1	Detection of transmissible viral proventriculitis (TVP) and Chicken proventricular necrosis
2	virus (CPNV) in the United Kingdom
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26 Abstract

Increasing evidence suggests that a new birnavirus, named *Chicken proventricular necrosis* 27 virus (CPNV), is the aetiological agent of transmissible viral proventriculitis (TVP). The 28 29 present work aimed to explore the possible presence of both TVP and CPNV in the UK. Fortyfour chickens showing TVP-compatible gross lesions were classified into 3 groups based on 30 the histological lesions: i) TVP-affected chickens: lymphocytic infiltration and glandular 31 necrosis (n=15); ii) lymphocytic proventriculitis (LP)-affected chickens: lymphocytic 32 infiltration without necrosis (n=18); and iii) without proventriculitis (WP): no lymphocytic 33 infiltration or necrosis (n=11). Nine proventriculi (7 out of 15 corresponding to TVP, and 2 out 34 of 11 corresponding to LP) were positive for CPNV by RT-PCR. These results support the 35 previously suggested idea of CPNV as causative agent of TVP. Moreover, this data shows that 36 37 CPNV can also be detected in a number of cases with LP, which do not fulfil the histological TVP criteria. Phylogenetic analysis of partial sequences of gene VP1 showed that British 38 CPNV sequences were closer to other European CPNV sequences and might constitute a 39 different lineage from the American CPNV. TVP cases with negative CPNV PCR results may 40 be due to chronic stages of the disease or to the reduced PCR sensitivity on formalin-fixed 41 paraffin embedded tissues. However, involvement of other agents in some of the cases cannot 42 totally be ruled out. As far as the authors are aware, this is the first peer-reviewed report of 43 TVP as well as of CPNV in the UK, and the first exploratory CPNV phylogenetic study. 44

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46 Keywords: Birnavirus; *Chicken proventricular necrosis virus* (CPNV); transmissible viral
47 proventriculitis (TVP); natural infection; poultry.

49 Introduction

Transmissible viral proventriculitis (TVP) is an infectious viral disease affecting chickens, 50 which is reported to have a significant economic impact on a global scale (Dormitorio et al., 51 52 2007; Guy et al., 2005). Affected chickens display non-specific clinical signs, which may include stunted growth, pallor and the presence of incompletely digested food in the faeces 53 54 (Goodwin et al., 1996). Flocks affected by TVP typically do not show a significant increase in 55 mortality (Hafner and Guy, 2013). TVP most commonly affects broiler chickens of four to five weeks of age (Bayyari et al., 1995). However, the disease has also been identified in both 56 57 broiler breeders and layer hens in the age range of nine to twenty weeks (Marusak et al., 2012). 58 TVP results in enlarged, fragile proventriculi which contain characteristic microscopic lesions (Goodwin et al., 1996; Hafner and Guy, 2013). Chicken flocks are diagnosed with TVP based 59 on the presence of histological lesions in the proventriculus: necrosis of oxynticopeptic cells, 60 61 lymphocytic infiltration and hyperplastic ductal epithelium which replaces the glandular epithelium (metaplasia) (Goodwin et al., 1996; Hafner and Guy, 2013). No specific control 62 measures or treatments are currently recommended for TVP (Hafner and Guy, 2013). 63

TVP was first described in the Netherlands almost 40 years ago (Kouwenhoven et al., 1978). 64 65 Since then, it has been reported in several countries in North-America (Guy et al., 2011b; Noiva et al., 2015), Europe (Grau-Roma et al., 2010; Marguerie et al., 2011) and Asia (Kim et al., 66 2015). Since the emergence of the disease, there has been considerable discussion as to the 67 aetiological agent responsible. Several viruses have been suggested including: an adenovirus 68 (Kouwenhoven et al., 1978), a reovirus (Jones, 2000), Infectious bronchitis virus (IBV) (Yu et 69 70 al., 2001), Infectious bursal disease virus (IBDV) (Huff et al., 2001), and a picornavirus (Kim et al., 2015). Few years ago, a new birnavirus, named Chicken proventricular necrosis virus 71 (CPNV), was detected in naturally and experimentally reproduced TVP-affected cases in USA 72 and proposed to be its cause (Guy et al., 2011a; Guy et al., 2011b). CPNV has subsequently 73

been detected in a few other studies in TVP-affected broiler chickens in France and USA
(Marguerie et al., 2011; Noiva et al., 2015).

