


## Article

# Orally Fed Recombinant *Lactococcus lactis* Displaying Surface Anti-Fimbrial Nanobodies Protects Piglets against *Escherichia coli* Causing Post-Weaning Diarrhea

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**Abstract:** Post-weaning diarrhea (PWD) and edema disease (ED), caused by enterotoxigenic and Shiga toxin producing *Escherichia coli* (ETEC and STEC) strains, are important diseases of newly weaned piglets worldwide. The objective of this study is to develop a passive immunization strategy to protect piglets against PWD and ED using recombinant *Lactococcus lactis* added to piglet diet at weaning. The Variable Heavy chain domains of Heavy chain antibodies (VHHs) or Nanobodies (Nbs), directed against the fimbrial adhesins FaeG (F4 fimbriae) and FedF (F18 fimbriae) of *E. coli* were cloned and expressed on the surface of *L. lactis*. In vitro, the recombinant *L. lactis* strains agglutinated and inhibited adhesion of cognate F4 or F18 fimbriae expressing *E. coli* to pig villous preparation. In vivo, the anti-F4 *L. lactis* protected weaned piglets against a challenge with an F4-positive ETEC strain. Piglets supplemented with oral anti-F4 *L. lactis* showed reduced fecal *E. coli* shedding. We concluded that the surface expressed Nanobodies on *L. lactis* neutralized the adhesins of targeted *E. coli* and abrogated gut colonization, the first step in disease pathogenesis. As a proof of concept, we demonstrated the potential of passive immunization with recombinant *L. lactis* as a viable alternative to antibiotic prophylaxis in preventing piglet-post-weaning diarrhea.

**Keywords:** post-weaning diarrhea; edema diseases; ETEC; STEC; VHH; Nanobodies; passive immunization; Lactococcus



**Citation:** Okello, E.; Moonens, K.; Erume, J.; De Greve, H. Orally Fed Recombinant *Lactococcus lactis* Displaying Surface Anti-Fimbrial Nanobodies Protects Piglets against *Escherichia coli* Causing Post-Weaning Diarrhea. *Agriculture* **2021**, *11*, 186. <https://doi.org/10.3390/agriculture11030186>

Academic Editors: Nicole Kemper and Grzegorz Woźniakowski

Received: 28 January 2021

Accepted: 19 February 2021

Published: 24 February 2021

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## 1. Introduction

Post-weaning diarrhea (PWD) and edema disease (ED) caused by enterotoxigenic and Shiga toxin producing *Escherichia coli* (ETEC and STEC) are a major cause of economic losses to pig farmers worldwide [1]. The losses result from mortality, morbidity, and the cost of treatment. The key virulence factors of these pathogenic strains are F4 and/or F18 fimbriae expressed in various combinations with heat labile toxin (LT), heat stable toxin (STa & STb) and Shiga toxin (Stx2e) [2–7]. Fimbriae are multi-subunit proteinaceous surface appendages that bind to specific receptors on enterocytes, enabling the bacteria to colonize the gut epithelium, the first step in disease pathogenesis [3]. Following colonization, the strains express toxins that induce secretory diarrhea or edema disease by interfering with water and electrolyte transport across epithelial cells, or toxic endothelial cell apoptosis, respectively [8–10].

Fimbriae are highly immunogenic and stimulate both serum and mucosal immune responses [11]. The peak antibody levels occur at 1 week (F4-specific IgA) and 3 weeks (F18-specific IgA) post infection respectively [12]. This immune response is dependent on

receptor recognition of the antigen, since the receptor negative piglets tolerate fimbriae as normal food antigens [13]. Vaccination of sows with fimbriae or fimbrial subunits is the current strategy for protecting neonatal piglets from ETEC infections through the induced lactogenic immunity [14]. However, active immunization inducing an effective intestinal mucosal immunity is required to protect post-weaning piglets following withdrawal of milk antibodies [15]. The experimental immunization of piglets orally with purified F4 fimbriae or the adhesive subunit (FaeG) induced protective F4-specific mucosal immune response in piglets [16,17]. Similarly, oral immunization with purified F18-fimbriae induced specific immune responses to the major subunit FedA, but the immune response against the FedF adhesin was weak and non-protective against F18+ *E. coli* challenge [18].

During the suckling period, antibodies in milk neutralizes oral vaccines and prevent the development of protective mucosal immunity. Since the diseases occur immediately after weaning, there is no sufficient time for the piglets to mount an immune response, and thus precludes the use of active immunization to prevent post-weaning diarrhea. Attempts to circumvent this problem using an encapsulated or enteric coated subunit and live vaccines resulted in improved immune response in mice models [19] and partial protection of challenged piglets [15,20].

Oral immunization of suckling piglets with *E. coli* or fimbriae, microencapsulated using spray dried poly (lactide-co-glycolide) microspheres, did not induce significant antibody levels 19 days after the booster dose, and the piglets were not protected from challenge post-weaning [21]. Experiments in mice showed that the microencapsulation of F4 and F18 fimbriae with thiolated eudragit microspheres for oral immunization resulted in increased antigen-specific IgA and IgG in serum and proliferation of immune cells in the spleen, lamina propria and the Peyer's patches [19]. Although the results were promising, the results were not validated in suckling piglets in the presence of milk antibodies and other inhibitory factors.

Passive immunization protects against gut colonization following withdrawal of milk antibodies at weaning, allowing the piglets to adapt to stressful changes during this period. Subsequent challenge with pathogenic strains would be better tolerated, since older piglets mount better immune responses and the fimbrial receptors diminish with age [22]. Passive immunization by the in-feed addition of serum-dried plasma antibodies or egg yolk antibodies from immunized chicken during the first two to three weeks after weaning was shown to prevent PWD [23,24]. However, the use of animal products in feed for farm animals is limited by the cost of production and regulatory restrictions, due to public health concerns [25]. Cost effective passive immunization can be achieved using recombinant antibodies expressed cheaply in microbial systems. *Lactococcus* is the microbial system of choice, due to its GRAS (Generally Regarded As Safe) status. Recent research studies have demonstrated the potential for use of engineered *Lactococcus* strains for passive immunization by secreting the antigens [26] or displaying subunit antigens and antibody fragments on the bacterial surface using anchor proteins [27–29].

The variable domains derived from llama Heavy chain antibodies (HCAb), also known as Nanobodies (Nbs), fused to Fc domains are suitable proteins for passive immunization, due to the ease of expression and scale up for bulk production in plant and yeast systems [30]. The HCAb consists only of the two heavy chains and lacks the two light chains found in conventional antibodies [31]. Nanobodies are derived from the variable domain of HCAb and can bind antigens despite having only VH domains. They represent a large diversity in their structural repertoire of antigen-binding sites and antigen-binding characteristics [32]. Nanobodies are small proteins (15 kDa) that are very soluble and very stable allowing for ease of expression in various heterologous systems. Nanobodies have recently gained wide application in animal production for research, diagnostics, therapy, and immunization. Studies reporting on diagnostic applications include the development of a specific nanobody for rapid and selective determination of *Salmonella enteritidis* in milk and a nanobody-based lateral flow assay to detect active *Trypanosoma congolense* infections in cattle [33,34]. Similarly, several studies have reported on the use of Nanobodies

for passive immunization, through systemic or oral administration, to protect animals against viral and bacterial infections such as foot and mouth disease, gastroenteritis, and *Clostridium difficile* [35–39].

The goal of this study is to develop a passive immunization strategy against post-weaning diarrhea in piglets using neutralizing Nanobodies, directed against fimbrial adhesins FaeG and FedF. All the Nanobodies used in this study were developed in our laboratory, with known structures and inhibitory properties [40,41]. Related studies had already demonstrated the neutralizing and protective effect of anti-FaeG Nb expressed in *Arabidopsis thaliana* and yeast cells, when added to feeds during weaning period [35,37]. These Nbs were expressed on *Lactococcus lactis* by surface display using the anchor domain of protein A of *Staphylococcus aureus* [42]. When added in feeds, the covalently linked Nbs on the *L. lactis* surface specifically recognized and neutralized the fimbrial adhesins on targeted ETEC and STEC strains and prevented intestinal colonization, the first step in disease pathogenesis.

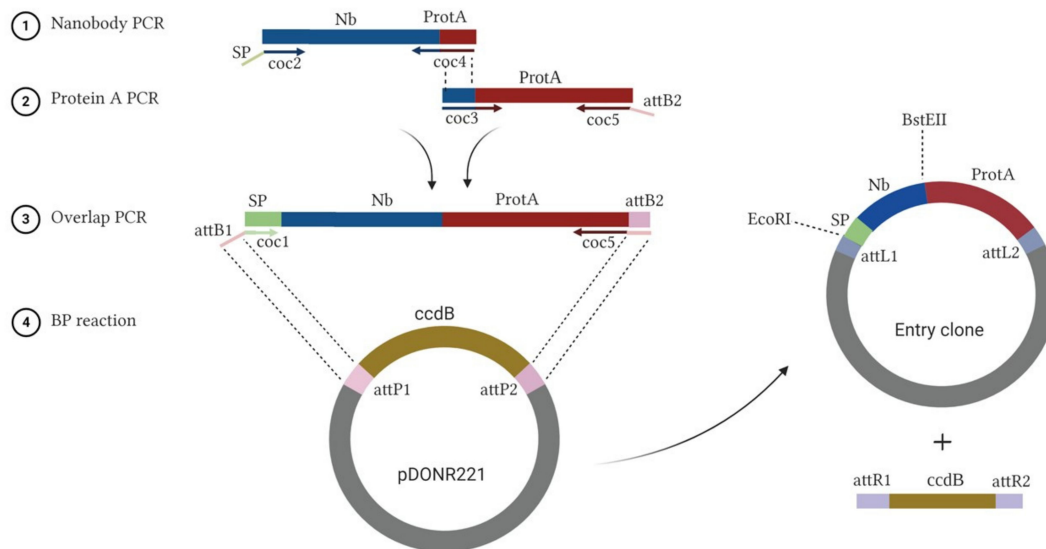
## 2. Materials and Methods

The fusion genes of different Nbs and protein A (ProtA) sequences from *S. aureus* were inserted into pTRKH3 shuttle vector [43] adapted in-house to the Gateway cloning system (Thermo Fisher Scientific) and transformed into the expression strain *L. lactis* MG1363 [44]. pTRKH3 is a shuttle cloning vector for *E. coli* and Gram-positive bacteria with a high copy number (45–85) in streptococcal and lactococcal hosts. The vector contains tetracycline and erythromycin genes; tetracycline resistance is only expressed in *E. coli*, while erythromycin resistance is expressed both in *E. coli* and Gram-positive bacteria.

