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Article

Chicken Interleukin-18 as a Vaccine Adjuvant in a Recombinant Fowlpoxvirus Vaccine for the Protection against Infectious Bursal Disease Virus

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Abstract: In mammals, the role of interleukin-18 (IL-18) in the immune response is to drive inflammatory and, normally therefore, anti-viral responses. IL-18 also shows promise as a vaccine adjuvant in mammals. Chicken IL-18 (chIL-18) has been cloned. The aim of this study was to investigate the potential of chIL-18 to act as a vaccine adjuvant in the context of a live recombinant Fowlpox virus vaccine (fpIBD1) against Infectious bursal disease virus (IBDV). fpIBD1 protects against mortality, but not against damage to the bursa of Fabricius caused by IBDV infection. The Fowlpox virus genome itself contains several candidate immunomodulatory genes, including potential IL-18 binding proteins (IL-18bp). We have knocked out (Δ) the potential IL-18bp genes in fpIBD1 and inserted ($::$) the cDNA encoding chIL-18 into fpIBD1 in the non-essential ORF030, generating five new viral constructs –fpIBD1::chIL-18, fpIBD1 Δ ORF073, fpIBD1 Δ ORF073::chIL-18, fpIBD1 Δ ORF214 and fpIBD1 Δ ORF214::chIL-18. The subsequent protection from challenge with virulent IBDV, as measured by viral load and bursal damage, given by these altered fpIBD1 strains, was compared to that given by the original fpIBD1. The results show that chIL-18 can act as a vaccine adjuvant, giving complete protection against challenge, with no detectable virus or damage in the bursa of Fabricius.

Keywords: chicken interleukin-18; vaccine adjuvant; recombinant Fowlpox virus FP9; fpIBD1

1. Introduction

Infectious bursal disease (IBD) was first recognised in Gumboro, Delaware, USA in 1962 [1], hence its alternative name, Gumboro disease. IBD causes considerable economic losses in the poultry industry throughout the world by inducing severe clinical signs, immunosuppression and a high mortality rate (may range from 1% to more than 50%) in infected chickens [2]. Before 1987, IBD was satisfactorily controlled by vaccination. Since 1987, however, vaccination failures have been described in different parts of the world, due to the emergence of variant and, later on, very virulent strains of IBDV. In the USA, the new strains are characterised by an antigenic variation that shows only a slight increase in virulence and are therefore called “variant” strains [3]. In Europe, IBDV strains still belong to classical serotype 1 strains but are characterised by a marked increase in virulence and are therefore called “hypervirulent” or “very virulent” IBDV (vvIBDV) strains [4].

Recombinant Fowlpox viruses have been used to express genes from a number of poultry pathogens, such as Newcastle disease virus [5,6], Avian influenza virus [7], Turkey rhinotracheitis virus [8], Marek's disease virus [9] and IBDV [10].

IBDV is a Birnavirus, characterised by having a bisegmented double stranded RNA genome [2]. IBDV encodes 5 proteins in which VP2 is a capsid protein. VP2 has been identified as the host-protective antigen [11]; hence it has been the focus of attempts to produce new vaccines by

recombinant DNA technology. VP2 was expressed in the FP9 strain of Fowlpox virus [12] to generate a recombinant vaccine, fpIBD1 [10]. Significant levels of protection were provided by vaccination with this recombinant. fpIBD1 afforded protection against mortality, but not against damage to the bursa of Fabricius [10]. In addition, the protective effect of the fpIBD1 vaccine was dependent on the titre of challenge virus and the major histocompatibility complex (MHC)-haplotype of the vaccinated chicken [13]. The current live attenuated vaccines in commercial use provide complete protection.

IL-18, also known as interferon-gamma-inducing factor (IGIF), is a pro-inflammatory cytokine that plays an important role in the development of T-helper type 1 (Th1) cells, which drive cell-mediated immune responses. As IL-18 is an inducer for Th1 response, it therefore seemed logical to investigate the efficacy of IL-18 as a vaccine adjuvant. Chicken IL-18 (chIL-18), originally identified in an EST database, was cloned and expressed [14].