Despite few cases with proventricular lesions compatible to TVP have previously been reported in the UK¹ (Randall & Reece, 1996), as far as the authors are aware, there is no peer-reviewed report indicating the presence of neither TVP nor CPNV in the United Kingdom (UK). Based on the recent description of both TVP (Grau-Roma et al., 2010; Marguerie et al., 2011) and CPNV (Marguerie et al., 2011) in Europe, the aims of this study were to determine whether TVP and CPNV are present in chickens in the UK, as well as to evaluate their association suggested previously.

83 Materials and methods

Study design. At the end of 2014, a prospective study was designed based on the collection 84 85 of samples from chicken post-mortems performed by poultry clinicians in different locations of the UK. Clinicians were asked to send proventricular samples fixed in 10% formalin to the 86 School of Veterinary Medicine and Science (SVMS) at the University of Nottingham when 87 TVP-compatible gross lesions were observed. These lesions included thickening of the 88 proventricular wall, dilation of proventriculus and/or evidence of white spots visible through 89 90 the proventricular serosa. Clinicians were also asked to report any other relevant abnormalities observed during the post-mortem examination. 91 In addition, chicken cases received at the Veterinary Pathology Service (VPS) of the SVMS 92 between January 2014 and June 2015 having a diagnosis of 'lymphocytic and/or necrotizing 93

94 proventriculitis' were included in the study as suspected TVP-affected cases.

¹ VLA (currently APHA) quarterly surveillance report: January-March, Volume 14, No. 1, page 14 (<u>http://www.thepoultrysite.com/articles/1765/uk-poultry-disease-quarterly-</u> surveillance-report-januarymarch-2010).

95 Histopathology. A complete cross-section on the central area of the proventriculus was performed from each chicken, routinely processed for histology and stained with haematoxylin 96 and eosin. Proventriculi were assessed for the three key histopathological findings which are 97 98 characteristic of TVP: (i) glandular lymphocytic infiltration; (ii) hyperplastic and metaplastic changes of ductal epithelial cells; and (iii) necrosis of oxynticopeptic cells. These parameters 99 were semi-quantified as follows: - (absence), + (>0 to 10% of the glands affected), ++ (>10 to 100 50% of the glands affected), +++ (>50% of the glands affected). In addition, the presence of 101 necrosis was also assessed as the percentage of glandular parenchyma affected in the most 102 103 affected gland, following the same percentages as described above. A mean of the 2 necrosis scores was then calculated, and mean scores between severity levels (e.g. +/++) were rounded 104 105 up to the higher level. The presence of inflammatory infiltrate within the lamina propria was 106 not taken into consideration, as it is reported to be a frequent finding in healthy birds (Kadhim *et al.*, 2011). 107

Based on the histopathological results, chickens were allocated a case status as follows: (i) transmissible viral proventriculitis (TVP)-affected chickens: lymphocytic infiltration and necrosis present in the proventriculus; (ii) lymphocytic proventriculitis (LP)-affected chickens: lymphocytic infiltration without necrosis present in the proventriculus; (iii) chickens without proventriculitis (WP): no lymphocytic infiltration or necrosis present in the proventriculus.

RNA extraction and RT-PCR. RNA was extracted from all formalin-fixed paraffin embedded (FFPE) proventriculi and tested subsequently by RT-PCR for CPNV. RNA extraction was done using four 25 µm-sections of each sample. Briefly, the extraction method used incubation with xylol (twice) followed by centrifugation. Pellet re-suspension was performed first with ethanol and second with a digestion buffer containing Proteinase K (Roche, Mannheim, Germany). After overnight incubation at 56°C and centrifugation, supernatant was mixed with TRIzol[®] Reagent (Invitrogen, 15596-018). Samples were then homogenized with chloroform (Sigma,

C2432) and centrifuged. The transparent phase was discharged and the pellet was mixed with
isopropanol (Sigma, I9516). Two more centrifugations followed by addition of cold ethanol
were performed. Finally, the pellet was left to dry and received 25 μl of warm RNase-free
water.

