



This document is a postprint version of an article published in Veterinary Immunology and Immunopathology © Elsevier after peer review. To access the final edited and published work see <https://doi.org/10.1016/j.vetimm.2021.110206>.

Document downloaded from:



1 **Assessment of three commercial ELISAs for the detection of antibodies against *Porcine***
2 ***epidemic diarrhea virus* at different stages of the immune response.**

3 Díaz I ^{1,2*}, Pujols J ^{1,2}, Cano E ^{1,2}, Cuadrado R ^{1,2}, Navarro N ^{1,2}, Mateu E ^{2,3,4}, Martín M ^{2,3,4}

4 ¹ IRTA, Centre de Recerca en Sanitat Animal (CReSA, IRTA-UAB), Campus de la Universitat
5 Autònoma de Barcelona, 08193 Bellaterra, Spain.

6 ² OIE Collaborating Centre for the Research and Control of Emerging and Re-Emerging Swine
7 Diseases in Europe (IRTA-CReSA), 08193 Bellaterra, Spain.

8 ³ Departament de Sanitat i Anatomia Animals, Universitat Autònoma de Barcelona (UAB), 08193
9 Bellaterra, Spain.

10 ⁴ Centre de Recerca en Sanitat Animal (CReSA, IRTA-UAB), UAB, Campus de la Universitat
11 Autònoma de Barcelona, 08193 Bellaterra, Spain.

12 * Corresponding author. Ivan.diaz@irta.cat

13

14 **Highlights**

- 15 • Commercial PEDV ELISAs showed significant differences in sensitivity.
- 16 • Five months after PEDV inoculation, the rate of detection ranged from 0% to 92%.
- 17 • The competitive ELISA was the only one that detected antibodies in a high percentage
18 of animals for up to five months.
- 19 • Differences between ELISAs could be related to the antigens and to the cut-offs.

20

21

22

23

24

25

26

27

28

29

30

31

32 **Abstract**

33 Three commercial ELISAs -two based on spike (E1 and E3) and one on nucleocapsid protein (E2)-
34 were used to analyze the development and persistence of antibodies against *Porcine epidemic*
35 *diarrhea virus* (PEDV). Seventy-five four-week-old PEDV-negative piglets were inoculated orally
36 with a European G1b PEDV (INOC) and fourteen were kept as controls (CTRL). After the
37 inoculation, E3 detected positive animals as soon as 7 days post inoculation (dpi), while the
38 earliest detection with E1 and E2 was at 14 dpi. All samples were positive at 21 and 28 dpi using
39 E1 and E3, respectively, while E2 failed to detect 23.3% of the inoculated pigs at any time point.
40 The percentages of positive samples were different through the study: E1 and E3 > E2 from 14
41 to 56 dpi; and E3 > E1 > E2 from 56 to 154 dpi (P<0.05). Five months after the inoculation, E3
42 still detected 92.0% (IC_{95%}=85.1-98.8%) of pigs as positive, while E1 and E2 detected only 27.0%
43 (IC_{95%}=16.0-37.9%) and 0%, respectively. The sensitivity for E2 never exceeded 0.62. Specificity
44 was 1 for all ELISAs. These different outcomes could be related to the ELISA strategies (indirect
45 versus competition), the antigens used, the cut-off, or to other intrinsic factors of each test. The
46 observed differences could be of importance when assessing whether older animals, such as
47 fatteners or gilts, had previously been in contact with PEDV.

48

49 **Keywords:** ELISA, *Porcine epidemic diarrhea virus* (PEDV), Spike, Nucleocapsid

50

51

52

53

54

55

56

57

58

59

60

61

62

63

64

65 **1. Introduction**

66 *Porcine epidemic diarrhea virus* (PEDV) is an enveloped, positive-sense, single-stranded
67 RNA virus belonging to the genus *Alphacoronavirus*, family *Coronaviridae*, order *Nidovirales*
68 (ICTV, 2011). In pigs causes a highly transmissible enteric disease characterized by acute watery
69 diarrhea that is especially severe in suckling piglets -case-fatality rate may reach 100% (Geiger
70 and Connor, 2013; Jung and Saif, 2015)-. PEDV was firstly described in UK and Belgium in the
71 1970s (Pensaert and de Bouck, 1978), causing diarrhea outbreaks with high mortality in
72 neonates. Afterwards, PEDV spread in many countries where it established endemically, with
73 sporadic outbreaks of varying severity (Carvajal et al. 2015, Martelli et al., 2008).

74 Based on their S gene sequences, PEDV strains can be differentiated into G1a, G1b, G2a
75 and G2b (Chen et al., 2012; Lee, 2015). Most of the PEDV originally present in Europe in the 70s
76 are classified as G1a, whilst the most recent ones -from 2010 onwards -, which can also be found
77 in Asia and North America, are G1b (Hanke et al., 2017; Kocherhans, et al., 2001; Theuns et al.,
78 2015). In 2013, a new introduction of PEDV in North America caused the death of millions of
79 piglets, being particularly devastating for the US and Canadian pig industries (Schulz and Tonsor,
80 2015; Weng et al., 2016). The strain causing those outbreaks in America could be traced back to
81 Asia (Huang et al., 2013). Thus, differences in pathogenicity of recent genogroups have been
82 demonstrated; G1b strains (known as S-INDEL) present low to moderate virulence, while G2b
83 (non S-INDEL), isolated only in Asia and North-America, show a higher virulence (Carvajal et al.
84 2015).

