

1 **EXPERIMENTAL BLUETONGUE VIRUS SUPERINFECTION IN**  
2 **CALVES PREVIOUSLY IMMUNIZED WITH BLUETONGUE VIRUS**  
3 **SEROTYPE 8**

4  
5 Ludovic Martinelle<sup>1</sup>, Fabiana Dal Pozzo<sup>1</sup>, Pierre Sarradin<sup>2</sup>, Willem Van Campe<sup>3</sup>, Ilse De Leeuw<sup>4</sup>, Kris  
6 De Clercq<sup>4</sup>, Christine Thys<sup>1</sup>, Etienne Thiry<sup>5</sup>, Claude Saegerman<sup>1\*</sup>

7  
8 <sup>1</sup> Research Unit in Epidemiology and Risk Analysis Applied to the Veterinary Sciences (UREAR-  
9 ULg), Center for Fundamental and Applied Research for Animal and Health (FARAH), Department of  
10 Infectious and Parasitic Diseases, Faculty of Veterinary Medicine, University of Liege, Quartier  
11 Vallée 2, Avenue de Cureghem 7A, B-4000, Liege, Belgium

12 <sup>2</sup> INRA UE 1277, Experimental Infectiology Platform, INRA – Research Centre of Tours, Nouzilly,  
13 France

14 <sup>3</sup> CODA-CERVA, Animal Experimental Centre, , Kerklaan 68 1830, Machelen, Belgium.

15 <sup>4</sup> CODA-CERVA, Unit Vesicular and exotic diseases, , Groeselenberg 99, B-1180, Uccle, Brussels,  
16 Belgium

17 <sup>5</sup> Veterinary Virology and Animal Viral Diseases, Center for Fundamental and Applied Research for  
18 Animal and Health (FARAH), Department of Infectious and Parasitic Diseases, Faculty of Veterinary  
19 Medicine, Quartier Vallée 2, Avenue de Cureghem 10, B-4000, Liège, Belgium

20 \*Corresponding author

21 Email addresses:

22 Ludovic Martinelle : [lmartinelle@ulg.ac.be](mailto:lmartinelle@ulg.ac.be)

23 Fabiana Dal Pozzo : [fdalpozzo@ulg.ac.be](mailto:fdalpozzo@ulg.ac.be)

24 Pierre Sarradin: [Pierre.Sarradin@tours.inra.fr](mailto:Pierre.Sarradin@tours.inra.fr)

25 Willem Van Campe : [willem.vancampe@coda-cerva](mailto:willem.vancampe@coda-cerva)

26 Ilse De Leeuw: [ilse.deleeuw@coda-cerva.be](mailto:ilse.deleeuw@coda-cerva.be)

27 Kris De Clercq : [kris.declercq@coda-cerva.be](mailto:kris.declercq@coda-cerva.be)

28 Christine Thys : [thys-ch@hotmail.com](mailto:thys-ch@hotmail.com)

29 Etienne Thiry : [etienne.thiry@ulg.ac.be](mailto:etienne.thiry@ulg.ac.be)

30 Claude Saegerman : [Claude.Saegerman@ulg.ac.be](mailto:Claude.Saegerman@ulg.ac.be)

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35 **Abstract**

36 The effect of a superinfection with bluetongue virus serotype 1 (BTV1) was evaluated on two groups  
37 of four calves. One group received a commercial inactivated BTV serotype 8 (BTV8) vaccine. This  
38 group and the non-vaccinated group of calves were challenged twice (4 months apart) with the  
39 European BTV8 strain isolated during the 2006-2007 epidemics. Calves were then infected with a  
40 BTV1 inoculum which was found to be unexpectedly contaminated by BTV serotype 15 (BTV15).  
41 BTV1 and BTV15 single infections were performed on two other groups of three BTV naïve calves.

42 A severe clinical picture was obtained after superinfection with BTV1/BTV15 in both vaccinated and  
43 non-vaccinated animals and after challenge with BTV8 in non-vaccinated animals. BTV1 and BTV15  
44 single infection caused only very slight clinical signs.

45 After superinfection and at the viraemic peak, there were an average of above 1000 times more  
46 BTV15 genomic copies than BTV1 ones. BTV1 RNA could be detected only in the spleen of one calf  
47 whereas BTV15 RNA was found in 15 organs of 7 different animals. BTV8 immunization whether it  
48 was acquired through vaccination and challenges or challenges alone did not change BTV1 or BTV15  
49 RNA detection in superinfected animals. However in these animals a partial cross neutralization  
50 between BTV8 and BTV1 might be involved in the lower BTV1 replication versus BTV15.

51 Infection with different serotypes can occur also in the field. Interference between virus strains,  
52 genetic reassortment and cross-protection were considered as mechanisms to explain the clinical  
53 outcomes and the other virological and immunological findings in the course of BTV1/BTV15  
54 superinfection.

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56 **Key words:** bluetongue virus, serotype 8, serotype 1, serotype 15, experimental superinfection.

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## 58 **Introduction**

59 Bluetongue (BT) is a non-contagious disease affecting ruminants and is caused by the bluetongue  
60 virus (BTV), the type species of the genus *Orbivirus*. BT is a World Organization for Animal Health  
61 reportable disease and is of considerable socioeconomic concern and of major importance in the  
62 international trade of animals and animal products [1]. Economic losses associated with BTV infection  
63 are caused directly through reductions in animal productivity and death, implementation of control  
64 measures, and more importantly, indirectly through trade losses due to animal movement restrictions  
65 [2].

66 Within each different *Orbivirus* species, several virus serotypes are identified, based on the specificity  
67 of reactions with the neutralizing antibodies generated by their mammalian host [3]. These reactions  
68 are dependent to a large extent to VP2 and also VP5, which are the most variable proteins in BTV;  
69 VP2 especially contains the most epitopes that drive neutralizing antibodies production and therefore  
70 is the main determinant of the serotype [4]. To date, 27 serotypes have been identified, including  
71 BTV25 identified in Switzerland in 2007 [5], BTV26 from Kuwait in 2010 [6] and BTV27 detected in  
72 goats in Corsica (France) in 2014 [7]. In addition, two putative new serotypes, respectively BTV28  
73 and BTV29 were recently detected [8]. Indeed serological cross-reactions between different serotypes  
74 are described [9] and evidences of possible heterologous cross-protection do exist [10, 11], but their  
75 influence on the epidemiology of the disease is not sufficiently understood.

76 From 2006 to 2015, seven BTV serotypes were detected in Western and Central Europe, namely  
77 BTV1, 6, 8, 11, 14, 25, and 27. Most of the economic losses have to be attributed to BTV8, and to a  
78 lesser extent, BTV1, with respectively over 27000 and 6000 holdings affected only in 2008. BTV8  
79 alone was responsible for the death of more than 20000 sheep in Belgium, which represents 5-10 % of  
80 the national flock [12]. The 2008 BTV8 epizootic in Northern Europe is believed to have caused  
81 greater economic damage than any previous single serotype BT outbreak [13]. By contrast re-  
82 emergence of BTV8 in France in 2015 was only of limited impact [14].

