ORIGINAL ARTICLE



Evaluation of the oral immunogenicity of M cell-targeted tetravalent EDIII antigen for development of plant-based edible vaccine against dengue infection

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Abstract

Four serotypes of the dengue virus that can cause severe disease in humans greatly increases the complexity of vaccine development. In this study, we evaluated an 'all-in-one' tetravalent dengue vaccine candidate as an alternative to the classical four viruses 'cocktail' vaccination approach. A single polypeptide chain of the tetra-EDIII comprising all four serotypes fused to the M-cell target ligand peptide (tEDIII-Co1) was constructed and expressed using a rice expression system under the control of a strong inducible promoter (rice amylase 3D: Ramy3D) and rice 3' untranslated region (UTR). The *tEDIII-Co1* was constructed using wild type or plant codon optimized gene sequence to compare the expression levels. The codon optimized gene combined with ER-localization showed higher and more stable expression than codon-optimization alone or the wild type. To test the oral immunogenicity of this construct, the transgenic rice cells containing 100 µg of tEDIII-Co1 were fed to mice and the antigen specific IgG and IgA antibodies analysed in sera or fecal extracts. The tEDIII fused to Co1 induced strong antibody responses in the absence of any adjuvants. We observed the strongest antigen-specific B and T cell responses in animals immunized with tEDIII-Co1. Taken together, this simple tetravalent EDIII-based fusion protein with the M cell ligand could has potential as an alternative dengue vaccine strategy against all four serotypes.

Keywords Oral immunogenicity · Tetravalent vaccine · Dengue · Transgenic rice · M cell ligand

Introduction

Mosquito-borne diseases caused by viruses (Dengue, Chikungunya, Yellow fever and Zika fever), parasites (malaria, filariasis and dirofilariasis) or bacteria (tularemia) create a

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A single tetravalent dengue antigen polypeptide fused to M-cell 1 target ligand (tEDIIICo1) was expressed in transgenic rice cells. ER targeting (SEKDEL) was shown to confer superior stability compared to the secreted protein. Orally administered tEDIIICo1 induced both systemic IgG and mucosal IgA antibodies while circumventing tolerance, thus suggesting that this could be a promising oral immunisation approach.

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² Department of Bioactive Material Sciences, Chonbuk National University, Jeonju, Republic of Korea big economic burden worldwide. In particular, dengue infection is estimated to affect 390 million people per year, in some cases leading to hospitalization due to severity of the diseases and various complications (Bhatt et al. 2013). Furthermore, 3.9 billion people across 128 countries are at risk of dengue infection (Brady et al. 2012). Member states in three WHO regions declared increasing infection rates (from 2.2 million in 2010 to 3.2 million in 2015) (http://www. who.int/news-room/fact-sheets/detail/dengue-and-sever e-dengue). An efficient vaccine to reduce the infection risk is urgently required while the complexity of the interactions between the host, vector and the virus makes it extremely difficult to control and manage this disease by other means.

So far, Dengvaxia® (also referred to as CYD-TDV, developed by Sanofi Pasteur) is the only approved vaccine and is licensed in 19 countries for use in 9–45 year olds. It confers reduction in severe dengue by 93% and dengue hospitalizations by 82% (Hadinegoro et al. 2015). However, its overall efficacy in preventing dengue infection is unsatisfactorily low and it is not recommended by WHO for children under the age of nine. Therefore, new efforts are needed to develop an alternative or additional vaccine. Currently, two leading vaccine candidates (TV003/TV005 and TDV developed by NIH/NIAID/Butantan Institute and Inviragen/Takeda, respectively) based on live attenuated chimeric tetravalent formulation are being evaluated in large Phase 3 trials, while other candidates, including inactivated whole virus (TDEN), subunit (V180), and DNA (D1ME100) vaccine candidates are in various stages of Phase I–II trials (Pang and Loh 2017).