### 2.1. Construction of Nb-ProtA Recombinant Fusion Genes

The Nb coding sequences were amplified from plasmids containing the inserts. These Nbs were generated in our laboratory and are well characterized [40,41]. The anchor domain of ProtA were amplified from total genomic DNA of *S. aureus* NTC8325 [45] using primer pair coc3 and coc5. The Nb coding sequences were amplified using primers coc2 and coc4 from miniprep DNA. The primers coc4 and coc3 were constructed with the respective 3' and 5' ends extension complementary to a segment of the Nb and ProtA sequences, and coc2 had 5' extension of Usp45 signal sequence. Nb and ProtA PCR fragments were joined in an overlap PCR reaction using primers pair coc1 and coc5. Primers coc1 and coc5 had, respectively, a 5' and 3' extension of attB sequence of the gateway system. The primer sequences and the PCR conditions are provided in Appendix A. The cloning scheme is shown in Figure 1.

The attB flanked Nb-ProtA fusion genes were cloned into the Gateway vector pDONR221 (Thermo Fisher Scientific, Merelbeke, Belgium) in a BP reaction according to the Gateway Technology manual (Thermo Fisher Scientific, Merelbeke, Belgium, Version E, Catalog nos.12535-019 and 12535-027). Briefly, the following reaction components were set up in 1.5 mL microcentrifuge tubes for each sample: 1.0 µL AttB-PCR product (66 ng/µL), 1.0 µL pDONR™ vector (150 ng/µL), and 8.0 µL TE Buffer pH 8.0. The product of BP reaction (5 µL) was used to transform CaCl<sub>2</sub>-competent DH5α cells [46] and transformed colonies selected on LB agar plates with kanamycin. Presumptive pENTR clones were screened by colony PCR using SeqLA1 (5'CTCTGCCGTTAACGCTAGCATGGAT3') and SeqLB (5'GTAACATCAGAGATTTTGAGACAC3') primers and positive clones confirmed by sequencing (Gene Sequencing Facility, University of Antwerp, Antwerp, Belgium). Sequence analysis of the pENTR clones showed variability in the ProtA sequence length. To standardize ProtA lengths, one short and long Nb-ProtA were selected and exchanged with each of the Nb-ProtA fragments of all the pENTR clones by Gibson assembly reactions (Appendix A).



**Figure 1.** Cloning of Nanobodies (Nb)-protein A (ProtA) fusion gene for surface display in *L. lactis*. The signal peptide (SP), the Nb genes and the anchor domain of ProtA were amplified in separate PCR reactions and the gene fragments were fused in an overlap PCR reaction introducing the attB sites of the Gateway system at both ends. Created in BioRender.

## 2.2. LR Reaction between pDONR221 Containing Nb-ProtA Insert and Gateway Adapted pTRKH3 Vector

The Nb-ProtA constructs in pDONR221 vector (pENTR) were transferred using a LR reaction into the Gateway-adapted pTRKH3 expression vector pHD669 carrying the *Lactococcus clpC* promoter upstream the Gateway cassette. The detailed procedure for adapting vector pTRKH3 to the Gateway system is provided in Appendix A. The LR reaction protocol was set up according to the Gateway Technology manual (Thermo Fisher Scientific, Merelbeke, Belgium, Version E, Cat. #12535-019). Briefly, the following components of the reaction solution was added to 1.5 microcentrifuge tube: 6  $\mu$ L of entry clone (Nb-ProtA in pDONR221), 2  $\mu$ L of Gateway-adapted pHD669, 4  $\mu$ L 5X LR Clonase reaction buffer, 4  $\mu$ L TE buffer, pH 8, and 4  $\mu$ L LR Clonase enzyme. The reaction solution was mixed well and incubated at 25  $^{\circ}$ C for 1 h, after which Proteinase K was added and incubated for 10 min at 37  $^{\circ}$ C to stop the reaction. Ten  $\mu$ L of the LR reaction product were transformed into DH5 $\alpha$  CaCl<sub>2</sub>-competent cells and selected on LB media supplemented with tetracycline. Miniprep DNA prepared from transformants were submitted for sequencing (Gene Sequencing Facility, University of Antwerp, Antwerp, Belgium) using primer pair pTRKH8 (5'CTAATAAAGCCGTAAGGAGACGGGTTC3') and pTRKH9 (5'CCCATCCTAACGGCCACG CATATG3'). Miniprep DNA prepared from expression clones in DH5 $\alpha$  were used to transform electro-competent *L. lactis* MG1363 [44]. Electroporation was conducted at 2.5 kV, 200  $\Omega$ , and 25  $\mu$ F. Positive clones were selected on M17 media supplemented with 0.5% glucose and 10  $\mu$ g/ mL erythromycin and confirmed by colony PCR.

## 2.3. Validation of Nb Surface Display on *L. lactis*

### 2.3.1. Slide Agglutination Assay

Recombinant *L. lactis* strains were grown on M17 broth (Oxoid<sup>®</sup>, Thermo Fisher Scientific, Merelbeke, Belgium) supplemented with glucose (0.5%) and erythromycin (5  $\mu$ g/) mL for 48 h at 30  $^{\circ}$ C. Cells were pelleted at 2000 rpm, washed once with 10 mL phosphate buffered saline (PBS), and re-suspended in PBS. Cell concentration was estimated by OD measurement and was adjusted to OD<sub>600</sub> = 3. High cell concentrations enabled the visualization and scoring of the agglutination effect. In the agglutination test, 15  $\mu$ L of the prepared *L. lactis* was added to an equal volume of *E. coli* cell suspension on an

agglutination plate and mixed by rocking for 1–3 min. Positive agglutination was observed visually against a light source and scored objectively on the basis of time of appearance and intensity. The scores ranged from 1–5 (5 = rapid uniform agglutination within 15 sec; 4 = rapid uniform agglutination within 30 sec; 3 = uniform agglutination after 30 sec–1 min; 2 = limited agglutination after 1 min; 1 = apparent agglutination seen after 1 min; 0 = no agglutination after 3 min).

### 2.3.2. In vitro Villous Adhesion Assay

In vitro villous adhesion assay was performed as previously described [22,47,48]. The cell concentration was adjusted to  $10^{10}$  cells/ mL for *L. lactis* and  $10^8$  cells/ mL for *E. coli* using OD estimates. The intestinal villi were prepared from the mid-jejunum of four-week-old piglets (provided by Prof. Dr. Eric Cox, Ghent University). The villi from each piglet were tested for the presence of the F4 (three variants F4ab, F4ac and F4ad) and the F18 receptors. Only villi from piglets positive for the F4 and F18 receptors were used in the adhesion assay. The intestinal sections were sliced, rinsed in PBS, and scraped into Krebs–Henseleit buffer containing 1% formaldehyde and stored at 4 °C. Before use, the thick villi suspension was washed twice with Krebs–Henseleit buffer without formaldehyde and re-suspended in PBS. For each assay, a mixture of 100 µL F4 or F18 *E. coli* with 100 µL of respective anti-F4 or anti-F18 *L. lactis* was incubated for 1 h at RT on a rotating plate (50 rpm), after which 50 µL prepared villous were added to the cell mixture and topped to 500 µL with PBS, and incubated for additional 1 h at RT. For the controls, anti-F4 and anti-F18 *L. lactis* were pre-incubated with 50 µg FaeG and FedF, respectively, for 15 min at RT on a slow rotating plate (50 rpm) before adding the *E. coli* strains and the subsequent steps were followed as above. To detect adherence, 12 µL of the prepared sample were observed under light microscopy and the number of *E. coli* cells adhering to 250 µm villous length was counted at  $\times 100$  magnification in oil immersion. Adhesion of <5, 5–30, and >30 *E. coli* cells per 250 µm brush border length was considered as negative, weak, and strongly positive, respectively. The mean difference in the number of *E. coli* cells adhered to 250 µm villous length between the test (adhesion inhibition assay) and controls (reversal of adhesion inhibition) were compared using paired t-tests.

