due to: a) locality close to a high hog density area, and b)  
94 surrounding herds have a history of ongoing PEDV. All C&D stations used a unidirectional flow  
95 pattern and recycled water for washing. For confidentiality, stations will be numbered/identified  
96 as 1, 2, and 3 (Table 1).

97

98 **Table 1:** Description of truck washes included in this study and corresponding truck types.

C&D station	Number of bays for C&D	Active disinfectant ingredient	Disinfectant dilutions studied	Truck types (month[s] of sampling)
1	1*	Accelerated Hydrogen Peroxide	1:32 and 1:40	Pigs-to-market (Nov-Feb); Pigs-to-farm (PED-) (Mar-May)
2	1	Quaternary Ammonium and Glutaraldehyde	1:100 and 1:128	Pigs-to-farm (PED+) (Nov-May)
3	3	Quaternary Ammonium and Glutaraldehyde	1:100 and 1:128	Pigs-to-farm (PED-) (Nov-Feb); Pigs-to-market (Mar-May); feed.

99 \* denotes there are three other bays at this station that utilized Quaternary Ammonium and Glutaraldehyde, but  
100 were not included in this study.  
101

102 The rationale for disinfectant choice was based on those currently used by the swine industry and  
103 field veterinarians, and what was available at the three selected C&D stations. The production  
104 company running the C&D stations typically disinfects using quaternary ammonium and  
105 glutaraldehyde (QAG) at a 1:256 dilution and advanced hydrogen peroxide (AHP) at a 1:32  
106 dilution. However, previous studies have compared different agricultural disinfectants and found  
107 discordant results regarding the efficacy of their ability to lower the concentration of PEDV RNA  
108 ([Bowman et al., 2015](#); [Holtkamp et al., 2016](#); [Houston et al., 2024](#)). Given this, we opted to test  
109 both disinfectants at two concentrations each. All three C&D stations rotated their disinfectant  
110 concentration between a higher and lower concentration for each collection period (Table 1). For  
111 QAG, the higher concentration was diluted at 1:100 and the lower concentration was diluted at

112 1:128. For AHP, the higher concentration was diluted at 1:32 and the lower was diluted at 1:40.  
113 While these were the stated concentrations, it was not possible to internally confirm a) if the  
114 working disinfectant solutions were prepared to these concentrations by truck wash workers  
115 (however, C&D station personnel verified the disinfectant concentration after preparing according  
116 to their validated procedure), and b) the volume of disinfectant administered to each truck. To  
117 measure the working concentrations of both disinfectants, test strips (QAG– Cid Lines, Ypres,  
118 Belgium; AHP– Virox Technologies Inc., Oakville, ON, Canada) were used to test the  
119 concentration of disinfectant after application to the back of the trailers of PED- pigs-to-farm and  
120 pigs-to market trucks.

121 The disinfectants in use at each of the C&D stations were as follows (Table 1): *Station 1* -  
122 QAG used in three bays (not sampled in this study) and AHP in the fourth bay per our request;  
123 *Station 2* - QAG in single bay; *Station 3* - QAG in all three bays, one of which was dedicated to  
124 feed truck disinfection. In March, the production company opted to move PED- pigs-to-farm trucks  
125 to C&D station 1 and pigs-to-market trucks to C&D station 3. Crew trucks were washed at the end  
126 of shifts at a drive-through car wash operated by the production company and therefore not  
127 subjected to the included disinfectants.

128

## 129 2.2 Sample Collection

130 For each of the four truck types, both an interior cabin sample and exterior truck/trailer sample  
131 were collected per vehicle. The exterior sample was collected at either the trailer or the tire (for  
132 vehicle types that did not carry live animals). Specific collection sites for each truck type were as  
133 follows: 1) *pigs-to-farm and market trucks*, the cabin floorboard underneath the brake pedal and  
134 the loading zone of the trailer and surrounding hinges; 2) *feed trucks*, the cabin floorboard

135 underneath the brake pedal and one tire from the feed-holding trailer; and 3) *crew trucks*, the cabin  
136 floorboard underneath the brake pedal and one tire from the truck (Figure 1). When collecting a  
137 sample, a Puritan Hydraflock swab (Puritan Medical Products, Guilford, ME, USA) was moistened  
138 by dipping into a 1.5 mL collection tube containing 500  $\mu$ L DNA/RNA Shield (Zymo Research  
139 Corporation, Irvine, CA, USA) and then subsequently wiped and rolled across approximately a  
140 20-by-20 cm area before placing the swab head back into the collection tube. The first truck for  
141 each truck type from each bi-weekly collection period was swabbed in triplicate; two samples  
142 served as collection duplicates and a third sample was used as a “processing positive control”  
143 (described below). Samples were kept on ice during collection and stored at  $-80^{\circ}\text{C}$  upon arrival at  
144 the laboratory. Samples were collected bi-weekly from 27 Nov 2023 to 31 May 2024 to include  
145 peak PEDV season which has a higher incidence from late fall to spring.

146 Additionally, to act as a positive control, samples were collected via swab from various  
147 locations at a farm confirmed to be PEDV+ (e.g. pen wall and floor, boot sole, and feed trough)  
148 and stored in 500  $\mu$ L of DNA/RNA Shield. Upon arrival at the laboratory, samples were frozen at  
149  $-80^{\circ}\text{C}$ .

150

### 151 *2.3 Detection of PEDV*

152 Reverse transcription-quantitative PCR (RT-qPCR) is a useful molecular biology method that  
153 allows researchers to detect specific sequences of RNA in a sample using custom nucleic acid  
154 primers that bind to and amplify a target sequence. Once amplified to a detectable level, a  
155 fluorescent dye can be measured which correlates to the quantity of the amplified RNA of the  
156 target sequence in the sample. Commercial RT-qPCR assays are available for the detection of  
157 many swine viruses, but are often expensive and not tailored to a specific sample type. This study

158 uses a custom in-house RT-qPCR assay that was designed to detect PEDV RNA specifically from  
159 environmental vehicle samples. Sections 2.3.1 and 2.3.2 detail the specific methods and reagents  
160 used in this assay.

161

### 162 *2.3.1 RNA Isolation*

163 The positive control samples collected from the PED+ farm were thawed at room temperature and  
164 isolated using the Quick-DNA/RNA Viral Kit (Zymo Research Corporation) according to the  
165 manufacturer's instructions. Eluates across multiple RNA isolations were pooled, separated into  
166 500  $\mu$ L aliquots, and stored at  $-80^{\circ}\text{C}$  until use.

167 The environmental samples collected from trucks were thawed to room temperature and  
168 swabs were placed in spin baskets set inside the collection tube (ChromeTech of Wisconsin Inc,  
169 Franklin, WI, USA) and centrifuged at 12,000 rpm for two minutes. This step was completed to a)  
170 ensure complete recovery of all liquid retained in the swab head, and b) facilitate the removal of  
171 large debris and particulates adhered to the swab. A total of 375  $\mu$ L of supernatant was transferred  
172 into a 1.5 mL clean tube. For the triplicate environmental samples collected to serve as "processing  
173 positive controls", 10  $\mu$ L of the pooled RNA eluate from the PED+ farm samples was spiked into  
174 the supernatant. This was completed to ensure that both the downstream RNA isolation processing  
175 and environmental materials associated with the samples did not impact PEDV RNA detection.  
176 The sample supernatants were frozen at  $-80^{\circ}\text{C}$  until RNA isolation could be completed in batches  
177 of 96.