A RT-PCR procedure was performed to amplify a 171 nucleotide (nt) sequence within the VP1
gene of CPNV using primers and protocols described previously (Guy et al., 2011b). FTA cards
with proventricular imprints from positive CPNV cases, kindly provided by Dr. Guerin
(National Veterinary School of Toulouse, France), were used as positive controls.

Sequencing of RT-PCR product and phylogenetic studies. The amplified products from the positive CPNV RT-PCR cases were purified using Mini Elute Gel Extraction Kit (Qiagen, Valencia, CA). Sequencing reactions were performed with ABI Prism BigDye Terminator Cycle Sequencing v.31 Ready Reaction (Applied Biosystems, Foster City, CA), and analysed using an ABI Prism model 3730 automated sequencer (Applied Biosystems, Foster City, CA). Positive and negative controls of extraction and amplification were added to each batch of samples tested.

Partial VP1 CPNV sequences obtained from British cases were compared with the sequence of 135 the American CPNV isolate R11/3 (Guy et al., 2011a) available in the Genbank 136 (http://www.ncbi.nlm.nih.gov, accession number HM038436.1), partial sequences obtained 137 from Spanish cases (Costa et al., submitted for publication) and the positive control (FTA card) 138 using MEGA6 (Molecular Evolutionary Genetics Analysis version 6.0) software (Tamura et 139 al., 2013). Sequences were aligned using ClustalW method. A nucleotide distance matrix 140 between sequences was computed to infer phylogenies and a Neighbor-joining (NJ) 141 phylogenetical tree was generated. The partial VP1 CPNV sequences reported in this work 142 have been deposited at GenBank under accession numbers KU933595 to KU933603. 143

Statistical analyses. Minitab version 17 was used for statistical analyses. Values -, +, ++ and +++ from the histological assessment were converted to 0, 1, 2 and 3, respectively, prior to the analyses. The distribution of variables was assessed using the Ryan-Joiner test. Kruskal-Wallis and Mann-Whitney U tests were used to assess for differences in variables between case statuses for nonparametric data.

149 **Results**

Epidemiological data and case status allocation. Forty-four chickens were included in this 150 study (Table 1). Thirty-nine chickens came from the prospective study (chickens with TVP-151 compatible gross lesions, chickens 1 to 39) and five chickens were received in the VPS and 152 were selected due to the presence of lymphocytic infiltration and/or necrosis of oxynticopeptic 153 154 cells (suspected TVP-affected cases, chickens 40 to 44). All the chickens were received between April 2014 and June 2015. The farm postcode was provided in 40 out of the 44 155 chickens, showing that the chickens came from 17 different farms located in England (n=16) 156 and Wales (n=1). The number of chickens received per farm ranged between 1 and 7. All the 157 received chickens were reported to have thickened and/or dilated proventriculus (Figure 1). 158 159 Most of the received cases corresponded to broiler chickens (42 out of 44), 1 corresponded to a layer hen (chicken 44), while the age of the chicken was not indicated in the remaining case. 160 Out of the 44 chickens studied, a total of 15 (34%) were classified as TVP-affected chickens, 161 18 (41%) as LP-affected chickens, and 11 (25%) as chickens without proventriculitis (WP) 162 (Table 1). Excluding the layer hen, the mean \pm SD age for each case status was: TVP=36 \pm 12, 163 LP=24±6 and WP=20±6 days. The only studied layer hen was 38 weeks-old, and was classified 164 within the TVP-affected chickens. In 8 out of the 15 TVP-affected chickens, submitted 165 166 veterinarians reported abnormal intestinal contents (including orange jejunal contents, intestinal dilation and loose caecal contents). 167

168 The TVP-affected chickens came from 9 different farms, all of them located in England. In these farms, the mean±SD number of chickens on a farm was 64,417±56,781, with a range of 169 20,000 to 203,000 chickens. The vaccination status was available for 5 out of these 9 farms 170 (from were 10 TVP-affected chickens came from, specifically cases 2, 3, 15, 17, 20 to 24 and 171 41). All of them were vaccinated with a live attenuated IBV vaccine variant strain 4-91, which 172 was combined with the virus strain IB Ma5 in 4 of the farms. In addition, the latter 4 farms also 173 used a live IBDV vaccine containing IBDV strain 228E. No other vaccines were used in these 174 farms. Data on monthly percentage mortality of flocks was available from 4 farms, which 175 176 ranged from 3.03% to 4.49%.