Traditional vaccines are mostly based on live attenuated organisms such as measles, mumps, rubella (MMR combined vaccine); varicella (chickenpox); influenza (nasal spray); rotavirus; yellow fever; smallpox etc. Also, inactivated/killed organisms (polio, hepatitis A, rabies) and toxoid vaccines (diphtheria, tetanus) are available. However, those types of vaccines generally have complex safety and storage requirements and there are issues with cost-effectiveness (Lee et al. 2012; Minor 2015). In contrast, subunit vaccines using only a part of the disease-causing virus could overcome both these concerns. Therefore, if they can be made to effectively stimulate the immune system they would be a very safe method of immunization for everyone, regardless of their health status. Nevertheless, subunit vaccines are less successful owing to short-lasting immunity against disease and requiring repeat administrations over specified periods. To increase their effectiveness, subunit vaccines are often engineered alongside various carriers or combined with adjuvants. Here, we genetically engineered a fusion protein that links the dengue antigen with a mucosal adjuvant and evaluated its potential as an oral vaccine in mice.

An ideal dengue vaccine should be safe, cheap and easily administered to young children in low-income countries such as South-East Asia and the Western Pacific regions (Mangold and Reynolds 2013) that show increasing incidence of the disease in the younger age group (Rodriguez-Barraquer et al. 2011). Some of these advantages could be realized by generating subunit vaccines in genetically modified plants (Daniell 2006) which have been shown to have a significant potential by many research groups (Chan and Daniell 2015; Huy and Kim 2017; Kim et al. 2016, 2015). For over 20 years, plant molecular engineering has been progressing to increase the yield of recombinant proteins, which is now comparable to other expression systems including the animal or insect cell cultures. The successful examples include 40% of TSP for HIV-1 antigen via a chloroplast expression (Gonzalez-Rabade et al. 2011); 3.7 mg/g fresh weight of GFP by transient expression using a double terminator (Yamamoto et al. 2018) and 150 µg/L of hGM-CSF using a Ramy3D expression system (Shin et al. 2003)-all of which proved to be high enough amounts to test their potential applications. Furthermore, plant-based vaccine candidates for human and animal diseases such as diarrhea caused by Norwalk virus, enterotoxigenic Escherichia coli (ETEC); Tuberculosis

caused by *Mycobacterium tuberculosis*; Gaucher disease caused by Taliglucerase alfa; Foot and mouth disease caused by foot and mouth disease virus (FMDV), have all been generated and evaluated in experimental models of infection (Chan and Daniell 2015). Likewise, many recombinant anti-cancer vaccines including tumor-associated colorectal cancer antigen, prostatic acid phosphatase (PAP) antigen, human papilloma virus 16 L1/L2 chimeras and the Non-Hodgkin's lymphoma (NHL) have also produced (Lee and Ko 2017) and some have advanced to or completed phases II and III trials (Faye and Gomord 2010).

This study aimed to generate a tetravalent dengue antigen comprising all four serotypes by using a rice amylase 3D expression system and endoplasmic reticulum (ER) targeting. We show that when fused with the M-cell targeting ligand Co1 as the mucosal adjuvant, this plant-based dengue vaccine candidate was immunogenic following oral delivery in mice.

Materials and methods

Construction of a binary vector encoding the tetravalent EDIII fusion Co1 gene

The wild type and synthetic tEDIII genes described in the previous paper (So-Chon Han 2018) were amplified using Ex Taq polymerase (TaKaRa, Korea) for fusion to M-cell ligand peptide (Co1, 12 aa) or further extension with ER targeting peptide (SEKDEL). The purified DNA fragment was subcloned into pGEM-T Easy vector and subsequently sequenced to confirm the correct sequence and in-frame gene (Cosmo Gene Tech, South Korea). The wild type tEDIII fused to Co1 (tEDIII-Co1) or synthetic *tEDIII-Co1* (*stEDIII-Co1*) was amplified with forward and reverse primer sets, Tetra1-F and tEDIII-Co1A(K)-R for wtEDIII-Co1; stEDIII(B)-F and Co1A(K)-R for stEDIII-*Col* (Table 1), respectively. Then, the plasmids digested by BamHI and KpnI of PCR product were cloned into a rice expression vector (So-Chon Han 2018), which contains two expression cassettes for foreign gene and selection marker in transgenic cells, yielding pMYV687 and pMYV690,