It was claimed that the ORF FPV073 (in the Fowlpox virus genome) was a homologue of human IL-18bp and an orthologue of IL-18bps from *Molluscum contagiosum* virus, Swinepox virus and Vaccinia virus [15]. It was later shown that ORF FPV214 is more likely to be the correct assignment [12]. Thus FPV214, but not FPV073, aligns with a conserved motif (97YWxxxxxFIEHL108 in humans) in the other IL-18bps. In contrast, FPV073 contains a GxGxxG nucleotide-binding motif and shows highest similarity to a tyrosine protein kinase. It seemed logical to delete IL-18bp from vector containing chIL-18 as an adjuvant. Therefore, both Fowlpox virus ORFs 073 and 214 were knocked-out, separately, from fpIBD1. ChIL-18 was then inserted into the non-essential [16] PC-1 gene (ORF FPV030) of fpIBD1. Then the protection provided by these new vaccines was compared to the original fpIBD1, in terms of clinical signs, bursal damage and viral loads in the bursa of Fabricius following challenge with virulent IBDV.

2. Materials and Methods

2.1. Chickens

Rhode Island Red (RIR) chicks were obtained from an unvaccinated flock maintained in isolation accommodation at the Institute for Animal Health, Compton, UK. The parents were confirmed to be free of antibodies to IBDV, chicken infectious anaemia virus, Marek's disease virus, reovirus and a number of other pathogens, so the chicks used in these experiments were deemed to be free of maternal antibodies against IBDV. The experiments met with local ethical guidelines as well as those of the UK Home Office.

2.2. Viruses

Fowlpox virus FP9 derivative fpIBD1 [10], expressing most of the IBDV F52/70 VP2 protein as a β -galactosidase fusion protein under the control of the Vaccinia virus p7.5 early/late promoter, from the BglII insertion site in ORF FPV002, was from laboratory stocks. fpIBD1 was grown on chicken embryo fibroblast (CEF) cells in the presence of 1X 199 medium (Sigma).

The vIBDV strain F52/70 [17] was used. The titre of virus stock was kindly determined by Dr Adriaan van Loon (Intervet BV, The Netherlands) [18]. Based on earlier studies, the dose of virus selected was 102.3 EID₅₀ vIBDV strain F52/70, which can overcome the protection provided by fpIBD1 and cause bursal damage, measured as the bursal lesion score in 2-3 week-old RIR chicks (Davison TF, personal communication).

2.3. Generation of novel fpIBD1 recombinants

fpIBD1 mutants carrying only deleted forms of the putative IL-18bp genes were isolated by trans-dominant selection [19] using selection for the *Escherichia coli* guanine phosphoribosyltransferase (gpt) gene and checked by PCR, essentially as described previously [16]. Deleted recombination constructs, containing 50 bp from either end of Fowlpox virus ORFs 073 and 214 as well as 500 bp flanking sequences, were, however, assembled by two-stage overlapping PCR before insertion into the BamHI and HindIII restriction sites of vector pGMR [16]. The constructs were then transfected, using Lipofectin (Invitrogen), into CEF infected with fpIBD1. After overnight

incubation at 37°C in 5% CO₂, the culture medium was discarded and replaced with 1X 199 medium + 2% new born bovine serum (NBBS) containing mycophenolic acid, xanthine and hypoxanthine (MXH) solution and reincubated for 2-4 days until a cytopathic effect was apparent. The virus was released by freeze/thawing the culture three times then plaque out under MXH selection.

Gpt⁺ recombinant clones were plaque-purified three times in MXH selective medium then further purified without selection until they became gpt⁻ (as determined by failure to plaque under MXH selective medium), that was confirmed by PCR.