178 The *Quick-DNA/RNA Viral 96 Kit* (Zymo Research Corporation) was used to isolate RNA  
179 according to the manufacturer's instructions, with the following modifications: 1) Viral  
180 DNA/RNA Buffer was left on the sample for 10 minutes to allow for an adequate lysis period for



181 environmental samples, 2) centrifuge steps were done at 3,250 rpm for 10 minutes, 3) a five-minute  
182 dry spin at 3,250 rpm was done before RNA elution, and 4) elution volume was increased to 18  
183  $\mu\text{L}$ . DNA digestion was performed during the isolation as recommended by the manufacturer to  
184 ensure pure RNA. Eluates were kept in 96-well plate format, sealed with a foil seal, and frozen at  
185  $-80^{\circ}\text{C}$  until further use.

186 To serve as a “RT-qPCR positive control,” fresh piglet intestines from PEDV+ diagnostic  
187 pathology cases were clipped to make six 50 mg sections of ileum. RNA was isolated from each  
188 ileum clipping using the *Quick-RNA* MiniPrep Plus Kit (Zymo Research Corporation) according  
189 to the manufacturer’s directions with the following modifications: 1) tissue clippings were each  
190 submerged in 600  $\mu\text{L}$  of PBS without calcium and magnesium and homogenized using 2.0 mm  
191 lysis beads in a high-speed bead-beater for three rounds of 20-second homogenization and one  
192 minute cooling on ice, 2) 60  $\mu\text{L}$  PK buffer and 30  $\mu\text{L}$  Proteinase K were added to each  
193 homogenized sample, allowed to incubate at room temperature for 30 minutes, and were  
194 centrifuged at 8,000 rpm five minutes to pellet any remaining tissue, and 3) RNA was eluted in  
195 two separate 50  $\mu\text{L}$  elutions. The first and second eluates from all six samples were pooled  
196 respectively, divided into 20  $\mu\text{L}$  aliquots, and stored at  $-80^{\circ}\text{C}$  until further use.

### 197 2.3.2 *RT-qPCR*

198 An 87 bp fragment of the PEDV N-gene was amplified and measured by RT-qPCR using the Go-  
199 Taq Enviro RT-qPCR System (Promega Madison, WI, USA). Each 20  $\mu\text{L}$  reaction consisted of 10  
200  $\mu\text{L}$  of 2X Go-Taq Enviro Master Mix, 0.4  $\mu\text{L}$  Go-Script Enzyme Mix, 0.8  $\mu\text{L}$  of 10  $\mu\text{M}$  forward  
201 primer (5’ - GAATTCCCAAGGGCGAAAAT - 3’) ([Ortiz et al., 2023](#)), 0.8  $\mu\text{L}$  of 10  $\mu\text{M}$  reverse  
202 primer (5’ - TTTTCGACAAATTCCGCATCT - 3’) ([Ortiz et al., 2023](#)), 0.2  $\mu\text{L}$  of 10  $\mu\text{M}$  probe  
203 (5’ - FAM-CGTAGCAGCTTGCTTCGGACCCA-BHQ1 - 3’) ([Ortiz et al., 2023](#)), 5.8  $\mu\text{L}$  of

204 nuclease-free water, and 2  $\mu$ L of isolated RNA. Additionally, every 96-well-plate of RT-qPCR  
205 included two controls: 1) a non-template control consisting of 2  $\mu$ L of DNA/RNA free water, and  
206 2) a “RT-qPCR positive control” consisting of 2  $\mu$ L of RNA isolated from the intestines of a  
207 PEDV+ piglet. All RT-qPCR reactions were run on the QuantStudio 5 (Applied Biosystems,  
208 Waltham, MA, USA) with the following cycling conditions: 45°C for 15 minutes, 95°C for 2  
209 minutes, and 40 cycles of 95°C for 15 seconds and 60°C for 1 minute. RT-qPCR results are  
210 reported as quantification cycle (Cq) values. When interpreting the resulting Cq value, an inverse  
211 relationship applies: a lower Cq value indicates a higher starting quantity of PEDV RNA in the  
212 sample, whereas a higher Cq indicates a lower starting quantity of PEDV RNA.

213 To confirm the accuracy of the in-house PEDV RT-qPCR assay, 10% of samples (n=254)  
214 across different truck types, collection periods, and collection sites (e.g. back-of-trailer, tire, and  
215 cabin) were additionally tested for PEDV, TGEV, and PDCoV using a commercial triplex kit  
216 (Indical Bioscience, Leipzig, Germany). Each 25  $\mu$ L reaction consisted of 18  $\mu$ L virotype® Mix  
217 +IC(TAMRA)-RNA, 2  $\mu$ L virotype® PEDV/TGEV/PDCoV Primers/Probes, and 5  $\mu$ L isolated  
218 RNA and were run on the QuantStudio 5 (Applied Biosystems) under the following cycling  
219 conditions: 50°C for 10 minutes, 95°C for 2 minutes, 40 cycles of 95°C for 5 seconds and 60°C  
220 for 30 seconds. Every 96-well-plate of RT-qPCR with the commercial triplex included two  
221 controls: 1) a non-template control consisting of 5  $\mu$ L of DNA/RNA free water, and 2) a positive  
222 control consisting of 5  $\mu$ L of the Indical virotype PEDV/TGEV/PDCoV positive control.

223

#### 224 *2.4 Statistical Analysis*

225 While the commercial (Indical) assay has clear interpretation guidelines (Cq  $\geq$  35 should be  
226 considered negative; Cq  $<$  35 should be considered positive), we did not assume those guidelines

227 would be appropriate for the results generated with the in-house assay. Thus, to determine an  
228 appropriate cutoff value for determining PEDV RNA positive and negative samples with the in-  
229 house assay, two separate approaches were carried out using the Cq values from the samples  
230 processed with both the in-house and commercial assay (n=254). Firstly, using data from the  
231 custom assay, the receiver operating characteristic (ROC) curve was plotted and the resulting area  
232 under the curve (AUC) values were determined for each potential cutoff value (every integer from  
233 25 to 36) using the commercial assay results with a cutoff of 35 as the reference. Secondly, the  
234 data was assumed to fit a Weibull distribution using a mixed models procedure. A No-U-turn  
235 sampler (NUTS) was used to estimate the posterior distribution of Cq values for positive samples  
236 to obtain the 95% credible interval. An agreement between these two methods was used to  
237 determine the cutoff value.

238 Using the statistically determined Cq threshold of 32 established for the in-house assay,  
239 study samples (n=2,004) were either deemed positive ( $\leq 32$ ) or negative ( $> 32$ ). Statistical analyses  
240 were completed using both positive/negative status and Cq value to assess the impact of study  
241 variables including truck type, disinfection status (e.g. before or after disinfection), disinfectant  
242 used, swab collection site, and month collected on the detection of PEDV RNA. For sample  
243 collection duplicates (n=190), Cq values were averaged. Any sample with a Cq $>40$  was omitted  
244 from the analysis (Supplementary Table 2). Summary statistics were calculated for the  
245 positive/negative status of samples. For Cq value comparisons, the Shapiro-Wilk test was used to  
246 determine normality. For normal data sets, pooled t-tests and Tukey–Kramer HSD tests were used  
247 with an  $\alpha = 0.05$ . For non-normal data sets, the Wilcoxon signed-rank test and Dunn tests with a  
248 Bonferroni correction were performed with an  $\alpha = 0.05$  to determine data significance. Details on  
249 the tests completed for each comparison are given in Supplemental Table 1.

