Double-stranded phosphodiester cytosine-guanine oligodeoxynucleotide complexed with calcium phosphate as a potent vaccine adjuvant for activating cellular and Th1-type humoral immunities

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Abstract: Conventional class B cytosine-guanine (CpG) (CpG-B) oligodeoxynucleotide (ODNs) consisting of a single-stranded (ss) phosphorothioate (PT) backbone (ss CpG-B-PT) is converted from a proinflammatory cytokine inducer to a type-I interferon (IFN) inducer when complexed with cationic materials. In this study, we designed ss CpG-B and double-stranded (ds) CpG-B ODNs with a phosphodiester (PD) backbone (ss CpG-B-PD and ds CpG-B-PD, respectively) that became type-I IFN inducers upon complexation with Lipofectamine 2000 (Lipo), a cationic liposome. The ds CpG-B-PD complex induced higher IFN-β expression in mouse macrophage-like RAW264 cells than ss CpG-B-PD and ss CpG-B-PT complexes. The fold induction of IFN-β increased with the number of CpG motifs in ds CpG-B-PD, and a complex of ds CpG-B-PD consisting of 72 base pairs with nine CpG motifs (ds CpG-B72-PD) and Lipo showed the highest capacity to induce IFN-β. The materials and method used for complexation influenced the degree of IFN-β induction: ds CpG-B72-PD entrapped by calcium phosphate (CaP) (ds CpG-B72-PD/CaP) showed a higher induction capacity than ds CpG-B72-PD adsorbed onto the CaP surface. Entrapment of ds CpG-B72-PD by CaP also enhanced the induction of the proinflammatory cytokine interleukin-12. Vaccinating mice with ds CpG-B72-PD/CaP in conjunction with ovalbumin (OVA) increased the ratios of OVA-specific CD8+ T cells in peripheral blood and of OVA-specific IgG2a associated with helper T (Th1) cells to OVA-specific IgG1 associated with Th2 cells. These results indicate that ds CpG-B72-PD/CaP is an effective vaccine adjuvant that can activate both cellular and Th1-type humoral immune responses.

Keywords: CpG oligodeoxynucleotide, calcium phosphate, vaccine adjuvant, cellular immunity, humoral immunity

Introduction

Synthetic single-stranded (ss) oligodeoxynucleotides (ODNs) containing unmethylated cytosine-guanine (CpG) dinucleotides have been studied as vaccine adjuvants for the prophylaxis or treatment of infectious diseases, cancers, and allergies,1–4 owing to their potential to activate humoral and/or cellular immunity.5,6 CpG ODNs are recognized by endosomal Toll-like receptor (TLR)9 and stimulate the production of proinflammatory cytokines and/or type-I interferon (IFN). In humans, TLR9 is expressed only in plasmacytoid dendritic cells (pDCs) and B cells,7–10 while in mice, it is also expressed in macrophages and myeloid DCs.11–13

CpG ODNs developed to date can be broadly divided into four classes based on differences in structure and immunostimulatory properties. Class A CpG (CpG-A) ODNs
(also known as D-type) have a palindromic CpG sequence consisting of a native phosphodiester (PD) backbone at the center of the sequence and a poly-G sequence with an artificial nuclease-resistant phosphorothioate (PT) backbone at either end.\textsuperscript{7,14,15} CpG-A activates pDCs to strongly induce type-I IFN but not B cell activation.\textsuperscript{15,16} Class B CpG (CpG-B) ODNs (also known as K-type) have a linear structure consisting entirely of a PT backbone containing multiple CpG motifs.\textsuperscript{7,14,17,18} CpG-B triggers B cells to activate and induce interleukin (IL)-6 and pDCs to mature and induce tumor necrosis factor (TNF)-\(\alpha\), while scarcely inducing type-I IFN.\textsuperscript{15,16,18} The other two classes, class C CpG (CpG-C) and class P CpG (CpG-P), have one and two palindromic CpG sequences, respectively, that consist entirely of a PT backbone, and both are known to activate B cells and pDCs.\textsuperscript{19–21} In addition, both CpG-C and CpG-P induce type-I IFN, although the latter has a higher capacity for induction.