For production of recombinants expressing chIL-18, a recombination vector was constructed. Briefly, pGEM-T::chIL-18 was digested with NotI, chIL-18 was then inserted into pEFgpt12S (containing S promoter). Sp+chIL-18 was then inserted into the PC-1 plasmid (pFPV-PC-1) to result in pFPV-PC-1::Sp+chIL-18, which was the target clone in which the chIL-18 gene, under the control of a synthetic early-late promoter, was inserted into the non-essential PC-1 gene (ORF FPV030) in a plasmid carrying the gpt gene under the control of the Vaccinia virus p7.5 promoter. CEF infected with parental fpIBD1, fpIBD1Δ073 and fpIBD1Δ214 viruses were then transfected with the pFPV-PC-1::S-promoter/chIL-18 plasmid. Gpt⁺ recombinants were selected and plaque purified three times. Insertion of the expression and selection cassette into the PC-1 gene was confirmed by PCR.

The six different recombinant viruses were titrated. Viruses carrying the gpt reporter gene (i.e. those that contain chIL-18) always had lower titres (Table 1).

Table 1. Titration of the six vaccines.

Virus	Final concentration (pfu/ml)
fpIBD1	1X10 ⁹
fpIBD1Δ073	2X10 ⁹
fpIBD1Δ214	1X10 ⁹
fpIBD1::IL-18	2X10 ⁸
fpIBD1Δ073::IL-18	2X10 ⁸
fpIBD1Δ214::IL-18	3X10 ⁸

2.4. Experimental design

Chicks received an initial vaccination at 1 week of age with 107 pfu fpIBD1 or manipulated fpIBD1 in a 50 µl volume. The inoculum was placed on the wing-web and the skin punctured 30 times over an area of 2 mm² with a 21-gauge hypodermic needle. The same procedure was repeated two weeks later to provide a booster vaccination. Chickens were challenged 10 days after final vaccination with 102.3 EID₅₀ IBDV strain F52/70 in a total volume of 100 µl by the intranasal route (50 µl in each nostril). For RNA preparation, blood samples (50 µl) were taken from a wing vein immediately into 350 µl RTL buffer every day after challenge (from the infected unvaccinated group). Five days after challenge, all birds were killed and the bursae removed for RNA, immunohistochemistry and H&E staining.

2.5. Sample processing

2.5.1. RNA extraction

Bursal tissue (~30 mg) was homogenised using a Bead mill (Retsch MM300). Briefly, the bursal tissue was placed in a 2 ml Safe-lock Eppendorf tube with 600 µl lysis buffer RLT. A 5mm stainless steel bead was added per tube. The tubes were placed in adaptors for the bead mill and run for 4 min at 20 Hz. Total RNA was then prepared from the homogenised tissues and the blood using the RNeasy mini kit (Qiagen) following the manufacturer's instructions. Purified RNA was eluted in 50 µl RNase-free water and stored at -70°C.

2.5.2. Frozen sections for immunohistochemical staining

Each bursal sample was put on a 2.5 cm² cork tile and covered with Tissue-Tek® O.C.T.M Compound. The samples were then snap-frozen in a dry-ice/iso-pentane bath and transferred to liquid nitrogen. Frozen blocks were then removed from the liquid nitrogen, wrapped in aluminium foil and stored at -70oC. Sections (6-8 m) were then cut from these blocks for immunohistochemistry staining using a cryostat, picked up onto glass slides, then fixed in acetone for 10 min and air-dried. Staining was then carried out using a Vectastain® ABC αmouse IgG HPR staining kit (Vector Laboratories, Burlingame, CA, USA), following the manufacture’s instructions. The monoclonal antibodies used were R63 [3] for IBDV and AV20 [20] for B cells.

2.5.3. H&E staining

Each bursal section from every bird was put into 40 ml formalin and stained with haematoxylin and eosin (H&E) to look for bursal damage.

2.6. Real-time quantitative RT-PCR

Quantitative real-time RT-PCR was carried out for IBDV and 28S [21]. IBDV-specific oligonucleotides were identified from genomic segment A [22]. The fluorescently labelled probes were labelled with the reporter dye 5-carboxyfluorescein (FAM) at the 5’ end and the quencher N,N,N’,N’-tetramethyl-6-carboxyrhodamine (TAMRA) at the 3’ end. Specific primers were designed to closely flank the probe. Primers and probes sequences are given in Table 2.

Table 2. Primers and probes for real-time quantitative RT-PCR.