250

### 251 3. Results

#### 252 3.1 Cq cutoff

253 The Cq cutoff value for determining positive and negative samples needed to be appropriate for  
254 the sensitivity level of the in-house RT-qPCR assay developed for this study. The maximum AUC  
255 of 0.8077 was found when using the cutoff Cq value of 33. The second-best AUC of 0.7974 was  
256 observed using a cutoff Cq value of 32. Both AUC values are considered “acceptable” ([Nahm et  
257 al., 2022](#)). The 95% credible interval for Cq values of positive samples determined using the NUTS  
258 method was 31.5-32.8. Based on the combined results from both methods, a Cq of  $\leq 32$  was selected  
259 as the optimal cutoff for classifying a sample as PEDV RNA positive using the custom RT-qPCR  
260 assay (Table 2).

261

#### 262 3.2 Prevalence of PEDV RNA on trucks

263 A total of 387 individual cabins, trailers, and tires (587 total cabins, trailers, and tires sampled due  
264 to repeat visits) were included in the study. The aim was to sample 30% of the production company  
265 fleet’s trucks for each truck type; based on availability, more than the 30% of each truck type was  
266 sampled for all except feed trucks (27%). The number of unique trucks included in the study per  
267 sample type is as follows: crew, 14 cabins/tires; feed, 52 trailers and 63 cabins; pigs-to-market, 72  
268 trailers and 48 cabins; pigs-to-farm, 80 trailers and 44 cabins (PED+ and PEV- combined). From  
269 these trucks, 2,004 environmental samples were collected and processed for PEDV RNA detection  
270 using RT-qPCR. A total of 92.07% of the “processing positive control” samples were deemed  
271 PEDV RNA positive, verifying that the sampling approach, storage, and RNA isolation procedures  
272 were appropriate for isolating PEDV RNA. Additionally, 190 samples were collection sample

273 duplicates; Cq values were averaged before further analysis. Of the remaining 1,650 samples,  
274 1,455 samples had detectable PEDV RNA (Cq < 40, 88.18%) and 625 were deemed positive for  
275 PEDV RNA (Cq ≤ 32, 37.88%).

276 Market trucks had the lowest median Cq value as well as the shortest median wash time of  
277 15 minutes (*pers. obs.*) and the highest percentage of positive results for samples collected before  
278 disinfection (Figure 2, Table 2, and Table 3). The highest Cq values and lowest percentage of  
279 positive results were observed from feed trucks (Figure 2, Table 2, and Table 3). PED+ and PED-  
280 pigs-to-farm truck median Cq values were within 1.40 cycles of each other when comparing  
281 samples collected before disinfection, as well as comparing samples disinfected with QAG (Table  
282 3). PED+ pigs-to-farm trucks had a median wash time of 60 minutes, while 45 minutes was the  
283 median wash time for PED- pigs-to-farm trucks (*pers. obs.*).

284 **Table 2.** Percentages of samples positive for PEDV RNA based on the in-house RT-qPCR results and a Cq cutoff value of  $\leq 32$

TRUCK TYPE	DISINFECTION STATUS						
	<i>Before Disinfection</i>			<i>After Disinfection</i>			
	<i>Location of sampling</i>			<i>Advanced Hydrogen Peroxide</i>		<i>Quaternary Ammonium/Glutaraldehyde</i>	
	<b>Cabin</b>	<b>Tire</b>	<b>Back-of-Trailer</b>	<b>1:40</b>	<b>1:32</b>	<b>1:128</b>	<b>1:100</b>
<b>PED+</b>	50.56% (45/89)	NA	48.86% (43/88)	NA	NA	25.00% (7/28)	36.96% (17/46)
<b>PED-</b>	47.25% (43/91)	NA	41.57% (37/89)	41.67% (5/12)	43.75% (7/16)	23.81% (5/21)	16.67% (3/18)
<b>Feed</b>	16.32% (31/190)	7.60% (13/171)	NA	NA	NA	26.92% (23/78)	3.19% (3/94)
<b>Market</b>	87.29% (103/118)	NA	79.17% (95/120)	100% (26/26)	97.22% (35/36)	70.37% (19/27)	84.84% (28/33)
<b>Crew</b>	67.65% (23/34)	30.00% (9/30)	NA	NA	NA	NA	NA

285 \*NA- Data not available as such samples were not collected from that vehicle type

286 \*Values are reported as follows: Percentage of PEDV RNA positive samples (number of positive samples/total number of samples).

287 \*Feed truck samples collected after disinfection were only collected from the tire. PED+, PED-, and market truck samples after disinfection were  
 288 only collected from the back of the trailer.

289

290 **Table 3.** Summary statistics for Cq values obtained from the in-house RT-qPCR results based on truck type, disinfection status, sampling  
 291 location, and disinfectant.

TRUCK TYPE	DISINFECTION STATUS						
	<i>Before Disinfection</i>			<i>After Disinfection</i>			
	<i>Location of sampling</i>			<i>Advanced Hydrogen Peroxide</i>		<i>Quaternary Ammonium/Glutaraldehyde</i>	
	<b>Cabin</b>	<b>Tire</b>	<b>Back-of-Trailer</b>	<b>1:40</b>	<b>1:32</b>	<b>1:128</b>	<b>1:100</b>
<b>PED+</b>	31.33 (23.58-37.82)	NA	30.46 (15.63-37.21)	NA	NA	32.14 (16.51-37.67)	33.45 (22.18-36.90)
<b>PED-</b>	31.69 (17.65-38.86)	NA	31.72 (21.59-37.03)	31.40 (15.87-37.25)	30.85 (16.07-36.99)	33.52 (27.00-37.49)	34.55 (28.94-39.58)
<b>Feed</b>	33.82 (17.39-39.91)	34.32 (31.21-37.52)	NA	NA	NA	33.41 (14.89-39.45)	34.89 (30.68-38.45)
<b>Market</b>	29.21 (23.86-35.19)	NA	28.83 (21.22-36.19)	24.49 (14.55-29.56)	26.05 (21.30-33.11)	29.78 (22.13-36.33)	29.43 (19.51-35.86)
<b>Crew</b>	29.08 (18.80-36.31)	32.92 (18.05-36.51)	NA	NA	NA	NA	NA

292 \*NA- Data not available as such samples were not collected from that vehicle type

293 \*Cq values are reported as follows: Median (minimum value – maximum value).

294 \*Feed truck samples collected after disinfection were only collected from the tire. PED+, PED-, and market truck samples after disinfection were  
 295 only collected from the back of the trailer

296        *3.3 Effect of disinfectants on PEDV RNA concentration*

297        When considering all back-of-trailer and tire samples and using Cq value as an assessment of  
298        cleanliness (i.e., more or less detectable PEDV RNA), neither disinfectant at the high nor low  
299        concentration left the trailers/tires significantly cleaner after disinfection (Figure 3, Supplementary  
300        Table 1). This is also observed when considering back-of-trailer and tire samples separately  
301        (Supplementary Figure 1 and Supplementary Figure 2, respectively). PED- pigs-to-farm trailers  
302        and pigs-to-market trailers were subjected to all four disinfection options; only QAG 1:128 left the  
303        PED- pigs-to-farm trailers significantly cleaner after disinfection (Figure 4A,  $p=0.024$ ) while  
304        neither disinfectant yielded significantly cleaner market trailers (Figure 4B,  $p>0.05$ ). Moreover,  
305        AHP 1:32 left market truck trailers significantly more contaminated with PEDV RNA after  
306        disinfection (Figure 4B,  $p=0.0197$ ). When considering after disinfection samples from each truck  
307        type, the Cq values observed from higher and lower disinfectant concentrations were never  
308        significantly different from each other (Figure 2, Figure 4, Supplementary Table 1). Additionally,  
309        when looking at market trailers after disinfection, the Cq values observed from trailers disinfected  
310        with AHP were significantly lower than what was observed for trailers disinfected with QAG  
311        (Supplementary Table 1).