Alum and emulsion are the preferred vaccine adjuvants for activation of humoral immunity, but new adjuvants that can induce cellular immunity have recently been used for prophylaxis or treatment of various diseases. Since type-I IFN induced by antigen-presenting cells including pDCs is involved in the activation of CD8\(^+\) T cells and helper T (Th)1 cells,\textsuperscript{22–24} CpG-A and CpG-P were considered as candidate vaccine adjuvants. However, these CpG ODNs form uncontrollable higher-order structures due to their palindromic and/or poly-G sequences,\textsuperscript{21,25,26} which precludes their clinical application.\textsuperscript{27} One way to overcome this problem is complexation of CpG-B with other materials such as cationic peptides, polycations, and polysaccharides,\textsuperscript{28–32} because multimerization of conventional CpG-B converts it from an inducer of proinflammatory cytokines to one of type-I IFN. However, complexation can be costly and complicated; this limits the use of CpG-B complexes as an adjuvant for vaccines with high coverage such as influenza vaccine. Furthermore, adverse drug reactions associated with the artificial nuclease-resistant PT backbone of CpG-B have been reported.\textsuperscript{33–40}

In this study, we demonstrate that CpG-B composed of ss native PD backbone (ss CpG-B-PD) can more potently induce IFN-\(\beta\) than conventional CpG-B with ss PT backbone (ss CpG-B-PT) when complexed with a commercially available cationic liposome, and that double-stranded (ds) CpG-B-PD is a more effective IFN-\(\beta\) inducer than the ss CpG-B-PD. We also found that ds CpG-B-PD entrapped by calcium phosphate (CaP; low-crystalline apatite) formed from inexpensive ingredients (calcium chloride and monosodium phosphate) could induce antigen-specific CD8\(^+\) T cells and IgG2a antibody to a degree similar to or higher than ds CpG-B-PD complexed with a cationic liposome. These results highlight the potential for using ds CpG-B-PD entrapped by CaP as an adjuvant for high-coverage vaccines.

**Materials and methods**

**ODNs**

Synthetic and fluorescein isothiocyanate (FITC)-labeled ODNs were obtained from Fasmac (Atsugi, Japan). B-form (B-) DNA was purchased from InvivoGen (San Diego, CA, USA). ds ODNs were prepared by combining an equivalent molar ratio of two synthetic ss ODNs in TES buffer (10 mM Tris–HCl [pH 8.0], 1 mM EDTA, and 0.25 mM NaCl) and incubating the mixture at 95°C for 10 min and at 30°C for 60 min. The ds ODNs were separated by polyacrylamide gel electrophoresis (E-R1020L; ATTO, Tokyo, Japan) at 20 mA and purified by excising the band of appropriate size from the gel in TES buffer followed by concentration with an ultracentrifugal filter unit (Amicon Ultra-0.5, Ultracel-3 membrane, 3 kDa; EMD Millipore, Billerica, MA, USA). The purified product was verified by polyacrylamide gel electrophoresis (Figure S1).

**Preparation of ODN complexes**

Lipofectamine 2000 (Lipo; Thermo Fisher Scientific, Waltham, MA, USA) and N-[1-(2,3-dioleoyloxy)propyl]-N,N,N-trimethylammonium methyl-sulfate (DOTAP) (Roche Applied Science, Penzberg, Germany) were used for the complexation of ODN with cationic liposomes. ODN-Lipo complexes were prepared by mixing ODN and Lipo in a 1:1 (w/w) ratio in Opti-MEM I Reduced Serum Medium (Thermo Fisher Scientific). ODN-DOTAP complexes were prepared by mixing ODN and DOTAP in a 1:6 (w/w) ratio in Tris–EDTA buffer.

ODN-CaP complexes were prepared by two different loading methods – namely, adsorption and entrapment. For the adsorption method, 164 \(\mu\)L of 11 mmol/L CaCl\(_2\) (pH 9) was added dropwise to 109 \(\mu\)L of 11 mmol/L Na\(_2\)HPO\(_4\) (pH 11) at 25°C to form CaP, and 17.5 \(\mu\)g of ODNs was added. After 1 h, the complex formed by adsorption of ODN onto the CaP surface was collected by centrifugation at 15,000 \(\times\) g and 4°C for 15 min. To estimate adsorption efficiency, the concentration of ODN in supernatant (ie, that was not adsorbed onto the CaP surface) was measured with a spectrophotometer (absorption wavelength =260 nm). For the entrapment method, 17.5 \(\mu\)g ODN was mixed with 164 \(\mu\)L of 11 mmol/L CaCl\(_2\) (pH 9), and the mixture was added dropwise to 109 \(\mu\)L of 11 mmol/L Na\(_2\)HPO\(_4\) (pH 11) at 25°C.
After 1 h, the complex resulting from the entrapment of ODN by CaP was collected by centrifugation. ODN concentration in the supernatant (ie, that was not entrapped by CaP) was measured as described above. The precipitated complexes were washed twice with H2O and resuspended in H2O at a concentration of 50 and 500 µg ODN/mL for in vitro and in vivo studies, respectively.