85 Diagnosis of PEDV outbreaks is mostly done by RT-PCR using feces (de Arriba et al.,
86 2002). On the contrary, the measurement of antibodies against PEDV by ELISA can be useful to
87 detect positive animals beyond the excretion period for different purposes, such as to know the
88 possible presence, persistence and spread of the virus within the farm, the screening of the
89 status of an animal source, to check the immune status of gilts and sows before farrowing, as
90 well as for immunological and epidemiological studies on PEDV.

91 ELISAs currently commercialized for the detection of antibodies against PEDV are based
92 on the spike or the nucleocapsid proteins. The aim of the present study was to compare the
93 performance of three PEDV commercial ELISAs -two based on spike protein (E1 and E3) and one
94 on nucleocapsid protein (E2)- using sera from animals experimentally infected with a G1b PEDV
95 strain in a long-term study.

96 **2. Material and methods**

97 *2.1 Experimental design*

98 Animals and samples used in this study were part of a larger project aimed to
99 characterize the duration of the immune response after the challenge with an European G1b
100 PEDV S-INDEL strain. Briefly, the present study used 89 three-week old piglets from a PEDV-
101 negative source determined by quantitative PCR (qPCR; VetMAX Swine Enteric Panel
102 TGEV/PEDV/PRV-A kit; Thermo Fisher Scientific Inc., Madrid, Spain) and ELISA (INgezim PEDV;
103 11.PED.K1; Eurofins INGENASA, Madrid, Spain). Animals were transported to experimental
104 facilities, ear-tagged and randomly distributed into two separated groups: inoculated (INOC,
105 n=75) and controls (CTRL, n=14). All animals were intramuscularly injected with Ceftiofur (3
106 mg/kg; EXCENEL, Zoetis, Hostalnou de Bianya, Spain) to prevent diarrhea by *E.coli*. After one
107 week of acclimation, namely at 0 days post-inoculation (dpi), piglets in INOC group were orally
108 inoculated using a gastric cannula with 2 mL of the G1b PEDV Calaf-1 (2014) (GenBank accession
109 number MT602520), at a dose of cycle threshold (Ct) =14.7, which corresponded to 10^{4.3} tissue
110 culture infective dose per mL (TCID₅₀/mL). Animals in CTRL group were kept mock-infected with
111 PBS.

112 All experiments involving pigs were done under the approval of Ethical Committee of
113 IRTA and authorized by the Catalan Government (Ref. CEO-H/9450). Animals were kept in
114 approved experimental facilities and were subjected to veterinary supervision for health and
115 welfare. Pig handling was done by veterinarians and trained personnel that fulfilled the Spanish
116 and European Union requirements.

117 2.2 Clinical follow-up and sampling

118 The appearance of individual feces was scored in a scale ranging from 0 (normal) to 3
119 (severe diarrhea). Hence, the maximum value for sum of scores at a given day was 225 (75
120 animals x 3). Individual serum samples were collected before virus inoculation and weekly until
121 day 42 and then at 56, 78, 105, 133 and 154 dpi. Individual fecal samples were also collected at
122 the same time points to assess the presence of PEDV.

123 2.3 PEDV genome detection in feces

124 The presence of PEDV in feces was determined by qPCR (VetMAX Swine Enteric Panel
125 TGEV/PEDV/PRV-A kit, Thermo Fisher Scientific Inc.), using the Path-ID Multiplex One-Step RT-
126 PCR kit (Thermo Fisher Scientific Inc.). Viral RNA was extracted using MagMAX pathogen
127 RNA/DNA (Thermo Fisher Scientific Inc.) and a BioSprint 96 workstation (Qiagen Iberia, Las
128 Matas, Spain), according to the manufacturer's instructions. Positive and negative controls were
129 included in each reaction and round of RNA extraction. Results were expressed as Ct.

130 2.4 ELISAs

131 Three commercial ELISAs for detection of anti-PEDV antibodies were considered: E1
132 (INgezim PEDV; 11.PED.K1; Eurofins INGENASA), E2 (ID Screen PEDV indirect; IDVet, Grabels,
133 France), and E3 (ID Screen PEDV Spike Competition; IDVet, Grabels, France). Both E1 and E2
134 were indirect ELISAs while E3 was a competition ELISA. E1 and E3 used as antigen spike
135 glycoprotein from G1b strains, while E2 was based on a G1b nucleocapsid protein.

136 All tests were performed and interpreted according to the manufacturer's instructions.
137 For E1, results were expressed as a sample-to-positive (S/P) ratio, namely a ratio of the optical
138 density (OD) of a given sample over the OD of the positive control provided with the kit. An S/P
139 ratio ≥ 0.35 was considered positive. For E2, S/P percentages (S/P%) were calculated: $[(DO$
140 $sample - DO\ negative\ control) / (DO\ positive\ control - DO\ negative\ control)] \times 100$. In this case,
141 the threshold to consider a sample positive was $> 60\%$. Since E3 was a competition ELISA, results
142 were expressed as a sample-to-negative (S/N) percentage $[(DO\ sample/DO\ negative\ control) \times$
143 $100]$; results lower than 50% were considered positive.