312        Test strips were used to determine the concentration of residual disinfectants after they  
313        were applied PED- and market trucks. Of the 49 trucks disinfected with AHP 1:32 and evaluated  
314        with test strips, seven had a reading of 1:32 or higher. Of the 33 trucks disinfected with AHP 1:40  
315        and tested, four had a reading of 1:32 or higher. Of the 28 trucks disinfected with QAG 1:100 and  
316        tested, five had a reading of 1:100. Lastly, of the 44 trucks disinfected with QAG 1:128, 12 had a  
317        reading of 0.5 or higher (equivalent to 1:200). In the majority of cases, the disinfectant remaining

318 on the truck was lower than the optimal concentration for effective disinfection. Notably, all of  
319 our test samples were collected from the back of the trailer.

320

### 321 *3.4 Impact of collection site on PEDV RNA detection*

322 Three different collection sites were sampled for this study: 1) back-of-trailer for the live-animal  
323 vehicles, 2) tires for the crew and feed trucks, and 3) the truck cabins for all vehicle types (Figure  
324 1). Market trucks were far more contaminated with PEDV RNA with the lowest median Cq values  
325 for the cabin and the back-of-trailer samples while crew truck tires and cabins had lower Cq values  
326 compared to feed trucks (Table 3, Figure 5A). When considering all pre-disinfection data, each of  
327 the three collection sites were significantly different from each other; the tire samples had a  
328 significantly higher Cq values than the cabin samples and the back-of-trailer samples had a  
329 significantly lower Cq values than the cabins (Figure 5B,  $p < 0.0001$ ). For each truck type, the  
330 cabins had a higher percentage of samples deemed PEDV RNA positive compared to the  
331 corresponding back-of-trailer or tire samples (Table 2). Additionally, for all live-haul vehicles,  
332 there was not a significant difference between the Cq values of the cabin and the back-of-trailer  
333 samples (Figure 5A, Supplementary Table 1).

334

### 335 *3.5 Impact of seasonality*

336 Samples were collected during 14 bi-weekly collection periods between November 2023 and May  
337 2024. Collection weeks one through seven took place during the winter period (November 2023-  
338 February-2024) and collection weeks eight through 14 were during the spring period (March 2024-  
339 May-2024). Four (57%) of the winter collection weeks had higher median Cq values after  
340 disinfection, whereas only three (42%) of the spring collection weeks had this result. However,



341 the median Cq values before disinfection for the winter collections were typically lower than the  
342 spring collections (Figure 6). Although Cq values for back-of-trailer and tire samples collected  
343 after disinfection were not significantly different between seasons, the Cq values observed for  
344 back-of-trailer and tire samples collected before disinfection were significantly lower in winter  
345 than in spring ( $p=0.016$ ).

346

#### 347 **4. Discussion**

348 By collecting samples from swine industry vehicles and trailers and utilizing an in-house RT-  
349 qPCR assay to detect PEDV RNA, we were able to examine the contamination level of vehicles  
350 with PEDV RNA before and after disinfection across different vehicle types when cleaned with  
351 different types and concentrations of disinfectants. The in-house assay, which is a more cost-  
352 effective alternative to commercial assays, was specifically optimized for use with environmental  
353 vehicle samples, which contain less genetic material compared to samples taken directly from  
354 organisms having clinical viral symptoms. We demonstrated that market trucks were a) the most  
355 contaminated with PEDV RNA, b) the most likely truck type to still be physically dirty after  
356 washing (*pers. obs.*), and c) washed for the shortest amount of time (*pers. obs.*) (Figure 2). This is  
357 most likely because these vehicles are only carrying pigs who are exiting the swine production  
358 system (e.g. the slaughterhouse) and no longer bear the risk of carrying PEDV to sow farms (e.g.  
359 [Mannion et al., 2008](#); [Galvis and Machado, 2024](#)). Pigs-to-farm trailers, which carry pigs from all  
360 production phases before being sent to the market, were more effectively washed and were  
361 significantly less likely to be contaminated with PEDV RNA compared to market trucks (Table 3,  
362 Figure 2; PED+ pigs-to-farm,  $p=0.0006$ , PED- pigs-to-farm,  $p<0.0001$ ) ([Galvis and Machado,](#)  
363 [2024](#)). Pigs-to-farm trucks from PED+ and PED- farms were never significantly different from

364 each other, despite being deliberately separated (e.g. washed at different C&D stations, driven by  
365 separate personnel, and sent to only farms with the same PEDV status) (Table 3, Supplementary  
366 Table 1). RNA can persist in environments which facilitate the stability of nucleic acids, such as  
367 in the presence of organic material, so it is plausible that trucks leaving PED- farms are testing  
368 positive due to residual PEDV RNA from previous trips. The most notable discrepancy between  
369 the PED+ and PED- pigs-to-farm trucks was that for PED+ farm trucks, 36.96% of the back-of-  
370 trailer samples disinfected with QAG 1:100 were PEDV RNA positive while only 16.67% of the  
371 back-of-trailer samples disinfected with QAG 1:100 from PED- farms were positive (Table 3,  
372  $p=0.196$ ). However, when comparing these samples disinfected with QAG 1:128, the percentages  
373 of PEDV RNA positive samples become much more comparable (Table 2,  $p=1.000$ ).

374         Given there was not a significant difference between the cabin and back-of-trailer pre-  
375 disinfection samples for all three truck types, it is likely that contaminants (possibly feces) are  
376 moving through fomites (e.g. boots, equipment) between the interior and exterior of the vehicles  
377 (Figure 5A, Supplementary Table 1). Since the crew and feed truck cabin samples were  
378 significantly more contaminated with PEDV RNA than the respective tire samples in this study,  
379 cabins could likely benefit from regular C&D similar to the exterior of the vehicles.

380         Based on PED- pigs-to-farm and market trucks, which were the only truck types disinfected  
381 with both disinfectants, QAG was the more effective disinfectant compared to AHP (Figure 4).  
382 QAG is on the approved list of disinfectants for decontaminating PEDV and has been shown to  
383 disrupt infectious PEDV in treated samples, but not PEDV RNA ([EQSP report 2015](#); [Bowman et](#)  
384 [al., 2015](#)). QAG was able to lower the percentage of PEDV RNA positive samples from before  
385 disinfection to after disinfection in 75% of the scenarios it was used, while AHP was not effective  
386 in this manner for either of the truck types it was used with (Table 3). Disinfectants were prepared

387 by production company staff at the beginning of every day and subsequently tested prior to use;  
388 our team used test-strips to evaluate the disinfectant concentration after vehicle application.  
389 Notably, the test-strip concentrations varied greatly and were typically low compared to the desired  
390 concentration. Resulting C<sub>q</sub> values were rarely significantly different from each other; only AHP  
391 1:32 saw significant differences in C<sub>q</sub> values when comparing across concentration readings  
392 (readings of 0 and 1:16, p=0.0083; all other pairs non-significant). The unreliable and low readings  
393 may be the consequence of applying disinfectant directly after washing. In such a scenario, the  
394 leftover water from washing is still running off the trailer and pooling on the surfaces during the  
395 disinfectant application, thus diluting the disinfectant to undetectable levels. The ideal solution for  
396 this would be to allow the truck to dry before applying the disinfectant, then allowing it to make  
397 contact for the manufacturer's recommended time; however, these may not be feasible due to  
398 company time constraints and the trailer's vertical surfaces (which the disinfectant cannot soak on  
399 [*pers. comm.*]). A possible alternative could be to apply the disinfectant at a high enough  
400 concentration that once it mixes with the residual water, it will dilute to a level that will effectively  
401 kill viral pathogens ([EQSP report 2015](#)). However, this needs to be examined more thoroughly as  
402 previous research rarely considers the concentration of the disinfectant after it is applied to a  
403 washed vehicle ([Bowman et al., 2015](#); [Boniotti et al., 2018](#)).