 Table 1
 List of primars to amplify Co1 fusion tEDIII genes and confirm the integrated genes using the genomic DNA PCR analysis

| Primer name | Primer sequence (5'–3') |
|------------------|---|
| Tertra 1-F | ggatccATGAAGGGGACGTCATAT |
| tEDIII-Co1A(K)-R | ggtaccTGGAAGTGGAGATCTAGCTGGAAG CTGATGAAAAGAAGAAGAACTTCCTTTCTT GAA |
| stEDIII(B)-F | ggatccATGAAGGGAACT |
| Co1A(K)-R | ggtaccTGGAAGTGGAGA |

respectively (Fig. 1a). For ER targeting of *stEDIII-Co1*, the *stEDIII-Co1* fragment digested by *BamHI/KpnI* was cloned into an intermediate rice expression vector pMYV800, which has ER retention peptide, SEKDEL, between *KpnI* and *SacI* restriction sites, generating pMYV804 (Fig. 1a).

Transformation and regeneration

To obtain the transgenic rice calli, the materials and cultivation protocol was as described in the previous paper (So-Chon Han 2018). Briefly, the embryogenic calli induced from the sterilized rice seeds (*Oryza sativa* L. cv. Dongin) were collected and transformed via a biolistic particle delivery system. The putative transgenic single cells were picked and propagated on the callus induction medium containing 50 mg/L of hygromycin B. For the development of transgenic calli shoots, the selected rice calli were incubated on Cambia medium supplemented with N6 salts with vitamins,

Analysis of the integration and transcription of target genes

sucrose (30 g/L), proline (0.5 g/L), glutamine (0.5 g/L)

casein enzymatic hydrolysate (0.3 g/L), NAA (0.5 mg/L),

BAP (3 mg/L) and hygromycin B (50 mg/l). MS basal medium plus Gelrite (2 g/L) was used to stimulate rooting.

To screen the transformed lines, genomic DNA PCR amplification and Northern blotting were performed as described previously (So-Chon Han 2018). Briefly, the genomic DNA extracted from non-transformed and transformed calli was amplified with gene-specific forward and reverse primers (Table 1) to verify the gene integration into the rice nuclear genome and total rRNA hybridized with α -³²P-labeled (Promega, USA) tEDIII-Co1 or stEDIII-Co1 DNA fragment to detect mRNA expression.

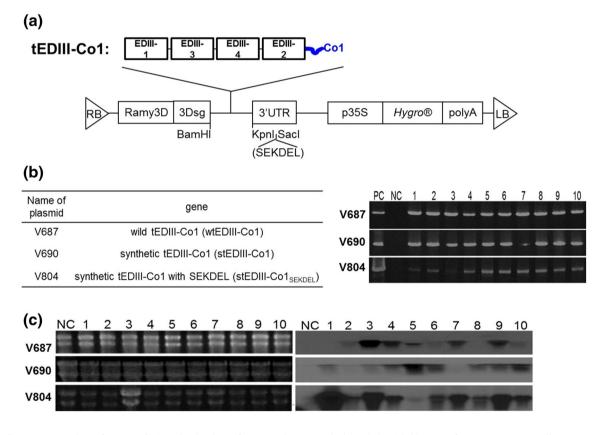


Fig. 1 Gene construction of tEDIII-Co1 and selection of transgenic cell lines by genomic DNA PCR and Northern blot analyses. **a** The wild type or synthesized tEDIII fused to Co1 gene constructed into a binary vector containing two expression cassettes: rice amylase 3D promoter (Ramy3D)—3D signal peptide sequence (3Dsg)—3' untranslated region (UTR); 35S promoter (p35S)—hygromycin resistant gene (Hygro®)—polyA terminator (polyA). SEKDEL peptide sequence was inserted between KpnI and SacI restriction sites within a binary vector for ER retention of tEDIII-Co1. **b** The gene integration of tEDIII-Co1 was confirmed by genomic DNA PCR

analysis and the 1.3 kb DNA fragment corresponding to tEDIII-Co1 detected in transgenic lines. **c** Total rRNA was fractionated on 1.2% formaldehyde-containing agarose gels (left) and the mRNA expression of tEDIII-Co1 was detected in the transgenic cells (right). PC, tEDIII-Co1 DNA amplified with plasmid used as positive control in PCR analysis. NC, the genomic DNA or rRNA extracted from non-transformed rice callus used as negative control in PCR or Norther blot analysis, respectively. Lanes 1–10, the genomic DNA or rRNA extracted from transformed rice callus lines