144 For any given ELISA, all samples yielding doubtful results were retested to discard any
145 potential error attributable to the laboratory processing. In order to minimize biases, all samples
146 were tested using the same ELISA batch.

147 *2.5 Statistical analyses*

148 Excel 2016 (Microsoft) was used to calculate descriptive statistics. Comparison of the
149 percentages of positive pigs between ELISAs at a given sampling day was done using StatsDirect
150 v3.2.8; firstly, a screening to assess if there were differences was done by Chi-square test for
151 multiple groups and then, the pairs were tested by the Fisher's exact test.

152 **3. Results and discussion**

153 PEDV infection beyond the first days of life is often mild and/or of short duration
154 (Crawford et al., 2015; Jung and Saif, 2015; Weng et al., 2016). In some cases, infection can
155 remain unnoticed making an indirect detection, such as the measure of antibodies a useful
156 screening tool. Several studies have evaluated both commercial and in-house ELISAs, either
157 spike-based or nucleocapsid-based ones (Gerber et al., 2016; Okda et al., 2015; Strandbygaard
158 et al., 2016). However, to our knowledge only one study has been published on the assessment
159 of PEDV antibodies during a long period (Knuchel et al., 1992). In that case, two in-house ELISAs
160 based on spike and nucleocapsid proteins were compared during six months after an
161 experimental infection. According to the authors, antibodies against spike protein remained
162 detectable for longer periods compared to the nucleocapsid ones. It is worth mentioning that
163 the experimental infection was done with a G1a PEDV strain (V215/78) already present in

164 Europe in the 70s (Pensaert and de Bouck, 1978). The aim of the present study was to assess the
165 dynamics of PEDV-specific production during five months, using three commercial ELISAs (two
166 spike-based ELISAs and one nucleocapsid-based ELISA), after a European G1b PEDV strain
167 experimental infection.

168 PEDV infection was demonstrated in all animals from INOC group, both by clinical
169 observations and PEDV genome detection in feces. Regarding clinical scores, presence of loose
170 stools/diarrhea was detected in all pigs. The maximum cumulative clinical score was 120 at day
171 4 dpi (out of a potential maximum of 225). No loose stools or diarrhea were observed beyond
172 21 dpi. Fecal excretion of PEDV was negative for all pigs at 0 dpi. All INOC piglets were positive
173 at 7 dpi (mean Ct=24.3±4.0). Afterwards, the percentage of positive samples and the average
174 amount of PEDV genome detected in positive animals declined until 42 dpi, when only one
175 animal was still positive (Ct=36.7). Afterwards, all animals remained negative until the end of
176 the study at 154 dpi. CTRL were all qPCR negative throughout the study.

177 Figure 1 summarizes the percentage of positive results detected by each ELISA. At the
178 beginning of the study, all pigs both in INOC and CTRL groups were negative by all ELISAs. In the
179 INOC group, 95.0% (IC_{95%}=91.5-100%) of the animals were detected as positive at 14 dpi with E1,
180 with all of them positive at 21 dpi. Using E2, the highest percentage of positive pigs was 62.2%,
181 detected at 21 dpi (IC_{95%}=51.1-73.2%). Importantly, with this E2, 19/75 INOC animals (23.3%;
182 IC_{95%}=16.3-36.9%) were never detected as positive (OD below the cut-off of the kit). When
183 samples were analyzed using E3, 50.7% of the pigs (IC_{95%}=39.4-62.0%) were classified as positive
184 as soon as 7 dpi; at 28 dpi all were positive and remained in that way until 56 dpi. From 14 to 56
185 dpi, the percentages of positive samples were always significantly higher in E1 and E3 than in E2
186 (P<0.05). Afterwards, significant differences were observed between all ELISAs, with E3 being
187 always the more sensitive (E3>E1>E2; P<0.05). Five months after the inoculation, E3 still
188 detected as positive 92.0% (IC_{95%}=85.1-98.8%) of the inoculated animals, while E1 and E2
189 detected only 27.0% (IC_{95%}=16.0-37.9%) and 0%, positive animals, respectively.

190 Regarding S/P, S/P% or S/N% means (figures 2a-2c), the maximum value in INOC group
191 was observed at 21 dpi for both E1 and E2, while the peak was seen at 28 dpi for E3. For E2,
192 mean S/P% was slightly above the cut-off at 28 dpi and from that day until 154 dpi was always
193 clearly below (figure 2b). The dynamics of the mean S/N% for E3 from 28 dpi onwards followed
194 similar mean values (figure 2c).

195 The sensitivity (Sn) for each ELISA at a given sampling day was also calculated (figure 3).
196 Sn was 1 for E1 at 21 dpi and for E3 at 28dpi; in contrast, for E2 Sn never exceeded 0.63. Later

197 on, Sn values declined for all ELISAs, although remained above 0.9 using E3. Mean S/N% values
198 and Sn for E3 remained highly constant during all the experiment, especially when compared
199 with E1 and E2. E3 was especially valuable for detecting individuals during longer periods,
200 maintaining the capability to detect >90% of positive animals after five months. Thus, after a
201 PEDV infection in the first days or weeks of life, E3 could still detect most animals as positive
202 until the finishing phase. In comparison, E1 seemed to have some limitations for long-term
203 detection and, therefore, it would be less useful to detect whether older animals, such as
204 finishers or replacement gilts, had been infected at young ages. In contrast, specificity (Sp) was
205 very high for all ELISAs (Sp=1).