404 Even though our feed trucks were relatively free of RNA contamination compared to the  
405 other truck types, research has shown that the only surfaces in a feed mill that had detectable RNA  
406 from porcine enteric viruses (including PEDV) were from the feed delivery system ([Elijah et al.,](#)  
407 [2021](#)). Previous studies have also shown that similar to our findings, feed delivery truck cabins  
408 were likely to be contaminated ([Elijah et al., 2021](#); [Houston et al., 2024](#); [Greiner, 2016](#)). Although  
409 we could not evaluate whether the drivers for any of the trucks were exiting their vehicles at their

410 destinations (e.g. the slaughterhouse, farm, feed mill), it has been shown that when drivers leave  
411 their cabin (or when other personnel step onto the trailers) the risk of contamination increases  
412 ([Lowe et al., 2014](#); [Greiner, 2016](#); [Houston et al., 2024](#); [Elijah et al., 2021](#)). This could explain  
413 why crew truck cabins were significantly more contaminated with PEDV RNA than the tires from  
414 the same vehicle, as it is very likely crew members are entering a contaminated area at their  
415 destination and then tracking PEDV RNA into their cabin when leaving.

416         Despite this study's attempt to maintain consistency by selecting C&D stations operated  
417 by the same production company, truck C&D was performed by company personnel and therefore  
418 could not be validated across sites; wash times, trailer cleanliness after disinfection, amount of  
419 disinfectant applied, and thoroughness of disinfectant application all varied between stations and  
420 between station personnel. Additionally, the recommended contact time for disinfectants could not  
421 always be followed, especially for vertical surfaces. Companies which manufacture QAG  
422 recommend a contact time of 5-12 minutes; however, due to the nature of the C&D stations, post-  
423 disinfection samples were often taken less than five minutes after the disinfectant was applied.  
424 Another limitation, concerning RT-qPCR results, was that not all of the "processing positive  
425 control" samples resulted in C<sub>q</sub> values  $\leq 32$  (which would be classified as PEDV RNA positive).  
426 While we would ideally want this to be 100%, these "processing positive controls" still confirmed  
427 that the storage solution (DNA/RNA shield; Zymo Research), the RNA isolation kit, and the in-  
428 house RT-qPCR were working effectively. Nonetheless, the main limitation of this study is that  
429 detecting PEDV RNA using RT-qPCR does not necessarily mean that there is an infectious virus  
430 present. Commercial disinfectants are designed to target viral capsids which often leave the RNA  
431 intact ([Bowman et al., 2015](#)). To study infectivity, the gold-standard approaches include growing  
432 the virus in cell culture or an animal model (bioassay) ([Puente et al., 2020](#)). Recent research

433 suggests that viability RT-qPCR assays can determine if there is viable virus in a sample ([Puente](#)  
434 [et al., 2020](#); [Balestreri et al., 2024](#)). However, these methods are unfeasible for pork producers due  
435 to time, validity, and material restrictions, especially during large outbreaks ([Bowman et al., 2015](#)).  
436 RT-qPCR allows the industry to determine if PEDV should be an immediate concern for their  
437 herds and to quickly address the outbreak.

438

## 439 **5. Conclusion**

440 The aim of this study was to determine how frequently industry vehicles are contaminated with  
441 PEDV RNA before and after C&D. We determined that pigs-to-market trucks are the most likely  
442 to be contaminated overall and that pigs-to-farm trucks from PED- farms have similar PEDV RNA  
443 loads to pigs-to-farm trucks from PED+ farms. We also identified that the cabins of all five truck  
444 types are not significantly cleaner than the exterior of the trucks. Based on these results, vehicle  
445 biosecurity measures involving the disinfection of truck interiors, as well as market truck  
446 disinfection as a whole, should be evaluated to prevent the spread of PEDV as trucks go back to  
447 farms after washing and as personnel exit and enter their cabins during vehicle visits. Additionally,  
448 this study identified QAG to be a more effective disinfectant than advanced hydrogen peroxide for  
449 decreasing PEDV RNA contamination of swine industry vehicles.

450

## 451 **Conflict of interest statement**

452 All authors confirm that there are no conflicts of interest to declare.

453

## 454 **Ethical statement**

455 The authors confirm the journal's ethical policies, as noted on the journal's author guidelines  
456 page. Ethics permits were unnecessary since this work did not involve animal sampling or  
457 questionnaire data collection by the researchers.

458

#### 459 **CRedit authorship contribution statement**

460 TBP: Data curation, formal analysis, investigation, methodology, project administration,  
461 validation, visualization, writing – original draft.

462 KAM: Conceptualization, data curation, funding acquisition, methodology, project  
463 administration, resources, supervision, writing – original draft.

464 GM: Conceptualization, data curation, funding acquisition, methodology, supervision,  
465 visualization, writing – review & editing.

466 MR: Supervision, writing – review & editing.

467 BSD: Investigation, visualization.

468 JBF: Conceptualization, data curation, funding acquisition, investigation, methodology, project  
469 administration, resources, supervision, writing – original draft.

470

#### 471 **Data availability statement**

472 The data supporting this study's findings are not publicly available and are protected by  
473 confidential agreements; therefore, they are not available.

474

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491

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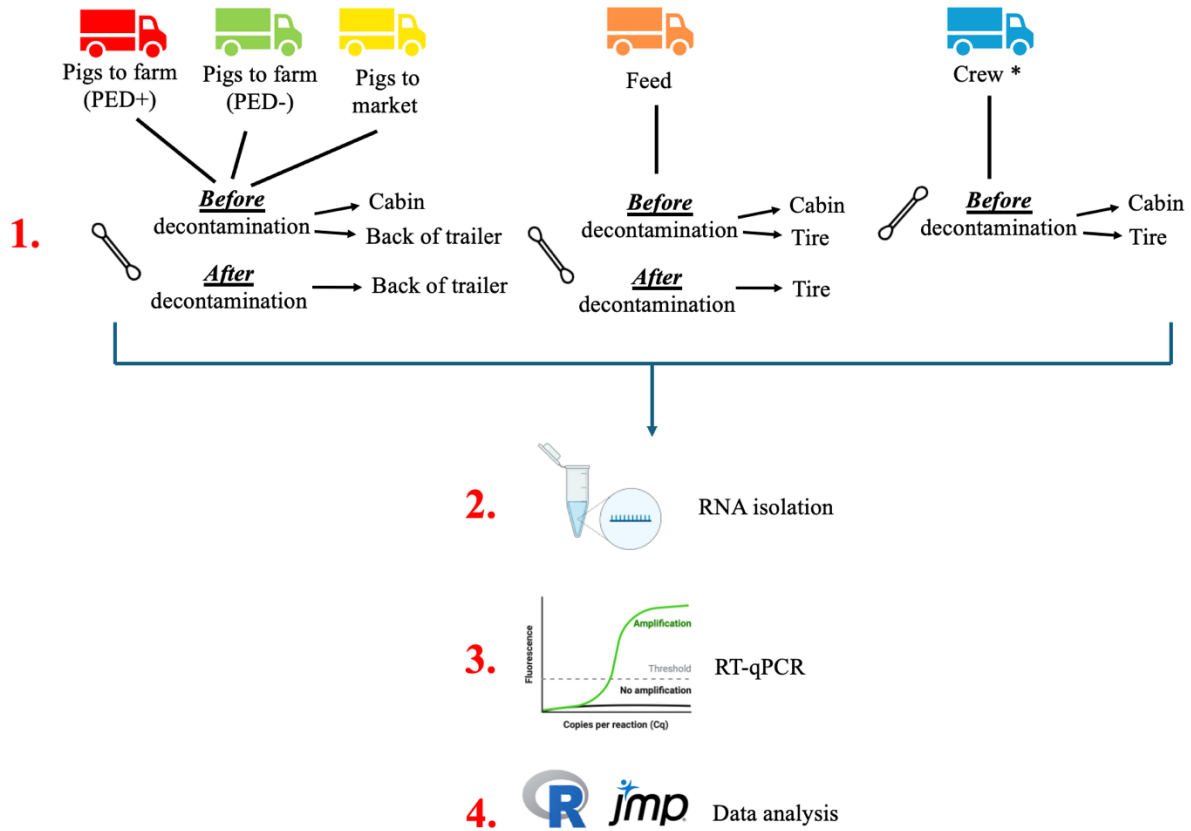