### 2.3.3. Fluorescence Microscopy of Recombinant *L. lactis*

Surface display of Nbs on *L. lactis* was visualized by fluorescence microscopy using specific Nb antigen (FaeG or FedF) labeled with FITC (Flouorescien-5-isothiocyanate). FaeG and FedF were conjugated with FITC according to the manufacturer's manual (Molecular Probes by Invitrogen; Cat.# F1906, Lot. 513684). Briefly, purified FaeG (8.1 mg/mL) and FedF (5.7 mg/mL) subunits in 20 mM Tris buffer were exchanged into 0.1 M sodium bicarbonate buffer (pH 8.5) using Zeba™ Spin Desalting Columns, 7K MWCO (Thermo Fisher Scientific, Lot# 203162, Merelbeke, Belgium) and concentrations adjusted to 5 mg/mL FITC dye were dissolved in DMSO to a final concentration of 10 mg/ mL, and 10 µL of the solution were added slowly with continuous vortexing to 200 µL FaeG and FedF preparations. The reaction mix was incubated at RT for 1 h on a slow rotating plate. Conjugated protein was purified using illustra™ NAP™-5 columns (GE Healthcare, Diegem, Belgium) to remove free dye. The *L. lactis* cells were prepared as described for the in vitro adhesion assay and resuspended in PBS to a concentration of  $1 \times 10^8$  cells/.mL The cells were blocked with 3% BSA for 20 min, labeled antigens added to final concentration of 2 µg/,mL incubated for additional 20 min, washed  $2 \times$  and resuspended in PBS. All washing steps were performed in PBS and centrifugation at 2000 rpm for 7 min at 4 °C. Fifteen microliters of labeled cell suspension was applied to a microscope slide and viewed under inverted fluorescence microscope (Nikon Eclipse TE2000-U, Groot-Bijgaarden, Belgium), at  $100 \times$  magnification, and image captured using attached camera.

#### 2.4. In vivo Piglet Challenge Experiments

A clinical trial was conducted to evaluate the protectiveness of the recombinant anti-F4 *L. lactis* in weaned piglets challenged with an F4<sup>+</sup> *E. coli*. Only anti-F4 *L. lactis* strains were tested in the challenge experiment based on the known challenge model for the F4-fimbriated ETEC strains [37].

##### 2.4.1. Animal Preparation

The research protocol was approved by the Uganda National Council for Science and Technology (UNCST) on 16 April 2013; Approval #HS 1361. Twenty piglets from 3 sows were weaned at 4 weeks and transported to the animal facility at the College of Veterinary Medicine, Makerere University. The piglets were drawn from 3 different litters and comprised of both males and females. The piglets were tagged and divided into 3 groups of 5, 5 and 10 animals (Table 1). The different groups were housed in disinfected, separate pens within the same housing unit. The different pens were separated by partial solid walls (1.25 m height) that precluded physical contact between piglet groups. Group 3 was split into 2 pens of 5 piglets each. To clear possible gut ETEC *E. coli* infection present at weaning, the piglets were treated with Enrofloxacin oral drench (5 mg/kg) for 3 consecutive days beginning 2 days prior to weaning. This was necessary because there were no known F4 or F18 *E. coli* negative farms. Enrofloxacin was chosen because all pathogenic *E. coli* strains isolated from pig farms in this region had tested susceptible to fluoroquinolones [49]. Piglets from the same litter were distributed equally in the different groups to account for cohort effect.

**Table 1.** Piglet groups and treatment.

Group	Number	Treatment	<i>L. lactis</i> Dose
1 (Negative control)	5	No F4 ETEC given	No <i>L. lactis</i> added
2 (Positive control)	5	F4 ETEC + MG1363 <i>L. lactis</i>	10 <sup>10</sup> cells/100g of feed
3 (Test)	10	F4 ETEC + Anti-F4 <i>L. lactis</i>	10 <sup>10</sup> cells/100g of feed

##### 2.4.2. Feeding and Challenge

All piglets were fed commercial weaner's meal supplied locally at a rate of 400 g/day divided in 2 daily rations (fed in the morning and evening), as per the supplier's recommendation. Weaners' meal comprised of maize bran base, protein and mineral supplements. For the treated group, the anti-F4 *L. lactis* suspension in 10 mL of PBS (10<sup>10</sup> cells/100 g of feed) was added and mixed in the feed with additional water to a uniform soft consistency. In the control groups, only PBS was added to the feed and additional water added to soft consistency as above. *L. lactis* strains were added to feeds starting two days before the challenge with the F4-positive strain and continued till day 18 after challenge.

Starting on day 0 (weaning day), the piglets were inoculated daily with a mix of *L. lactis* strains in feed at a dose of 10<sup>10</sup> cells/100 g of feed. Cell concentration was estimated from OD measurements. On day 2 and 3, the piglets (test and positive control groups) were challenged with 10<sup>9</sup> F4<sup>+</sup> *E. coli* strain (C95-72) by oral drenching. The *E. coli* cells were harvested from an overnight culture incubated at 37 °C. *L. lactis* strains were grown in M17 medium supplemented with 0.5% glucose (final concentration) at 30 °C for 48 h.

##### 2.4.3. Monitoring Response

The piglets were weighed daily and monitored daily for any signs of diarrhea or edema disease. Fecal samples were also collected daily from each piglet for total bacterial counts. F4<sup>+</sup> *E. coli* isolates in fecal samples were confirmed by slide agglutination test using anti-F4 *L. lactis* on 10 selected clones per piglet. Blood samples were taken from the jugular vein on days 0, 7, and 14 post-challenge to evaluate immune response by. On day 15, the piglets were sacrificed and jejunal villi isolated to confirm F4 receptor status

by in vitro adhesion assay, as previously described [22]. The Enzyme-Linked Immuno Sorbent Assay (ELISA) protocol used was previously described by the authors [49]. Briefly, we ran indirect ELISA serial dilutions of a known positive and negative serum samples, and all the test samples, to establish the optimum dilution that would allow an optimal comparison. Subsequently, we chose a single point dilution (1:10) in serum dilution buffer (PBS + 0.05% (v/v) Tween20 + 3% w/v BSA) and ran all the samples again in duplicate plates concurrently under identical conditions to minimize variations. The serum samples were pre-treated with kaolin before dilution. The plates (Nunc MaxiSorp, Thermo Fisher Scientific, Merelbeke, Belgium) were coated with 1.5 µg/ mL of purified recombinant FaeGac in coating buffer (100 mM bicarbonate/carbonate buffer; pH 9.6) for 3 h at 37 °C and blocked overnight at 4 °C with PBS + 0.2% (v/v) Tween80. Serum samples were applied, and plates incubated at 37 °C for one hour. Rabbit anti-pig IgG HRP conjugated secondary antibody (1:10,000 dilution) was then added and incubated at 37 °C for 1 h, followed by the substrate (3,3',5,5'-tetramethylbenzidine (TMB, Invitrogen) for 20 min at RT, and the reaction was stopped with an equal volume of 1 N HCl. Optical density (OD) was read at 450 nm. The plates were washed twice with PBS between each incubation step.

#### 2.4.4. Statistical Analysis

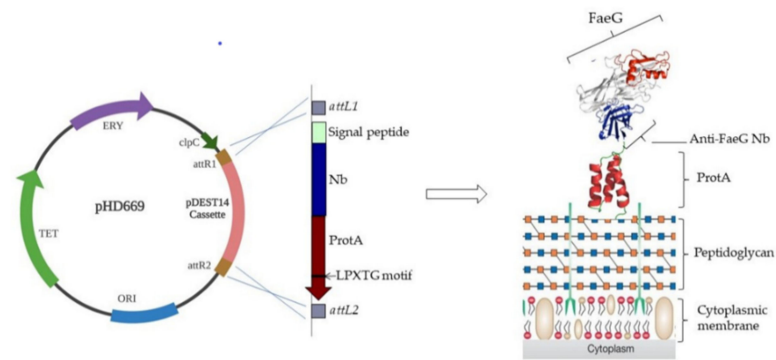
Experimental data were recorded in Microsoft Excel and analyzed using Stata IC 15 (Stata Corp LLC, College Station, TX, USA) and GraphPad Prism 9.0 statistical software. Repeated measures of body weight and fecal bacterial shedding were summarized using descriptive statistic means for continuous variables. Uncertainty measures including standard errors and 95% Confidence Intervals (CI) were also reported. The analysis of variance (ANOVA) test was used to compare the differences in weight gain, fecal bacterial shedding, and immune response between the different treatment groups. Mixed effects logistic regression models were used to predict changes in weight gain, immune response, and fecal bacterial shedding; treatment, experiment day, and interaction between treatments and experiment day were the fixed factors, and individual animals were the random factors. Checks on data normality distribution showed that the measure on fecal bacteria shedding was skewed to the right and was hence transformed to decadic logarithm for the final analysis.

### 3. Results

#### 3.1. Cloning Nb for Surface-Display on *L. lactis*

The current study was initiated by the design of *L. lactis* recombinant strains that covalently present anti-FedF or anti-FaeG (respective adhesive subunits of F18 or F4 fimbriae) Nanobodies on their cell surface. The Nanobodies were cloned in pTRKH3 shuttle vector adapted to the Gateway system (pHD669) for expression in *L. lactis*. The vector pHD669 contain the clpC promoter from *L. fermentum* BR11. This promoter displayed high activity in all *Lactobacillus* strains tested when present as a transcriptional fusion with the *gusA* gene [50]. The fusion gene constructs consisting of a G-positive signal peptide, the Nb, and the protein A anchor domain derived from *S. aureus* sequences were expressed from this promoter (Figure 2).

We used both the long (163 amino acid residues) and short (63 amino acid residues) ProtA anchor domain to construct different recombinant *L. lactis* strains (Table 2) and compared the effect of the anchor length on biological activity. Complete protein sequence alignment of the fusion gene constructs for anti-FaeG and anti-FedF Nanobodies are shown in Supplementary Figure S1 and Figure S2, respectively.