206 The rationale behind the differences observed among the Sn of the three ELISAs are
207 unclear. Regarding the antigens, E2 is based on the nucleocapsid, which is well-conserved among
208 PEDV strains (Lin et al., 2015), is the most abundant viral protein and induces a strong humoral
209 response (Li, 2015; Sturman and Holmes, 1983; Utiger et al., 1995). However, it has been
210 suggested that the spike protein has a remarkably higher immunogenicity compared to
211 nucleocapsid, and that spike-based ELISAs, such as E1 and E3, are more specific and sensitive
212 than nucleocapsid-based ones (Hou et al., 2007; Knuchel et al., 1992; Okda et al., 2015).
213 Moreover, when spike and nucleocapsid-based ELISAs have been compared in the same PEDV-
214 infected pigs, antibodies against spike have remained detectable in sera for longer periods
215 (Knuchel et al., 1992). Interestingly, sera from PEDV-infected pigs not only could reach higher
216 titers in spike-based ELISAs, but also these antibodies could closely link with neutralization
217 activity and be, therefore, relevant for protection (Paudel et al., 2014).

218 The abovementioned facts may partially explain the presence of false negatives and the
219 lowest Sn seen for E2 compared to E1 and E3. Conversely, some authors have claimed that
220 differences among assays could be more related to intrinsic factors, rather than to the PEDV
221 antigen used, as demonstrated using a combination of different antigens and different ELISA
222 strategies (Gerber et al., 2016). Accordingly, differences observed between E1 and E3 (both
223 spike-based ELISA) could be due to ELISA strategies (indirect versus competition) or to other
224 intrinsic factors: cut-off values; coating with a recombinant versus purified spikes, or
225 alternatively, to spike heterogeneity (Lin et al., 2015).

226 To determine if the low Sn calculated for E2 was due to the cut-off value ($S/P\% > 60\%$),
227 a new one was calculated based on the results obtained from truly negative pigs in the present
228 study [(n=243 observations: all samples in INOC group at day 0 dpi (n=75) plus all samples from
229 0 to 154 dpi in CTRL group (n= 14 pigs x 12 sampling days = 168)]. For this, mean $S/P\% + 5xSD$

230 was calculated and set as a potential cut-off (99.9% of values for negative samples). In regards
231 of Sn, figure 3 (E2', dotted gray line) shows the ability of E2' to detect positive animals using the
232 new threshold ($S/P\% > 17\%$). The magnitude and pattern for the new Sn (E2') was like the one
233 obtained for E1 and clearly higher than that calculated using the original threshold (E2). Using
234 the new cut-off, all animals yielded positive results at 21 dpi, percentages of positives remained
235 at 88.0% ($IC_{95\%}=80.6-95.3\%$) at 56 dpi and 19.3% ($IC_{95\%}=15.4-23.3\%$) at 154 dpi. Applying $S/P\% >$
236 17% as cut-off, only one sample was detected as false positive ($Sp=99.6\%$). It cannot be
237 discarded that the original cut-off of E2 was higher because the field samples used during the
238 standardization and validation process showed a different behavior in the ELISA, compared to
239 experimental samples.

240 In summary, this study demonstrates the existence of significant differences between
241 three commercial PEDV-specific ELISAs. As an example, E2 showed a very low Sn, with a 25% of
242 false-negative animals throughout all the study. Differences were particularly marked during
243 long-term phases after the infection. Thus, E3 still detected as positive more than 90% of the
244 animals as late as five months after the infection; while, at that time, E1 and E2 only detected
245 27% and 0%, respectively. The differences could be due to ELISA strategies (competition versus
246 indirect), the antigen used and/or other intrinsic factors of each ELISA. This fact could be a
247 challenge when monitoring the PEDV immune status in older animals, such as pigs during the
248 final phase of fattening or gilts, if they were infected some months before.

249 **Declaration of Competing Interest**

250 The Authors declare no conflict of interest with respect to the research, authorship,
251 and/or publication of this article.

252 **Acknowledgements**

253 We would like to thank Eurofins INGENASA and IDVet for kindly providing the ELISA kits
254 used in this study. Also, we thank Field Trials Unit (IRTA) for technical support and assistance.
255 CERCA Programme from the Generalitat de Catalunya is also acknowledged.

256 **Funding**

257 This work was supported by the National Institute of Research and Agricultural and Food
258 Technology (INIA, reference E-RTA2015-0003-C02-01).

259 **References**

260 Carvajal, A., Argüello, H., Martínez-Lobo, F.J., Costillas, S., Miranda, R., G de Nova, P.J., Rubio, P., 2015.
261 Porcine epidemic diarrhoea: new insights into an old disease. *Porcine Health Manag.* 1, 12.

262 Chen, J., Liu, X., Shi, D., Shi, H., Zhang, X., Feng, L., 2012. Complete genome sequence of a porcine epidemic
263 diarrhea virus variant. *J. Virol.* 86, 3408.

264 Crawford, K., Lager, K., Miller L., Opriessnig, T., Gerber, P., Hesse, R., 2015. Evaluation of porcine epidemic
265 diarrhea virus transmission and the immune response in growing pigs. *Vet. Res.* 46, 49.