177 **Histopathology.** The histopathological results of each chicken are detailed in Table 1.

The mean necrosis of oxynticopeptic cells score was mild (+) in 8 out of the 15 TVP-affected cases (53%), and moderate (++) in the remaining 7 (47%). Necrotic cells showed hypereosinophilia, fragmentation and karyorrhexis, karyolysis and/or pyknosis, and were usually seen as small clusters within the lumen of dilated proventricular alveoli, often within the edge of the lobule (Figure 2). Collecting ducts (secondary ducts) were often dilated and filled with necrotic debris and sloughed cells. No inclusion bodies were observed in any of the cases.

Thirteen out of the 15 TVP-affected chickens (87%) had severe lymphocytic infiltration, while in the remaining 2 chickens (13%) the lesion was moderate. Lymphocytic infiltration was usually multifocal, located within the interstitium of the proventricular glands (Figure 3). In some cases, lymphocytic cells formed nodular aggregates. Unaffected glands were usually intermingled with affected ones. The median of the lymphocytic infiltration score in the TVPaffected group (3) was significantly higher than in the LP-affected group (1.5) (p=0.002). Although the inflammation in TVP-affected cases was predominantly lymphocytic, few plasmacells, macrophages and occasional heterophils were also present.

Finally, all the 15 TVP-affected chickens (100%) had severe hyperplasia and metaplasia of ductal epithelial cells (Figure 2 and 3), whereas it was present in 11 out of 18 (61%) LP-affected cases and in 2 out of the 11 (18%) chickens within the WP group. The median score for TVPaffected group (3.0) was significantly higher than for LP-affected group (1.5) (p<0.001) and for WP (0.0) (p<0.001). However, trend only was observed between the median score for LP and WP (p=0.053).

CPNV RT-PCR and phylogenetic studies. Nine chickens gave positive results for CPNV
RT-PCR in the proventriculus (Table 1). Seven out of these 9 chickens (78%) belonged to the
TVP-affected group and the remaining 2 (22%) to the LP-affected group. When looking at
each group, 7 out of the 15 TVP-affected cases (47%) and 2 out of the 18 LP-affected
chickens (11%) were positive for CPNV RT-PCR. None of the proventriculi from chickens
belonging to the WP group gave positive results.

All RT-PCR CPNV positive cases corresponded to broiler chickens. The only analysed layer
hen, which was histologically classified as TVP, gave a negative CPNV RT-PCR results.

The VP1 gene of the 9 positive CPNV cases was partially sequenced, including a fragment of 207 171 nucleotides (nt). Figure 4 shows a phylogenetical tree including these 9 sequences, 1 208 sequences obtained from the French CPNV RT-PCR positive control, 2 Spanish CPNV 209 sequences and 1 American case (Genbank accession number: HM038436.1). All the 9 British 210 211 CPNV sequences were closer to each other than to sequences obtained from other countries. Their percentage of similarity was below 90% only when compared with the American and 212 213 one of the Spanish sequences. Specifically, the cases from the UK showed 99.4-100% nucleotide similarity between the sequences. When compared with the sequences retrieved 214

from the other countries, the % similarity was 92.4- 92.8%, 88.3-94.2% and 89.5-90.1%, with

the French, Spanish and American sequences, respectively (Table 2).

217 Discussion

Cases of chickens with lymphocytic and necrotising proventriculitis, consistent with TVP, have
been reported in the USA (Guy et al., 2011b), South Korea (Kim et al., 2015) and several
countries in Europe (Grau-Roma et al., 2010; Kouwenhoven et al., 1978; Marguerie et al.,
2011). However, as far as the authors are aware, this is the first peer-reviewed description of
TVP as well as of CPNV in the UK.