**Figure 2.** Cloning scheme for Nb surface display on *L. lactis* cell wall using ProtA anchor domain. Left: The pTRKH3 shuttle vector was adapted to the Gateway cloning system by inserting the pDEST14 cassette flanked by the attR1 and attR2 recombination sites, downstream of the cplC promoter sequence, in the ApaLI restriction site to form plasmid pHD669. The fusion gene comprised the UspA signal peptide sequence for Gram-positive bacteria (green), Nb (blue) sequence, and protein A anchor domain sequence (red). The C-terminus of protein A contained the LPXTG motif (X is any amino acid) that is recognized by sortase, a transaminopeptidase enzyme, that cleaves the protein A between the threonine and glycine residues of the motif and links it covalently to the pentaglycine peptide of the cell wall peptidoglycan. The pentaglycine peptide is an interbridge between the tetrapeptides that cross-link the N-acetylglucosamine (NAG) and N-acetylmuramic acid (NAM) polysaccharide chains to form the lattice-like structure of the cell wall. On the right is a schematic presentation of the anti-FaeG Nb (blue) in the complex with the FaeG subunit from F4 fimbriae (in grey is the immunoglobulin like core and in red is the receptor binding domain) [41] anchored on the cell wall [51] by protein A (red) [52].

**Table 2.** Different recombinant *L. lactis* strains expressing Nanobodies against FaeG and FedF anchored on the cell surface via a short or long protein A protein linker.

Anti-FaeG <i>L. lactis</i> Strains		Anti-FedF <i>L. lactis</i> Strains	
Long Anchor ProtA	Short Anchor ProtA	Long Anchor ProtA	Short Anchor ProtA
pEXP366 (V2)	pEXP444 (V2)	pEXP362-1 (NbFedF6)	pEXP447 (NbFedF9)
pEXP368 (V4)	pEXP445 (V3)	pEXP364-3 (NbFedF7)	pEXP451 (NbFedF12)
pEXP441 (V1)	pEXP455 (V1)	pEXP459 (NbFedF9)	pEXP449 (NbFedF11)
pEXP457 (V4)		pEXP463 (NbFedF12)	pEXP365 (NbFedF7)
		pEXP461 (NbFedF11)	pEXP363 (NbFedF6)

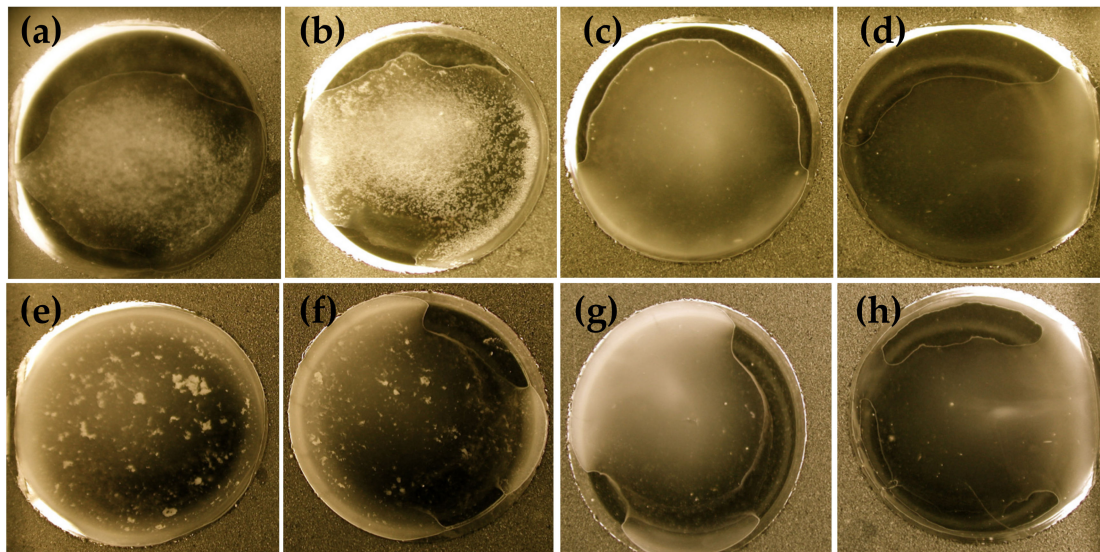
### 3.2. In vitro Assay to Validate Nanobodies Surface Display on *L. lactis*

We demonstrated the expression and functional surface display of the Nbs on *L. lactis* using slide agglutination test, immunofluorescence, and flow cytometric analysis.

#### 3.2.1. Agglutination Assay

To cause agglutination, the surface Nbs on *L. lactis* needs to specifically recognize FedF or FaeG subunits of F4 or F18 fimbriae present on *E. coli*. In our agglutination assay, overnight cultures of *L. lactis* and *E. coli* strains were harvested, washed, and re-suspended in PBS at OD<sub>600</sub> 3. Equal volumes of *L. lactis* and the *E. coli* suspension were added into wells of agglutination plates, mixed by gentle rocking, and observed agglutination against light source. The agglutination was scored on the basis of the time of appearance and the extent of granulation. Cross-linking of several *E. coli* and *Lactococcus* cells created a mesh forming macroscopic granules. The result of the agglutination assay is shown in Figure 3. The rest of the *L. lactis* constructs was tested for agglutination with different *E. coli* strains, and the results are summarized in Supplementary Tables S1 and S2.





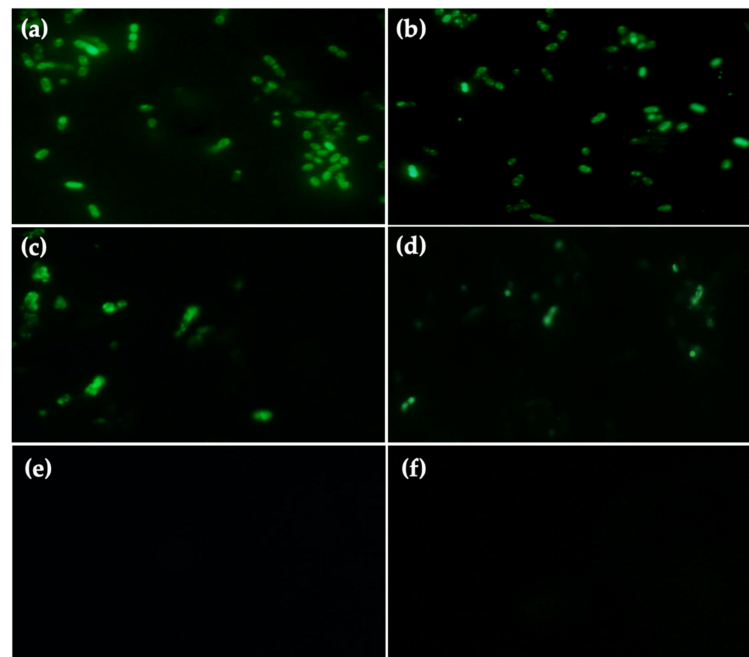
**Figure 3.** Agglutination of F4<sup>+</sup> and F18<sup>+</sup> *E. coli* by recombinant *L. lactis* displaying surface anti-F4<sup>+</sup> and anti-F18<sup>+</sup> Nb, respectively, anchored on the cell wall by protein A. Top panel: (a) positive agglutination of F4<sup>+</sup> *E. coli* strain C1065-79 and (b) F4<sup>+</sup> *E. coli* strains C544-79 by anti-F4 *L. lactis* (pEXP368;  $\alpha$ F4 V4 on long protein A anchor). In the control tests, (c) no agglutination was seen between *E. coli* C1065-79 and control *L. lactis* (pEXP362;  $\alpha$ F18 NbFedF6 on long anchor ProtA) or (d) C1065-79 in PBS. Bottom panel: agglutination of F18<sup>+</sup> *E. coli* strain by anti-F18 Lactococcus strains, (e) F18<sup>+</sup> *E. coli* K514  $\Delta$ fim (pIH120) that contain the gene cluster for F18 fimbriae biogenesis [8] and *L. lactis* (pEXP362), (f) F18<sup>+</sup> *E. coli* K514  $\Delta$ fim (pIH120) and *L. lactis* (pEXP364 (NbFedF7 on long protein anchor). In the control tests, no agglutination is seen between (g) K514  $\Delta$ fim (pIH120) and non-F18 *L. lactis* (pEXP368,  $\alpha$ F4 V4) or (h) K514  $\Delta$ fim (pIH120) in PBS.

### 3.2.2. Effect of Heat Treatment on Viability and Agglutination Capacity

To test the possibility of using inactivated *Lactococcus* strains in piglets, the strains were heated at different temperatures and times in a water bath and used in the agglutination test. Successful inactivation was evaluated by colony count (CFU/) mL of heat-treated samples plated on M17 (Oxoid) agar supplemented with glucose and erythromycin. Complete inactivation with a full agglutinating capacity was achieved at 60 °C for 20 min. Inactivation at higher temperatures abolished agglutination. The same strains were also treated with 5 mM dithiothreitol (DTT) for 20 min to reduce the internal disulfide bond present in the Nbs anchored to the surface of the recombinant *L. lactis* strains. This treatment abolished agglutination, but the strains remained viable. The results for heat and DTT treatments are summarized in Supplementary Table S3. Non-specific binding was blocked by 3% bovine serum albumin in PBS.

### 3.2.3. Immunofluorescence Microscopy

To demonstrate the surface display of the anti-FaeG and anti-FedF Nbs by recombinant *L. lactis* using FITC fluorescence dye, FITC conjugated FaeG and FedF were mixed with the respective anti-FaeG and anti-FedF Nbs expressing *L. lactis* strains and observed under fluorescence microscope (Figure 4). The anti-FaeG and anti-FedF *L. lactis* strains specifically recognized the respective antigen conjugates, FaeG-FITC and FedF-FITC, as detected by the positive fluorescence signals.