Detection and quantification of antigen proteins

For detection and quantification of the expressed protein, four transgenic lines showing a strong mRNA expression of each gene construct were selected and lyophilized for further experiments. Total soluble protein (TSP) was extracted with a protein extraction buffer (200 mM Tris-HCl, pH 8.0, 100 mM NaCl, 400 mM sucrose, 10 mM EDTA, 14 mM 2-mercaptoethanol, 1 mM phenylmethylsulfonyl fluoride (PMSF), and 0.05% Tween 20) and used for Western blot analysis after probing by anti-dengue monoclonal antibody (Serotech, UK). TSP of non-transformed callus was used as the negative control in this study. The protocol for generating recombinant tEDIII or cEDIII and the immunoblot analysis was described previously (Kim et al. 2016, So-Chon Han 2018). To measure the amount of tEDIII-Co1 protein produced by each construct, indirect enzyme-linked immunosorbent assay (ELISA) was conducted. Briefly, 3 µg/mL of TSP was loaded in the top wells of a 96-well microtiter plate (NUNC, USA) following two-fold serial dilution in bicarbonate buffer (15 mM Na₂CO₃ and 35 mM NaHCO₃), pH 9.6 and then coated overnight at 4 °C. The wells were blocked by adding 300µL/well of 1% bovine serum albumin (BSA) in PBS and incubating at 37 °C for 2 h, followed by three washes using PBST buffer (PBS containing 0.05% Tween20). The wells were loaded with 100µL/well of a 1:2000 dilution of mouse anti-dengue antibody (Serotech, UK), and incubated for 2 h at 37 °C, followed by washing three times with PBST buffer. The plate was then incubated with 100µL/well of the secondary antibody, a 1:7000 dilution of anti-mouse IgG conjugated with alkaline phosphatase (Sigma-Aldrich, USA) for 2 h at 37 °C and washed three times with PBST buffer. The Optical density (OD) was measured at 405 nm in an ELISA reader (Tecan, Austria) after 15 min incubation with 100µL/well of phosphatase substrate (Sigma-Aldrich, USA).

Oral immunization

The 6 weeks old Balb/c female mice were purchased from Charles River Laboratories through Orient Bio, Inc. (Sungnam, Korea) and maintained under general specific pathogen-free conditions. The experimental procedures involving laboratory animals was approved by the Institutional Animal Care and Use Committee of the Chonbuk National University (Approval Number: CBU 2013-1-0021). One hundred µg of tEDIII-Co1_{SEKDEL} protein was administered orally to mice with or without 10 µg of Cholera Toxin (sigma, USA) or 2.5 mg of Saponin Quillaja sp (Sigma, USA). For oral delivery, a total of 500 mg of dried rice cells, either wild type or producing tEDIII-Co1, was suspended in 500 µL of PBS buffer and administrated by gavage needle to five mice in each group. Equal amount of non-transgenic lyophilized rice cells was used as a control. Oral administration was performed at weeks 0, 2, and 4, and serum and feces from the immunized mice collected 4 days later to analyze the antigen specific antibody induction as depicted in Fig. 3b. Before culling the mice, 100 μ g of bacterial tEDIII-Co1 protein formulated with 100 μ L of incomplete Freund's adjuvant was injected subcutaneously to test for immune tolerance.

Analysis of antibody induction

To test the antibody induction in immunized sera or feces, the recombinant tEDIII protein (80 ng/well) was coated and ELISA performed as described above. 1:200 of serum or 1:10 dilution of fecal extracts were used as the starting concentration and diluted serially. Alkaline phosphatase (AP) conjugated anti- mouse IgG or IgA (Sigma, USA) was used as the detection antibody. For serotype specific cross-reactivity assay, *E. coli*—produced EDIII protein representing each serotype (1–4) was used.

