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Document downloaded from:

<http://hdl.handle.net/10459.1/64978>

The final publication is available at:

<https://doi.org/10.1016/j.virusres.2014.03.004>

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26 **ABSTRACT**

27 In the present work, we studied the infection caused by the low virulence strain Cat01 in
28 domestic pigs, which is responsible for the 2001- 2002 CSFV outbreak in the Catalonia
29 region in Spain, in comparison with the infection caused by a CSFV virulent strain
30 responsible for inducing the acute form of CSF in the field. We focused on assessing the
31 impact of CSFV virulence on the kinetics of different cytokines, such as IFN- α (innate
32 immunity) and IFN- γ (adaptive immune response), during the first weeks after infection.
33 In addition, we evaluated the impact of CSFV virulence on the induction of the humoral
34 immune response and its relation to the course of infection and the RNA CSFV viral
35 load. IFN- α levels in the serum samples from pigs infected with the low virulent strain
36 were lower than those detected in pigs infected with the virulent one. After the infection
37 with the low virulent strain, some IFN- γ response against CSFV was detected. Likewise,
38 the humoral response was detected only in pigs infected with the low virulent strain.
39 Noteworthy, the lowest load of CSFV RNA was detected in the serum samples of these
40 animals. Similarly, the lowest viral load levels were detected in the tonsils of these pigs.
41 Both the T cells and the humoral response generated in most of the pigs infected with
42 strain Cat01 could be related to the protection in the symptom progression of CSF
43 against this CSFV strain. However, despite the levels of protection observed, this strain
44 still impaired the adaptive immunity against CSFV, as not all animals developed a
45 detectable T-cell response and adequate levels of neutralising antibodies that are capable
46 of neutralising the virus at both the serum and the tissue levels. Thus, RNA viral
47 detection was observed in the tonsils from all Cat01-infected pigs. This fact can
48 encourage the spread of the virus and its persistence after infection with CSFV low viral
49 strains. These types of results explain in part why CSFV are capable of producing an

50 outbreak that goes unnoticed during the first months of infection and how difficult its
51 resulting eradication is in endemic countries today.

52 **INTRODUCTION**

53 Classical swine fever (CSF) is a highly contagious and deadly viral disease caused by a
54 small enveloped single-stranded RNA virus that belongs to the genus Pestivirus within the
55 family *Flaviviridae* (Heinz et al., 2004). Today, CSF represents one of the leading threats
56 to the pig industry worldwide (Wang et al., 2008). The virus is known to have tropism for
57 vascular endothelial and immune system cells, mainly those derived from the monocyte-
58 macrophage lineage (Summerfield et al., 1998, Summerfield et al., 2001). It is assumed
59 that the infection of these cells plays an important role in the disease epidemiology and
60 pathogenesis, for instance for the spread and the persistence of this virus, as well as for
61 immunosuppression.

62 Depending on the virulence of the strain and the immunological status of the infected
63 animals, pigs can have as much as 90% of their total T cells depleted in the final stages of
64 the disease (Pauly et al., 1998). This effect can be observed as early as one day after
65 infection, even before viraemia has been established (Summerfield et al., 1998).
66 Immunosuppression can be detected much earlier than seroconversion and the presence of
67 clinical signs, which is relevant for early diagnosis and for the study of viral pathogenesis
68 (Ganges et al., 2008, Pauly et al., 1998).

69 Type-I interferon (IFN) has antiviral and immunomodulatory effects and is a
70 component of the innate immune response against viruses. It is attributed a beneficial
71 effect in viral infections, restricting the viral dissemination and promoting
72 immunopathological events when released at high levels over longer periods
73 (Summerfield et al., 2006). However, CSFV exacerbates the IFN- α response that is
74 detected in the serum of infected pigs, and this has been suggested to relate to disease

75 severity rather than to protective immune responses (Summerfield et al., 2006; Tarradas
76 et al., 2010; Summerfield, 2011).

77 Depending on the virulence of the strain, varying degrees of disease severity can
78 be observed, ranging from acute to subclinical or chronic forms (van Oirschot 1988;
79 2004). Interestingly, whereas infections with virulent strains result in an acute
80 hemorrhagic disease, the infection caused by less virulent isolates can become chronic.
81 Due to the severity of the disease, the acute form of the infection has been widely
82 studied, whereas there is very little information about the pathogenesis of these types of
83 CSFV strains and their relation with the disease progress. Curiously, a trend toward
84 milder, chronic presentations of CSF has been observed as a consequence of years of
85 several CSF outbreaks in different endemic countries (Pérez et al., 2012; Shen et al.,
86 2011). In addition, pigs infected with low-virulence strains are known to shed virus
87 continuously or intermittently for months, representing a major threat not only for
88 countries endemically infected but also worldwide (Pérez et al., 2012, Weesendorp et al.,
89 2011).

90 In spite of the potential involvement of the T cellular response against CSFV vaccines,
91 the principal role of IFN- γ is to control the infection (Tarradas et al., 2010; Graham et
92 al., 2012). The immune response against CSFV infection with strains of various levels of
93 virulence is poorly understood.

94 In the present work, we studied the infection caused by the low virulent strain
95 Cat01 in domestic pigs, which was responsible for the 2001- 2002 CSFV outbreak in the
96 Catalonia region in Spain, in comparison with the infection caused by a CSFV virulent
97 strain responsible for inducing the acute form of CSF in the field. We focused on
98 assessing the impact of CSFV virulence on the kinetics of various cytokines, such as IFN-
99 α (inmate immunity) and IFN- γ (adaptive immune response), during the first weeks after

100 infection. In addition, we evaluate the impact of CSFV virulence on the induction of
101 humoral immune response and its relation to the course of infection and the viral load of
102 CSFV RNA.

103 **MATERIALS AND METHODS**

104 **Cells and viruses**

105 The porcine kidney PK15 cell line was used to grow and titre the viral stocks. The cells
106 were cultured in DMEM media (Invitrogen, USA) supplemented with 10% fetal bovine
107 serum (FBS) at 37°C in 5% CO₂. PK15 were infected with a 0.1 TCID₅₀/cell using DMEM
108 media with 2% FBS, and the virus was harvested at 48 h post-infection. Stock titration
109 was performed using the peroxidase-linked assay (PLA) (Wensvoort et al., 1986),
110 following the statistical method previously described (Reed and Muench, 1938). The
111 Catalunya 01 (Cat01) viral strain used in this study was isolated from CSF Spanish
112 epizootic in 2000-2001. This isolate was included by sequence analysis into the CSFV 2.3
113 genogroup (Pérez et al., 2012). The course of the infection at the farm, where this virus
114 was isolated, was found to be mild (Allepuz et al., 2007). The virulent Margarita strain
115 belongs to the 1.4 genotype, and has been used in Cuba since 1965 for a vaccine potency
116 test (Díaz de Arce et al., 1999; Ganges et al., 2005).