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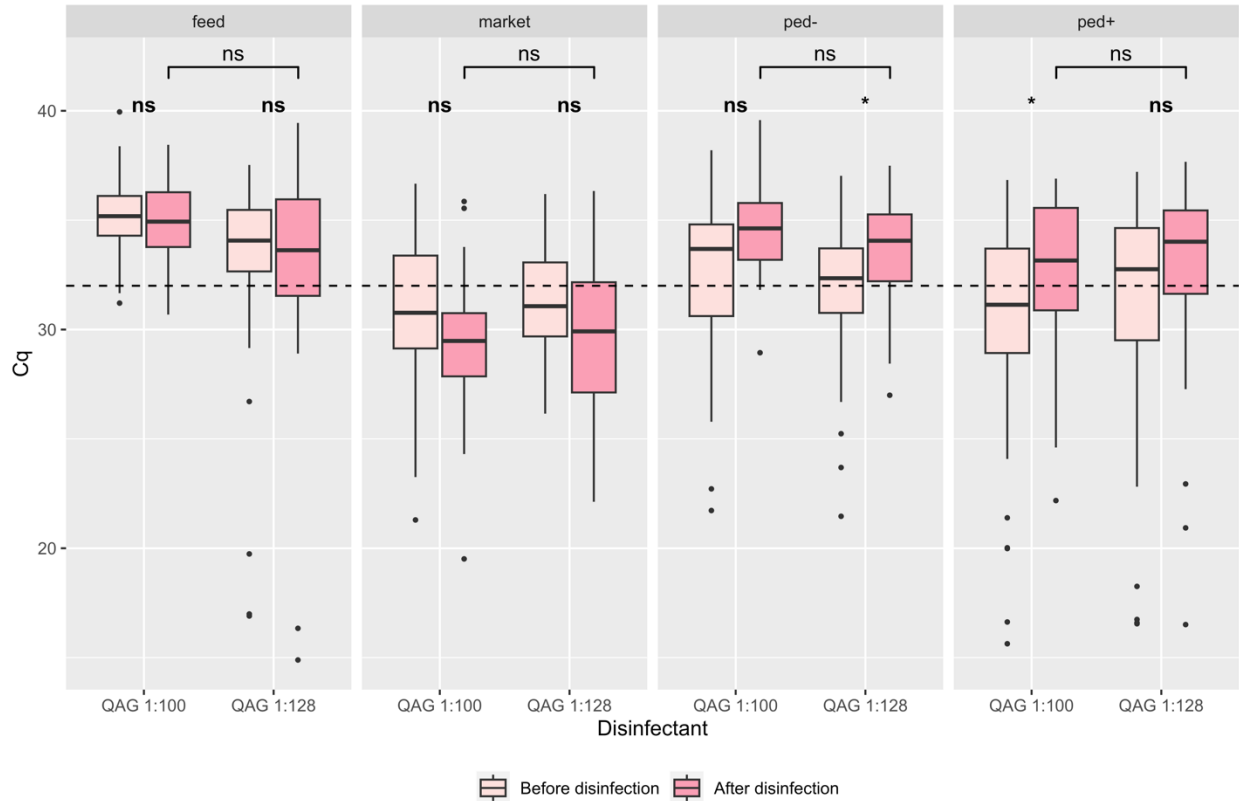
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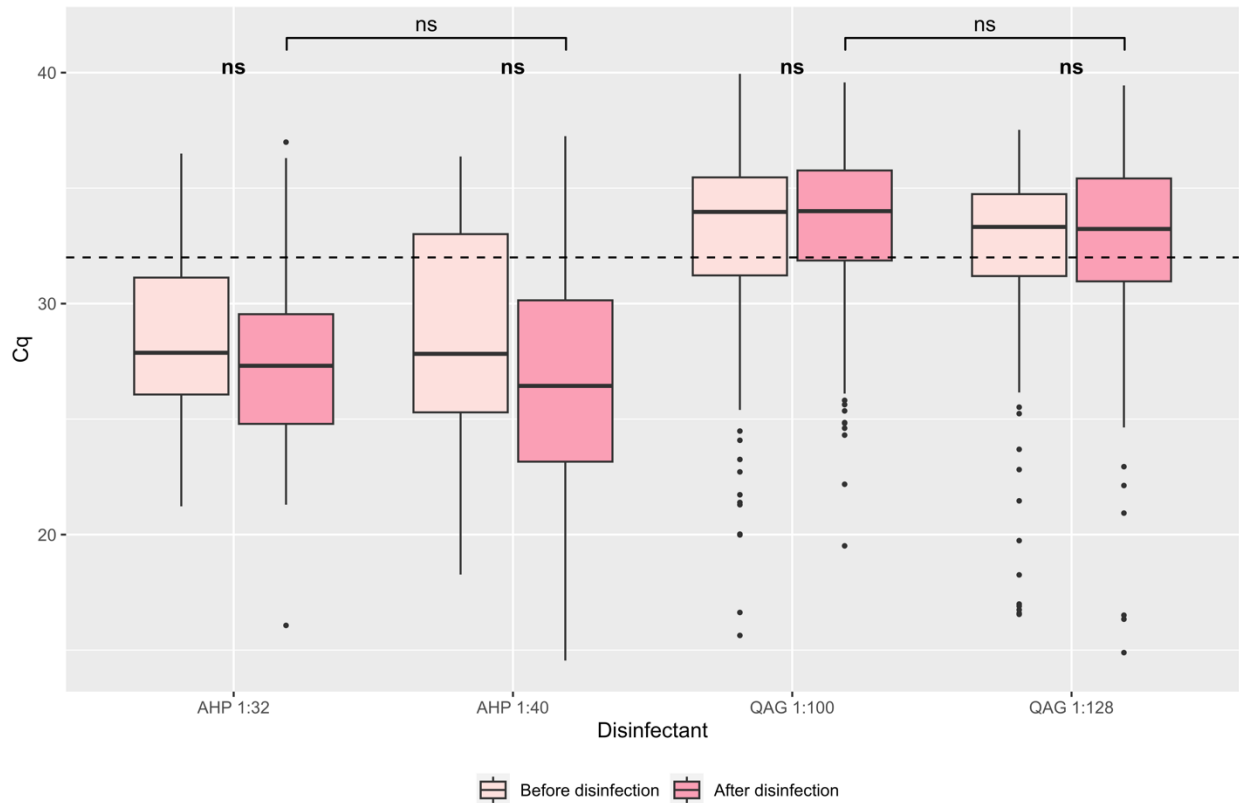
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**Figure 1. Sampling and processing workflow used in this study.** \* denotes that crew trucks were not decontaminated at C&D stations, therefore “after decontamination” samples were not collected.