83 On the other hand, BTV1 that circulated contemporarily in Southwest Europe, was described as a  
84 virus leading to subclinical or mild disease in cattle [15]. As a consequence of this epidemiologic

85 context, domestic ruminants in the field could be sequentially infected by these two serotypes, as it  
86 was reported in France and Spain [16].

87 In this paper the results of a 9 month-long experiment are shown. Calves were originally divided in  
88 two groups, with one group being vaccinated against BTV8, and were subsequently both challenged  
89 with a homologous BTV8 European strain. In order to mimic the occurrence of repetitive infections  
90 according to studies reporting several peaks of vector activity during the course of the year [17], the  
91 same calves were infected a second time with the same BTV8 strain and later with BTV serotype 1  
92 (BTV1) (superinfection). The aim of this study was to analyse the outcome of these successive  
93 challenges, taking into account the influence of vaccine immunity as well as natural post-infection  
94 immunity. The BTV1 inoculum appeared to be contaminated with BTV15 [18]. In order to evaluate  
95 any *in vivo* cross-protection, the consequences of a BTV1 and BTV15 single infection in BTV naïve  
96 calves were also considered.

## 97 **Materials & methods**

### 98 **BTV8 successive infections**

99 Ten Holstein female calves, about 6-7 months old, were used. All the animals were tested seronegative  
100 and non viraemic for BTV and bovine viral diarrhoea virus (BVDV), and seronegative for bovine  
101 herpesvirus 1 (BoHV1). A thorough general clinical examination was carried out on all the animals by  
102 a veterinarian before including them in the study, to confirm their asymptomatic state, in accordance to  
103 physiological standards [19].

104 The calves were housed in an insect-secured BSL3 zone at the Experimental Infectiology Platform  
105 (PFIE) of the INRA centre of Tours (Nouzilly, France). The local ethical committee approved the  
106 experimental protocol (dossier n.2011-10-1). Three groups were created: a group of four non-  
107 vaccinated calves (group NV, calves 1-4), a group with four vaccinated calves (group V, calves 5-8),  
108 and an environmental control group with two calves (group C, calves 15 and 16). Vaccination against  
109 BTV8 was performed using the inactivated commercial vaccine BTVPUR AISAP 8 (Merial, Lyon,  
110 France) following manufacturer's instructions.

111 Calves of group V and NV were infected twice, four months apart, using the same BTV8 inoculum  
112 described in a previous experimental infection [20]. Briefly, a calf inoculated with a BTV-8 strain  
113 passaged twice in baby hamster kidney fibroblasts (BHK-21) cells (BEL2006/01 BHK-21 P2), was  
114 blood sampled at the viraemic peak, showing clinical signs. The first infection took place 50 days after  
115 the second vaccine shot, and the second challenge 120 days later (Figure 1A). Each time, half of the  
116 dose was inoculated via the jugular vein and half subcutaneously. For each one of these challenges, all  
117 8 infected animals were administered a total of 15 ml of blood, corresponding to a titre of  $10^3$  embryo  
118 lethal dose 50 (ELD<sub>50</sub>).

119 The animals were daily examined and sampled for EDTA-blood and whole blood on dry tubes during  
120 the 3 first weeks after infection, then twice a week until superinfection. For clarification purpose, days  
121 post infection regarding BTV8 successive infection are mentioned as dpi<sub>BTV8</sub>, with day of first  
122 challenge as dpi<sub>BTV8</sub> 0.

### 123 **BTV superinfection**

124 About 5 months (160 days) after the first BTV8 infection, and 40 days after the second one, calves of  
125 groups NV and V were challenged with BTV1 infectious blood, kindly provided by the Friedrich-  
126 Loeffler Institute. Each animal received  $10^{6.15}$  tissue culture infective dose 50% (TCID<sub>50</sub>) of virus, half  
127 intravenously and half subcutaneously. An incidental contamination of the inoculum with BTV15 was  
128 discovered during the course of the study [18].

129 The animals were daily examined and sampled for EDTA-blood and whole blood on dry tubes until  
130 the end of the experiment.

131 Calves were euthanized 21 days after the superinfection and necropsied (Figure 1A-B). Days post  
132 infection for superinfection is mentioned as dpi<sub>BTV1/15</sub>.

### 133 **BTV1 and BTV15 single infections**

134 Eight Holstein male calves, about 6-7 months old, were housed in the BSL3 facility, at CODA-  
135 CERVA's (Veterinary and Agrochemical Research Centre, Uccle, Belgium) experimental centre  
136 (Machelen, Belgium). These animals fulfilled the same inclusion criteria of the BTV8 successive

137 infections and were naïve for BTV, BVDV and BoHV-1. The experimental protocol was reviewed by  
138 the competent authority (Ethical Committee of the Institute of Public Health-Veterinary and  
139 Agronomical Research Centre) and subsequently approved (ref. 110228-01 RT 10/10  
140 BLUETONGUE).

141 After an acclimatization period of two weeks, calves were divided in three groups (BTV1: calves 9-11,  
142 and BTV15: calves 12-14, single infections; mock-infection: four calves (calves 17-20)). The  
143 infection was performed with a volume of 1 ml of virus diluted in Dulbecco's Modified Eagle Medium  
144 (DMEM, Lonza BioResearch, Belgium), half intravenously and half subcutaneously.

145 The BTV1 strain has been provided by CODA-CERVA, from sub-saharian origin, derived from the  
146 European Community Reference Laboratory for bluetongue at the Pirbright Laboratory, United-  
147 Kingdom) collections, and subsequently passaged 2 times in BHK-21 cells. The infection was  
148 performed with  $10^6$  TCID<sub>50</sub> per animal.

149 BTV15 was provided by CODA-CERVA, derived from the European Community Reference  
150 Laboratory for bluetongue at the Pirbright Laboratory, United Kingdom) and was then passaged twice  
151 on BHK21 cells at CODA-CERVA. Calves were infected with  $10^4$  TCID<sub>50</sub>.

152 Mock-infected calves were inoculated with sterile DMEM following the same routes and volume.

153 The animals were daily examined and sampled for EDTA-blood and whole blood on dry tubes until  
154 the end of the experiment. Calves were euthanized 35 dpi and necropsied. Days post infection for  
155 single infections are mentioned as dpi<sub>single</sub>.

#### 156 **Clinical and post-mortem examination**

157 After each challenge, the individual body temperature and the clinical signs were monitored for three  
158 weeks. The severity of the infection was quantified by calculating clinical scores on a daily basis,  
159 leading to overall cumulative clinical scores by groups and animal. For this purpose, a standardised  
160 clinical form adapted from Saegerman *et al.* was used [21]. As BTV1 and BTV15 single infections

161 only involved 3 animals whereas other groups had 4 calves each, total clinical score was pondered to  
162 allow direct comparison of clinical scores from different experiments.