266 de Arriba, M.L, Carvajal, A., Pozo, J., Rubio, P., 2002. Isotype-specific antibody-secreting cells in systemic
267 and mucosal associated lymphoid tissues and antibody responses in serum of conventional pigs
268 inoculated with PEDV. *Vet. Immunol. Immunopathol.* 84, 1-16.

269 Geiger, O., Connor, J.F., 2013. Porcine epidemic diarrhea, diagnosis and elimination. Web available
270 <https://www.aasv.org/aasv%20website/Resources/Diseases/PED/13-05-29PEDWhitePaper.pdf>.
271 [Accessed on 08.06.20.](#)

272 Gerber, P.F., Lelli, D., Zhang, J., Strandbygaard, B., Moreno, A., Lavazza, A., Perulli, S., Bøtner, A., Comtet,
273 L., Roche, M., Pourquier, P., Wang, C., Opriessnig, T., 2016. Diagnostic evaluation of assays for detection
274 of antibodies against porcine epidemic diarrhea virus (PEDV) in pigs exposed to different PEDV strains.
275 *Prev. Vet. Med.* 135, 87-94.

276 Hanke, D., Pohlmann, A., Sauter-Louis, C., Höper, D., Stadler, J., Ritzmann, M., Steinrigl, A., Schwarz, B.A.,
277 Akimkin, V., Fux, R., Blome, S., Beer, M., 2017. Porcine epidemic diarrhea in europe: in-detail analyses of
278 disease dynamics and molecular. *Viruses* 9, 177.

279 Hou, X.L., Yu, L.Y., Liu, J., 2007. Development and evaluation of enzyme-linked immunosorbent assay
280 based on recombinant nucleocapsid protein for detection of porcine epidemic diarrhea (PEDV) antibodies.
281 *Vet. Microbiol.* 123, 86-92.

282 Huang, Y., Dickerman, A.W., Piñeiro, P., Li, L., Fang, L., Kiehne, R., Opriessnig, T., Meang XJ, 2013. Origin,
283 evolution, and genotyping of emergent porcine epidemic diarrhea virus strains in the United States. *MBio.*
284 4:e00737-13.

285 International Committee on Taxonomy of Viruses, 2011. [https://talk.ictvonline.org/ictv-](https://talk.ictvonline.org/ictv-reports/ictv_9th_report/positive-sense-rna-viruses-2011/w/posrna_viruses/222/coronaviridae)
286 [reports/ictv_9th_report/positive-sense-rna-viruses-2011/w/posrna_viruses/222/coronaviridae](https://talk.ictvonline.org/ictv-reports/ictv_9th_report/positive-sense-rna-viruses-2011/w/posrna_viruses/222/coronaviridae).
287 [Accessed on 01.06.20.](#)

288 Jung, K., Saif, L.J., 2015. Porcine epidemic diarrhea virus infection: Etiology, epidemiology, pathogenesis
289 and immunoprophylaxis. *Vet J.* 204, 134-143.

290 Kocherhans, R., Bridgen, A., Ackermann, M., 2001. Completion of the porcine epidemic diarrhoea
291 coronavirus (PEDV) genome sequence. *Virus Genes* 23, 137-144.

292 Knuchel, M., Ackermann, M., Müller, H.K., Kihm, U., 1992. An ELISA for detection of antibodies against
293 porcine epidemic diarrhoea virus (PEDV) based on the specific solubility of the viral surface glycoprotein.
294 *Vet. Microbiol.* 32, 117–134.

295 Lee, C., 2015. Porcine epidemic diarrhea virus: an emerging and re-emerging epizootic swine virus. *Virology*
296 *J.* 12, 193.

297 Li, F., 2015. Receptor recognition mechanisms of coronaviruses: A decade of structural studies. *J. Virology*
298 89, 1954-1964.

299 Lin, C., Gao, X., Oka, T., Vlasova, A.N., Esseili, M.A., Wang, Q., Saif, L.J., 2015. Antigenic relationships among
300 porcine epidemic diarrhea virus and transmissible gastroenteritis virus strains. *J. Virology* 89, 3332-3342.

301 Martelli, P., Lavazza, A., Nigrelli, A.D., Merialdi, G., Alborali, L. G., Pensaert, M.B., 2008. Epidemic of
302 diarrhoea caused by porcine epidemic diarrhoea virus in Italy. *Vet. Rec.* 162, 307-310.

303 Okda, F., Liu, X., Singrey, A., Clement, T., Nelson, J., Christopher-Hennings, J., Nelson, E.A., Lawson, S.,
304 2015. Development of an indirect ELISA, blocking ELISA, fluorescent microsphere immunoassay and
305 fluorescent focus neutralization assay for serologic evaluation of exposure to North American strains of
306 Porcine Epidemic Diarrhea Virus. *BMC Vet. Res.* 11, 180.

307 Paudel, S., Park, J.E., Jang, H., Shin, H.J., 2014. Comparison of serum neutralization and enzyme-linked
308 immunosorbent assay on sera from porcine epidemic diarrhea virus vaccinated pigs. *Vet. Q.* 34, 218-23.

309 Pensaert, M.B., de Bouck, P., 1978. A new coronavirus-like particle associated with diarrhea in swine. *Arch.*
310 *Virology* 58, 243-247.

311 Schulz, L.L., Tonsor, G.T., 2015. Assessment of the economic impacts of porcine epidemic diarrhea virus
312 in the United States. *J. Anim. Sci.* 93, 5111-5118.