Analyses of lymphocytes by ELISPOT (enzyme-linked immunosorbent spot) and proliferation assay

The splenocytes isolated from the immunized mice were used to measure the number of IgG-secreting cells (SCs) by ELISPOT or to perform lymphocyte proliferation assays as described previously (Kim et al. 2016). Briefly, for ELIS-POT assay, the recombinant tEDIII antigen protein was precoated onto the cellulose ester membrane-lined microtiter plates (Millipore, USA) overnight at 4 °C and incubated with the isolated lymphocytes for 36 h at 37 °C. The cells were then washed and incubated with alkaline phosphatase (AP) conjugated anti-IgG antibody and reaction developed using the BICP/NBT reagent. The spots were counted under a microscope and the mean number of spots from the triplicates for each dilution was calculated. The tEDIII-specific lymphocyte proliferation assay was performed by ³[H] -thymidine incorporation assay as described previously (Kim et al. 2016). Stimulation indices were calculated by dividing the tritium incorporation (cpm) in cells treated with tEDIII by that induced by medium alone.

Statistical analysis

Five animals were used per group for immunization experiments. All ELISA assays were performed in triplicates and the values (from a representative experiment of typically three performed) are shown as the mean +/- standard deviation. For cellular response assay, *t*-test was used to determine statistically significant differences. Exell Microsoft and GraphPad Prism v.7.04 software used for statistical analysis.

Results

Construction of a binary vector encoding the tetravalent EDIII fusion Co1 gene

A single chain tetrameric EDIII (*tEDIII*) covering the all four serotypes has been generated and fused to Co1 peptide (SFHQLPARSPLP) which is a known mucosal adjuvant (Kim et al. 2010). For the extension of *tEDIII* with Co1 at the carboxyl-terminus, a plant codon optimized Co1 was designed for in-frame fusion and synthesized using tEDIII-Co1A(K)-R reverse primer (Table 1). Co1 was added to a wild type or synthetic *tEDIII* by PCR amplification and cloning into a binary vector harboring rice 3D amylase promoter with its signal sequence at the upstream and amylase 3' UTR at the downstream to control an expression of a foreign gene. A strong promoter (Ramy3D) combining with plant codon optimization and/or targeting to subcellular localization (ER) was employed to achieve improved production of tEDIII-Co1 in this study.

Integration and transcription of target genes

Three constructs including pMYV687, pMYV690 and pMYV804 were used to transform rice embryogenic calli and gene integration into the nuclear genome and mRNA expression verified by DNA PCR and Northern blot analyses, respectively. 1.3 kb PCR fragment corresponding to *tEDIII-Co1* was detected in the putative transgenic cell lines, as shown in Fig. 1b, but not in the negative control. Most of the cell lines growing in the hygromycin medium had detectable mRNA expression with somewhat stronger expression detected for synthetic than the wild type gene (Fig. 1c).

Expression of tEDIII-Co1 fusion protein

Four strongest mRNA expressing cell lines were subjected to analysis by anti-dengue monoclonal antibody under reducing

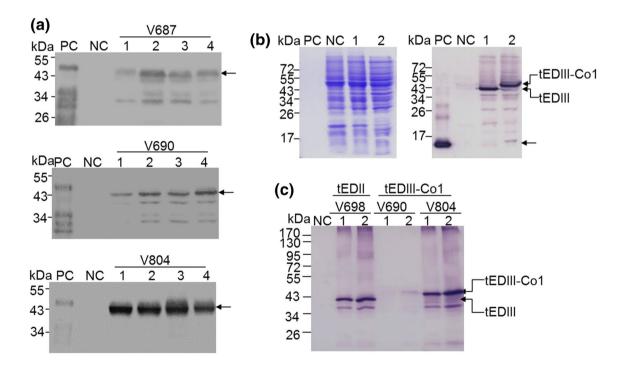


Fig. 2 Detection of tEDIII-Co1 by Western blot analysis. **a** Protein extracts transformed with each construct were analysed to detect tEDIII-Co1 fusion protein. Immunoblotting was performed under reducing (**a**) or non-reducing (**b** and **c**) condition and probing by anti-dengue monoclonal antibodies. The expected size of 47 kDa was detected in all constructs although some truncated fragments were detected in secreted (top and middle) but not ER-targeted protein (bottom). **b** The tEDIII-Co1 cell line showing the highest expression was selected for propagation. 30 µg of total soluble proteins containing tEDIII (lane 1) or tEDIII-Co1 (lane 2) were separated on 12% acrylamide gel (left) and analysed by Western blotting (right)—lane 1: tEDIII; lane 2, tEDIII-Co1. **c** The dried calli expressing tEDIII