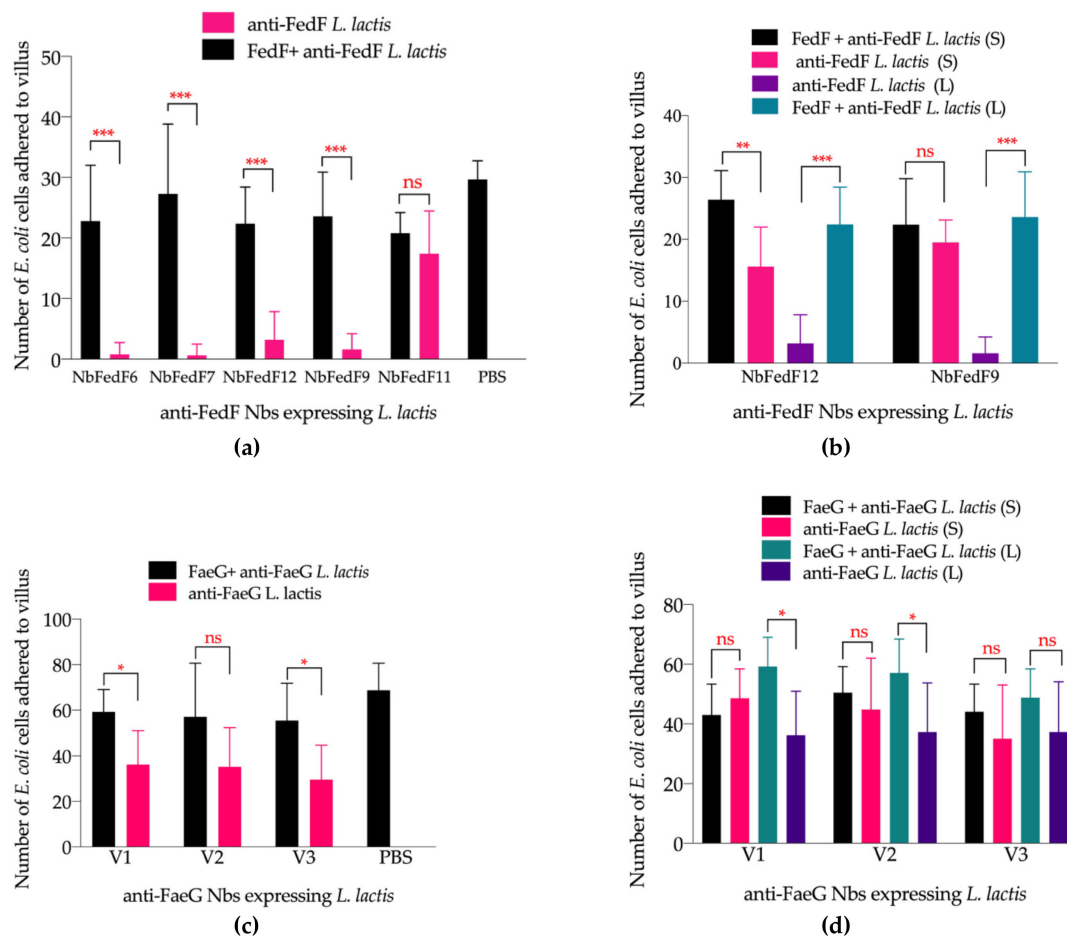
**Figure 4.** Fluorescence of recombinant *L. lactis* labeled with (Flourescien-5-isothiocyanate) FITC-conjugated antigens. The anti-FaeG and anti-FedF *L. lactis* strains specifically recognized FITC labeled FaeG and FedF antigens, respectively, as shown by fluorescence microscopy. Slides: (a) anti-FaeG Nb *L. lactis* (pEXP368) labeled with FaeG-FITC; (b) anti-FaeG *L. lactis* (pEXP366) labeled with FaeG-FITC; (c) anti-FedF *L. lactis* (pEXP364) labeled with FedF-FITC; (d) anti-FedF *L. lactis* (pEXP362) labeled with FedF-FITC; (e) non FaeG/FedF *L. lactis* (pEXP242) labeled with FaeG-FITC; (f) non FaeG/FedF *L. lactis* (pEXP242) labeled with FedF-FITC. Image captured at X100 magnification under oil immersion.

### 3.2.4. In vitro Adhesion Assay

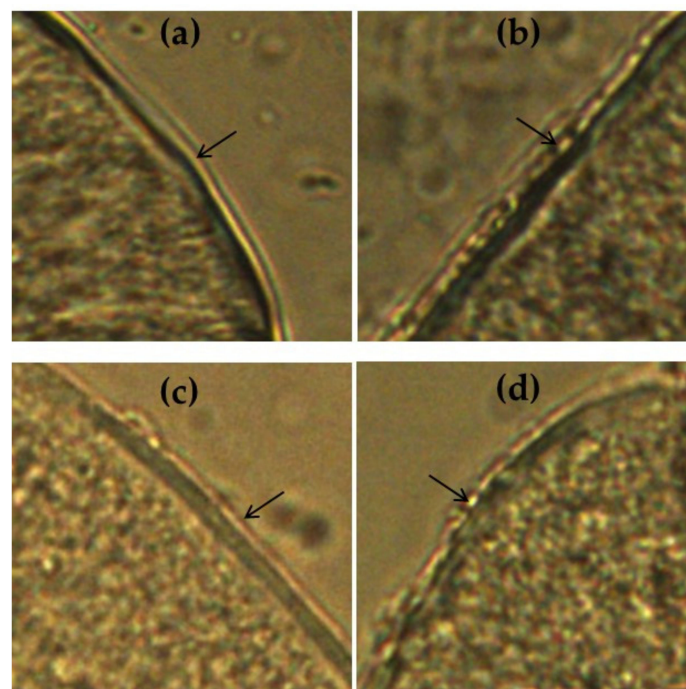
The anti-F4 and anti-F18 Nbs expressing *L. lactis* strains were tested for their inhibitory effect on the respective F4 or F18 *E. coli* adhesion to piglet intestinal villi in an in vitro adhesion test [22,48].

#### *Lactococcus lactis* Expressing anti-FedF Nanobodies

The anti-FedF *L. lactis* strains (Table 2) inhibited F18<sup>+</sup> *E. coli* (strain 107/86) adhesion to the F18R<sup>+</sup> piglet villi, as measured by the number of *E. coli* cells adhering to the piglet villi. Adhesion inhibition was reversed by pre-incubating anti-FedF Nb expressing *L. lactis* with FedF, the tip adhesin of F18 fimbriae (Figures 5a and 6a). Paired t-test showed significant difference in the mean number of F18<sup>+</sup> *E. coli* adhered to the villi, following the adhesion inhibition by anti-FedF *L. lactis* strains and the reversal of the inhibitory effect by pre-incubating the *L. lactis* strains with FedF adhesin ( $p < 0.001$ ). The optimal anti-FedF Nb expressing *L. lactis* concentration for maximal inhibition was  $10^{10}$  cells/.mL Inhibitory effect of *L. lactis* strains expressing short (83 residues) or long (163 residues) protein A anchor domains was tested in the in vitro assay (Figures 5b and 6b). All the *L. lactis* strains with long protA anchor had a strong inhibitory effect on binding of F18+ *E. coli* that was reversed by pre-incubation of the villi with FedF adhesin ( $p < 0.001$ ), while *L. lactis* with Short ProtA anchor showed only partial inhibition and adhesion inhibition reversal effects by pre-incubation of the anti-FedF *L. lactis* strains with purified FedF (Figure 5a) that was significant for NbFedF12 *L. lactis* strain ( $p = 0.003$ ) and non-significant for NdFedF9 *L. lactis* strain.



**Figure 5.** Inhibition of adherence of the F18<sup>+</sup> and F4<sup>+</sup> *E. coli* strains to prepared piglet microvilli by *L. lactis* expressing anti-FaeG and anti-FedF Nb. **(a)** Inhibition of adherence of the F18<sup>+</sup> *E. coli* 107/86 to microvilli in vitro by *L. lactis* expressing anti-FedF Nbs on the cell surface. All the *L. lactis* strains showed strong inhibition except for the positive control Nb242 (non-neutralizing anti-FedF Nb) and *E. coli* strain 107/86 alone in PBS. Adherence inhibition was reversed by pre-incubating *Lactococcus* strains with purified FedF protein. **(b)** Effect of protein A anchor domain length on the inhibition of F18<sup>+</sup> *E. coli* 107/86 adherence to microvilli by anti-FedF Nb expressing *L. lactis*. A short ProtA anchor domain (S) resulted in partial inhibition (magenta), whereas a long ProtA (L) caused nearly complete inhibition (purple). This positive adherence inhibition was reversed by pre-incubation of the anti-FedF *L. lactis* strains with FedF (black and blue bars). **(c)** Partial inhibition of F4<sup>+</sup> *E. coli* (C95-72; K88ac) adherence to prepared piglet microvilli by *L. lactis* expressing surface anti-FaeG Nbs (magenta). This inhibition was reversed by pre-incubation of these *Lactococcus* strains with purified FaeG (black). All the anti-FaeG *L. lactis* strains showed partial inhibition. **(d)** Effect of protein A anchor domain on the inhibition of F4<sup>+</sup> *E. coli* C95-72 adherence to microvilli. A short linker (S) did not inhibit adherence (magenta), whereas a long linker (L) resulted in partial inhibition (purple). Pre-incubation of the anti-FaeG *Lactococcus* strains with FaeG (black and blue) allowed adherence. Adherence was measured as counts of *E. coli* cells attached to 250  $\mu$ m villus length. Level of significance:  $p > 0.05$  (ns),  $p \leq 0.05$  (\*),  $p \leq 0.002$  (\*\*), and  $p \leq 0.001$  (\*\*\*)



**Figure 6.** Inhibition of F18<sup>+</sup> and F4<sup>+</sup> *E. coli* adhesion to F18R<sup>+</sup> and F4R<sup>+</sup> piglet microvilli by anti-FedF or anti-FaeG Nbs expressing *L. lactis*. (a) Complete adhesion inhibition of F18<sup>+</sup> *E. coli* (107/86) to F18R<sup>+</sup> microvilli by *L. lactis* (pEXP362) expressing anti-FedF NbFedF6 on its surface. (b) Inhibitory effect of anti-FedF Nb expressing *L. lactis* was blocked by pre-incubating with 50 µg of FedF that neutralized surface displayed NbFedF6. (c) Partial inhibition of F4<sup>+</sup> *E. coli* (C95-72) adhesion to F4R<sup>+</sup> microvilli by *L. lactis* (pEXP368) expressing anti-FaeG Nb V4. (d) Adhesion inhibition was reversed by pre-incubating *L. lactis* with 50 µg of FaeG. The arrow points to the villi surface without bound *E. coli* (a,c) and with bound *E. coli* cells (b,d).