117 **Experimental design**

118 To evaluate the pathogenesis induced by both strains, we used three groups of
119 domestic pigs. Each group included six Landrace x Large White healthy pigs of ten weeks
120 of age, free of porcine circovirus type 2, porcine reproductive respiratory syndrome virus
121 and Pestivirus. The groups were: group 1 or the Margarita group (numbered 1-6), group 2
122 or the Cat01group (7-12), and group 3 or the control, non-infected group (13-18). The
123 animals were experimentally infected by i.m. injection in the neck with 10⁵ TCID₅₀ of

124 CSFV Margarita, Cat01strains, and PBS, respectively. After infection, serum samples and
125 nasal swabs were collected at 0, 7, 13 and 17 days post-infection (DPI); whole blood
126 samples at 0, 7 and 13 DPI and tonsils samples were taken at the time of necropsy (for
127 Margarita group) or at 18 DPI (for Cat01 and control group). A trained veterinarian
128 recorded the rectal temperature and the clinical signs of disease daily in a blind manner.
129 Clinical signs compatible with a CSFV infection were anorexia, conjunctivitis, diarrhoea,
130 constipation, abdominal petechiae, and prostration. In addition, the presence or absence of
131 nervous symptoms was recorded independently. The clinical status of the animals were
132 scored from 0 to 7, rating fever, clinical signs (not including nervous symptoms) and
133 nervous symptoms as follows: 0 - no signs; 1 - mild pyrexia; 2 - pyrexia plus mild clinical
134 signs; 3 - mild-moderate clinical signs but absence of nervous disorders; 4 - slight nervous
135 disorders and moderate rest of clinical signs; 5 - moderate nervous disorders and
136 moderate–severe rest of clinical signs; 6 - severe clinical signs (including nervous
137 disorders); 7 - death. For ethical reasons, the animals were euthanised either when the
138 clinical score reached 5 or higher or when showing prostration and moderate–severe
139 nervous disorders. After euthanasia, an exhaustive necropsy was conducted in which the
140 presence of pathological signs in different organs and tissues was evaluated. The
141 experiments were approved by the Ethics Committee for Animal Experiments of the
142 University Autonomous of Barcelona (UAB) according to existing national and European
143 regulations.

144 **Detection of CSFV RNA**

145 The samples from sera, nasals swabs and tissue homogenates from nasal swabs and
146 tissues were processed following the methodology described previously (Díaz de Arce et
147 al., 1998; Pérez et al., 2011). The RNA was extracted from sera, nasals swabs and tissue
148 homogenates using the RNA viral isolation kit Nucleospin II according to the

149 manufacturer's instructions (Macherey-Nagel). In all cases, an initial volume of 150 μ L
150 was used to obtain a final volume of 50 μ L of RNA, which was stored at -80°C . The real-
151 time RT-PCR was performed using the Quantitect SYBR Green PCR kit (Qiagen GmbH,
152 Hilden, Germany). Specific CSFV detection was performed in serum, nasals swabs and
153 tonsil samples by using primer pairs targeting a region corresponding to the NS5B protein
154 (Díaz de Arce et al., 1998). The serum and nasals swab samples were collected at 0, 7, 13
155 and 17 DPI. The improved primer set concentration and thermocycling conditions have
156 been recently described (Pérez et al., 2011). The limits of detection used were 1, 10 and
157 10^2 gene copies/ μ l for nuclease-free water, serum and tissue homogenate, respectively.

158 **Antibody detection**

159 To evaluate the humoral response, we followed the methodology as described previously
160 (Ganges et al., 2005; Tarradas et al., 2010; Tarradas et al., 2011). Serum samples were
161 collected from all animals at 0, 7, 13 and 17 DPI. The serums were tested for neutralising
162 antibodies using a neutralisation peroxidase-linked assay (NPLA) (Terpstra et al., 1984),
163 and titres were expressed as the reciprocal dilution of serum that neutralised 100 TCID₅₀
164 of Catalunya and Margarita strains, respectively, in 50% of the culture replicates.
165 Detection of E2 specific antibodies was performed using a commercial ELISA
166 (CEDITEST; Lelystad); samples were considered to be positive when blocking
167 percentages $\geq 40\%$, following the manufacturer's recommendations.

168 **ELISA for IFN- α and IFN- γ detection in serum samples**

169 Anti-IFN- α monoclonal antibodies (K9 and K17) and IFN α recombinant protein (PBL
170 Biomedical Laboratories, Piscataway, New Jersey) were used in the ELISA assay
171 (Nowacki and Charley et al., 1993; Guzylack-Piriou et al., 2004). For IFN- γ detection,
172 mAbs P2G10, P2C11 and IFN- γ recombinant protein SD066 were used (BD Biosciences),

173 (De Antonio et al., 1998). The 3,3',5,5'- tetramethylbenzidine (TMB) was from Sigma.
174 For each ELISA test, the cut-off value was calculated as the average of the optical density
175 of negative controls (blank and negative serums before CSFV infections) plus three
176 standard deviations. Cytokine concentrations in the serum were determined using a
177 regression line consisting of the optical densities of the cytokine standards used in each
178 test.

179 **PBMCs and Elispot assay for CSFV-specific IFN- γ detection**

180 Peripheral blood mononuclear cells (PBMCs) were obtained from total blood collected
181 at 0, 7 and 13 DPI and were separated by density-gradient centrifugation with Histopaque
182 1077 (Sigma). The PBMC number and the viability of the cells were obtained by staining
183 with Trypan Blue (Ganges et al., 2005). The Elispot assay to detect CSFV-specific IFN γ
184 cells was performed as previously described (Tarradas et al., 2010). In brief, plates (Costar
185 3590, Corning) were coated overnight with 5 μ g/ml of capture antibody (P2G10,
186 Pharmigen). Detection was performed using a biotinylated antibody (P2C11,
187 Pharmigen). A number of samples with 5×10^5 PBMC/well were plated in triplicate at 0.1
188 multiplicity of infection (MOI) of the CSFV Cat01 and Margarita strains, respectively.
189 Controls were incubated in the absence of virus or with phytohaemagglutinin (PHA) (10
190 μ g/ml). The counts of spots in the media for mock-stimulated wells were considered to be
191 the baseline for the calculation of antigen-specific frequencies of IFN- γ producing cells.

192 **Statistical Analysis**

193 All statistical analyses were performed using SPSS 15.0 software (SPSS Inc., Chicago,
194 IL, USA) using "pig" as the experimental unit. The significance level (α) was set at 0.05,
195 with statistical tendencies reported when $P < 0.10$. Throughout the trial, a non-parametric
196 test (Mann–Whitney) was chosen to compare values obtained from the immunological and

197 the clinical parameters between groups. This non-parametric analysis was chosen due to
198 the number of animals used in each experimental group.

199 **RESULTS**

200 **Development of clinical signs**

201 The clinical score results showed that no pigs infected with the Cat01 strain reached
202 level 3 in the clinical score value (grading from 0 to 7), and moreover, level 2 (pyrexia
203 plus mild clinical signs) was not observed until days 11 to 15 post-infection (Fig. 1 and
204 2). Animals infected with the Margarita strain developed pyrexia (rectal temperature
205 above 40°C) after day 2 days post-infection, with peaks reaching 42° C. From day 3 until
206 the end of the trial, the rectal temperature was always lower in the Cat01 group than in
207 the Margarita group ($p < 0.05$) (Fig. 1). In Margarita-strain infected pigs, moderate and
208 severe clinical signs (≥ 4 points in score value), such as anorexia, conjunctivitis,
209 diarrhoea, constipation, abdominal petechiae, severe nervous signs, and prostration were
210 observed from day 5 up to day 12 -13 post-infection, when animals were euthanised for
211 ethical reasons after reaching level 5 or more in the clinical score valuation (Figure 1 and
212 2). The difference in the clinical score values for the Cat01 and the Margarita strains
213 were statistically significant from day 3 until the end of the trial ($p < 0.05$) (Fig. 2).

214 **Virus detection in serum, nasals swabs and tonsils**

215 Viral RNA in the serum and the nasals swabs was detected from day 7 post-infection in
216 all infected animals (Fig 3a, 3b). The RNA viral load was higher for the Margarita-strain
217 infected pigs in the serum and the nasals swabs than for the Cat01-strain infected pigs in
218 all samples evaluated ($p < 0.05$). Three out of the six animals infected with the Margarita
219 strain were tested at 13 DPI and showed similar RNA loads to the ones previously
220 determined at 7 DPI. The other three animals were either dead or were euthanised before

221 13 DPI since they reached a clinical score of 5 (Fig 2). At day 17 PI, only pigs infected
222 with the Cat01 strain could be evaluated. Interestingly, only one pig (number 11) could
223 clear the RNA virus in the serum and the nasals swabs (Fig. 3a, 3b). However, all Cat01-
224 infected pigs were positive to RNA viral detection in the tonsil samples. Nonetheless, the
225 RNA viral load was significantly higher for the Margarita-strain infected pigs than for
226 those infected with Cat01 ($p < 0.0005$) (Fig 3c).