313 Strandbygaard, B., Lavazza, A., Lelli, D., Blanchard, Y., Grasland, B., Poder, S.L., Rose, N., Steinbach, F., van
314 der Poel, W.H.M., Widén, F., Belsham, G.J., Bøtner, A. 2016. Inter-laboratory study to characterize the
315 detection of serum antibodies against porcine epidemic diarrhoea virus. *Vet. Microbiol.* 197, 151-160.

316 Sturman, L.S., Holmes, K.V., 1983. The molecular biology of coronaviruses. *Adv. Virus. Res.* 28, 35-112.

317 Theuns, S., Conceicao-Neto, N., Christiaens, I., Zeller, M., Desmarets, L.M., Roukaerts, I.D., Acar, D.D.,
318 Heylen, E., Matthijssens, J., Nauwynck H.J., 2015. Complete genome sequence of a porcine epidemic
319 diarrhea virus from a novel outbreak in Belgium, January 2015. *Genome Announc.* 3:e00506–15.

320 Utiger, A., Tobler, K., Bridgen, A., Suter, M., Singh, M., Ackermann, M., 1995. Identification of proteins
321 specified by porcine epidemic diarrhoea virus. *Adv. Exp. Med. Biol.* 380, 287-90.

322 Weng, L., Weersink, A., Poljak, Z., de Lange, K., von Massow, M., 2016. An economic evaluation of
323 intervention strategies for porcine epidemic diarrhea (PED). *Prev. Vet. Med.* 134, 58-68.

324 **Figure 1. Percentage of positive results in each ELISA.** Bars indicate the percentage of positive
325 results in INOC group for E1 (black bar) for E2 (grey bar), and for E3 (white bar) at a given
326 sampling day. Bars with a letter above show significant differences among ELISAs at a given
327 sampling day (Fisher's exact test; <0.05).

328 **Figure 2. Longitudinal serological profiles.** Lines indicate the mean \pm SD results at a given
329 sampling day. In figure 2a, means are shown as mean of the S/P ratios (E1); in figure 2b, means
330 of the S/P% (E2); and in 2c, means of the S/N% (E3). Black lines correspond to INOC group and
331 dashed grey lines to CTRL group. The dotted black lines show the threshold value for each ELISA.

332 **Figure 3. Sensitivity (Sn) for each ELISA throughout the study.** Lines indicate the Sn values of
333 each ELISA at different days: black line (E1), dashed grey line (E2) and grey line (E3). Dotted grey
334 line (E2') indicates the Sn for E2 using a recalculated cut-off.