#### *Lactococcus lactis* Expressing anti-FaeG Nanobodies

Anti-FaeG *Lactococcus* partially inhibited the in vitro adhesion of F4<sup>+</sup> *E. coli* to the F4R<sup>+</sup> villi, and this effect was reversible when the strains were pre-incubated with FaeG, the specific antigen for the Nanobodies displayed by *L. lactis* strains. Inhibition effect was concentration dependent, with only partial inhibition at 10<sup>10</sup> cells/ mL for *L. lactis* (Figures 5c and 6d). The difference between mean number of F4<sup>+</sup> *E. coli* adhered to F4R<sup>+</sup> villi in the adhesion inhibition and reversal of adhesion inhibition assay was only significant for the V1 ( $p = 0.004$ ) and V3 ( $p = 0.004$ ) *L. lactis* strains, but it was not significant for V2 *L. lactis*. Strains expressing short (83 residues) and long (163 residues) ProtA anchor domain sequences were compared in the in vitro assay. The result showed that constructs with a short ProtA anchor domain did not inhibit adhesion, whereas long links resulted in partial agglutination that was reversible on pre-incubation of the *Lactococcus* strains with purified FaeG. Among the long ProtA anchor *L. lactis* strains, the mean difference in the number of F4<sup>+</sup> *E. coli* adhered to the F4R<sup>+</sup> villi during the inhibition adhesion and inhibition reversal assays were significant for V1 ( $p = 0.004$ ) and V2 ( $p = 0.005$ ) *L. lactis* strains but not significant for V3 ( $p = 0.106$ ) *L. lactis*. The results are shown in Figures 5d and 6d.

The in vitro adhesion assays thus demonstrated a clear effect of the length of the ProtA anchor domain for the anti-FedF Nbs directed against the tip adhesin FedF (1 adhesin per fimbria) of F18 fimbriae. In comparison, the effect of the length of the ProtA anchor domain inhibition of anti-FaeG Nbs directed against the major subunit and adhesin FaeG (1000 adhesins per fimbria) of F4 fimbriae only resulted in a small difference in the inhibition of adhesion.

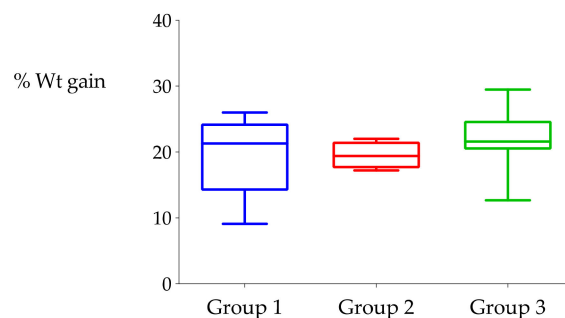
### 3.3. In vivo Piglet Challenge Assay

In a preliminary clinical trial, twenty piglets were weaned at four weeks and randomly divided into three groups for the infection challenge experiment. The F4<sup>+</sup> *E. coli* strain C95-72 (F4ac) was used in experimental infection challenge. To control F4<sup>+</sup> *E. coli* infection, the treated group received passive immunization through feeding diet supplemented with a mixture of recombinant anti-FaeG *L. lactis* (pEXP366(V2) and pEXP368(V4)); the positive control group diet was supplemented with laboratory strain *L. lactis* MG1363 [46], while the negative control group received no diet supplement (Table 1).

The piglets were monitored for clinical signs of diarrhea daily. The fecal bacterial shedding, weight gain and immune response were measured over a two-week period after which the piglets were sacrificed to assess the individual F4R receptor status by in vitro villous adhesion assay. One piglet from Group 2 and two piglets from Group 3 were found to be F4R negative and were removed from subsequent analysis.

#### 3.3.1. Weight Gain

All the piglets were weighed on days 0, 7, and 14 (before sacrificing). Average weight of piglets assigned to the three experimental groups, measured at the three time points, is shown in Figure 7 and Supplementary Table S4. Since the piglets were drawn from different litters and varied in weights on day 0, percentage weight gain during the 2-week study period was computed to compare growth rate between the different groups. The boxplot in Figure 7 depicts the quartile distribution of the average percent weight for each treatment group. The mean percent weight gain over the study period (day 0–14) was comparable between the three groups. Numerically, the mean percentage weight gain was highest in Group 3 piglets (infected and treated), as shown in Supplementary Table S5, followed by Group 1 (negative control) and least in Group 2 (positive control). Prediction of weight gain using mixed effects logistic regression showed no significant association between treatment group and weight gain (Supplementary Table S6). All interactions between treatment groups and experiment days were not significant.

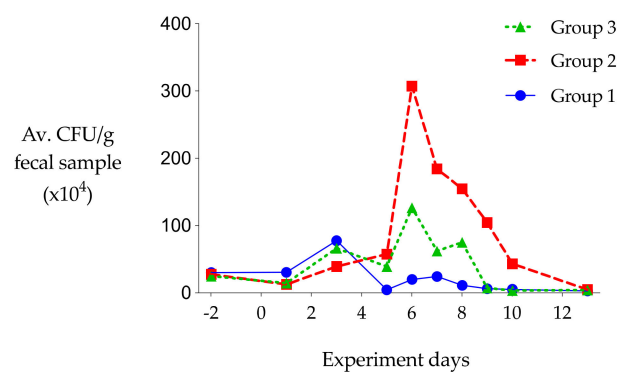


**Figure 7.** Average weight gain post weaned piglets following challenge with F4 ETEC strain. Group 1, negative group (not infected); Group 2, positive group (not treated and challenged with F4<sup>+</sup> ETEC); Group 3; test group (treated with anti-FaeG *L. lactis* added to the feed from day 0 to day 8 and challenged with F4<sup>+</sup> ETEC and). The edges of the boxes are drawn at 25th and 75th percentiles, the middle line is the median, and the lines are drawn at 1.5 IQR (inter quartile range).

#### 3.3.2. Fecal Bacterial Shedding and Diarrhea

Fecal samples were taken on day −2, 1, 3, 5, 6, 7, 8, 9, 10, and 13 post-challenge. Ten-fold serial dilutions of 10 g fecal in PBS was plated on MacConkey media for *E. coli* enumeration. The CFUs per gram of the fecal sample for each piglet was determined and the average for each group computed. The negative control group (Group 1) was used as a reference basal CFUs counts to compare the other groups. The results (Figure 8) showed the infected and treated piglets (Group 3) had lower CFU counts compared to the infected and non-treated piglets (Group 2). Both groups registered the peak CFU counts on day 6 (3 days post-challenge), followed by a progressive decline thereafter. The infected and

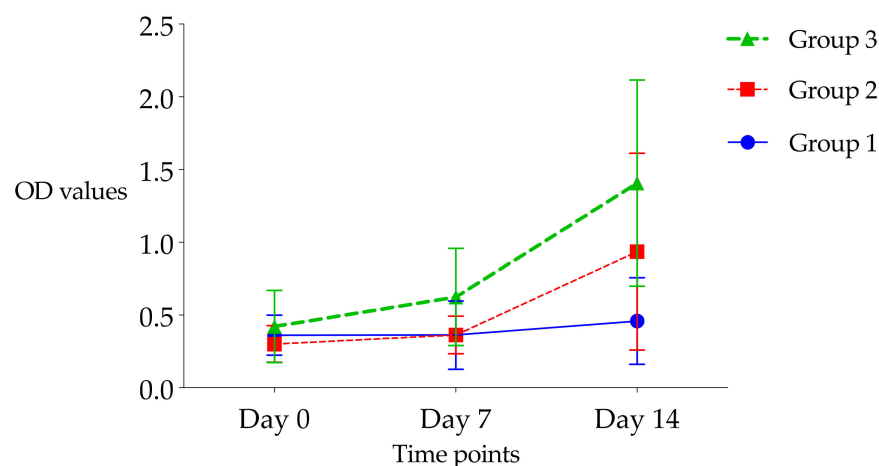
treated piglets (Group 3) however returned to basal level of bacterial shedding 5 days earlier than the infected non-treated piglets (Group 2). An ANOVA test comparing the bacterial shedding between groups concluded that the mean CFU was not significantly different for all groups ( $F = 2.92$ ,  $p < 0.071$ ). None of the piglets in Group 1 (non-infected) and Group 3 (infected and treated) developed clinical diarrhea during the study period. Among the piglets in Group 2 (infected and not treated), two piglets (out of 5) developed clinical diarrhea starting day 3 and 4 post challenge. The diarrhea lasted for 2–3 days and was self-limiting. Mixed effects logistic regression analysis of Log<sub>10</sub> CFU/g fecal bacteria shedding by experimental piglets, with treatment and experiment days as fixed factors and animal ID as random factor, estimated 0.24 and 0.28 logs decrease in fecal bacterial shedding, by groups 2 and 3 respectively, although these decreases were not significant. Analysis of the interaction between treatment groups and experiment day estimated a >1 log increase in fecal bacterial shedding in group 2 on days 5–10 compared to reference group 1, and this increase was significant ( $p < 0.05$ ). In group 3, significant increase in fecal bacterial shedding ( $p < 0.05$ ) was observed on days 3–6, but a >1 log increase was estimated on day 5 only (Supplementary Table S7).