223 The mean mortality of broiler flocks in the UK has previously been reported at 4.1% (Dawkins et al., 2004). This percentage is close to the mortality recorded in the TVP-affected farms in 224 this study. These results are therefore in line with earlier studies that state there is no significant 225 226 increase in mortality in flocks affected by TVP (Hafner and Guy, 2013). All the TVP-affected chickens came from farms located in counties across England. TVP-affected broilers were in 227 the age range of 21 to 49 days old, which is consistent with previous reports of the disease 228 (Bayyari et al., 1995; Hafner and Guy, 2013). The majority of the chickens submitted to the 229 study were Ross 308 broiler chickens, with only 1 case corresponding to a layer hen. There 230 may be 2 reasons for this: (i) TVP affects mainly broiler chickens; (ii) broilers chickens equate 231 to 80% of the chicken post-mortems carried out by the poultry clinicians submitting the 232 samples. The studied layer hen studied was histologically classified as TVP, becoming the 233 second report of this disease in layer hens in peer-reviewed literature (Marusak et al., 2012). 234 Marusak et al. (2012) diagnosed TVP in broiler breeder and commercial layer hens ranging 235 from 9 to 20 weeks of age. Therefore, the hen included here is the oldest chicken reported to 236 be affected by TVP, although it was negative by CPNV RT-PCR. 237

238 The detection of CPNV by RT-PCR in almost 50% (7 out of 15) of TVP-affected chickens together with the negative results in all the chickens within the WP group supports the idea that 239 CPNV is the cause of TVP (Guy et al., 2011a; Guy et al., 2011b). All the TVP-affected cases 240 241 showed moderate to severe lymphocytic infiltrates and severe tubular hyperplasia and metaplasia, which are features of chronicity as demonstrated in experimentally reproduced 242 TVP-affected cases (Guy et al., 2011b). This experimental infection also showed that CPNV 243 was only detectable by RT-PCR from 1 to 14 days post exposure (PE), while the microscopic 244 lesions were present from 5 to 35 days PE (Guy et al., 2011b). Therefore, the TVP cases with 245 246 negative RT-PCR CPNV in the present study may correspond to chronically CPNV infected chickens, where the virus is not detectable further within the lesions. In addition, the well-247 known reduced sensitivity of RT-PCR on FFPE tissues compared to fresh tissues might account 248 249 for a number of these negative RT-PCR results (Lewis et al., 2001). Finally, it can not however be ruled out that other infectious or non-infectious agents, alone or together with CPNV, may 250 be involved in some of the TVP-affected chickens presented here (Huff et al., 2001; Dormitorio 251 et al., 2007; Kim et al., 2015). 252

253 It has been reported that proventriculitis can be caused by a number of different factors, including infectious (viruses, bacteria, fungi or parasites), mycotoxins and nutritional factors 254 (Dormitorio et al., 2007). It seems likely that, in this study, the group of LP-affected chickens 255 correspond to a mixture of cases with different aetiologies. Amongst them, a number of LP-256 affected chickens with negative CPNV RT-PCR results may correspond to chronically affected 257 TVP cases, where the virus is not detectable (Guy et al., 2011b). Interestingly, 11% of LP-258 affected chickens gave positive results by CPNV RT-PCR. These chickens likely correspond 259 to chronic TVP-affected chickens, where necrosis of oxynticopeptic cells is not observed, but 260 where the virus is still detectable. This finding indicates that TVP should still be suspected in 261 cases of LP without glandular necrosis and that the CPNV RT-PCR can have diagnostic value 262

in these cases. It is worth mentioning that areas of necrosis may have been present within other
 non-examined areas of the proventriculi, since this study was performed on a single complete
 cross-section of each proventriculus.

The % of nucleotide similarities showed the highest value similarity when comparing the 266 267 sequences obtained from within the UK. Moreover, the phylogenetical tree of nucleotide sequences suggested that the UK CPNV sequences may be more similar to the other European 268 sequences than to the American sequence. These geographical variations may be due to 269 270 mutations as a result of different selection pressures between countries and continents, as previously suggested for other virus such as IBDV (Jackwood and Sommer-Wagner, 2007). 271 Although the current study included short nucleotide sequences (171 nt) within the VP1 gene, 272 it must be acknowledged that this gene encodes for the RNA-dependent RNA-polymerase, 273 which is known to be a well-conserved gene in cellular organisms as well as in viruses (Pan et 274 275 al., 2007). For this reason, based on the results obtained, it could be hypothesized that two distinct lineages of CPNV, i.e. European and American, are present in the two continents. It 276 would, however, be necessary to perform larger studies, increasing the number of sequences 277 278 and their length, to try to confirm these initial observations