227 **E2-specific antibodies and neutralising activity at 13 and 17 DPI detected in pigs** 228 **infected with Cat01 CSFV**

229 The serum Samples from the infected animals and the control pigs were analysed for the
230 presence of E2-specific antibodies (ELISA) and for neutralising antibodies (NPLA); no
231 antibodies were detected before 13 DPI (data not shown). Interestingly, the presence of
232 E2-specific antibodies detected by ELISA was observed in almost all infected animals
233 with the Cat01 strain from day 13 to 17 PI (Fig 4). This observation could not be made for
234 pigs infected with the Margarita strain since these animals died or were euthanised at
235 approximately 13 DPI, but at the time of death, there were no detectable antibody levels.
236 Neutralising antibodies were detected in two pigs infected with the Cat01 strain at 13 DPI
237 and in four pigs at 17 DPI (Table 1), particularly in animals number eight and eleven,
238 which had high neutralising antibody titrations (>32).

239 **Detection of levels of IFN- α and IFN- γ in the serum samples of Cat01-infected pigs**

240 The IFN- α cytokine levels in the serum samples obtained from Cat01- and Margarita-
241 strain infected animals were compared. The results showed significantly lower IFN- α
242 levels in all Cat01-infected pigs at day 3 ($p < 0.003$), and at day 7 post-infection ($p = 0.01$)
243 (see Fig 5a) than those observed in animals infected with the Margarita strain. Levels of
244 IFN- γ in the serum samples were determined by the ELISA technique (Fig 5 b). No

245 cytokines were detected for the Margarita-strain infected pigs (pig 1 to 6) or for the
246 control pigs. Interestingly, for the Cat01 strain, significant amounts of the cytokine were
247 detected in all samples. The ELISA results for IFN- γ were positive for five animals at
248 day 7 PI and for one animal at day 13 PI. The IFN- γ serum concentration was
249 significantly higher for Cat01- ($p=0.01$) than for Margarita-strain infected pigs (no
250 detectable levels) at day 7 post-infection (Fig. 5b).

251 **CSFV-specific IFN- γ producing cells after infection**

252 The Elispot results showed that the CSFV-specific IFN- γ producing cells response was
253 not detectable in pigs infected with the Margarita strain. Therefore, the response detected
254 in pigs infected with the Cat01 strain was statistically significant ($p=0.0025$) at 7 and 13
255 DPI (Fig. 5c).

256 **DISCUSSION**

257 The severity of the acute form of CSF responsible for the high mortality rate, the
258 auto-limiting nature of the virulent virus strains, and the pathogenesis of these forms have
259 been the subject of many studies (Hüsser et al., 2012; Summerfield et al., 2009; Ganges et
260 al., 2005; Sánchez-Cordon et al., 2005; Knoetig et al., 1999). Nevertheless, some animals
261 are likely to develop a chronic, or worse, unapparent form of the disease, which can be
262 misdiagnosed because of the confusing pathological picture and can be the source of virus
263 dissemination for an extended period of time (Weesendorp et al., 2010; Pérez et al., 2012,
264 Rout et al., 2012).

265 In most cases in the outbreak of early 2000 in Spain, the animals presented nonspecific
266 clinical signs consisting only of fever and apathy (Allepuz et al., 2007). In the present
267 work, we studied the infection caused by the Cat01 strain in domestic pigs, responsible for
268 the 2001- 2002 CSFV outbreak in the Catalonia region in Spain in comparison with the

269 infection caused by the Margarita strain, responsible for inducing the acute form of CSF in
270 the field (Diaz de Arce et al., 1999; Ganges et al., 2005). We focused on assessing the
271 impact of CSFV virulence on the kinetics of various cytokines involved in the innate and
272 the adaptive immune responses to infection as well as the humoral response generated and
273 its relation to the course of infection and the RNA CSFV load during the first weeks after
274 CSFV infection. In accordance with the data previously collected in the field (Allepuz et
275 al., 2007), the Cat01 strain caused a very mild form of the disease with almost unapparent
276 CSF clinical signs in the experimentally infected pigs (Fig. 1 and 2). Nevertheless, the
277 viral RNA load was detected in the majority of the samples tested, including in the tonsils
278 of Cat01-infected pigs at 18 DPI, although the detected RNA loads were significantly
279 lower than those detected in pigs infected with the virulent strain (Fig. 3). The results of
280 this study are in agreement with those suggesting that the RNA CSFV load detected in the
281 serum, the nasal swabs and the tonsils has a direct correlation between strain virulence and
282 the clinical signs developed (Donahue et al., 2012, Uttenthal et al., 2003, Weesendorp et
283 al., 2010).

284 Type I IFN responses are usually associated with the antiviral defences of the host. They
285 are induced by viral infection or double-stranded RNA (dsRNA), a by-product of viral
286 replication, and lead to the production of a broad range of antiviral proteins and
287 immunoactive cytokines (Haller et al., 2006). However, in accordance with previous
288 reports, in our in vivo experiment, we detected high levels of IFN- α in the sera of pigs
289 infected with the CSFV virulent strain after infection (Tarradas et al., 2010; Renson et al.,
290 2010; Rosen et al., 2013). In contrast, pigs infected with the Cat01 strain, which developed
291 a very mild or unapparent CSF disease, expressed delayed and lower levels of IFN- α (Fig.
292 5a), associating the IFN- α levels with the severity of the disease after infection. Similar
293 findings were obtained for the influenza virus strains, where low pathogenic strains could

294 be related to low levels of type-I IFN (Moulin et al., 2011). Our results support the
295 previously reported correlation between the exacerbation of the type-I IFN production in
296 the host by virulent CSFV strain, the degree of severity of the illness, and the level of
297 replication and detection of the CSFV (Fig. 1, 2 and 5a) (Renson et al., 2010).
298 Additionally, type-I IFN is known to promote apoptosis in infected and non-infected cells
299 (Sivaraman et al., 2011, Tanaka et al., 1998, Thyrell et al., 2002). It has been reported that
300 there is a very strong correlation between high IFN- α levels in pigs with the acute form of
301 CSF and lymphopenia caused by bystander apoptosis (Renson et al., 2010, Summerfield et
302 al., 2009, Sun et al., 2010). Likewise, the surviving animals infected with the Margarita
303 strain showed very severe clinical signs and leucopenia at 13 DPI (Figure 1, 2 and Table
304 2). Conversely, pigs infected with the less virulent strain (CAT01) showed a lower degree
305 of impact on the number of PBMC after infection (Table 2). The severe lymphocyte
306 depletion observed in the acute CSF reflects the failure of the adaptive response in pigs
307 infected with the Margarita strain (Ganges et al., 2005, Renson et al., 2010, Summerfield
308 et al., 2006).

309 In general, for highly virulent hemorrhagic fever viruses, the failure of the adaptive
310 immune responses can play a crucial role in its lethality (Bray et al., 2005). For example,
311 data from Ebola outbreaks indicate that failure to develop specific antibodies by the
312 second week of illness is predictive of death (Ksiazek et al., 1999, Sanchez et al., 2004).
313 Our results confirm the ineffectiveness of highly virulent CSFV strains to induce
314 neutralising antibodies after 13 days post infection (Ganges et al., 2005, Tarradas et al.,
315 2010).