RNA target	Primer/Probe*	Sequence (5’-3’)
28S	F	GGC GAA GCC AGA GGA AAC T
	R	GAC GAC CGA TTT GCA CGT C
	Probe	AGG ACC GCT ACG GAC CTC CAC CA
IBDV (VP2)	F	GAG GTG GCC GAC CTC AAC T
	R	AGC CCG GAT TAT GTC TTT GAA G
	Probe	TCC CCT GAA GAT TGC AGG AGC ATT TG

RT-PCR was carried out using reagents from the TaqMan® EZ RT-PCR kit (PE Applied Biosystems). Amplification and detection of specific products were undertaken using the ABI PRISM™ 7700 Sequence Detection System with the following cycle profile: 1 cycle of 50°C for 2 min, 1 cycle of 96°C for 5 min, 1 cycle of 60°C for 30 min, 1 cycle of 95°C for 5 min and 40 cycles of 94°C for 20 sec and 59°C for 1 min. Quantification was based on the increased fluorescence detected by the ABI PRISM™ 7700 Sequence Detection System (PE Applied Biosystems) due to hydrolysis of the target-specific probes by the 5’ nuclease activity of the rTth DNA polymerase during PCR amplification. A passive reference dye ROX (present in the EZ reaction buffer), which is not involved in amplification, was used to correct for fluorescent fluctuations resulting from changes in the reaction conditions for normalisation of the reporter signal. Results are expressed in terms of the threshold cycle value (Ct), the cycle at which the change in the reporter dye (ΔRn) passes a significance threshold.

2.7. Construction of standard curves for quantitative PCR and RT-PCR assays

To generate standard curves for the 28S rRNA-specific reaction, total RNA, extracted from stimulated splenocytes, was serially diluted in sterile RNase-free water and dilutions made from 10⁻¹ to 10⁻⁵. To generate standard curves for the IBDV, total RNA was extracted from 50 l IBDV stock and serially diluted similarly. Regression analysis of the mean values of 6 replicate RT-PCRs for the log₁₀ diluted RNA was used to generate standard curves.

3. Results

3.1. Protection from IBDV challenge by the recombinant vaccine fpIBD1 and the new viral constructs

Within 2 days, birds vaccinated with parental fpIBD1 or manipulated fpIBD1 developed pocks, at the site of inoculation, which disappeared by 10 days. Protection from virulent IBDV challenge by the parental recombinant vaccine fpIBD1 and the altered fpIBD1 was measured by the appearance of IBD clinical signs, using the bursal damage scoring index of Muskett et al. [23] (Table 3 & Figure 1), and viral loads (IBDV) in the bursa, using immunohistochemical staining and real-time quantitative RT-PCR, at 5 days post-infection (dpi).