163 Samples of spleen, thymus, prescapular and mesenteric lymph nodes were collected from infected and  
164 control calves and stored at  $-80^{\circ}\text{C}$  for virus detection.

### 165 **Serology**

166 Neutralizing antibodies (Abs) were titrated by seroneutralization (SNT). Two-fold serial dilutions of  
167 the sera (1:10-1:1280) were tested in the presence of 100 TCID<sub>50</sub> of virus, as previously described  
168 [22]. The neutralizing antibody titre was defined as the reciprocal of the serum dilution causing a 50%  
169 reduction in cytopathic effect. Serum samples with a titre  $<20$ ,  $=20$  and  $>20$  were considered negative,  
170 doubtful and positive, respectively.

171 In order to identify *in vitro* cross neutralization between BTV1, BTV8 and BTV15, the serum of the  
172 calf infected with BTV8 and showing the highest anti-BTV8 antibody titre was tested in the presence  
173 of BTV1 or BTV15. Similarly, the serum of the calves infected with BTV1 and BTV15 following  
174 single serotype infections and showing the highest neutralizing antibody titres against the  
175 correspondent virus were tested against heterologous serotypes. *In vitro* cross neutralization was  
176 measured using the percentage of neutralization obtained using heterologous serotypes with immune  
177 serum and compared to homologous neutralization as reference (100 %).

178 In the course of the two BTV8 challenges, and after BTV1 and BTV15 single infections,  
179 seroconversion against VP7 antibodies was also evaluated using a commercial competitive ELISA kit  
180 (ID Screen® Bluetongue Competition ELISA kit, ID Vet, France). Results were expressed as % of  
181 negativity (PN) compared to the negative kit control and transferred to a positive, doubtful or negative  
182 result according to the cut-off settings provided by the manufacturer (PN  $\leq 35$  is positive;  
183  $35 < \text{PN} \leq 45$  is doubtful; PN  $> 45$  is negative).

### 184 **BTV RNA detection**

185 Viral RNA extraction from the blood was achieved using the QIAamp Viral RNA Mini Kit (Qiagen,  
186 Germany). Viral RNA denaturation and reverse transcription followed by qPCR were performed as

187 previously described [23]. BTV RNA was detected by serotype specific RTqPCR, using a fragment of  
188 BTV segment 2 as the target. Serial dilutions of *in vitro* constructed plasmids (pGEM®-T Easy  
189 Vector, Promega, The Netherlands) carrying the target part of the segment 2, specific for each  
190 serotype, allowed the absolute quantification of the viral cDNA equivalent in samples. Quantification  
191 was expressed in cDNA copy number/ml of blood. RTqPCR cycling conditions, primers and probes  
192 were similar to the ones described by Vandenbussche et al. (2009) for BTV1 and BTV8  
193 (RTqPCR\_BTV1\_S2 and RTqPCR\_BTV8\_S2, respectively), and Eschbaumer and al. (2011) for  
194 BTV15 (RTqPCR\_BTV15\_S2). In all the RTqPCR of this study, bovine beta-actin was  
195 contemporaneously amplified as internal control (RTqPCR\_ACT) [24].

196 BTV RNA detection was performed on all the collected organs starting from approximately 100 mg of  
197 tissue, which was processed using TRI reagent according to the manufacturer's instructions (Life  
198 Technologies Europe BV, Gent, Belgium). BTV and bovine beta-actin detection were performed by  
199 RTqPCR as described above.

#### 200 **Viral growth assay**

201 *In vitro* replication of BTV1, BTV8 and BTV15 were compared on VERO and Bovine Pulmonary  
202 Endothelial cells (BPAEC). BTV8 was the same as in the BTV8 successive infections, and BTV1 and  
203 BTV15 were the same as in single infections. These viruses were used for a growth assay following a  
204 protocol previously described [25]. Briefly, VERO and BPAEC confluent cells in 24 wells plates were  
205 inoculated with a multiplicity of infection (M.O.I.) of 0.05, and after 0, 8, 24, 48 and 120 hours post  
206 infection (hpi), supernatant was removed and stored at -80°C. Each virus underwent three independent  
207 assays on each cell type. Supernatants were then titrated by end-point dilution and titres expressed as  
208 Log<sub>10</sub> of TCID<sub>50</sub>/ml.

#### 209 **Statistical analysis**

210 Mean cumulative clinical scores were analysed using linear mixed model, with calf as random effect.  
211 Viraemia and serological results were compared using two-way ANOVA with repeated measures.  
212 RNA detection in organs at necropsy, frequencies and proportions were compared with Fisher's Exact



213 Test for count data [26]. For all tests,  $P$  values  $< 0.05$  were considered significant. In case of multiple  
214 comparisons, a Bonferroni correction was applied to reduce the risk of type I error (conservative  
215 approach) and a Holm correction was applied when more than 4 comparisons had to be tested.  
216 Statistical analyses were performed using the R software/environment (R-3.1.2, R Foundation for  
217 Statistical Computing, <http://www.r-project.org/>) and SAS software, Version 9.3 TS level 1M2 of the  
218 SAS System for Unix, and SAS University Edition (SAS Institute, Cary, NC).

## 219 **Results**

### 220 **BTV8 successive infections**

#### 221 *Clinical examination*

222 From the beginning to the end of the experiment, control calves and vaccinated animals (V) did not  
223 show any clinical signs that could be related to BTV infection. After the first challenge, clinical signs  
224 showed by NV calves were slight, mostly consisting in ocular lesions and to a lesser extent by oral  
225 lesions. Clinical signs consistent with BTV8 infection could be reported from 7 to 21 dpi<sub>BTV8</sub>. After  
226 the second challenge no clinical manifestations or temperature rise could be detected in any animal.

#### 227 *Serology*

228 VP7 and anti-BTV8 neutralizing Abs in control calves could not be detected at any tested time points.  
229 First vaccination of the calves did not induce detectable neutralizing Abs, which could only be  
230 detected 7 days after the booster vaccination. In NV group, neutralizing Abs were first detected at 18  
231 dpi<sub>BTV8</sub> (Figure 1A). The titre of anti-BTV8 neutralizing Abs of the V group was significantly higher  
232 between -43 dpi<sub>BTV8</sub> (thus 35 days after the first vaccine shot) and 27 dpi<sub>BTV8</sub> ( $P < 0.005$ ), and then Abs  
233 titres of both NV and V groups followed a similar trend until the end of the experiment (Figure 1A).  
234 Following the second BTV8 challenge, neutralizing Abs titres underwent a boost in both NV and V  
235 groups until 33 dpi<sub>BTV8</sub> (Figure 1A).