316 It is noteworthy that previous studies indicate the role of specific IFN- γ
317 producing cells, mainly the CD4⁺CD8 α ^{low} double-positive T cells (memory), in the
318 elicited protection against CSFV (Suradhat et al., 2001; Tarradas et al., 2010; Graham et

319 al., 2012; Franzoni et al., 2013). After the infection with the low virulent strain Cat01 at
320 7 and 13 DPI, and despite the slight cell depletion detectable in PBMC (Table 2), the
321 PBMC in pigs infected with the low virulent Cat01 strain were capable of being activated
322 and produced some levels of IFN- γ response against CSFV (Fig. 4). Alongside the T-cell
323 response, the humoral response was detected in some of the low virulent infected pigs
324 (Cat01) among the 13 and 17 DPI, mainly in pigs 8, 10 and 11 (Figure 3 and Table 1).
325 Curiously, in the serum samples of these animals were detected the lowest load of
326 CSFV RNA, achieving clearance of the virus in one of them at 17 DPI. Similarly, in the
327 tonsils of two of these animals (pigs 10 and 11) the lowest viral load levels were
328 detected. Both the T cells and the humoral response generated in most of the infected
329 pigs with strain Cat01 could be related to the protection in the symptom progression of
330 CSF against this CSFV strain (Fig 1 and 2).

331 However, despite of the levels of protection that were observed, the Cat01 strain still
332 impaired the adaptive immunity against CSFV because not all animals developed a
333 detectable T-cell response and, in parallel, the adequate levels of antibodies that are
334 capable of neutralising the virus in both the serum and the tissues. Thus, RNA viral
335 detection was observed in the tonsils from all Cat01 infected pigs, corroborating the role
336 of this lymphoid organ in the persistence of CSFV (Ganges et al., 2008), principally in
337 pigs infected with a low virulence strain. Thus, pigs with nearly unapparent CSF
338 symptoms, and even negative for CSFV RNA in the serum that are positive in lymphoid
339 tissues, and with a chance of excreting the virus (even though in low levels), can hamper
340 the diagnosis and the control of this viral infection. This fact can encourage the spread of
341 the virus and its persistence in endemic countries. These types of results explain in part
342 why CSFV of low virulence strains are capable of producing an outbreak that goes

343 unnoticed during the first months of infection (Allepuz et al., 2007) and how difficult its
344 resulting eradication is in endemic countries today.

345

346 **Author Disclosure Statement**

347 None

348 **ACKNOWLEDGMENTS**

349 We thank Iván Cordón and David Solanes for their help in the animal facilities. This
350 research was supported by grant AGL2012-38343 from Spanish government.

351 **REFERENCES**

352 Allepuz A, Casals J, Pujols J, Jové R, Selga I, Porcar J, Domingo M. 2007. Descriptive
353 epidemiology of the outbreak of classical swine fever in Catalonia (Spain),
354 2001/02. *Vet. Rec.* 12:398-403.

355 Alvarez B, Sánchez C, Bullido R, Marina A, Lunney J, Alonso F, Ezquerra A,
356 Dominguez J. 2000. A porcine cell surface receptor identified by monoclonal
357 antibodies to SWC3 is a member of the signal regulatory protein family and
358 associates with protein-tyrosine phosphatase SHP-1. *Tissue Antigens.* 55:342-51.

359 Bray M. 2005. Pathogenesis of viral hemorrhagic fever. *Curr. Opin. Immunol.* 17:399-
360 403.

361 Cheville NF, Mengeling WL. 1969. The pathogenesis of chronic hog cholera (swine
362 fever). Histologic, immunofluorescent, and electron microscopic studies. *Lab.*
363 *invest.* 20: 261-74.

364 Denham S, Zwart RJ, Whittall JT, Pampusch M, Corteyn AH, Bianchi AT, Murtaugh
365 MP, Parkhouse RM, Tlaskalova H, Sinkora J, Sinkora M, Rehakova Z. 1998.
366 Monoclonal antibodies putatively identifying porcine B cells. *Vet. Immunol.*
367 *Immunopathol.* 60:317-28.

368 Diaz de Arce H, Nuñez JI, Ganges L, Barreras M, Frías MT, Sobrino F. 1998. An RT-
369 PCR assay for the specific detection of classical swine fever virus in clinical
370 samples. *Vet. Res.* 29:431-40.

371 Diaz de Arce H, Nuñez JI, Ganges L, Barreras M, Teresa Frías M, Sobrino F. 1999.
372 Molecular epidemiology of classical swine fever in Cuba. *Virus Res.* 64:61-7.

373 De Antonio, E.M., Husmann, R.J., Hansen, R., Lunney, J.K., Strom, D., Martin, S.,
374 Zuckermann, F.A. 1998. Quantitative detection of porcine interferon-gamma in
375 response to mitogen, superantigen and recall viral antigen1. *Veterinary*
376 *Immunology and Immunopathology*, 61: 265-277.

377 Donahue BC, Petrowski HM, Melkonian K, Ward GB, Mayr GA, Metwally S. 2012.
378 Analysis of clinical samples for early detection of classical swine fever during
379 infection with low, moderate, and highly virulent strains in relation to the onset of
380 clinical signs. *J. Virol. Methods.* 179:108-15.

381 Ezquerra A, Revilla C, Alvarez B, Pérez C, Alonso F, Domínguez J. 2009. Porcine
382 myelomonocytic markers and cell populations. *Dev. Comp. Immunol.* 33:284-98.

383 Franzoni G, Kurkure NV, Edgar DS, Everett HE, Gerner W, Bodman-Smith KB, Croke
384 HR, Graham SP. 2013. Assessment of the phenotype and functionality of porcine
385 CD8 T cell responses following vaccination with live attenuated classical swine
386 fever virus (CSFV) and virulent CSFV challenge. *Clin Vaccine Immunol.* 10:1604-
387 16

388 Ganges L, Barrera M, Nuñez JI, Blanco I, Frias MT, Rodríguez F, Sobrino F. 2005. A
389 DNA vaccine expressing the E2 protein of classical swine fever virus elicits T cells
390 response that can prime for rapid antibody production and confer protection upon
391 viral challenge. *Vaccine.* 23: 3741-52.

392 Ganges L, Núñez JI, Sobrino F, Borrego B, Fernández-Borges N, Frias-Lepoureau MT,
393 Rodríguez F. 2008. Recent advances in the development of recombinant vaccines
394 against classical swine fever virus: cellular responses also play a role in protection.
395 *Vet. J.* 177: 169-177.

396 Gerner W, Hammer SE, Wiesmüller KH, Saalmüller A. 2009. Porcine T lymphocytes
397 and NK cells--an update. *Dev Comp Immunol.* 33:310-20.

398 Graham SP, Haines FJ, Johns HL, Sosan O, La Rocca SA, Lamp B, Rümenapf T,
399 Everett HE, Crooke HR. , 2012. Characterisation of vaccine-induced, broadly
400 cross-reactive IFN- γ secreting T cell responses that correlate with rapid protection
401 against classical swine fever virus. *Vaccine.* 30:2742-8.

402 Guzylack-Piriou L, Balmelli C, McCullough KC, Summerfield A. 2004. Type-A CpG
403 oligonucleotides activate exclusively porcine natural interferon-producing cells to
404 secrete interferon-alpha, tumour necrosis factor-alpha and interleukin-12.
405 *Immunology.* 112:28-37.

406 Haller O, Kochs G, Weber F. 2006. The interferon response circuit: Induction and
407 suppression by pathogenic viruses. *Virology.* 344: 119-130.

408 Haverson K, Zuckermann F, Saalmüller A, Lipp J, Aasted B, Stokes CR. 1998.
409 Summary of workshop findings for porcine adhesion molecule subgroup. *Vet.*
410 *Immunol. Immunopathol.* 60: 351-65.

411 Heinz FX, Stiasny K, Allison SL. 2004. The entry machinery of flaviviruses. *Arch.*
412 *Virol. Suppl.* 18:133-7.

413 Hüsser L, Ruggli N, Summerfield A. 2012. N(pro) of classical swine fever virus prevents
414 type I interferon-mediated priming of conventional dendritic cells for enhanced
415 interferon- α response. *J. Interferon Cytokine Res.* 32:221-9.

416 Knoetig SM, Summerfield A, Spagnuolo-Weaver M, McCullough KC. 1999.
417 Immunopathogenesis of classical swine fever: role of monocytic cells.
418 Immunology. 97:359-66.