(synthesized by plasmid pMYV698) and the two tEDIII-Co1 constructs (synthesized by pMYV690 and pMYV804) were analyzed by Western blotting following 1 year storage at -70 °C; 10 µL and 20 µL of TSP loaded in lane 1 and lane 2, respectively. Recombinant tEDIII-Co1 or EDIII protein produced in E. coli were used as the positive control (Lane PC) and non-transformed rice callus as the negative control (Lane NC) in (**a**), (**b**), and (**c**). Lanes 1–4 in (**a**) represent the transgenic cell lines producing tEDIII-Co1 fusion proteins synthesized by the plasmids pMYV687, pMYV690 and pMYV804, respectively. Molecular weight markers (M) in kDa are shown on the left of each panel (Fig. 2a) or non-reducing (Fig. 2b, c). The main protein band at 47 kDa corresponding to tEDIII-Co1 and some truncated bands appeared in the rice cell lines transformed with constructs pMYV687 (wild type) and pMYV690 (synthetic). Similarly, the recombinant tEDIII-Co1 produced in E. coli appeared as one dominant and a couple of smaller bands - the slightly higher molecular weight of the main expression product than the plant-expressed protein due to the presence of the N-terminal His tag. However, tEDIII-Co1_{SEKDEL} (transformed with pMYV804) appeared as a single band with increased stability (Fig. 2a, bottom panel). No protein was detected in non-transgenic cells for any of the constructs. This result indicates that the higher expression of intact protein is dependent on ER targeting more than codon optimization. The tEDIII-Co1 fusion protein was slightly bigger than the tEDIII alone (Fig. 2b, right panel) due to the presence of Co1 peptide (1.3 kDa). The size of the fusion protein was calculated using the ProParam tool (https://www.expasy.org/proteomics). The quantity of proteins was determined by densitometric analysis of immunoblots by comparing to the known amounts of the bacterial EDIII as described previously (So-Chon Han 2018). The results showed that tEDIII-Co1_{SEKDEL} (V804) expressed 2.4 times higher than tEDIII-Co1 (V690), with TSP estimated to be 3.6% and 1.5%, respectively. However, similar to our previous studies with ER-targeted tEDIII (V698) (So-Chon Han 2018), the transgenic cells, when lyophilized and stored at -70 °C for 1 year showed stable expression of tEDIII-Co1 (V804)] only when ER-targeted, while in the absence of ER targeting (V690) the expression significantly decreased (Fig. 2c). The amount of tEDIII-Co1_{SEKDEL} (V804) was determined to be 316.7 µg per gram of dry weight, which was sufficiently high to be feasible for mouse oral studies.

Oral immunogenicity of tEDIII-Co1 fusion protein

Sera and fecal extracts from mice immunized three times with tEDIII-Co1 alone showed antigen specific-IgG and IgA responses, respectively, (Fig. 3b). Combining with CT or saponin generally did not significantly enhance the mean responses but somewhat reduced variations between individual mice, at either 2 or 4 weeks after the final immunisation. Therefore, neither adjuvant gave significant advantage over Co1 fusion protein alone. Mock immunised mice did not show any antigen specific antibodies. Orally administered tEDIII-Co1 induced a balanced Th1/2 response as indicated by high titre of IgG1 (Th2) and IgG2a and IgG2b (Th1) antibody subtypes (Fig. 3c).

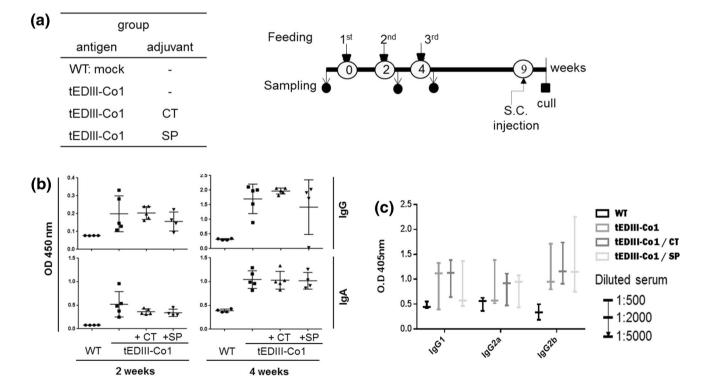


Fig. 3 Oral immunization and antibody responses in sera and feces. **a** Dried cells containing 100 µg of tEDIII-Co1 were gavaged to mice. Four groups including non-transgenic dried cells (WT), tEDIII-Co1 with/without additional adjuvants such as cholera toxin (CT) or saponin (SP) were tested. Three oral administrations to test oral immuno-