236 The use of an ELISA allowed the detection of anti-VP7 Abs in all the vaccinated animals as soon as 3  
237 days after second vaccine injection (Figure 2). Then PN of vaccinated animals did not evolve

238 significantly until the end of the measures at 180 dpi<sub>BTV8</sub> (additional file 1). Non-vaccinated calves  
239 were confirmed seropositive between 10 and 19 dpi<sub>BTV8</sub>, with no significant variations until the end of  
240 the measures at 180 dpi<sub>BTV8</sub> (additional file 1,  $P > 0.05$ ).

#### 241 *BTV RNA detection*

242 BTV8 RNA was never detected in the EDTA-blood samples of control and vaccinated calves during  
243 the course of the two BTV8 infections. In the NV group of calves, BTV RNA could be detected  
244 starting from 5 dpi<sub>BTV8</sub>. After the viraemic peak (11-15 dpi<sub>BTV8</sub>) a progressive decrease in BTV8 RNA  
245 was measured (Figure 1A-B), until the end of the experiment.

#### 246 **BTV superinfection**

##### 247 *Clinical and post-mortem examination*

248 In the NV as well as in the V group, lesions following BTV1/BTV15 superinfection were mainly  
249 conjunctivitis with serous to purulent discharge. Erosions of the muzzle, erosions and ulcerations of  
250 the gums and the dental pad and later crusts on the muzzle or on the cutaneous-mucous junction were  
251 commonly notified. Reddening and swelling of the coronal margin and interdigital space were also  
252 mentioned. Conjunctivitis and congestion of the lower limb were mild to severe and respectively less  
253 severe and absent in the previous BTV8 infection. At the end of the experiment cumulative clinical  
254 score of the NV BTV1/BTV15 superinfected group was higher than in NV BTV8 and V  
255 BTV1/BTV15 groups, however the difference was not significant ( $P > 0.4$ , additional file 2).

256 The necropsy revealed petechial haemorrhages of limited extent in prescapular and submandibular  
257 lymph nodes, and thymus, at least in one of these organs in all the superinfected calves. BTV8 RNA  
258 could be detected in prescapular lymph node of one calf and in the spleen of another one, both from  
259 NV group; BTV1 RNA could only be detected in the spleen of one vaccinated calf and BTV15 RNA  
260 could be detected with a significant higher frequency ( $P < 10^{-5}$ ) in 15 organs belonging to 7 different  
261 calves (5/15 in NV group and 10/15 in V group; Figure 3).

262 No BTV RNA could be detected in any of the tested organs from control animals.

263 *Serology*

264 After the superinfection, high levels of residual neutralizing antibodies against BTV8 were found  
265 throughout the experiment in all the infected animals, with a roughly steady level (Figure 1A and B).

266 There were no significant differences between NV and V groups through time in anti BTV8  
267 neutralizing antibodies (two-way ANOVA with repeated measures, group effect:  $P=0.78$ , group time  
268 interaction:  $P=0.84$ ). By contrast, BTV1 only gave rise to very low titres of neutralizing antibodies.

269 BTV15 neutralizing antibody titre increased regularly after infection, with a positive titre detectable in  
270 most of the animals at 9 dpi<sub>BTV1/15</sub> (Figure 1B). BTV15 neutralizing Abs followed an increasing trend  
271 until the time of euthanasia; in previously vaccinated animals a higher and significant earlier raise of  
272 neutralizing antibody was found compared to non-vaccinated ones (two-way ANOVA with repeated  
273 measures, group effect:  $P<0.02$ , group time interaction:  $P<2\times 10^{-4}$ ) (Figure 1B).

274 *BTV RNA detection*

275 At the time of the superinfection, a residual BTV8 RNAemia was detected in the NV group of calves.  
276 BTV8 RNAemia decreased through the time of the experiment, but was still detectable in all the NV  
277 animals by the time of euthanasia (180 days after the first BTV8 challenge – Figure 1).

278 The superinfection inoculum contained respectively  $10^{6.8}$  and  $10^{7.6}$  copies of segment 2 cDNA per ml  
279 of blood, for BTV1 and BTV15.

280 After superinfection in the NV and the V groups, BTV1 could only be detected inconstantly, from one  
281 to 3 days amongst all the tested day-points and at lower copy number than BTV15 (Figure 1B). On the  
282 contrary, BTV15 could be easily detected in both groups and among all the infected calves. RNAemia  
283 through time was significantly different between BTV15 and BTV1 whichever the considered  
284 vaccination status of the animals ( $P<10^{-8}$ ). Mean copy number at viraemic peak was  $10^{5.4}$  (+/- 0.7 Log)  
285 cDNA copy number / ml of blood for BTV15 and  $10^{2.4}$  (+/- 1.8 Log) for BTV1. BTV15 could be  
286 detected until the end of the experiment (Figure 1B). Between V and NV groups, detection of BTV1  
287 and BTV15 was not significantly different ( $P=0.18$  and  $P=0.86$  for BTV1 and BTV15 respectively).

288 No viral RNA could be detected in control animals (data not shown).

289 **BTV1 and BTV15 single infections**

290 *Clinical and post-mortem examination*

291 During BTV1 single infection, one calf underwent sporadic hyperthermia and all the three calves of  
292 the group had mild oral and ocular lesions. No systemic impact was reported in any of these animals.  
293 During BTV15 single infection, infected animals showed very mild clinical conditions compatible  
294 with BT, including congestion and crusts on the nostrils and oral mucosa. One calf showed  
295 hyperthermia at 7 dpi (39.6°C) and at 14 (40°C) and 15 dpi (40.7°C), with no other lesion throughout  
296 the experiment (additional file 2). No hyperthermia was recorded in any of the other cattle at any stage  
297 of the experiment.

298 Overall, the sum of clinical scores in BTV1 and BTV15 single infected groups were not significantly  
299 different when compared to control animals or to the V group after BTV8 challenge ( $P>0.14$ ), but  
300 were significantly lower when compared to first BTV8 challenge in NV calves and the superinfection  
301 with BTV1/15 inoculum in both V and NV groups ( $P<0.002$ ).

302 Necropsy for BTV1 single infection revealed moderate petechial haemorrhages in mesenteric and  
303 mediastinic lymph nodes in one calf, prescapular and mediastinic lymph nodes in another one and no  
304 lesions in the last one.

305 In the calves with BTV15 single infection, no BTV specific lesions could be found at necropsy.  
306 Petechial haemorrhages were reported on the thymus and the prescapular lymph node of two calves,  
307 respectively, in both cases on a limited amount.

308 All the three BTV1 infected calves had a positive BTV1 detection in prescapular lymph nodes. In  
309 addition, viral RNA could be detected in the thymus and spleen of two other calves, respectively  
310 (Figure 3).

311 RTqPCR revealed BTV15 positive detection in the spleen and the prescapular lymph nodes of all of  
312 the three BTV15 infected calves. In addition, thymus and mesenteric lymph node were shown to be  
313 positive in two calves, respectively (Figure 3). The frequency of positive detection in organs was not  
314 different between BTV1 and BTV15 (Fisher's Exact Test for count data,  $P>0.4$ ).