419 Ksiazek TG, Rollin PE, Williams AJ, Bressler DS, Martin ML, Swanepoel R, Burt FJ,
420 Leman PA, Khan AS, Rowe AK, Mukunu R, Sanchez A Peters CJ. 1999. Clinical
421 virology of Ebola hemorrhagic fever (EHF): virus, virus antigen, and IgG and IgM
422 antibody findings among EHF patients in Kikwit, Democratic Republic of the
423 Congo, 1995. J. Infect. Dis. 179:177-87.

424 Li MO, Flavell RA. 2008. Contextual regulation of inflammation: a duet by transforming
425 growth factor-beta and interleukin-10. Immunity. 28:468-76.

426 Lohse L, Nielsen J, Uttenthal A. 2012. Early pathogenesis of classical swine fever virus
427 (CSFV) strains in Danish pigs. Vet. Microbiol., in press.

428 Lunney JK. 1994. Current status of the swine leukocyte antigen complex. Vet. Immunol.
429 Immunopathol. 43:19-28.

430 Mittelholzer C, Moser C, Tratschin JD, Hofmann MA. 2000. Analysis of classical swine
431 fever virus replication kinetics allows differentiation of highly virulent from
432 avirulent strains. Vet. Microbiol. 74:293-308.

433 Moennig V. 2000. Introduction to classical swine fever: virus, disease and control
434 policy. Vet. Microbiol. 73: 93-102.

435 Mosser DM, Zhang X. 2008. Interleukin-10: new perspectives on an old cytokine.
436 Immunol. Rev. 226:205-18.

437 Moulin HR, Liniger M, Python S, Guzylack-Piriou L, Ocaña-Macchi M, Ruggli N,
438 Summerfield A. 2011. High interferon type I responses in the lung, plasma and
439 spleen during highly pathogenic H5N1 infection of chicken. Vet. Res. 42: 6-11.

440 Nielsen J, Lohse L, Rasmussen TB, Uttenthal A. 2010. Classical swine fever in 6- and
441 11-week-old pigs: Haematological and immunological parameters are modulated in
442 pigs with mild clinical disease. *Vet. Immunol. Immunopathol.* 138: 159-173.

443 Nowacki, W., Charley, B. 1993. Enrichment of coronavirus-induced interferon-
444 producing blood leukocytes increases the interferon yield per cell: A study with pig
445 leukocytes. *Research in Immunology*, 144:111-120.

446 Pauly T, König M, Thiel HJ, Saalmüller A. 1998. Infection with classical swine fever
447 virus: effects on phenotype and immune responsiveness of porcine T lymphocytes.
448 *J. Gen. Virol.* 79: 31-40.

449 Pérez LJ, Díaz de Arce H, Perera CL, Rosell R, Frías MT, Percedo MI, Tarradas J,
450 Dominguez P, Núñez JI, Ganges L. 2012. Positive selection pressure on the B/C
451 domains of the E2-gene of classical swine fever virus in endemic areas under C-
452 strain vaccination. *Infect. Genet. Evol.* 12:1405-1412.

453 Pérez LJ, Díaz de Arce H, Tarrades J, Rosell R, Perera CL, Muñoz M, Frías MT,
454 NuñezJI, Ganges L. 2011. Development and validation of a novel SYBR Green
455 real-time RT-PCR assay for the detection of classical swine fever virus evaluated
456 on different real time PCR platforms. *J. Virol. Methods.* 174:53-9.

457 Pescovitz MD, Lunney JK, Sachs DH. 1984. Preparation and characterization of
458 monoclonal antibodies reactive with porcine PBL. *J. Immunol.* 133:368-75.

459 Reed LJ, Muench H. 1938. A simple method of estimating fifty percent endpoints. *Am.*
460 *J. Hyg.* 27:493-497.

461 Renson P, Blanchard Y, Le Dimma M, Felix H, Cariolet R, Jestin A, Le Potier MF.
462 2010. Acute induction of cell death-related IFN stimulated genes (ISG)
463 differentiates highly from moderately virulent CSFV strains. *Vet. Res.* 41: 7-22.

464 Rout M, Saikumar G. 2012. Virus load in pigs affected with different clinical forms of
465 classical swine fever. *Transbound. Emerg. Dis.* 59:128-33.

466 von Rosen T, Lohse L, Nielsen J, Uttenthal A. 2013. Classical swine fever virus
467 infection modulates serum levels of INF- α , IL-8 and TNF- α in 6-month-old pigs.
468 *Res Vet Sci.* 95:1262-7.

469 Saalmuller A, Aasted B, Canals A, Dominguez J, Goldman T, Lunney JK, Maurer S,
470 Pescovitz MD, Pospisil R, Salmon H, *et al.* 1994. Analyses of mAb reactive with
471 porcine CD8. *Vet. Immunol. Immunopathol.* 43:249-54.

472 Sanchez A, Lukwiya M, Bausch D, Mahanty S, Sanchez AJ, Wagoner KD, Rollin PE.
473 2004. Analysis of human peripheral blood samples from fatal and nonfatal cases of
474 Ebola (Sudan) hemorrhagic fever: cellular responses, virus load, and nitric oxide
475 levels. *J. Virol.* 78:10370-7.

476 Sánchez-Cordon PJ, Núñez A, Salguero FJ, Pedrera M, Fernández de Marco M, Gómez-
477 Villamandos JC. 2005. Lymphocyte apoptosis and thrombocytopenia in spleen
478 during swine fever:role of macrophages and cytokines. *Vet. Pathol.* 42:477-88.

479 Shen H, Pei J, Bai J, Zhao M, Ju C, Yi L, Kang Y, Zhang X, Chen L, Li Y, Wang J,
480 Chen J. 2011. Genetic diversity and positive selection analysis of classical swine
481 fever virus isolates in south China. *Virus Genes.* 43:234-42.

482 Sivaraman V, Zhang L, Su L. 2011. Type I Interferon contributes to CD4⁺ T cell
483 depletion induced by infection with HIV-1 in the human thymus. *J. Virol.* 85:
484 9243-6.

485 Summerfield A, Alves M, Ruggli N, de Bruin MG, McCullough KC. 2006. High IFN-
486 alpha responses associated with depletion of lymphocytes and natural IFN-
487 producing cells during classical swine fever. *J. Interferon Cytokine Res.* 26:248-55.

488 Summerfield A, Hofmann MA, McCullough KC. 1998. Low density blood granulocytic
489 cells induced during classical swine fever are targets for virus infection. *Vet.*
490 *Immunol. Immunopathol.* 63:289-301.

491 Summerfield A, Knötig SM, McCullough KC. 1998. Lymphocyte apoptosis during
492 classical swine fever: implication of activation-induced cell death. *J. Virol.*
493 72:1853-61.

494 Summerfield A, McCullough KC. 2009. The porcine dendritic cell family. *Dev. Comp.*
495 *Immunol.* 33:299-309.

496 Summerfield A, McNeilly F, Walker I, Allan G, Knoetig SM, McCullough KC. 2001.
497 Depletion of CD4⁺ and CD8^{high+} T-cells before the onset of viraemia during
498 classical swine fever. *Vet. Immunol. Immunopathol.* 78: 3-19.

499 Summerfield 2006

500 Sun J, Shi Z, Guo H, Tu C. 2010. Changes in the porcine peripheral blood mononuclear
501 cell proteome induced by infection with highly virulent classical swine fever virus.
502 *J. Gen. Virol.* 91: 2254-62.

503 Suradhat S, Intrakamhaeng M, Damrongwatanapokin S. 2001. The correlation of virus-
504 specific interferon-gamma production and protection against classical swine fever
505 virus infection. *Vet. Immunol. Immunopathol.* 83:177-89.

506 Suradhat S, Sada W, Buranapraditkun S, Damrongwatanapokin S. 2005. The kinetics of
507 cytokine production and CD25 expression by porcine lymphocyte subpopulations
508 following exposure to classical swine fever virus (CSFV). *Vet. Immunol.*
509 *Immunopathol.* 106: 197-208.