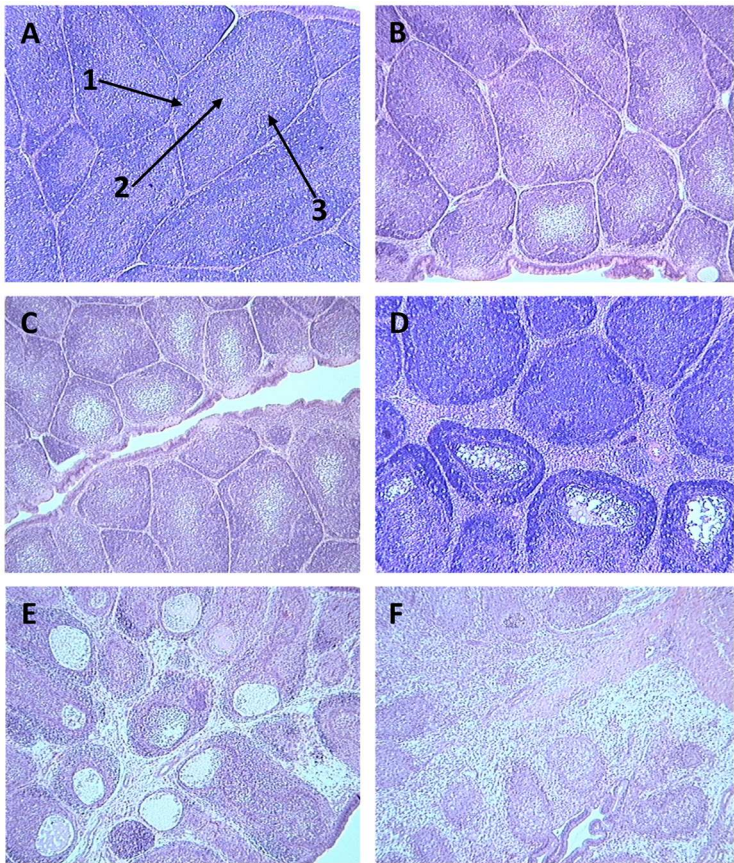


Figure 1. Sections of bursae stained with H&E from birds infected with vIBDV strain F52/70, showing different bursal damage scores as detailed in Table 3. A = bursal damage score of 0, B = bursal damage score of 1, C = bursal damage score of 2, D = bursal damage score of 3, E = bursal damage score of 4, F = bursal damage score of 5. Arrows indicate the cortex (1), medulla (2) and cortico-medullary junction (3) of a bursal follicle.

Table 3. Bursal damage scores for different groups after infection with vIBDV based on the histological scoring system* of Muskett et al. [23].

Vaccination	IBDV challenge	Bursal Damage Score					
		0	1	2	3	4	5
-	-	5	-	-	-	-	-
-	+	-	-	-	1	3	1
fpIBD1	+	1	3	1	-	-	-
fpIBD1Δ073	+	2	2	1	-	-	-
fpIBD1Δ214	+	4	1	-	-	-	-
fpIBD1::IL-18	+	5	-	-	-	-	-
fpIBD1Δ073::IL-18	+	5	-	-	-	-	-
fpIBD1Δ214::IL-18	+	5	-	-	-	-	-

Severe bursal damage was observed for all infected, unvaccinated birds (Table 3). Four out of five (80%) fpIBD1-vaccinated birds were not protected. Three birds out of five (60%) vaccinated with fpIBD1 Δ 073 were not protected. Only one bird out of five (20%) vaccinated with fpIBD1 Δ 214 was not protected. Interestingly, no bursal damage was seen for all birds vaccinated with viral constructs containing IL-18.

Massive depletion of B cells was observed in the bursae of infected, unvaccinated birds. Some B cell depletion was seen in the bursae of birds vaccinated with fpIBD1. Very little B cell depletion was seen in the bursae of birds that were vaccinated with the knockout viruses 073 and 214 and no B cell depletion was observed in the bursae of birds that were vaccinated with the viruses that contained chIL-18 (Figure 2).