In conclusion, this study identified the presence of TVP amongst chicken populations in the UK. Results indicate that CPNV can often be detected within the proventriculus of TVPaffected chickens, and in a number of chickens with LP. Moreover, preliminary phylogenetic studies on partial CPNV sequences indicate that European and American chicken populations may have 2 different CPNV genetic lineages. Additional investigations, encompassing larger sample sizes, are needed in order to determine the incidence and prevalence of TVP in the UK and globally.

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376 Tables

Table 1. Microscopic proventricular lesion scores^a, RT-PCR results and case status^b for each
 chicken. Genbank accession numbers for positive *Chicken proventricular necrosis virus* (CPNV) RT-PCR cases are detailed.

Chicken	Mean necrosis score ^c Interstitial lymphocyti c infiltration		Ductal epithelial hyperplasia and metaplasia	Case Status	CPNV RT- PCR	Identifier	Accession number		
1	-	-	-	WP	Negative				
2	+	+++	+++	TVP	Negative				
3	+	+++	+++	TVP	Negative				
4	-	-	-	WP	Negative				
5	-	-	-	WP	Negative				
6	-	+	-	LP	Negative				
7	-	+++	+++	LP	Negative				
8	-	+	-	LP	Negative				
9	-	++	++	LP	Negative				
10	-	+	-	LP	Negative				
11	-	-	-	WP	Negative				
12	_	_	-	WP	Negative				
13	_	+	-	LP	Negative				
14	_	+++	++	LP	Negative				
15	++	+++	+++	TVP	Positive	CPNV-UK-1	KU933597		
16	-	++	++	LP	Positive	CPNV-UK-2	KU933598		
17	++	+++	+++	TVP	Negative				
18	-	+	++	LP	Negative				
19	-	-	++	WP	Negative				
20	++	+++	+++	TVP	Negative				
21	+	+++	+++	TVP	Positive	CPNV-UK-3	KU933599		
22	++	++	+++	TVP	Positive	CPNV-UK-4	KU933600		
23	++	+++	+++	TVP	Positive	CPNV-UK-5	KU933601		
24	++	+++	+++		Positive Nogetive	CPNV-UK-0	KU955002		
25	+	++	+++		Nogativo				
26	+	+++	+++		Negative				
27	+	+++	+++		Negative				
28	-	++	-	LP	Negative				
29	-	+++	++	LP	Negative				
30	-	-	-	WP	Negative				
31	-	+	++	LP	Negative				
32	-	-	-	WP	Negative				
33	-	+	+	LP	Negative				
34	-	-	++	WP	Negative				
35	-	+	-	LP	Negative				
36	-	-	-	WP	Negative				
37	-	+	+	LP	Negative				
38	-	-	-	WP	Negative				
39	-	++	-	LP	Negative				
40	-	+++	+++	LP	Negative				

Microscopic proventricular lesion scores

	41	+	+++	+++	TVP	Positive	CPNV-UK-7	KU933603
	42	-	+++	+++	LP	Positive	CPNV-UK-8	KU933595
	43	++	+++	+++	TVP	Positive	CPNV-UK-9	KU933596
	44	+	+++	+++	TVP	Negative		
380	^a -: absence	; +: >0 to 1	0% of the gla	inds affected; -	++: >10 to	50% of th	ne glands aff	fected, +++:
381	>50% of th	ie glands af	fected					
382	^D TVP: trar	nsmissible	viral provent	riculitis; LP:	lymphocyt	ic proven	triculitis; W	P: without
383	proventricu	ılitis.						
384	^c Mean nec	rosis score	is the mean	combined scor	e of necro	osis of ox	ynticopeptic	cells in all
385	glands and	necrosis of	f glandular pa	renchyma in th	e most aff	ected glan	d.	
386	C		0 1	2		U		
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Table 2. Percentage of homology between the studied *Chicken proventricular necrosis virus* (CPNV) partial VP1 sequences. Sequences

included are American (CPNV-USA, n=1), French (CPNV-Fr, n=1), Spanish (CPNV-Sp, n=2) and British (CPNV-UK, n=9).