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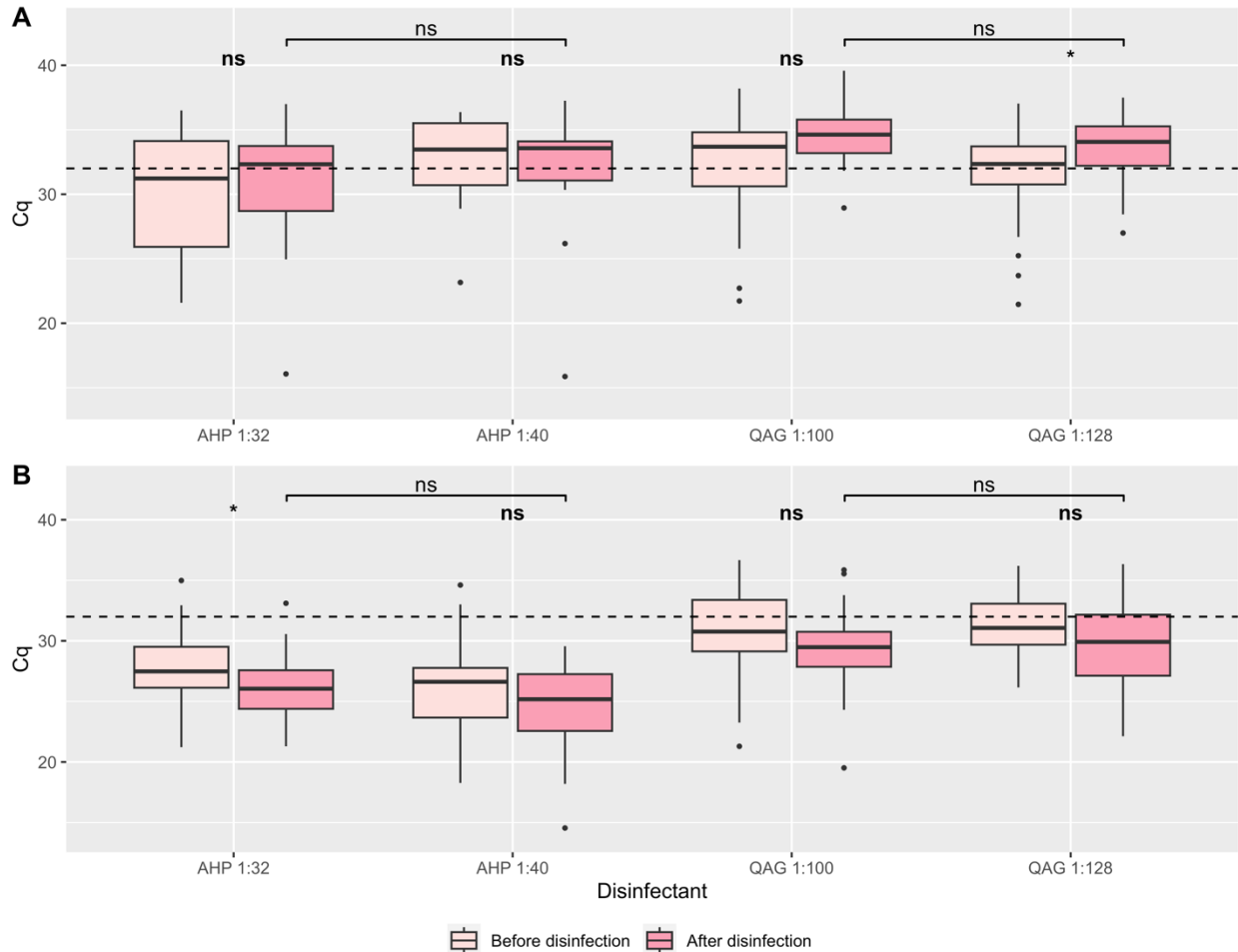
**Figure 2. Comparing disinfectant efficacy in removing PEDV RNA from vehicle exteriors (back-of-trailer and tire samples) across truck types (separated by facet).** Light pink box and whisker plots show the Cq values for samples before disinfection. Darker pink box and whisker plots show the Cq values for samples after disinfection. Lines indicate the median; the box indicates the 1<sup>st</sup> and 3<sup>rd</sup> quartiles; and the whiskers indicate the 1.5 x interquartile range. Individual dots represent outliers. The dashed horizontal line spanning all plots denotes a Cq value of 32, used as the threshold for determining PEDV positive (<32) and negative (≥32) samples. Abbreviations are as follows: Cq, quantification cycle; QAG, Quaternary Ammonium, and Glutaraldehyde; ns, no significance. \*p<0.05 (Supplementary Table 1).



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**Figure 3. Comparing disinfectant efficacy in removing PEDV RNA from vehicle exteriors (back-of-trailer and tire samples) across pigs-to-farm, pigs-to-market, and feed trucks.**

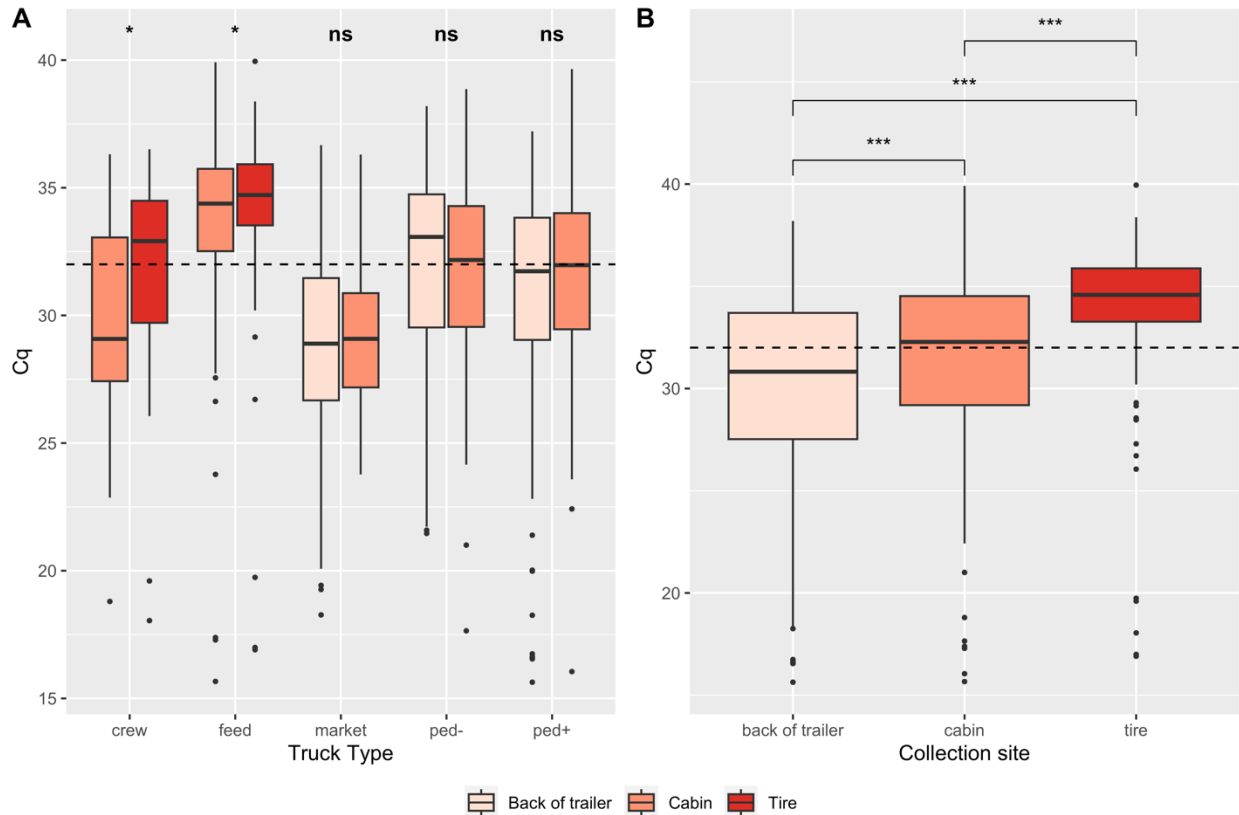
Light pink box and whisker plots show the Cq values for combined back-of-trailer and tire samples before disinfection. Darker pink box and whisker plots show the Cq values for combined back-of-trailer and tire samples after disinfection. Lines indicate the median; the box indicates the 1<sup>st</sup> and 3<sup>rd</sup> quartiles; and the whiskers indicate the 1.5 x interquartile range. Individual dots represent outliers. The dashed horizontal line spanning all plots denotes a Cq value of 32, used as the threshold for determining PEDV positive (<32) and negative (≥32) samples. Abbreviations are as follows: Cq, quantification cycle; AHP, Accelerated Hydrogen Peroxide; QAG, Quaternary Ammonium, and Glutaraldehyde; ns, no significance (Supplementary Table 1).