510 Tanaka N, Sato M, Lamphier MS, Nozawa H, Oda E, Noguchi S, Schreiber RD,
511 Tsujimoto Y, Taniguchi T. 1998. Type I interferons are essential mediators of
512 apoptotic death in virally infected cells. *Genes Cells.* 3:29-37.

513 Tarradas J, Argilaget JM, Rosell R, Nofrarias M, Crisci E, Cordoba L, Pérez-Martin E,
514 Diaz I, Rodríguez F, Domingo M, Montoya M Ganges L. 2010. Interferon-gamma
515 induction correlates with protection by DNA vaccine expressing E2 glycoprotein
516 against classical swine fever virus infection in domestic pigs. *Vet. Microbiol.* 142:
517 51-58.

518 Terpstra C, Bloemraad M, Gielkens AL. 1984. The neutralizing peroxidase-linked assay
519 for detection of antibody against swine fever virus. *Vet. Microbiol.* 9:113-120.

520 Thyrell L, Erickson S, Zhivotovsky B, Pokrovskaja K, Sangfelt O, Castro J,
521 Einhorn S, Grandér D. 2002. Mechanisms of Interferon-alpha induced apoptosis in
522 malignant cells. *Oncogene.* 21:1251-62.

523 Uttenthal A, Storgaard T, Oleksiewicz MB, de Stricker K. 2003. Experimental infection
524 with the Paderborn isolate of classical swine fever virus in 10-week-old pigs:
525 determination of viral replication kinetics by quantitative RT-PCR, virus isolation
526 and antigen ELISA. *Vet. Microbiol.* 92:197-212.

527 van Oirschot JT, De jong D, Huffels ND. 1983. Effect of interations with swine fever
528 virus on immune functions. II. Lymphocyte response to mitogens and enumeration
529 of lymphocyte subpopulations. *Vet Microbiol.* 8:81-95.

530 Wang Z, Shao Y, Guo Y, Yuan J. 2008. Enhancement of peripheral blood CD8⁺ T cells
531 and classical swine fever antibodies by dietary B-1,3/1,6-glucan supplementation in
532 weaned piglets. *Transbound. Emerg. Dis.* 55: 369-76.

533 Weesendorp E, Willems EM, Loeffen WL. 2010. The effect of tissue degradation on
534 detection of infectious virus and viral RNA to diagnose classical swine fever virus.
535 *Vet. Microbiol.* 141:275-81.

536 Weesendorp E, Backer J, Stegeman A, Loeffen W. 2011. Transmission of classical
537 swine fever virus depends on the clinical course of infection which is associated
538 with high and low levels of virus excretion. *Vet. microbiol.* 147: 262-73.

539 Wensvoort G, Terpstra C, Boonstra J, Bloemraad M, Van Zaane D. 1986. Production of
540 monoclonal antibodies against swine fever virus and their use in laboratory
541 diagnosis. *Vet. Microbiol.* 12:101-8.

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561 TABLES

562 Tarradas et al.,

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564 TABLE 1. Detection of neutralising antibodies titres after infection with the Cat01 strain
565 of CSFV

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568 TABLE 2. Total number of live PBMCs obtained per ml of blood at days 0 (T0) and 13
569 (T13) post CSFV infection (DPI)

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576 FIGURE 1. Clinical signs after CSFV infection. (a) Individual rectal temperature values
577 detected during the experiment. Pigs infected with the Margarita strain were euthanised
578 between 7 and 13 DPI. Pigs infected with the Cat01 strain or the control (uninfected)
579 pigs lived until 18 DPI, which was the end of the experiment. A temperature above 40°C
580 was considered fever.

581 (b) Mean and standard deviation of the rectal temperature recorded after CSFV infection
582 with the Margarita and the Cat01 strains. An asterisk indicates statistical significance
583 between the two groups from day 3 until day 12 ($p < 0.05$), and the Margarita strain score
584 is significantly higher.

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586 FIGURE 2. Clinical signs after CSFV infection. Mean of the clinical scores after CSFV
587 infection with the Margarita and Cat01 strains. An asterisk indicates statistical
588 significance between the two groups from day 3 until day 13 ($p < 0.05$), and the
589 Margarita strain score is significantly higher. The symbol † indicates euthanasia for
590 human reasons or death. Scores are defined in the material and methods section 2.

591

592 FIGURE 3. Detection of CSFV RNA through real time RT-PCR in serum, nasal swabs
593 and tonsils obtained from pigs infected with the CSFV Margarita and Cat01 strains. (a)
594 Shows the serum viral loads from pigs 1-6 infected with the Margarita strain at day 7 PI,
595 for all animals, and at day 13 for 3 animals because the other three were either
596 euthanised or dead. Serum viral loads from pigs 7-12 were analysed at days 7, 13 and
597 17pi and were found to be significantly lower than those obtained in Margarita infected
598 pigs ($p < 0.05$).

599 (b) Shows the nasal swab viral loads from pigs 1-6 infected with the Margarita strain at
600 day 7 PI, for all animals, and at day 13 for 3 animals because the other 3 were either
601 euthanised or dead. Nasal swabs viral loads from pigs 7-12 were analysed at days 7, 13
602 and 17pi and were found to be significantly lower than those obtained in Margarita
603 infected pigs ($p < 0.05$). (c) Depicts the tonsils viral loads of Margarita- and Cat01-
604 infected animals and highlights the differences between both groups ($p < 0.0005$).

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606 FIGURE 4. Antibody response against E2 glycoprotein detected by ELISA (in blocking
607 %) from pigs infected with the Cat01 strain at 13 and 17 DPI. Values above 40% were
608 considered positive.

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611 FIGURE 5. Serum levels of IFN- α , and IFN- γ and induction of CSFV-specific-IFN- γ
612 producing cells from pigs infected with the CSFV Margarita and Cat01 strains. (a) IFN-
613 α levels at days 0, 3, 7 and 13 PI. These levels are significantly higher in animals 1-6
614 (Margarita group) than in 7-12 (Cat01 group) at days 3 and 7 PI ($p=0.003$ and $p=0.01$,
615 respectively). (b) IFN- γ levels in serum can only be detected in pigs from the Cat01
616 group ($p=0.01$ at day 7 PI). (c) Lack of response to IFN- γ in the Elispot assay in pigs
617 infected with the Margarita strain and the detection of a response in some of the pigs
618 infected with the Cat01 strain at 7 and 13 DPI ($p<0.0025$).

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Table 1:

Neutralizing antibodies vs Cat 01 strain		
Pig	13 DPI	17 DPI
7	1/5	1/10
8	1/160	1/160
9	1/5	N
10	1/5	1/20
11	1/80	>1/640
12	N	1/20

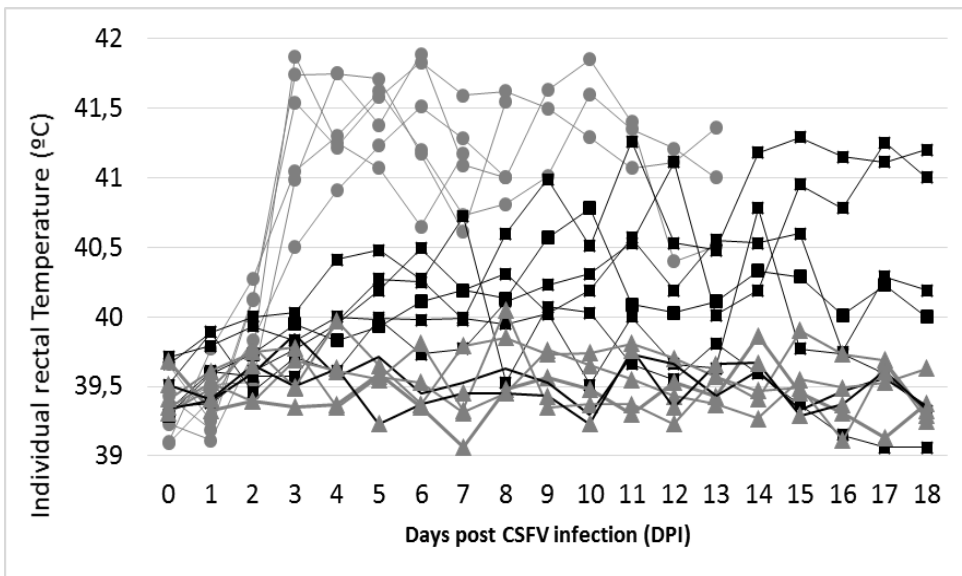
Table 2:

Pig	Live cells ^(a)	
	T0	T13
1	3.5	0.12
2	4.8	0.18
3	6.5	0.15
4	1.8	†
5	5.5	†
6	4.1	†
7	6.5	2.8
8	5.5	3.5
9	6.1	1.7
10	2.5	1.6
11	1.7	2.0
12	6.4	3.1
13	4.5	5.5
14	6.0	5.4

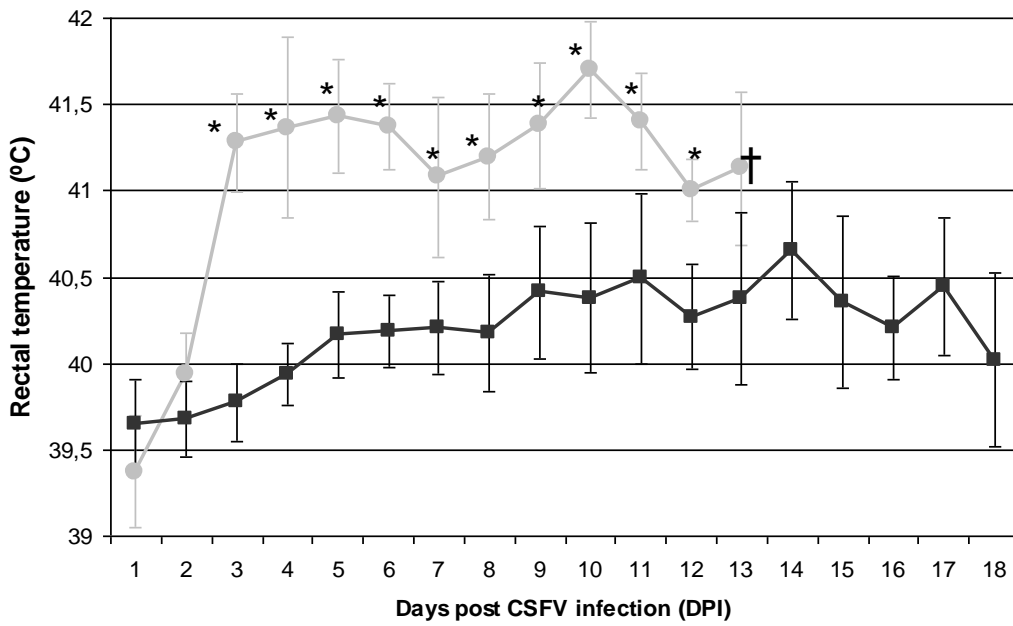
^a Total number of live PBMCs ($\times 10^6$) obtained per ml of blood at days 0 (T0) and 13 (T13) pi (DPI)
Pigs 1 to 6 infected with Margarita strain.
Pigs 7 to 12 infected with Cat01 strain.
Pigs 13 and 14 belong to the control group.

Figure 1:

a



b



● Pigs infected with Margarita strain

■ Pigs infected with Cat01 strain

▲ Control pigs

Figure 2:

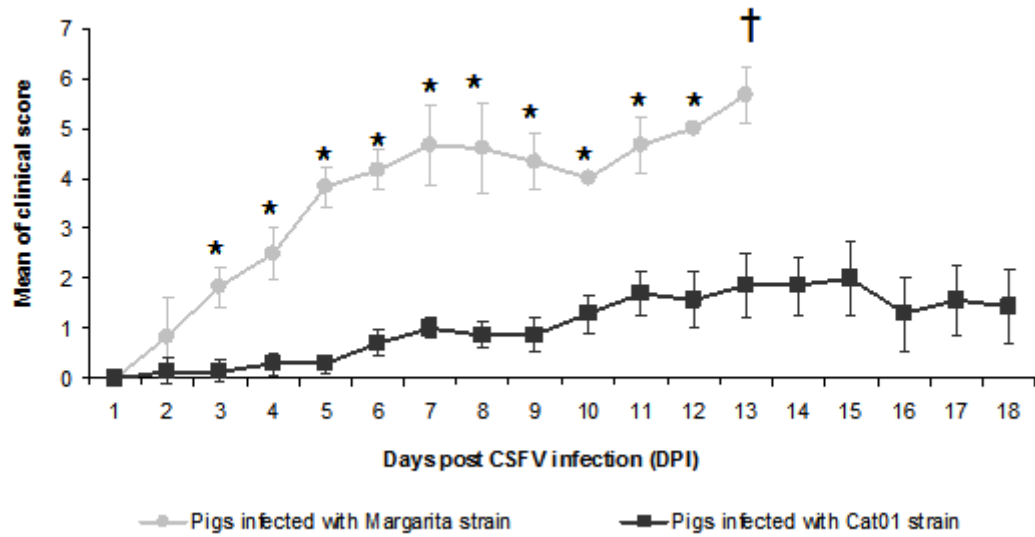
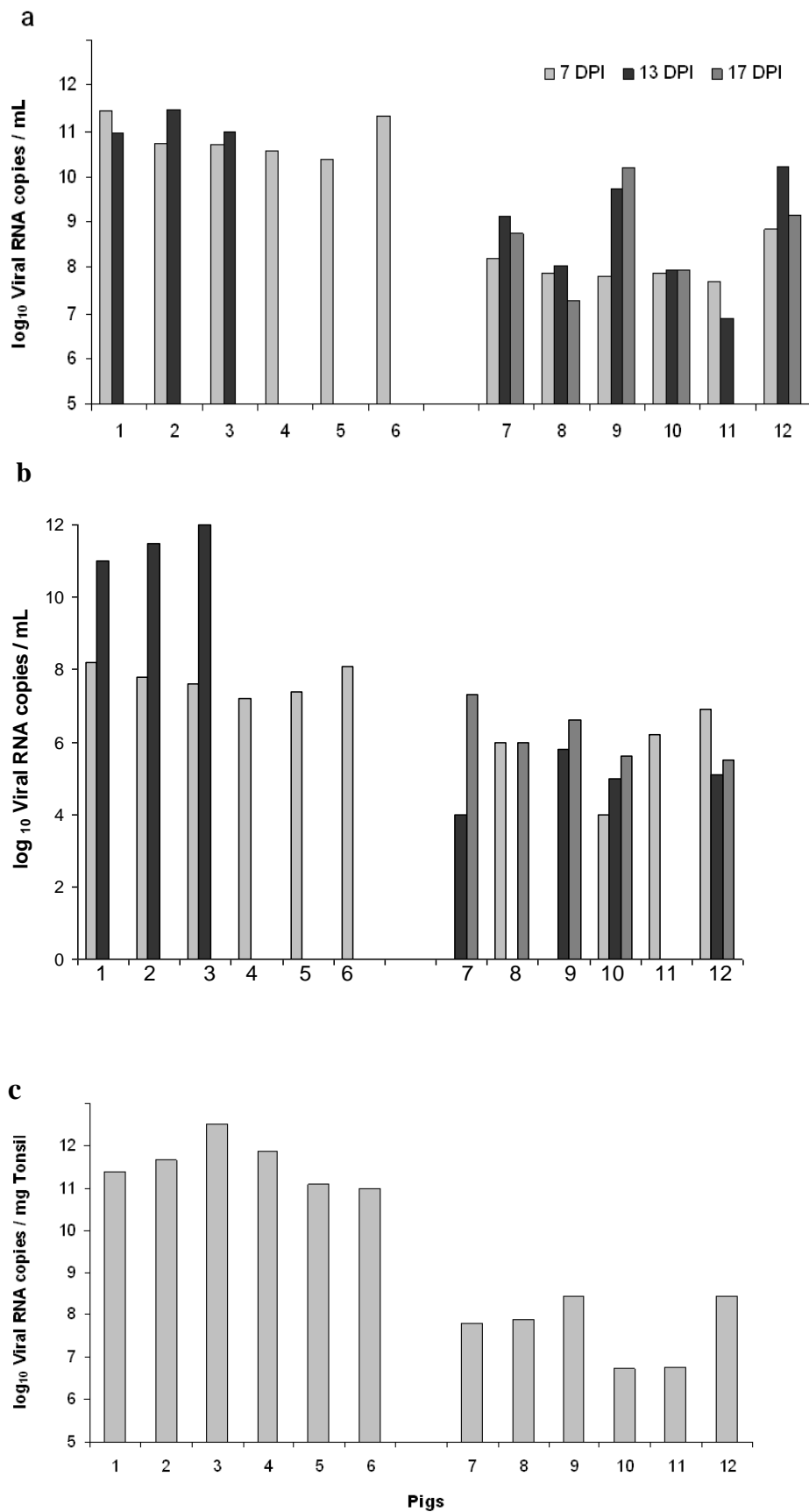
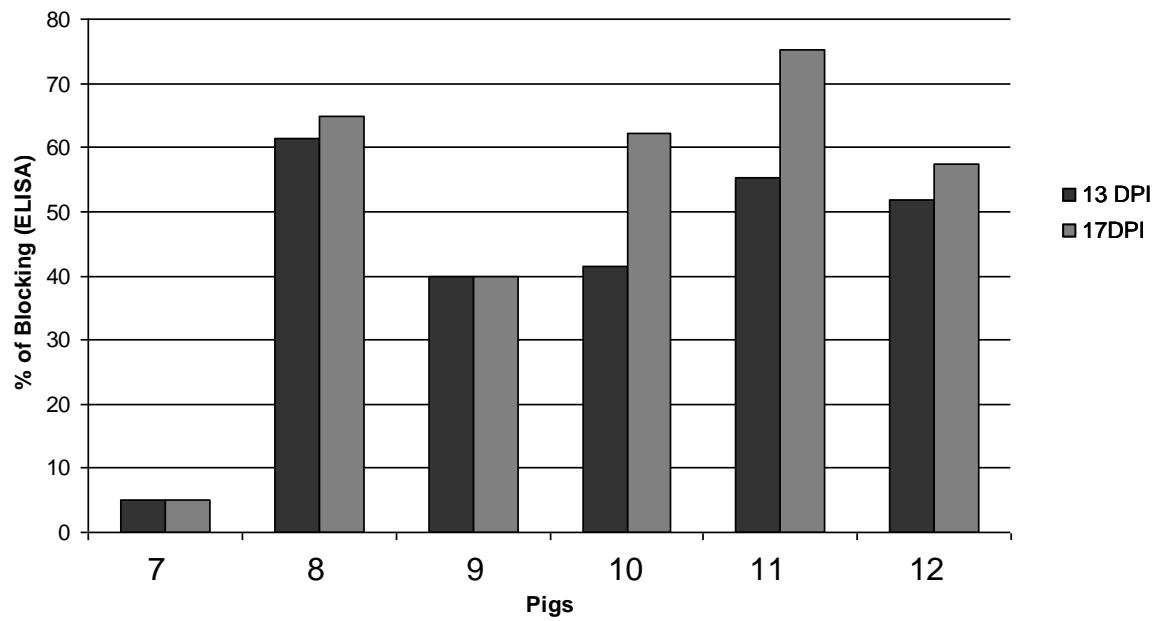