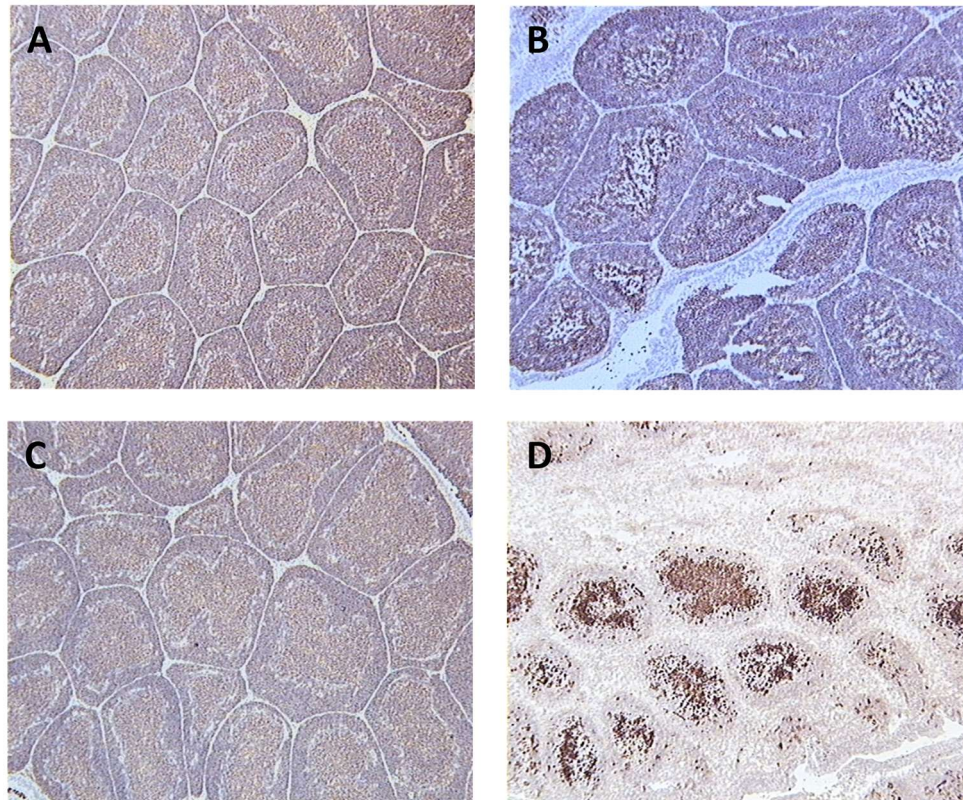


Figure 2. Bursal tissue sections from different groups taken at 5 dpi. with the vIBDV strain F52/70. The sections were stained with the anti-B cell monoclonal antibody, AV20, and counterstained with haematoxylin. A = unvaccinated and unchallenged (-ve control), B = vaccinated with fpIBD1 and challenged, C = vaccinated with fpIBD1 Δ 214::IL-18 and challenged, D = unvaccinated and challenged (+ve control).

3.2. Detection of IBDV using immunohistochemistry

The bursae of unvaccinated and challenged chickens were swamped with the virus. Massive destruction of the bursae had taken place. In contrast, no virus was detected in the bursae of vaccinated chickens (Figure 3).

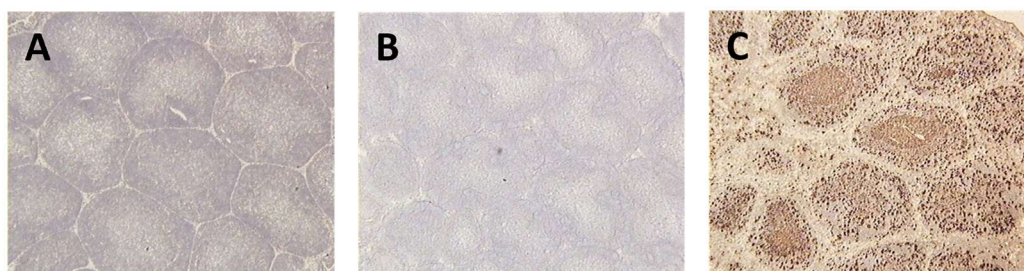


Figure 3. Bursal tissue sections stained with the anti-IBDV monoclonal antibody, R63, and counterstained with H at 5 dpi. A = Unvaccinated and unchallenged, B = vaccinated with fpIBD1 and challenged, C = unvaccinated and challenged. Challenge was with $10^{2.3}$ EID₅₀ vIBDV strain F52/70.

3.3. Detection of IBDV using real-time quantitative RT-PCR

Viral RNA in 50 μ l of whole blood extracted from infected chickens (unvaccinated) over the first 5 dpi with vIBDV strain F52/70 (every 24 h) was quantified. The total number of birds in this group was 7 birds. 5 birds came to the clinical end-point by 3 dpi and 2 birds survived until the end of the experiment (5 dpi). The results show that IBDV levels in the blood reach a peak at 3 dpi. The two surviving birds had lower levels of IBDV in the blood compared to the other birds (Figure 4).