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	CPNV-												
	USA-1	Fr-1	Sp-2	Sp-1	UK-8	UK-9	UK-1	UK-2	UK-3	UK-4	UK-5	UK-6	UK-7
CPNV-USA-1	100.00	90.06	89.47	91.81	89.47	89.47	90.06	90.06	90.06	90.06	90.06	90.06	90.06
CPNV-Fr-1	90.06	100.00	94.15	92.40	92.40	92.40	92.98	92.98	92.98	92.98	92.98	92.98	92.98
CPNV-Sp-2	89.47	94.15	100.00	94.74	93.57	93.57	94.15	94.15	94.15	94.15	94.15	94.15	94.15
CPNV-Sp-1	91.81	92.40	94.74	100.00	88.30	88.30	88.89	88.89	88.89	88.89	88.89	88.89	88.89
CPNV-UK-8	89.47	92.40	93.57	88.30	100.00	100.00	99.42	99.42	99.42	99.42	99.42	99.42	99.42
CPNV-UK-9	89.47	92.40	93.57	88.30	100.00	100.00	99.42	99.42	99.42	99.42	99.42	99.42	99.42
CPNV-UK-1	90.06	92.98	94.15	88.89	99.42	99.42	100.00	100.00	100.00	100.00	100.00	100.00	100.00
CPNV-UK-2	90.06	92.98	94.15	88.89	99.42	99.42	100.00	100.00	100.00	100.00	100.00	100.00	100.00
CPNV-UK-3	90.06	92.98	94.15	88.89	99.42	99.42	100.00	100.00	100.00	100.00	100.00	100.00	100.00
CPNV-UK-4	90.06	92.98	94.15	88.89	99.42	99.42	100.00	100.00	100.00	100.00	100.00	100.00	100.00
CPNV-UK-5	90.06	92.98	94.15	88.89	99.42	99.42	100.00	100.00	100.00	100.00	100.00	100.00	100.00
CPNV-UK-6	90.06	92.98	94.15	88.89	99.42	99.42	100.00	100.00	100.00	100.00	100.00	100.00	100.00
CPNV-UK-7	90.06	92.98	94.15	88.89	99.42	99.42	100.00	100.00	100.00	100.00	100.00	100.00	100.00

401 Figures

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Figure 1. Proventriclulus and gizzard, broiler chicken, chicken 22, Transmissible viral proventriculitis (TVP)-affected case. The proventricular wall is severely and diffusely thickened. Multifocal and small (up to 0.5 cm) circular areas of congestion and haemorrhage are present within the proventricular mucosa generating a pattern that highlights the proventricular mucosal papillae (arrowhead). The proventricular wall shows prominent glandular lobules (asterisk). Gizzard shows no gross lesions.



Figure 2. Proventriculus, broiler chicken, chicken 17. Transmissible viral proventriculitis (TVP)-affected case. Photomicrograph showing severe proventricular interstitial lymphocytic infiltration and moderate replacement of glandular epithelium by hyperplastic ductal epithelium (ductal epithelial metaplasia). Inset: Higher magnification of the same proventriculus, showing the predominantly lymphocytic interstitial infiltration as well as the tubular epithelial hyperplasia and metaplasia. Haematoxylin and eosin.

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Figure 3. Proventriculus, broiler chicken, chicken 20. Transmissible viral proventriculitis
(TVP)-affected case. Photomicropraph showing multifocal aggregates of necrotic cells
(arrowhead) within the lumen of proventriculuar alveoli located at the periphery of a
proventricular lobule. Areas with replacement of glandular epithelium by hyperplastic ductal
epithelium (ductal epithelial metaplasia) are also present (asterisk). Haematoxylin and eosin.



449 **Figure 4**. Phylogenetic tree based on the NJ method for 13 partial (171 nucleotides) VP1

450 CPNV sequences. Sequences originate from 4 different countries: USA (CPNV-USA-1),

451 France (CPNV-Fr-1), Spain (CPNV-Sp-1 and 2), and UK (CPNV-UK-1 to 9). Numbers

452 along the branches refer to the percentages of confidence.