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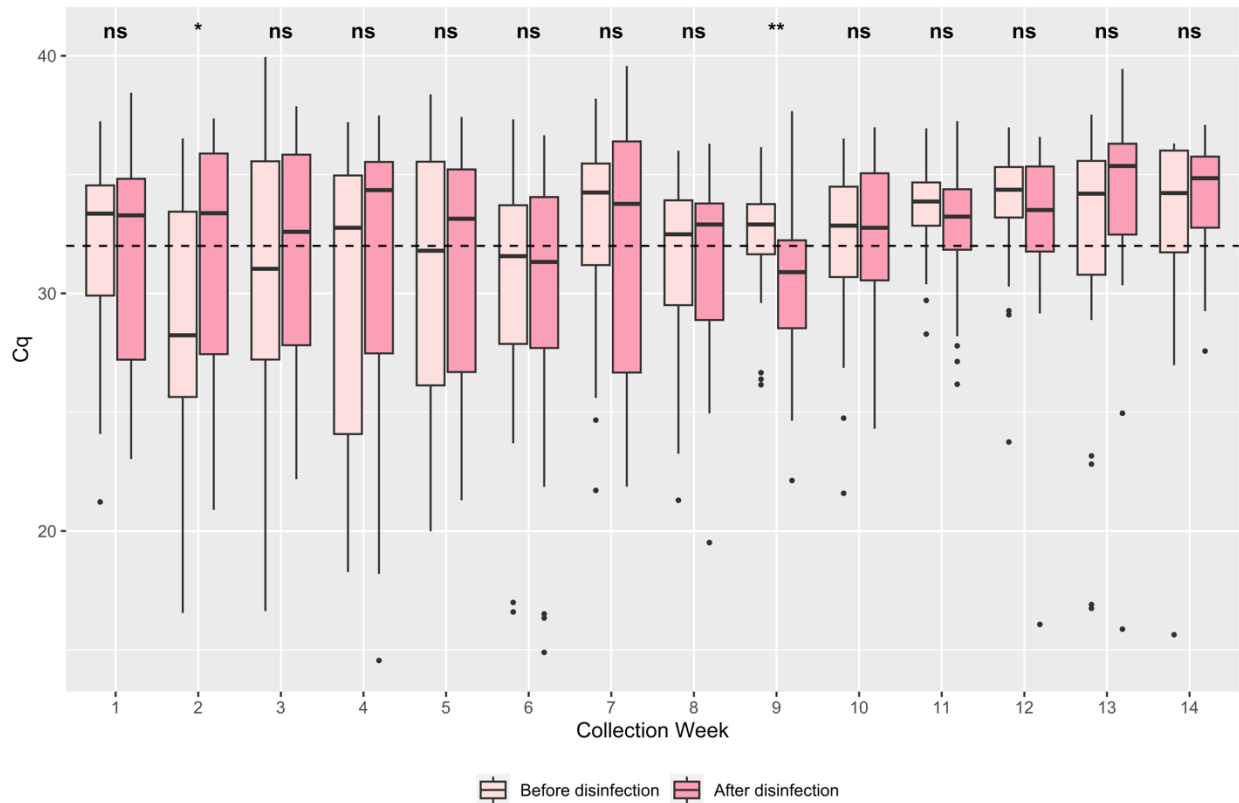
**Figure 4. Comparing disinfectant efficacy across (A) back-of-trailer samples from PED-pigs-to-farm trucks and (B) back-of-trailer samples from pigs-to-market trucks.** Light pink box and whisker plots show the Cq values for samples before disinfection. Darker pink box and whisker plots show the Cq values for samples after disinfection. Lines indicate the median; the box indicates the 1<sup>st</sup> and 3<sup>rd</sup> quartiles; and the whiskers indicate the 1.5 x interquartile range. Individual dots represent outliers. The dashed horizontal lines spanning all plots denote a Cq value of 32, used as the threshold for determining PEDV positive (<32) and negative (≥32) samples. Abbreviations are as follows: Cq, quantification cycle; AHP, Accelerated Hydrogen Peroxide; QAG, Quaternary Ammonium, and Glutaraldehyde; ns, no significance. \*p<0.05 (Supplementary Table 1).





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**Figure 5. Comparing pre-disinfection Cq values across (A) collection sites with respect to truck type and (B) collection sites across all truck types.** Light peach box and whisker plots show the Cq values for samples collected from the back-of-trailer. Orange box and whisker plots show the Cq values for samples collected from the cabin. Red box and whisker plots show the Cq for samples collected from the tire. Lines indicate the median; the box indicates the 1<sup>st</sup> and 3<sup>rd</sup> quartiles; and the whiskers indicate the 1.5 x interquartile range. Individual dots represent outliers. The dashed horizontal line spanning both plots denotes a Cq value of 32, used as the threshold for determining PEDV positive (<32) and negative (≥32) samples. Abbreviations are as follows: Cq, quantification cycle; ns, no significance. Wilcoxon tests used to determine statistical significance; \*p<0.05, \*\*\*p≤0.0001



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644 **Figure 6. Comparing disinfectant efficacy in removing PEDV RNA from pigs-to-farm,**  
645 **pigs-to-market, and feed truck exteriors (back-of-trailer and tire samples) across collection**  
646 **weeks.** Seasonality is as follows: Winter, November-February, weeks 1-7; Spring, March-May,  
647 weeks 8-14. Light pink box and whisker plots show the Cq values for samples before  
648 disinfection. Darker pink box and whisker plots show the Cq values for samples after  
649 disinfection. Lines indicate the median; the box indicates the 1<sup>st</sup> and 3<sup>rd</sup> quartiles; and the  
650 whiskers indicate the 1.5 x interquartile range. Individual dots represent outliers. The dashed  
651 horizontal line denotes a Cq value of 32, used as the threshold for determining PEDV positive  
652 (<32) and negative (≥32) samples. Abbreviations are as follows: Cq, quantification cycle; ns, no  
653 significance. Wilcoxon tests used to determine statistical significance; \*p<0.05, \*\*p<0.001.