315 No specific lesions or BTV RNA detection could be found in control animals. A few non-specific  
316 abscesses could be found in the lung of one of these calves.

### 317 *Serology*

318 During BTV1 single infection anti-BTV1 neutralizing antibodies could be detected for the first time at  
319 the 16 dpi<sub>single</sub> tested time point and then increased regularly until the end of the experiment (Figure  
320 2A). Similarly, also anti-BTV15 neutralizing antibodies were measured starting from 16 dpi<sub>single</sub> in the  
321 course of BTV15 single infection and the titres increased regularly until the time of euthanasia (Figure  
322 2A).

323 There was no significant difference between SNT titres of BTV1 and BTV15 single infection and  
324 BTV8 after first infection in NV group during the first 35 days ( $P>0.24$ ).

325 BTV1 infected calves seroconverted regarding anti-VP7 antibodies between 7 and 16 dpi<sub>single</sub>, and  
326 were still all seropositive at the end of the experiment.

327 Anti-VP7 Abs in BTV15 infected animals clearly increased between 10 and 15 dpi<sub>single</sub> in all infected  
328 calves (Figure 2B). However, only 1/3 calves seroconverted at the end of the experiment at 35 dpi<sub>single</sub>,  
329 whereas the two other calves remained slightly out the positivity limit (mean PN =38+/- 5.7, Figure  
330 2B). At 35 dpi<sub>single</sub> PN of BTV15 infected group was significantly higher when compared to BTV1  
331 single infection group and BTV8 groups at 35 dpi<sub>BTV8</sub> ( $P<0.007$ ).

332 *In vitro* cross neutralization assay only showed limited cross reactivity between BTV8 immune serum  
333 against BTV1 virus (18 % +/- 7.4 % of the BTV8 immune serum homologous neutralization, Figure  
334 2C). However, this cross reactivity was significantly higher than the one measured between BTV1  
335 versus BTV15 immune serum, BTV8 versus BTV1 and BTV15 immune serums ( $P<0.02$ ). BTV1 and  
336 BTV8 immune serum elicited a limited cross reactivity toward BTV15 (10 % +/- 4.5 % and 10 % +/-  
337 1.8 %, respectively).

### 338 *BTV RNA detection*

339 In the course of BTV1 single infection BTV RNA could be detected as soon as 1 dpi in all the calves  
340 of the group, but then could only be detected again at 2 and 3 dpi for respectively 1 and 2 calves.

341 During BTV15 single infection viral RNA could be detected starting from 7 dpi in all three calves.  
342 BTV15 RNA could be detected until the end of the experiment (Figure 2A). The levels of RNA  
343 peaked in the blood of both BTV1 and BTV15 groups between 9 and 11 dpi<sub>single</sub>. BTV1 cDNA copy  
344 number detected in single infected calves was significantly higher than in BTV1 superinfected ones,  
345 no matter their vaccination status ( $P < 10^{-4}$ ). By contrast, there were no significant differences in cDNA  
346 copy numbers between BTV15 superinfected (V and NV groups), BTV1 and BTV15 single infected  
347 calves ( $P > 0.2$ ). Moreover, BTV1 and BTV15 cDNA copy numbers were not significantly different  
348 from BTV8 cDNA copy number during first infection in NV group ( $P > 0.13$ ).

349

### 350 **Viral growth assay**

351 From 0 to 24-48 hpi, BTV1 showed a faster replication, however not significant, whichever the  
352 considered cell line (Figure 4). In VERO cells BTV15 grew less efficiently than BTV1 from 0 to 48  
353 hpi ( $P < 10^{-5}$ ), but finally reached by 120 hpi similar titres to BTV1 and BTV8 in VERO cells ( $P > 0.14$ )  
354 and to BTV1 in BPAEC ( $P > 0.9$ ). Homologous viral growth was not significantly different between  
355 cell types ( $P = 0.15$ ), and there was no significant differences between serotypes in BPAEC ( $P > 0.3$ ).

### 356 **Discussion**

357 The influence of the existent active immunity towards the European BTV8 strain on the outcomes of a  
358 superinfection with BTV1 was evaluated. The BTV1 inoculum appeared to be contaminated with  
359 BTV15 thus the animals were actually infected with a mixed BTV1-BTV15 inoculum. BTV8 active  
360 immunity was evaluated either by vaccination followed by infectious challenges or by infectious  
361 challenges alone Two successive infections with the same BTV8 strain were realized 4 months apart.  
362 In line with field data [9], vaccination only elicited the production of neutralizing Abs detectable after  
363 the second vaccine boost. Vaccinated animals underwent a significantly earlier detection of  
364 neutralizing antibodies after the first BTV8 challenge when compared to non-vaccinated calves. In the  
365 NV BTV8 group, the detection of BTV8 RNA lasted until 180 dpi<sub>BTV8</sub> (end of the experiment), which  
366 is consistent with currently existing literature data [27, 28]. Non-vaccinated calves infected with

367 BTV8 showed a slight to mild clinical picture. A moderate impact of the disease caused by BTV8 on  
368 cattle is not unusual, in experimental infections [22, 29] as well as in the field [30-32].

369 After the unexpected contamination of the BTV1 inoculum with BTV15, the influence of both viruses  
370 on the outcome of the infection was investigated. In the inoculum used for the superinfection, the copy  
371 number of BTV1 segment 2 cDNA per ml of inoculum was about 10 fold lower than BTV15. After  
372 superinfection BTV1 could be found irregularly and only at a few tested time-points in the blood of  
373 the calves while BTV15 was detected with high levels of RNAemia until the end of the experiment in  
374 both V and NV groups. The overwhelming replication of BTV15 *versus* BTV1 is in line with results  
375 reported by Eschbaumer *et al.*, (2011). Domination of one serotype on another during mixed infection  
376 has been previously reported [33] and the same authors observed that about 5 % of progeny viruses  
377 were actually reassortants. Any genome segment can be involved in reassortment which is readily  
378 generated, as demonstrated by Shaw *et al.* [34]. As in the current study viral RNA was based on  
379 segment 2 quantification, it is not possible to rule out that some of the segment 2 detected by RTqPCR  
380 being actually part of reassortant viruses. This was also one of the hypotheses brought to light by Dal  
381 Pozzo *et al.* (2013) to explain the predominance of BTV8 on BTV1 and BTV15 in the course of a  
382 triple co-infection. Another possible explanation is the viral interference, occurring between two or  
383 more viruses infecting simultaneously the same host.

384 BTV8 was recently used to study underlying IFN-I control mechanisms by the virus [35]. To  
385 investigate whether the different tested serotypes would also show different replication patterns *in*  
386 *vitro* or not, growth curves of BTV1, BTV8 and BTV15 were established in two common cell lines,  
387 BPAEC and VERO cells. VERO cells are deficient for IFN-I production [36]. BTV15 did not replicate  
388 as much as BTV1 in VERO cells during the first 48 hours. Nevertheless, in IFN-I competent cells such  
389 as BPAEC primary line, both BTV serotypes replicated following a similar pattern through time.