**Figure 8.** Bacterial load in fecal samples measured as CFU. The piglets were challenged on days 2 and 3. Peak fecal bacterial load was recorded on day 6 in groups 2 and 3, after which there was a progressive decline. Group 2 had lower CFU counts than group 3 and resolved quickly to the basal level (Group 1) on day 9. Group 1, negative control group (not infected); Group 2, positive group (infected and not treated); Group 3; test group (infected with F4<sup>+</sup> ETEC and anti-FaeG Nanobodies on the surface of *L. lactis* added to the meals from day 0 to day 8).

### 3.3.3. Immune Response

Blood samples were taken on days 0, 7, and 14, and a change in IgG levels of these blood samples were determined using an anti-FaeG ELISA (Figure 9). Immune response was measured as OD (optical density) in the ELISA method. The calculated inter- and intra-assay coefficient of variation (CV) were 5.7% and 6.2% respectively. Increased anti-FaeG IgG level was observed in the infected and treated piglets (Group 3) compared to the challenged and non-treated piglets (Group 2), while the negative control group had negligible change in the IgG levels (Figure 9). Prediction of immune response using mixed effects logistic regression estimates showed no significant effects of treatment group and experiment days on immune response, although the analysis indicated a lower and increased immune response in groups 2 and 3, respectively. A similar pattern was observed for experiment days 7 and 14. However, analysis of the interaction between treatment group and experiment day estimated a significant increased immune response for group 3 on day 14 ( $p < 0.001$ ) (Supplementary Table S8).



**Figure 9.** Mean IgG level in sera samples taken on days 0, 7, and 14. IgG level was measured using an anti-FaeG ELISA and depicted as optical density (OD) at 450 nm. The error bars indicate standard deviation. Group 1, negative group (not infected); Group 2, positive group (infected and not treated); Group 3; test group (infected with F4 ETEC and anti-F4 *L. lactis* added to the meals from day 0 to day 8).

#### 4. Discussion

Despite several research efforts to develop vaccines against post-weaning diarrhea in piglets, there is still no effective vaccine available on the market. Previous researches focusing on oral immunization with live or subunit vaccines failed to give protective immunity, due to the limited time available for the piglets to mount an immune response after weaning, given the fact that lactogenic immunity neutralizes vaccines during the suckling period [15]. Passive immunization is currently considered the most viable option in preventing post-weaning diarrhea in piglets [37]. The major challenge is however to design a cost-effective method for mass production for commercial application. *L. lactis* is a normal resident of the intestinal flora and is a suitable system due to the GRAS (generally regarded as safe) status [27,29,53]. In this study, we grafted the llama Heavy chain antibody binding domains (Nanobodies or Nbs) on protein A anchor domain derived from *S. aureus* for surface display on *L. lactis* through covalent linkage to the cell wall. These *L. lactis* producing the fused Nb on ProtA anchor domain were subsequently used for passive immunization.

Successful Nanobody display in active form was evaluated in vitro by slide agglutination, immunofluorescence, flow cytometry, and villous adhesion inhibition assay. All tests confirmed correct display in biologically active form. The recombinant *L. lactis* strains specifically recognized F4 or F18 *E. coli* strains in a slide agglutination assay and abrogated bacterial adhesion to F4 or F18 receptor positive villous preparations in vitro. Whereas adhesion inhibition of F18 strains was complete, the F4 strains were only partially inhibited possibly due to the fact that the targeted FaeG adhesin subunit is also the major fimbrial subunit that exists in multiple copies [54]. FedF tip adhesin of F18 strains on the other hand is a single copy fimbrial tip adhesin that can be easily neutralized. We compared the inhibitory effect of the Nbs with varied ProtA anchor domain length and could demonstrate that a longer linker peptide improved the inhibition of villous adhesion for both anti-F4 and anti-F18 producing *L. lactis* strains. This could be attributed to the improved degree of mobility and increased display exposure outside the bacterial surface that enables a wider sweep area for the attached Nanobodies to locate and bind their cognate receptors on *E. coli* cells.

In a challenge experiment, only the anti-FaeG *L. lactis* was tested on the basis of the known experimental results when expressed in the plant system [36]. Anti-FedF *L. lactis* will be tested in future challenge experiments. A minimum dosage of *Lactococcus* ( $10^{10}$  cells/100 g of feed for of the three anti-FaeG Nanobodies) was used on the basis of inhibitory concentrations in the in vitro assay. We were able to culture a low concentration of the

anti-FaeG *L. lactis* in the fecal samples of the piglets from the treated group, indicating these strains could remain viable through the gastrointestinal tract. Since the *L. lactis* strains retained activity following heat inactivation, future experiments should test the protective effect of heat inactivated strains in vivo to address any concerns regarding the use of live organisms. Such experiments should evaluate several dosage regimes.

Results of the challenge experiment showed that passive immunization by in-fed addition of the recombinant *Lactococcus* strains protected the piglets against F4 ETEC colonization, as shown by reduced fecal bacterial shedding following challenge in the treated group compared to the non-treated. The slightly elevated bacterial counts in the treated group were also quickly reduced to the basal level with 6 days after challenge. In addition, none of the challenged and treated piglets developed diarrhea, while two in five piglets in the challenged and non-treated group had diarrhea, indicating the potential of anti-FaeG *L. lactis* to protect piglets against post-weaning diarrhea. This confirms the results previously reported using the same Nanobodies expressed in *Arabidopsis thaliana* [37]. However, use of a microbial system is a much more viable option, due to the ease and cost effectiveness of scaling up production for commercial application. There was a marginal improved weight gain in the treated group compared to the positive control, indicating that the reduced bacterial load impacted positively on the growth rate. Since the piglets varied in the weaning weight, percentage weight gain was a preferred method for comparison. Future experiments should use piglets of similar weight for a more accurate growth rate comparison. The result of the ELISA showed that the low anti-FaeG IgG level (OD values) among the non-infected piglets (Group 1) remained low throughout the follow up period; this was an expected outcome in the absence of F4-positive ETEC challenge. Among the other groups, increased OD values were observed in the infected and treated piglets (Group 3) compared to the challenged and non-treated piglets (Group 2). Group 2 piglets were untreated but challenged, resulting in massive shedding of the F4-positive ETEC at days 6, 7, and 8, while Group 3 (treated and challenged) only shed very low levels of the F4-positive ETEC strain. We hypothesize that the F4 immunity in the piglets was primed during the suckling period, as shown by the high OD values in all the three groups on day 0. The low amount of challenge F4+ ETEC boosted the immune response in both Group 2 and 3 piglets. However, the immune response in Group 2 piglets was initially suppressed while shedding large numbers of ETEC bacteria but increased later when ETEC shedding reduced. Group 3 had a significantly higher OD value on day 14 compared to the other groups, which could be explained by the ETEC immune boost that was not impeded since the *L. lactis* added in diet was protective against colonization and limited ETEC shedding. This hypothesis is supported by the observed low OD values in the F4R negative piglets in Groups 2 and 3, that remain at the levels observed for the non-infected group. It is already known for a long time that the immune response to oral administered F4 fimbriae is F4R receptor-dependent [55]. Hence, F4R-negative piglets will not develop an immune response. On the other hand, an increased immunity is a desired outcome that ensures protection following withdrawal of the probiotic supplement. Additionally, the improved immune response in the treated group could also be possibly explained by non-specific immunomodulatory effect of lactic acid bacteria [56]. At weaning, the microbial population structure was shown to change from predominantly *Lactobacilli* species at neonatal stage to emergence of *Clostridia* and *E. coli* during the early weaning stage [57]. Addition of *L. lactis* at weaning possibly stabilized the gut microbiota and improved immune response [58].

At the end of the challenge experiment, three piglets from two treatment groups were shown to be F4R negative and excluded from the analysis. Prescreening for the F4R receptor is not possible, as only postmortem in vitro adhesion assay can ultimately determine the F4R status of piglets. Genotypic detection of *mucin4* genotype polymorphism have been considered a possible prescreening method for F4R status but is not an accurate predictor for the susceptibility phenotype [59].



## 5. Conclusions

Passive immunization of piglets with recombinant *Lactococcus* strains could be a cost-effective alternative to control post-weaning diarrhea and edema disease. Therefore, we constructed recombinant *L. lactis* displaying anti-FaeG (F4 fimbriae) Nbs or anti-FedF (F18 fimbriae) Nbs on the bacterial surface. The presence of the Nbs on the surface of recombinant *L. lactis* was shown by agglutination and immunofluorescence microscopy. These recombinant *L. lactis* also inhibit the in vitro adhesion of the corresponding ETEC bacteria to microvilli. The recombinant *L. lactis* expressing the anti-FaeG (F4 fimbriae) Nbs was also used to evaluate the in vivo protection of weaned piglets against F4 ETEC infections. We could demonstrate the potential use of recombinant *Lactococcus* strains for passive immunization against F4 ETEC infections causing piglet post-weaning diarrhea. If the anti-FedF Nbs expressing *L. lactis* strains are also protective in vivo, combining both strains in feed would offer complete protection against ETEC and STEC infections during the weaning period.