**Cell culture**

The mouse macrophage-like RAW264 cell line (RCB0535) and mouse fibrosarcoma L929 cell line (RCB1451) were purchased from RIKEN BioResource Center (Tsukuba, Japan). The human monocyte-like THP-1 cell line (ATCC® TIB-202™) was purchased from American Type Culture Collection (Manassas, VA, USA). Minimum Essential Medium supplemented with 10% fetal bovine serum (FBS) (Sigma-Aldrich, St Louis, MO, USA), 1% penicillin/streptomycin (P/S; Thermo Fisher Scientific), and 1% nonessential amino acid solution (Wako Pure Chemical Industries, Osaka, Japan) was used for RAW264 cell cultures. Dulbecco’s Modified Eagle’s Medium (Sigma-Aldrich) supplemented with 5% FBS was used for L929 cell cultures. Roswell Park Memorial Institute 1640 medium (Thermo Fisher Scientific) supplemented with 10% FBS and 1% P/S was used for THP-1 cell cultures. Cells were seeded in a 96-well plate at a density of 1×10^4 cells/well (5×10^5 cells/mL). After 18 h, free ODN (final concentration: 50 µg/mL) or ODN complex (final concentration: 5 µg/mL) was added to the culture medium, and RAW264 and THP-1 cells were cultured for another 6 h while L929 cells were cultured for another 24 h to evaluate the induction of IFN-β and IL-12 transcripts. Cell viability was assayed with Cell Counting Kit-8 (Dojindo Laboratories, Kumamoto, Japan).

**Analyses of relative transcript levels and transcription product levels of cytokines**

For analysis of relative transcript levels, total RNA was extracted from cells and purified using Isogen (Nippon Gene, Tokyo, Japan). The mRNA was converted to cDNA with reverse transcriptase (Takara Bio, Kusatsu, Japan) after DNase I treatment. IFN-β and IL-12p40 transcript levels were measured by quantitative real-time PCR (LightCycler 2.0; Roche Applied Science) with SYBR Green and the primers listed in Tables S1 and S2, and normalized to that of glyceraldehyde 3-phosphate dehydrogenase.

To analyze the transcription product levels, we collected the culture medium and diluted it with water. The transcription product level of IFN-β in the diluted culture medium was assayed using a VerKine™ Mouse Interferon-beta ELISA Kit (PBL Assay Science, Piscataway, NJ, USA).

**Confocal fluorescence microscopy**

Cells stimulated for 1 or 4 h with either free or complexed FITC-labeled ODNs were fixed for 15 min with 4% (w/v) paraformaldehyde at 25°C, washed twice with phosphate-buffered saline (PBS), and then permeabilized with 0.2% Tween-20 for 5 min at 25°C before blocking with 3% bovine serum albumin for 1 h at 4°C. The cells were then incubated overnight at 4°C with rabbit anti-ly-sosome associated membrane protein (LAMP)-1 antibody (Abcam, Cambridge, UK) followed by Alexa 555-labeled goat anti-rabbit IgG (Thermo Fisher Scientific) for 1 h at 25°C. The cells were washed twice with PBS and then treated with an Antifade kit (Thermo Fisher Scientific). Fluorescence imaging was performed on a TCS SP5 confocal laser scanning microscope (Leica Microsystems, Wetzlar, Germany).

**Characterization of complexes**

Hydrodynamic size and zeta potential of complexes were measured using a laser diffraction particle size analyzer (SALD-2100; Shimadzu, Kyoto, Japan) and laser electrophoresis zeta-potential analyzer (ELSZ-1000; Otsuka, Tokyo, Japan), respectively.

For scanning electron microscopy (SEM), CaP and ds CpG-B-PD consisting of 72 base pairs (bp) entrapped by CaP (ds CpG-B72-PD/CaP) were resuspended in ethanol and sonicated for 2 min. The suspension was added dropwise onto a silicon wafer that was desiccated at room temperature and then coated with platinum for SEM analysis (JSM-7000F; JEOL, Tokyo, Japan) at an accelerating voltage of 7 kV. Elemental analysis was carried out by energy-dispersive X-ray (EDX) spectroscopy (JED-2200; JEOL) combined with SEM.