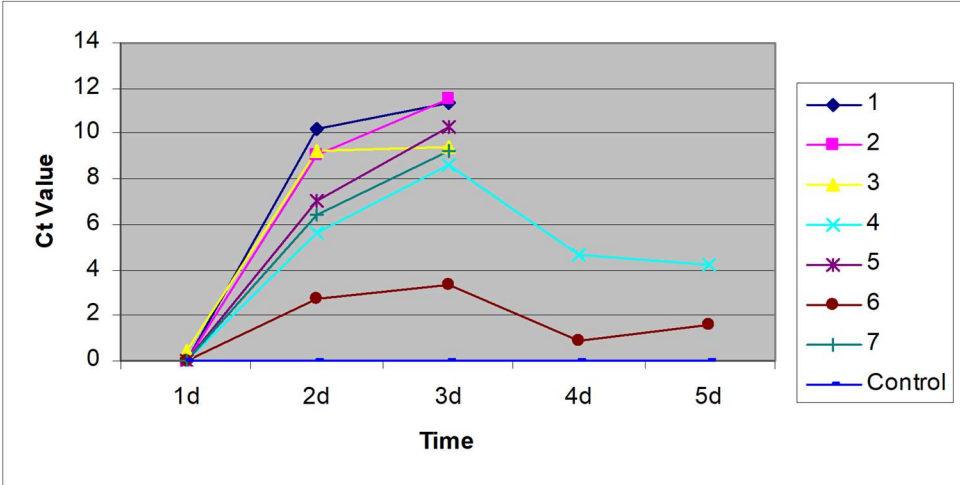


Figure 4. IBDV mRNA levels in the blood of an unvaccinated birds (-ve control) and birds challenged with IBDV $10^{2.3}$ EID₅₀ strain F52/70 were quantified by real-time quantitative RT-PCR. Birds nos 4 and 6 survived until the end of the experiment, whereas the other birds reached the clinical end-point by 3 dpi. Results are expressed as 40-Ct values.

IBDV was detected in the bursa at very high levels in infected, unvaccinated birds (Figure 5). There was a low level of IBDV in the bursae of birds vaccinated with fpIBD1 or with fpIBD1 Δ 073 and lower levels still in the bursae of birds vaccinated with fpIBD1 Δ 214. Interestingly, hardly any IBDV was detected in the bursae of birds vaccinated with fpIBD1::IL-18, and even more interestingly, no virus was detected in the bursae of birds vaccinated with either fpIBD1 Δ 073::IL-18 or fpIBD1 Δ 214::IL-18 (Figure 5). The experiment was repeated and similar results were obtained.

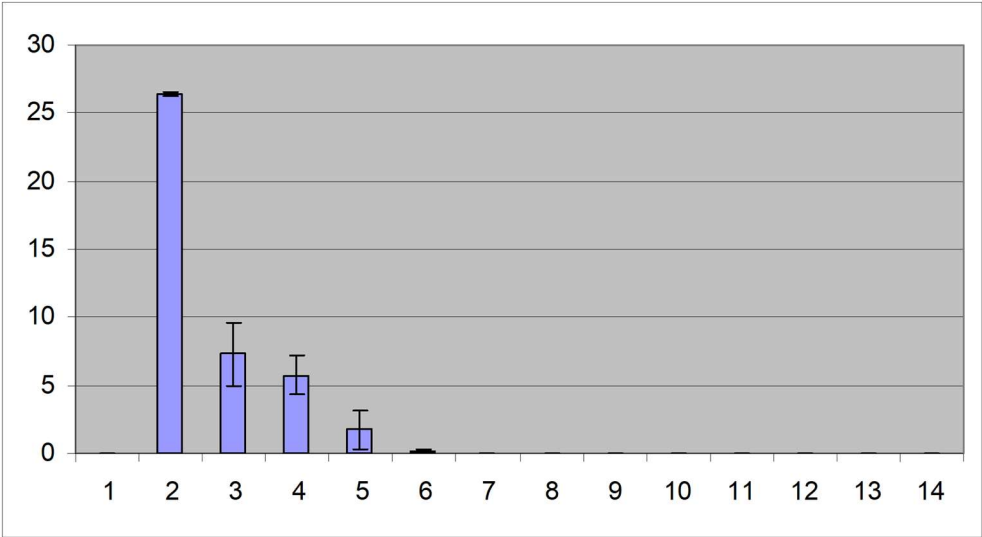


Figure 5. Viral loads (vIBDV strain F52/70) in the bursa 5 d.p.i. (1): -ve control; (2): +ve control (vIBDV); (3): fpIBD1; (4): fpIBD1Δ073; (5): fpIBD1Δ214; (6): fpIBD1::IL18; (7): fpIBD1Δ073::IL18; (8): fpIBD1Δ214::IL18; (9): fpIBD1; (10): fpIBD1Δ073; (11): fpIBD1Δ214; (12): fpIBD1::IL18; (13): fpIBD1Δ073::IL18; (14): fpIBD1Δ214::IL18; (Groups 2-8): Challenged with vIBDV; (Groups 9-14 and group 1): No challenge.