**Supplementary Materials:** The following are available online at <https://www.mdpi.com/2077-0472/11/3/186/s1>, Figure S1: Sequence alignment of the fusion protein (Signal peptide-FaeG Nanobody-ProtA) expressed in *Lactococcus lactis*, Figure S2: Sequence alignment of the fusion protein (Signal peptide-FedF Nanobody-ProtA) expressed in *Lactococcus lactis*, Table S1: F18 *L. lactis* agglutination assay, Table S2: F4 *L. lactis* agglutination assay, Table S3: Effect of heat treatment on viability (CFUs/mL) and agglutination properties of *L. lactis* strains, Table S4: Average weight of piglets assigned to the 3 experimental groups, Table S5: Meant percent weight gain between day 0–14, Table S6: Mixed effects logistic regression estimates of weight gain in experimental piglets with treatment and experiment days as fixed factors, and animal ID as random factor, Table S7: Mixed effects logistic regression estimates of Log<sub>10</sub> CFU/g fecal bacteria shedding by experimental piglets, with treatment and experiment days as fixed factors and animal ID as random factor, Table S8: Mixed effects logistic regression estimates of immune response in experimental piglets with treatment and experiment days as fixed factors, and animal ID as random factor.

**Author Contributions:** H.D.G. conceived the study. E.O. made the recombinant *L. lactis* strains. E.O. and K.M. carried out the in vitro characterization of the recombinant *L. lactis* strains and analyzed the results. E.O. and J.E. did the in vivo pig studies and the analysis of the challenge data. E.O. and H.D.G. interpreted all the results and wrote the manuscript. All authors have read and agreed to the published version of the manuscript.

**Funding:** This research was funded by VLIR-UOS (Vlaamse Interuniversitaire Raad) under the Interuniversity Development Cooperation program (Belgium).

**Institutional Review Board Statement:** “The study was conducted according to the guidelines of the Declaration of Helsinki and approved by the Institutional Review Board (or Ethics Committee) of the Uganda National Council for Science and Technology (UNCST) on 16 April 2013; Approval #HS 1361.”

**Informed Consent Statement:** Not applicable.

**Data Availability Statement:** Not applicable.

**Acknowledgments:** The authors thank F. Deboeck for excellent technical assistance in constructing the Gateway shuttle vectors. We also thank Eric Cox (Ghent University) for the intestinal villi from the mid-jejunum of four-week-old piglets, positive for the F4 or F18 receptor. Finally, we thank the School of Veterinary Medicine, Makerere University, for providing animal housing facilities and care for the piglets.

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

## Appendix A Detailed Experiment Procedures

### Appendix A.1 Primers Sequences

Primer Name	Nucleotide Sequence (5'-3')
<b>coc1</b>	GGGGACAAGTTTGTACAAAAAAGCAGGCTGAATTCAGAAAGGAGA TATACGCATGAAAAAAAAGATTATCTCAGCTATTTTAATG
<b>coc2</b>	AGATTATCTCAGCTATTTAATGTCTACAGTGATACTTTCTGCTGCAGC CCCGTTGTCAGGTGTTACGCTCAGGTGCAGCTGCAGGAGTCTG
<b>coc3</b>	CTATTGGGGCCAGGGGACCCAGGTCACCGTCTCCTCAGGAC CAAAAGAGGAAGACAATAACAAGCCT
<b>coc4</b>	AGGCTTGTTATTGTCTTCTCTTTTGGTCTGAGGAGACGG TGACCTGGGTCCCCTGGCCCAATAG
<b>coc5</b>	GGGGACCACTTTGTACAAGAAAGCTGGGTAAAGCTTTTATAGTTCG CGACGACGTCCAGCTAATA
<b>coc19</b>	CTTTGTACAAAAAAGCAGGCTGAATTCAGAAAGGAGATATACGCATG AAAAAAAAGATTATCTCAGCTATTTAATG
<b>coc20</b>	GACCCAGGTCACCGTCTCCTCAGGACCAAAAAGAG
<b>Aida9</b>	GCGAAATTAATACGACTCACTATA
<b>pETrev</b>	GGTTATGCTAGTTATTGCTCAGCG
<b>pTRKH8</b>	CTAATAAAGCCGTAAGGAGACGGGTTC
<b>pTRKH9</b>	CCCATCCTAACGGCCACG CATATG

### Appendix A.2 Construction of Nb-ProtA Recombinant Fusion Genes

The Nb coding sequences were amplified from plasmids containing the inserts. These Nbs were generated in our laboratory and are well characterized [40,41]. The anchor domain of ProtA were amplified from total genomic DNA of *S. aureus* NTC8325 [45] using primers pair coc3 and coc5. The Nb coding sequences were amplified using primers coc2 and coc4 from miniprep DNA. The primers coc4 and coc3 were constructed with the respective 3' and 5' ends extension complementary to a segment of the Nb and ProtA sequences, and coc2 had 5' extension of Usp45 signal sequence. Nb and ProtA PCR fragments were joined in an overlap PCR reaction using the primer pair coc1 and coc5. Primers coc1 and coc5 had, respectively, a 5' and 3' extension of attB sequence of the gateway system. The primer sequences are outlined in the table above.

All PCR reaction volumes were 50 µL PCR: 5 µL 10x ExTakara buffer, 4 µL 2.5 mM dNTPs mix, 1 µL each 20 µM primers, 0.2 µL ExTakara DNA polymerase, 10 µL DNA template, and 14.4 µL dH<sub>2</sub>O. The cycling parameters were: initial denaturation at 94 °C for 55 sec, denaturation at 94 °C for 10 sec, annealing at 55 °C for 30 sec, extension at 68 °C for 2 min, and final extension at 72 °C for 7 min.

### Appendix A.3 Adapting pTRKH3 Vector to the Gateway System

To adapt the pTRKH3 to the Gateway system, the vector was restricted by EcoRI and ApaLI restriction enzymes in two separate reactions and the ends of the linear plasmid blunted by T4 polymerase. The reaction mix comprised of 15 µL restricted plasmid DNA, 2 µL dNTPs (2.5 mM each), 2 µL H<sub>2</sub>O and 1 µL T4 DNA polymerase. The pDEST14 cassette was amplified with the primers Aida9 and pETrev (same PCR protocol as above; primer sequences are outlined in the table above), and the PCR fragment was blunted with T4 polymerase for subsequent ligation with a linearized and blunted pTRKH3 vector, using Boehringer Quick ligation kit. The blunting reaction mix comprised of 15 µL Qiaquick purified PCR fragment, 2 µL of 10x restriction buffer or 10x T4 DNA polymerase buffer, 2 µL of dNTPs (2.5 mM each), and 1 µL T4 DNA polymerase. The blunting reactions were incubated at 15 °C for 15 min and inactivated at 65 °C for 10 min. The subsequent ligation

reaction (Boehringer Rapid Ligation kit) comprised of 5.5  $\mu$ L ApaLI-linearized vector (blunted), 2.5  $\mu$ L pDEST14 cassette PCR fragment (blunted), 2.0  $\mu$ L 5X DNA ligation buffer, 10  $\mu$ L 2X T4 DNA ligation buffer and 1.0  $\mu$ L T4 DNA ligase. The reaction was incubated at 16 °C for 5 min, and the product was immediately transformed into CSH50 gyrA462 cells [60,61] and plated on LB plates supplemented with chloramphenicol. Miniprep DNA prepared (Qiaquick Miniprep DNA kit) from selected transformed colonies were analyzed for the presence and nature of the inserts by EcoRI and HindIII double digest, and agarose gel electrophoresis.

#### Appendix A.4 Gibson Assembly Exchange Reaction

Initial screening and sequencing of pENTR clones showed variability in the ProtA sequence length. To standardize ProtA length, one long and short Nb (V1)-ProtA fusion gene (pENTR-V1L and pENTR-V1S) were selected and confirmed by sequencing. The other Nbs against F4 (V2, V3, V4), and F18 (the inhibiting Nb's NbFedF6, NbFedF7, NbFedF9 or NbFedF12 and the non-inhibiting Nb NbFedF11) were exchanged with V1 of pENTR-V1L and pENTR-V1S in a Gibson assembly reaction [62] to give constructs of uniform ProtA length. Briefly, the Nb genes were amplified sequentially with coc2/coc20 and coc19/coc20 (primer sequences outlined in the table above) to introduce the Usp45 signal peptide and end homology with EcoRI/BstEII restricted pENTR-V1 constructs. The fusion genes and linearized vector were joined in a Gibson Assembly<sup>®</sup> reaction according to the manufacture's manual. The reaction products were transformed into DH5 $\alpha$  CaCl<sub>2</sub>-competent cells and correct insert confirmed by colony PCR and sequencing. Confirmed Nb-ProtA constructs were transferred into destination vector pHD669 in a LR reaction (as previously described). The LR product was then transformed into DH5 $\alpha$  CaCl<sub>2</sub>-competent cells and selected on LB media supplemented with tetracycline and confirmed by PCR and sequencing primer pair pTRKH8 and pTRKH9 (primer sequences are outlined in the table above). Miniprep DNA prepared from expression clones in DH5 $\alpha$  were used to transform electro-competent *L. lactis* MG1363 [44]. Electroporation conditions were conducted at 2.5 kV, 200  $\Omega$ , and 25  $\mu$ F. Positive clones were selected on M17 media supplemented with 0.5% glucose and 10  $\mu$ g/ mL erythromycin and confirmed by colony PCR.

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