The crystalline phase composition of CaP was examined by X-ray diffraction (XRD) (MiniFlex 300/600; Rigaku, Tokyo, Japan) using an X-ray source with the potential set at 40 kV and current set at 15 mA. The specimens were scanned in a range from 5° to 60° at a speed of 0.02°/s. Patterns were analyzed using a model auto-matched to the International Centre for Diffraction Data database. The structural characteristics and vibration bonds of CaP were analyzed by Fourier transform infrared (FTIR) spectroscopy (IRTracer-100; Shimadzu). For the analysis, CaP was prepared as a finely ground powder along with potassium bromide (2%–10%, w/w) and then formed into a transparent pellet using a crimping machine. Pellets were dried under vacuum for 30 min.
Degradation of CaP at various pH

The pH of PBS (originally pH 7.4) was adjusted with 1 mol/L HCl. CaP collected by centrifugation was resuspended in solutions with different pH values. After 1 h, the optical density (OD) at 320 nm was measured with a spectrophotometer (NanoDrop 2000; Thermo Fisher Scientific). The degradation of CaP was calculated with the formula (OD_{various \, pH} / OD_{pH \, 7.4}) \times 100\%. For visualization, FITC-labeled ODNs entrapped by CaP were resuspended in PBS (pH 7.4) and 10 mmol/L acetic acid (pH 5.0). After 1 h, the FITC signal was detected by confocal microscopy.

Immunization

Animal studies were conducted with approval from the Animal Care and Use Committee at the National Institute for Materials Science according to the Guidelines for Proper Conduct of Animal Experiments established by the Science Council of Japan. C57BL/6j mice (5 weeks old) were obtained from Charles River Laboratories Japan (Yokohama, Japan). A 200 µL volume of vaccine solution containing 200 µg ovalbumin (OVA) (Sigma-Aldrich) alone or mixed with 50 µg CpG ODNs with Lipo or CaP was subcutaneously injected into the back of 6-week-old mice twice at a 1-week interval. One week after the second injection, mice were euthanized with overdose of isoflurane (DS Pharma Animal Health, Osaka, Japan), and peripheral blood was collected from the caudal vena cava.

Estimation of OVA-specific CD8^+ T cell abundance and OVA-specific antibodies

A portion of the collected peripheral blood was treated with EDTA/3K. A 10 µL volume of T-Select H-2K^b OVA Tetramer-SINFEKL-phycocerythrin (PE) (TS-5001-1C; Medical & Biological Laboratories, Nagoya, Japan) was added to 50–100 µL EDTA-treated blood, and the mixture was allowed to stand at 25°C for 20 min before adding 1 µL of mouse anti-CD3 PE-cyanine 5-conjugated antibody (Thermo Fisher Scientific) and 1 µL of mouse anti-CD8 FITC-conjugated monoclonal antibody (KT15) (Medical & Biological Laboratories). After 20 min, 500 µL Optilyse C (Beckman Coulter, Brea, CA, USA) was added to the blood to induce hemolysis, and the mixture was allowed to stand for 10 min before adding 500 µL PBS. The cell suspension was purified on a cell sorter (SH800; Sony, Tokyo, Japan). To identify OVA-specific CD8^+ T cells, lymphocytes were gated on the basis of a forward scatter (FSC)-vs-side scatter (SSC) density plot, and then developed to SSC-vs-CD3 density plot. The CD3^+ population was further developed into an OVA tetramer-vs-CD8 density plot.

To obtain serum, peripheral blood was centrifuged at 1,800 \times g for 15 min at 4°C. The levels of OVA-specific IgG1 and IgG2a were assayed using mouse anti-OVA IgG1 and IgG2a assay kits (both from Chondrex, Redmond, WA, USA).

Statistics

Statistical significance for in vitro data was evaluated with the two-tailed unpaired Student’s t-test. For multiple comparisons, the p-value for significance was corrected with the Bonferroni method. For in vivo data, all groups were compared by analysis of variance followed by Tukey’s honestly significant difference post hoc test and Ryan’s method. A p-value <0.05 was considered statistically significant.