Figure 3:



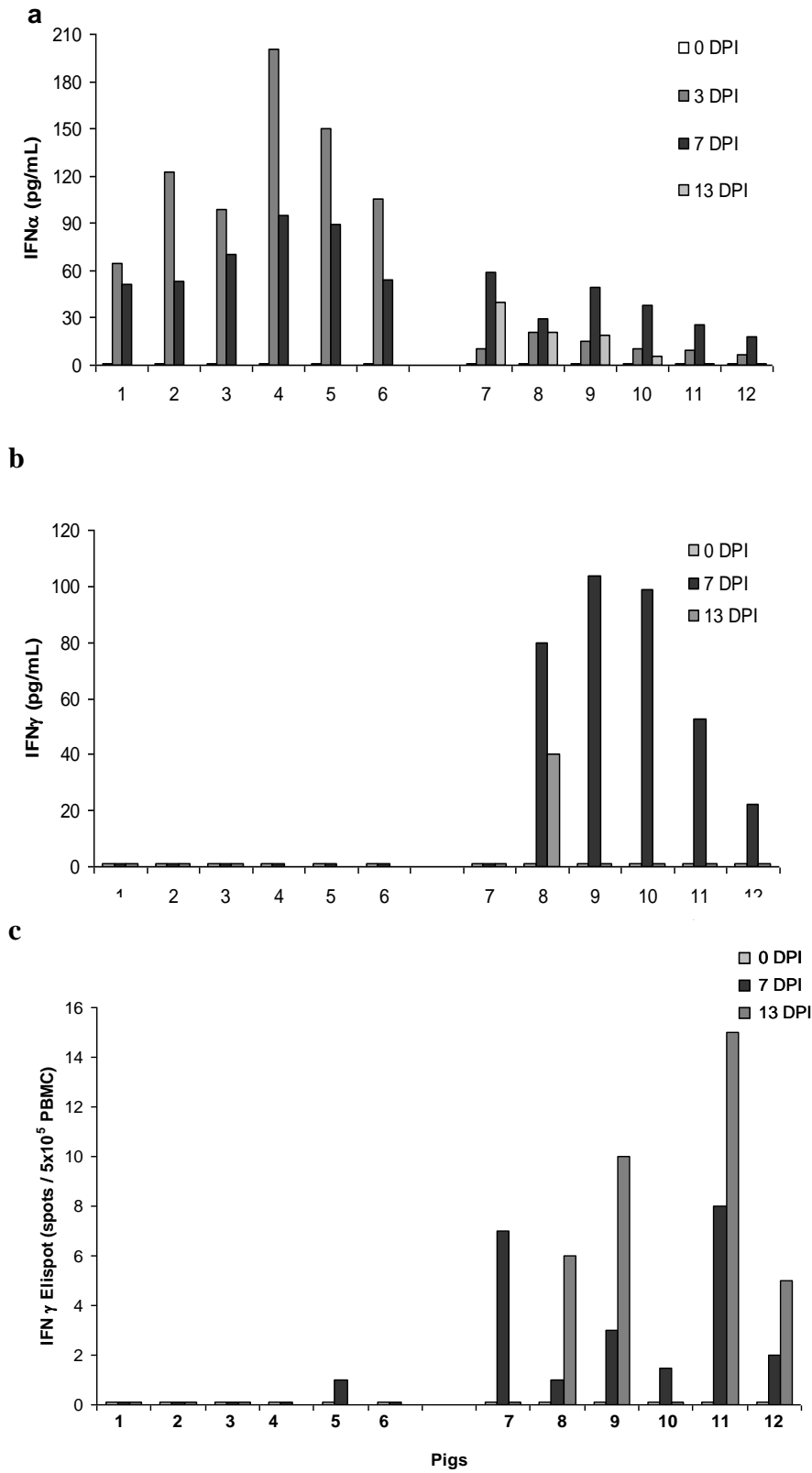
(Pigs 1 to 6: infected with Margarita strain / Pigs 7 to 12: infected with Cat01 strain)

Figure 4:



(Pigs 7 to 12: infected with Cat01 strain)

Figure 5:



(Pigs 1 to 6: infected with Margarita strain
 Pigs 7 to 12: infected with Cat01 strain)