335

Figure 1

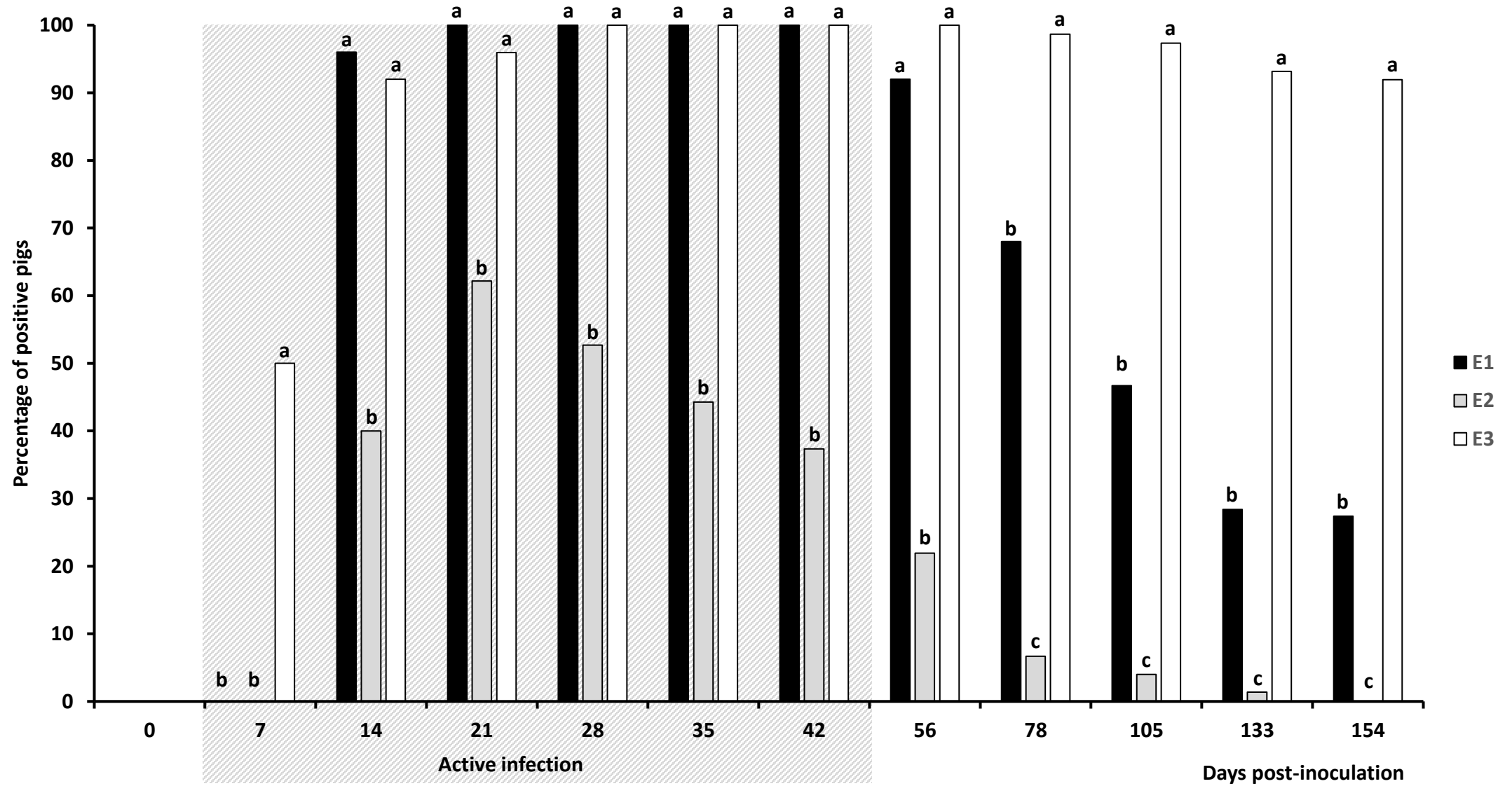


Figure 2a

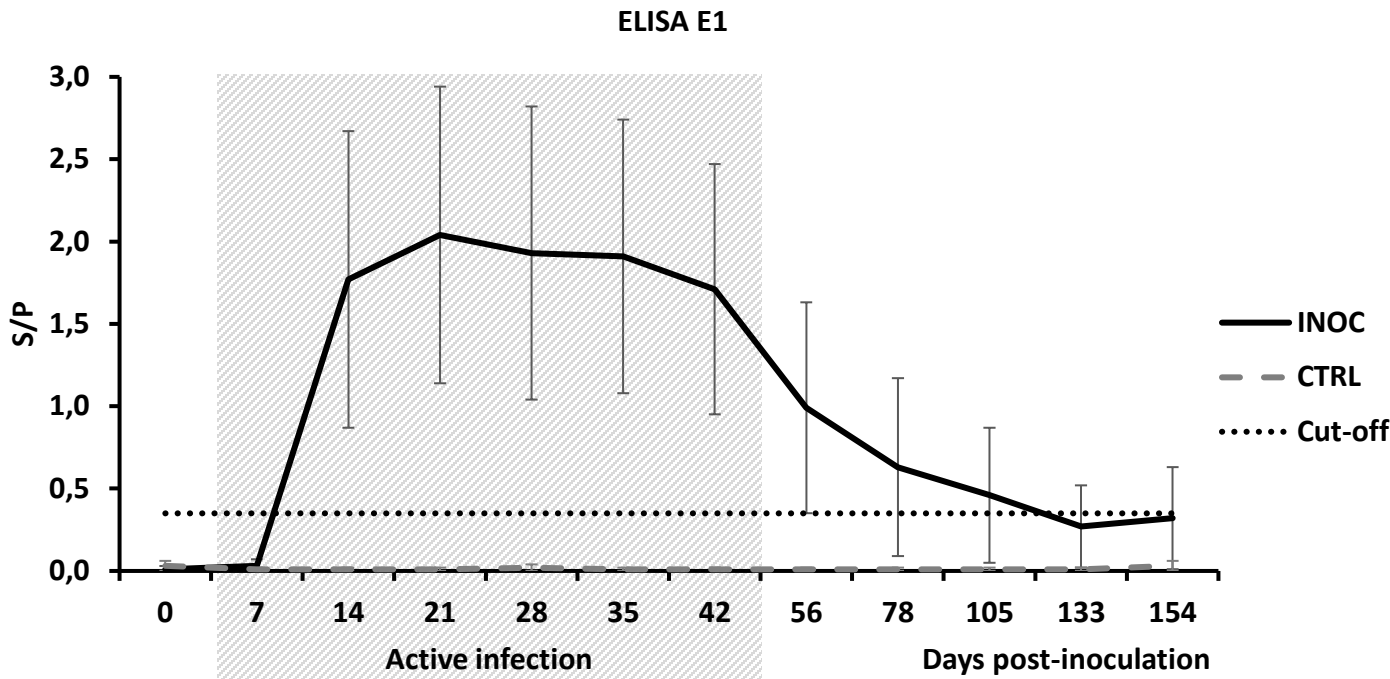