4. Discussion

In previous studies, the recombinant vaccine fpIBD1 protected outbred Rhode Island Red chickens against mortality induced by virulent (F52/70) and very virulent (CS89) strains of IBDV, but not against damage to the bursa of Fabricius [10]. Successful vaccination with fpIBD1 is dependent on the titre of challenge virus, for high titres of challenge virus were able to overcome protection induced by fpIBD1, whereas challenge with a low titre of virus did not [13]. Hence, we decided to use a high titre (102.3 EID₅₀) of IBDV strain F52/70 to overcome the protection provided by the original fpIBD1 and to cause bursal damage.

Cell mediated immunity is involved in protection against challenge with IBDV after vaccination with fpIBD1, as there are no detectable antibodies against IBDV after vaccination and before challenge with IBDV, although there are high levels of antibodies against Fowlpox virus [10,13]. IBDV strain F52/70 induces a Th1 response following infection. IL-18 is an inducer of the Th1 response. It therefore seemed logical to investigate the efficacy of IL-18 as a vaccine adjuvant with fpIBD1. Including chIL-18 in the fpIBD1 vaccine could result in more rapid clearance of the vaccine from the host, again with alteration in the magnitude of the immune response to the vaccine.

The Fowlpox virus genome contains several immunomodulatory genes, including a postulated IL-18bp [12,15]. It seemed logical to delete this gene from vectors containing chIL-18 as an adjuvant. However, Fowlpox virus presumably uses its vIL-18bp as part of a strategy to avoid the host immune response. Deleting the vIL-18bp gene might, therefore, have had an adverse effect on the persistence of the fpIBD1 vaccine in the host and could, therefore, have altered the magnitude of the immune response to the vaccine.

The data indicate that chIL-18 can act as a vaccine adjuvant. Despite the use of a challenge dose of IBDV high enough to overcome protection and cause bursal damage in birds vaccinated with fpIBD1, there was no bursal damage in birds vaccinated with fpIBD1::IL-18, fpIBD1Δ073::IL-18 and fpIBD1Δ214::IL-18 (Figure 1). Furthermore, no IBDV was detected in the bursae of birds vaccinated with fpIBD1Δ073::IL-18 and fpIBD1Δ214::IL-18 (Figure 5). The results also indicate that, as suggested [12], ORF214 is the better candidate for IL-18bp, as fpIBD1Δ214 showed significantly better protection than fpIBD1 or fpIBD1Δ073 (Figure 5 & Table 3).

5. Conclusions

In conclusion, we believe our data indicate that chIL-18 can act as a vaccine adjuvant with fpIBD1 when challenging with a virulent strain of IBDV. It will be interesting to investigate if chIL-18 can act similarly when challenging with a very virulent strain of IBDV using the same experimental model.

Author Contributions: Conceptualization, I.E. and P.K.; methodology, I.E.; validation, I.E., L.R., M.S. and P.K.; investigation, I.E.; data curation, I.E., L.R.; writing—original draft preparation, I.E.; writing—review and editing, I.E., L.R., M.S. and P.K.; supervision, P.K. All authors have read and agreed to the published version of the manuscript.

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Institutional Review Board Statement: The study was conducted in accordance with Animals (Scientific Procedures) Act 1986. The animal study was reviewed and approved by the ethics committee of the Institute for Animal Health and the United Kingdom Government Home Office.

Data Availability Statement: All data supporting the findings of this study are available within the manuscript. Any additional data are available from the corresponding author upon reasonable request.

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Conflicts of Interest: The authors declare no conflict of interest.

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