390 These results are not inconsistent with the hypothesis of BTV15 to be better adapted to IFN-induced  
391 state when compared to BTV1, possibly explaining the relatively more efficient replication of BTV15  
392 *in vivo* or *in vitro* in IFN-competent cell lines. Another hypothesis to explain the difference between  
393 BTV1 and BTV15 RNAemia after superinfection might be related to the influence of BTV8 immunity

394 on these two serotypes. The level of heterologous reactivity as assessed by SNT between BTV8  
395 immune serum and BTV serotypes 1 and 15 was low, yet higher for BTV1 (18+/- 7.4 % versus 10 +/-  
396 1.8 % for BTV1 and BTV15, respectively; Fig. 2 C). This result is in line with Hund *et al.* (2012),  
397 which reported partial cross neutralization between BTV8 positive serum and BTV1, despite the  
398 genetic distance between these serotypes [9].

399 In addition, no significant difference was reported between RNAemia of BTV15 superinfected calves  
400 (from V and NV groups, both immunized against BTV8) and RNAemia of BTV15 single infected  
401 calves. Thus this low *in vitro* humoral cross reactivity between BTV8 immune serum and BTV15  
402 seemed to have no significant influence on BTV15 RNAemia *in vivo* in contrast to the BTV1  
403 RNAemia.

404 BTV1/BTV15 superinfection led to clinical disease in both V and NV animals. On the contrary,  
405 during BTV1 and BTV15 single infection the calves had very low clinical scores. The reason of this  
406 difference remains uncertain; however individual variability could be part of the explanation.

407 After superinfection, BTV1 neutralizing Abs only reached very low levels, as a consequence of the  
408 very low BTV1 RNA detection. By contrast, BTV15 neutralizing Abs extended to high titres, either in  
409 BTV1/BTV15 superinfected animals or in BTV15 single serotype infected calves.

410 Despite high neutralizing Abs detection following BTV15 single serotype infection, ELISA detecting  
411 VP7 Abs showed a mean PN at 35 dpi just above the positivity threshold. This is consistent with  
412 previous reports showing that significant immunological differences exist between BTV15 and other  
413 BTV serotypes and that monoclonal antibodies raised against BTV1 VP7 failed to react with BTV15  
414 VP7 [37]. When assessing diagnostic tools aiming at non-serotype specific detection, it would be  
415 therefore advisable to include distantly related strains in the test panel to cover most of the genetic  
416 variability displayed by BTV proteins.

417 Vandenbussche *et al.* suggested to use the ID Vet cELISA kit with a cut off of 66 PN instead of 35 for  
418 BTV8, as recommended by the manufacturer, to achieve optimal accuracy for both screening and



419 diagnostic [38]. Taking into account this suggestion, all of the three calves would have been  
420 considered as seropositive by day 21 post infection with BTV15.

#### 421 **Conclusion**

422 Unlike the European BTV8, known for its increased virulence in bovine, BTV1 and BTV15 have been  
423 associated with subclinical or very mild disease in this species. Numerous factors are known to  
424 influence the severity of BT in individual ruminants; nutritional status, immune status and age, breed,  
425 environmental stresses such as high temperature and ultraviolet radiation [39]. . In this study, the  
426 accidental co-infection with BTV1 and BTV15 and the obtained severe clinical outcome underlined  
427 the potential higher pathogenicity of a co-infection.

428 The main objective of this study was to observe the outcomes of a superinfection in calves previously  
429 immunized with the European BTV8. BTV1 or BTV15 RNA detection in superinfected animals was  
430 not different whether BTV8 immunization was acquired through vaccination and challenges or  
431 challenges alone. Furthermore, a low cross neutralization was measured between BTV8 and BTV1,  
432 and between BTV8 and BTV15. Taken all together in the context of the European BT epidemiological  
433 situation, the results could suggest that an infection or a vaccination with the European BTV8 strain  
434 would not efficiently protect the bovines from a superinfection with the BTV1 or BTV15 strains used  
435 in the study.

#### 436 **Competing interests**

437 The authors declare that they have no competing interest.

#### 438 **Author's contributions**

439 CS, ET, FDP, LM, KDC, and PS conceived and designed the experiments. LM, FDP, PS and WVC  
440 performed the experiments *in vivo*. LM, FDP, CT and IDL performed the experiments *in vitro*. LM,  
441 FDP and CS analysed the data. LM, FDP and CS wrote the paper and all the authors approved the  
442 final manuscript.

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453

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577

## 578 **Figures**

### 579 **Figure 1. Neutralising antibodies and viral RNA detection following BTV8 challenges and** 580 **BTV1/BTV15 superinfection.**

581 (A) Mean neutralising antibodies titres and mean copy number of VP2 cDNA are shown for  
582 vaccinated (V) and non-vaccinated (NV) calves, after vaccinations and challenges with BTV8 and  
583 BTV1/BTV15 superinfection. Standard deviation is shown as error bars. The two vaccine injections  
584 are represented as arrows followed by Vacc1 and Vacc2 respectively. Infectious challenges are  
585 represented as arrow labelled with the corresponding serotypes. \* above braces: time period with  
586 neutralising Abs titre of V group significantly higher than NV group;  $P < 0.05$ . (B) Focus on the  
587 BTV1/BTV15 superinfection experiment.

588 **Figure 2. Serology, RNA detection and cross neutralization results following single infections**  
589 **with BTV1, BTV15 and BTV8.**

590 (A) Results of RTqPCR and SNT in V and NV groups of calves, after BTV1 and BTV15 single  
591 infection challenges compared with BTV8 first challenge in NV group. Dashed lines represent SNT  
592 results. (B) cELISA % of negativity following BTV1 and BTV15 single infection, BTV8 vaccination  
593 in V group and first BTV8 challenge in NV group. Dotted line represents positivity threshold, with %  
594 of negativity <35 considered as positive. (C) Cross neutralization results, with BTV1, BTV8 and  
595 BTV15 compared to the respective heterologous immune serum. Results are expressed as a percentage  
596 of the highest homologous neutralization titre. \*:  $P < 0.05$ ; \*\*  $P < 0.01$ .

597 **Figure 3. BTV RNA detection in organs.**

598 (A) BTV1, BTV8 and BTV15 detection in organs following BTV1/BTV15 superinfection. (B) BTV1  
599 and BTV15 detection in organs following single infections. Results are expressed as  $\text{Log}_{10}$  copy  
600 number/100 mg of tissue. Mesent. LN: Mesenteric lymph node. Prescap. LN: Prescapular lymph node.