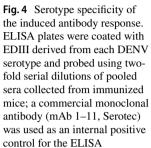
genicity and then a subcutaneous injection 5 weeks later to confirm overcoming oral tolerance, were performed (right). **b** Antigen specific IgG (sera) or IgA (feces) responses, **c** IgG1, IgG2a and IgG2b sub-type responses in pooled sera of immunized mice detected in different serum dilutions

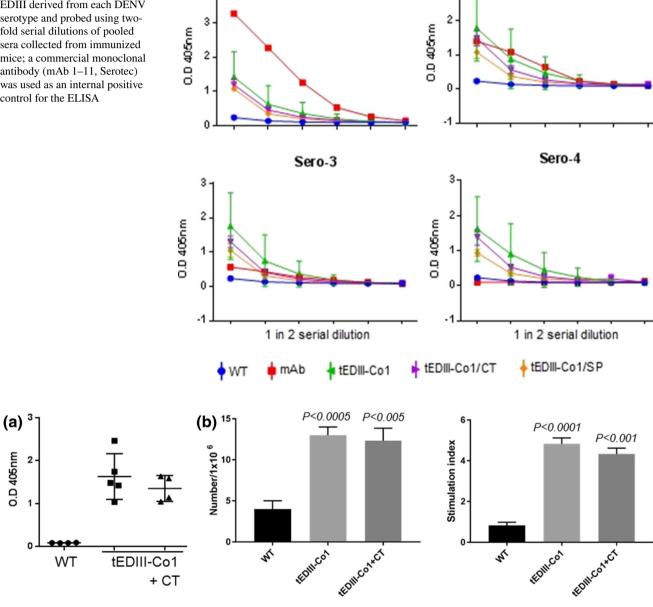
(Fig. 4). Successful oral priming and overcoming of the immunological tolerance was also demonstrated by a 'remote' subcutaneous injection (5 weeks later) that was still capable of boosting the antibody response, as shown in Fig. 5a. Furthermore, antigen-specific B and T cells could be detected in splenocytes from tEDIII-Co1 but not mock immunised mice (Fig. 5).

3

Sero-2

The IgG antibodies induced by tEDIII-Co1 in all groups could react to each serotype specific EDIII antigen and so could be expected to protect against all four serotypes





Sero-1

Fig. 5 Test for oral tolerance and splenic lymphocyte responses in immunized mice. a The IgG antibody showed after subcutaneous injection with recombinant tEDIII protein in the immunized group with tEDIII-Co1 W/O CT adjuvant and the test for oral tolerance confirmed. b B cell ELISPOT, showing the mean number of antibodysecreting cells from triplicate cultures; WT induced low frequency antibody-secreting cells but tEDIII-Co1 induced a statistically significant increase (middle). T cell proliferation measured by 3[H]-thymidine incorporation and presented as the stimulation indices (right)

Discussion

The fastest growing human arboviral disease, dengue, has become a global life-threatening public health concern, with 2.5 billion individuals in more than 100 countries at risk of infection (Brady et al. 2012). To reduce this threat, new antiviral drugs, better vector (mosquito) control and development of an efficient and affordable dengue vaccine are all needed (Hasan et al. 2016). However, the existence of four antigenically different serotypes (DEN-1, DEN-2, DEN-3, and DEN-4) of dengue virus makes the latter task very difficult and complex. Natural immunity to dengue virus infection elicits long-lasting protection only to the same serotype and only short-term (months) protection against other serotypes. Furthermore, severe cases such as dengue hemorrhagic fever (DHF) and dengue shock syndrome (DSS) can develop by sequential infection with different dengue virus serotypes (Halstead 1988; Kurane 2007; Snow et al. 2014). It is for this reason that the WHO strongly recommends that the new dengue vaccine (Dengvaxia®) is given only to people who were previously infected, since giving it to unexposed young children under the age of 9 could lead to more severe subsequent infections (Hadinegoro et al. 2015).