Figure 2b

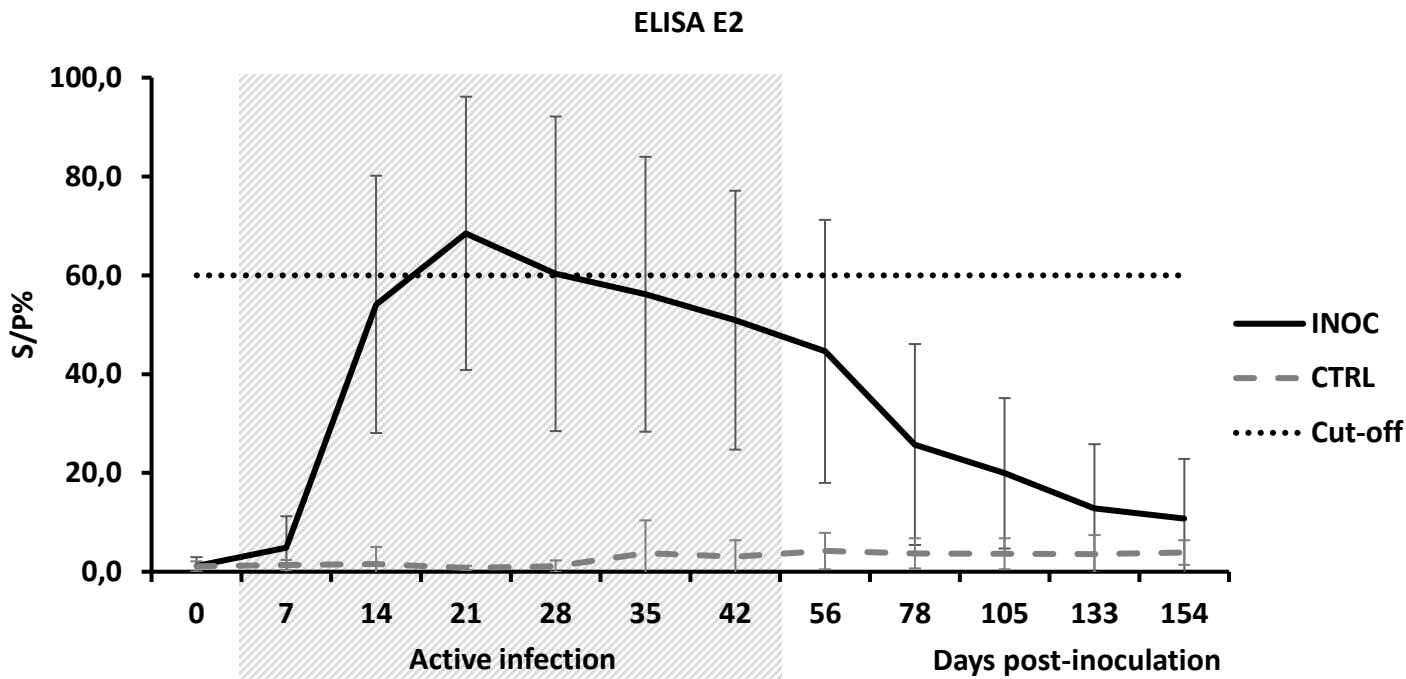


Figure 2c

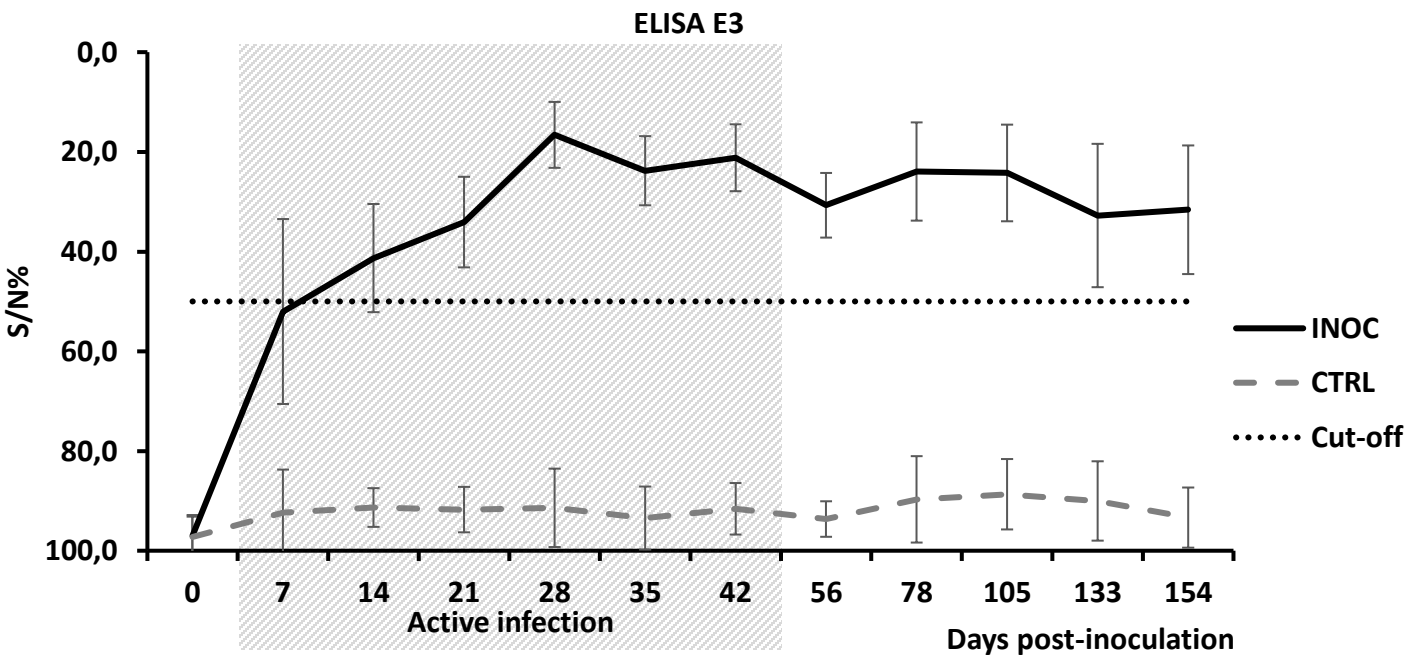


Figure 3