601 **Figure 4. *In vitro* growth kinetics of BTV1, 8 and 15 on VERO and BPAEC cells.**

602 VERO cells (A) and BPAEC (B) were infected at a multiplicity of infection (M.O.I.) of 0.05 and  
603 supernatants collected at 8, 24, 48, and 120 h post infection (hpi). Viruses were the same as used for  
604 single infection (BTV1 and BTV15) and BTV8 was the same as in the BTV8 successive infection  
605 experiment. Supernatants were then titrated on VERO cells by end-point dilution and the virus titres  
606 expressed as  $\text{log}_{10}$  TCID<sub>50</sub>/ml. Three different assays were performed independently, each time  
607 involving the three serotypes, tested on the different cell lines. Standard deviations are displayed as  
608 error bars.

609 **Additional files**

610 Additional file 1.tif



611 **BTV group specific anti-VP7 antibodies, after vaccination against BTV8 and BTV8 challenges,**  
612 **for non-vaccinated, control and vaccinated calves**

613 BTV group specific anti-VP7 antibodies as the % of negativity. Dashed line represents the cut off  
614 value. A % of negativity under the cut off (35 %) is considered positive. NV: non vaccinated; V:  
615 vaccinated. The two vaccine injections are represented as arrows labelled Vac1 and Vac2 respectively,  
616 and first and second BTV8 challenges are represented as green arrows. Standard deviations are  
617 represented as error bars.

618

619 Additional file 2.csv

620 **Cumulative clinical scores**

621 Cumulative clinical scores after the first BTV8 challenge in NV and V animals, BTV1/BTV15  
622 superinfection and BTV1 and BTV15 single infections. Sd: Standard deviation.

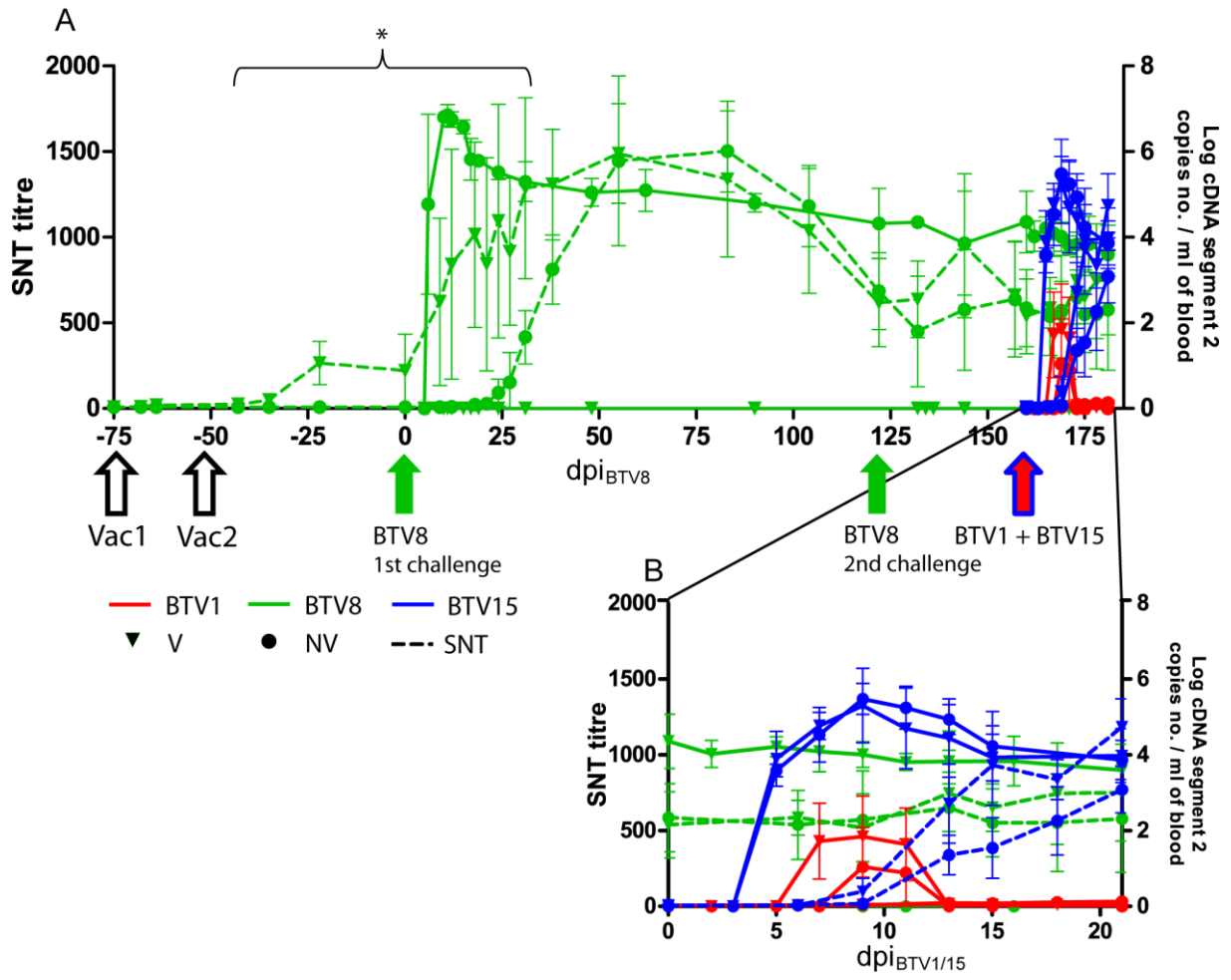
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626 Fig. 1