Most of the leading vaccine candidates are formulated with four live-attenuated viruses (Pang and Loh 2017), requiring multiple-dose immunization regimens (Kanesathasan et al. 2001). Therefore, more effort and diverse approaches are required to identify the most promising dengue vaccination strategy. This almost certainly will require detailed scrutiny and comparisons of vaccineinduced immune responses to those induced in natural infection. Different types of vaccine candidates may need to be considered including the so-called 'self-adjuvanting' subunit vaccines, for safety and cost-effectiveness reasons. We reported previously on one such vaccine candidate, a multimeric IgG Fc fusion protein targeting antigen presenting cells and inducing high-level humoral and cellular immune responses (Kim et al. 2018a, 2017). The others include the modified immune complex platform (Kim et al. 2015) and cholera toxin B subunit fusion constructs (Kim et al. 2016). An alternative strategy based on targeting the gut microfold (M) cells was explored in this study is. M cells have an important role in antigen sampling without degradation, which can facilitate antigen-specific immune responses (Cerutti and Rescigno 2008). Kim et al. (2010) isolated the M cell-specific peptide ligands (identified as Co1, Co2 and Co3) and characterized the function of Co1 as a ligand for complement C5a receptors expressed on M cells, thus having a mucosal adjuvant activity (Kim et al. 2010). They demonstrated the uptake of Co1 fusion antigen protein (produced in E. coli) by M cells and induction of antigen-specific B and T cells after mucosal vaccination (Kim et al. 2018b, 2010). More recently, we demonstrated the immunogenic potential of two types of tetravalent dengue vaccine candidates (cEDIII-Co1 and Tet-EDIII-Co1) expressed in plant and yeast cells, respectively (Kim et al. 2013; Nguyen et al. 2015). Furthermore, we and others also tested an animal vaccine candidate based on Co1 fusion with porcine epidemic diarrhoea virus (PEDV) antigens and expressed in transgenic rice or lactobacillus bacteria (Huy et al. 2012; Ma et al. 2018). These studies highlighted the potential application of C5a receptor as a target for mucosal vaccines.

The major target for antibody and T cell responses against flaviviruses such as dengue virus is the envelope (E) protein. In comparison, non-structural (NS) protein 2 is predominantly the target for T cell responses and only a weak antibody response, while NS3 induces only T cell response (Rothman 2004). E domain III of dengue virus is a suitable vaccine target because it is involved in the host cell receptor-binding, can induce neutralizing antibodies and is implicated in the pathogenesis of DHF/DSS. However, EDIII has only a very low intrinsic potential for inducing protective antibodies (Chin et al. 2007; Crill and Roehrig 2001; Simmons et al. 2001, 1998). Etemad et al. (2008) generated the tetravalent chimeric EDIII protein derived from four DEN virus serotypes in *Pichia pastoris* and the purified protein elicited neutralizing antibodies against all four serotypes after intraperitoneal injection with different adjuvants (Etemad et al. 2008). In the present study, oral delivery of crude rice cells expressing tetrameric EDIII-Co1 protein within the ER compartment induced systemic and mucosal immune responses to the dengue antigen. This is the proofof-principle of the feasibility of this approach to develop a cheap and safe vaccine that does not require an exogenous adjuvant. The utility of the M cell targeting by incorporation of the Co1 peptide was underscored by the absence of an immune response by the orally administered tEDIII alone in our previous (unpublished) studies.

In summary, a single tetravalent dengue antigen polypeptide fused to M-cell target ligand (*tEDIII-Co1*) and expressed in transgenic rice cells showed to be a promising oral vaccine candidate. ER targeting (SEKDEL) showed to confer a superior stability and slightly enhanced expression profile compared to the secreted protein. Only 100 μ g of tEDIII fused to Co1 was sufficient to induce both systemic IgG and mucosal IgA antibodies and circumvent oral tolerance. We therefore conclude that this simple tetravalent dengue fusion protein with Co1 could be exploited as an alternative subunit dengue vaccine strategy to complement the existing vaccine approaches.

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Author Contributions M-Y K. conceived and designed the overall study. M-Y K. and B-Y K performed the experiments and co-wrote the manuscript.

Compliance with ethical standards

Conflict of interest The authors declare no conflict of interest.

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