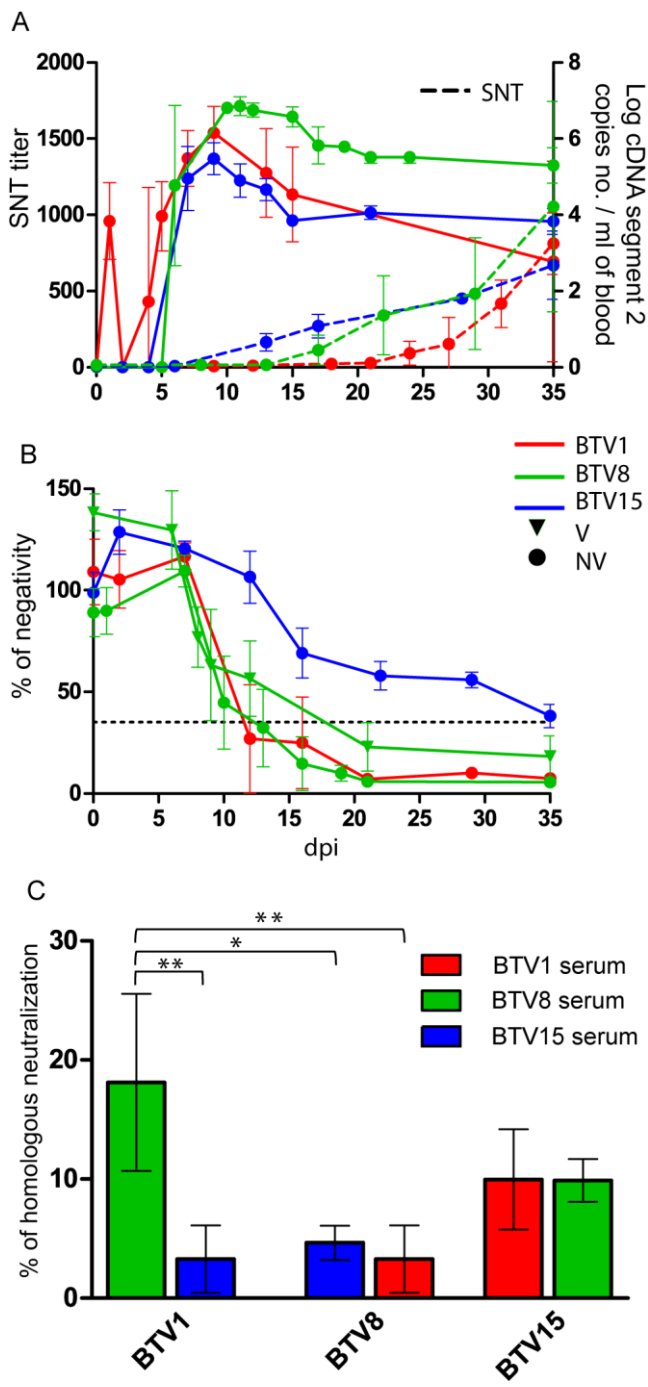
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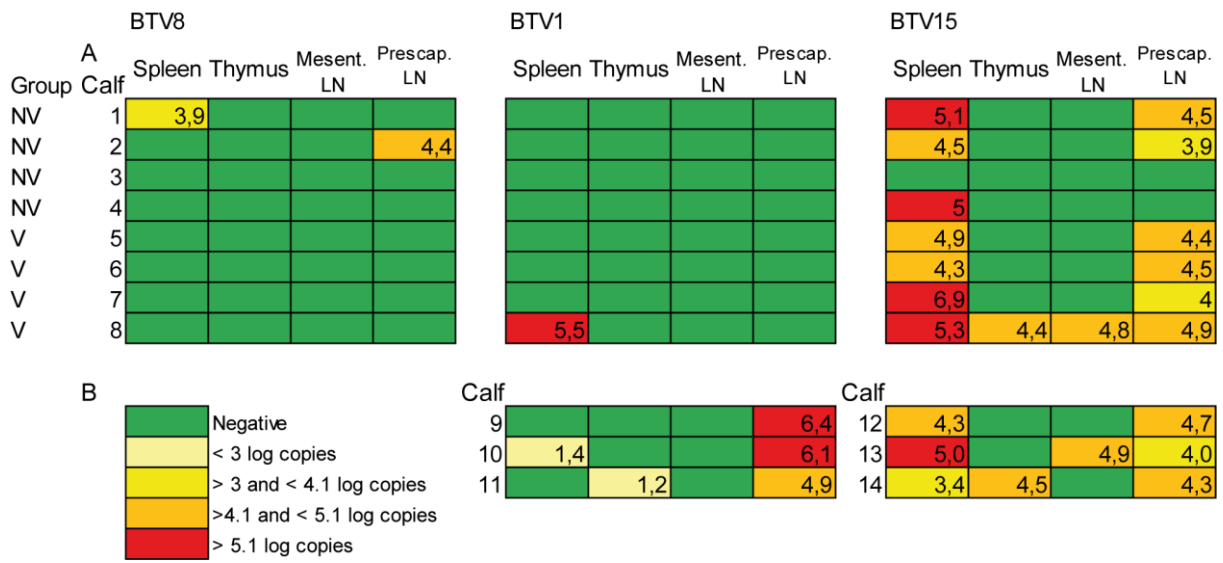
630 Fig. 2



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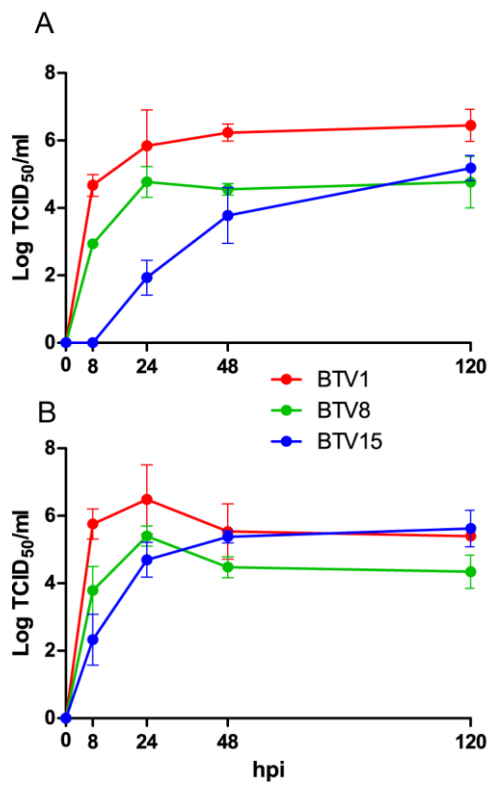
633 **Fig. 3**



634

635

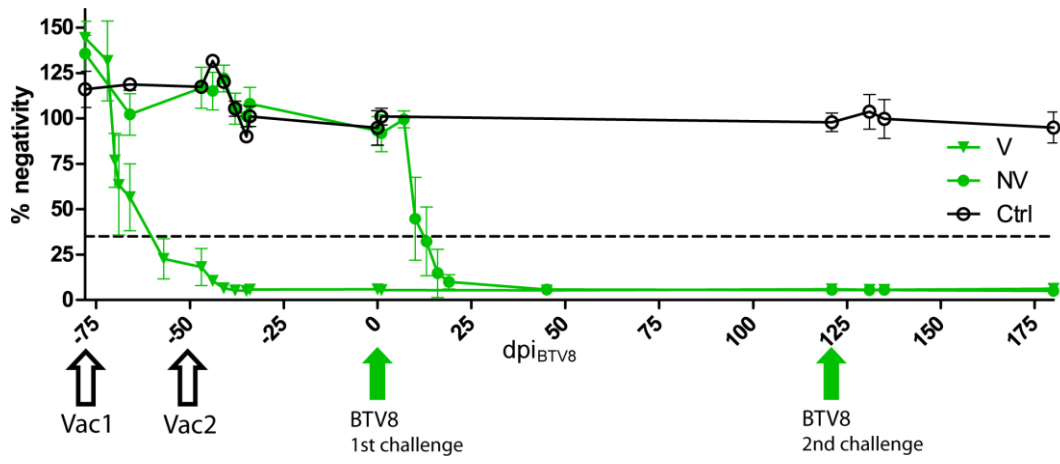
636 Fig. 4



637

638

639 **Supplementary material**



640