Results

ds CpG-B-PD complex is a potent type-I IFN inducer

ODN sequences used in this study are shown in Box 1. ss ODN2006 (hereafter referred to as ss CpG-B24-PT) is a conventional CpG-B ODN consisting of 24 PT nucleotides (nt) and three CpG motifs (GTCGTT). Although complexation of CpG-B with cationic materials reportedly converts CpG-B from proinflammatory cytokine inducer to type-I IFN inducer,29-31 there was no induction of IFN-β when mouse macrophage-like RAW264 cells expressing TLR9 were stimulated with ss CpG-B24-PT complexed with Lipo, a commercially available cationic liposome (Figure 1A). On the other hand, the complex of ss CpG-B24-PD – which has a PD backbone – with Lipo showed a high capacity for inducing IFN-β as compared to the ss CpG-B24-PT complex (Figure 1A), while the complex of ds CpG-B24-PD and Lipo had a greater capacity to induce IFN-β than the ss CpG-B24-PD complex (Figure 1B).

ds CpG-B24-PD consists of 24 bp with three CpG motifs. We prepared ds CpG-B48-PD and ds CpG-B72-PD, which have two and three linked ds CpG-B24-PD sequences, respectively (ie, six and nine CpG motifs, respectively) (Box 1). IFN-β induction in the ds CpG-B-PD-Lipo complexes increased with the number of CpG motifs (Figure 1C). We observed IFN-β induction by ds CpG-free24-PD, ds CpG-free48-PD, and ds CpG-free72-PD complexes that
were 24, 48, and 72 bp ODNs, respectively, without CpG motifs, although the induction level was lower than that by the complex of B-DNA, which is a ligand of cytosolic DNA receptors (Figure 1C). In this experiment, we also analyzed the level of IFN-β secreted into the culture medium and confirmed the correlation between the transcript level and secretion level in RAW264 cells (Figure S2). IFN-β induction by CpG-free ODN complexes may be mediated by cytosolic DNA receptors that recognize ds PD DNA in a CpG-independent manner. IFN-β levels induced by the ds CpG-B24-PD, ds CpG-B48-PD, and ds CpG-B72-PD complexes were higher than those induced by ds CpG-free ODN complexes, suggesting that IFN-β induction by the former is highly dependent on the presence of CpG motifs. We evaluated IFN-β induction potential of ds CpG-B-PT (ds CpG-B24-PT and ds CpG-B72-PT) complexed with Lipo but did not observe any significant induction (Figure 1D).

We previously reported that free ss CpG-B72-PD is a proinflammatory cytokine inducer. Free ds CpG-B72-PD induced IL-12 in RAW264 cells, although the level was markedly lower than that induced by free ss CpG-B72-PD (Figure S3). On the other hand, free ds CpG-B72-PD showed no capacity to induce IL-12. These results imply that free ds CpG-B72-PD is also a proinflammatory cytokine inducer, and that this activity depends on the CpG motif.

CaP is an effective material for ds CpG-B-PD complexation

We tested another commercially available cationic liposome, DOTAP, for complexation with ds CpG-B72-PD. The level of IFN-β induced by ds CpG-B72-PD complexed with DOTAP was significantly lower than that induced by ds CpG-B72-PD complexed with Lipo (ds CpG-B72-PD-Lipo) (Figure 2A). This suggests that the material used for ds CpG-B-PD ODN complexation influences the degree of IFN-β induction. Although effective, Lipo is expensive and can only be used in laboratory settings. Important considerations for adjuvants – especially those for high-coverage vaccines – are manufacturing cost as well as adjuvant effect. We therefore examined whether CaP can be used for complexation of ds CpG-B-PD, since the ingredients required for CaP preparation are inexpensive and production is simple and readily scalable. We complexed ds CpG-B72-PD with CaP by entrapment of ds CpG-B72-PD by CaP (ds CpG-B72-PD/CaP) and by adsorption of ds CpG-B72-PD onto the CaP surface (ds CpG-B72-PD-CaP) (Figure S4A and B). The complexation efficiency of ds CpG-B72-PD was
approximately 100% in both the entrapment and adsorption methods. Assuming that all Ca\(^{2+}\) was used for CaP formation, the amount of ds CpG-B72-PD loaded was estimated to be around 68 and 65 \(\mu\)g/mg CaP, respectively. The mean hydrodynamic size of naked CaP was 1,115 ± 332 nm, whereas the mean sizes of ds CpG-B72-PD/CaP and ds CpG-B72-PD-CaP were 437 ± 135 and 717 ± 183 nm, respectively (Figure S5).

The zeta potential was −5.46 mV for naked CaP, −16.1 mV for ds CpG-B72-PD/CaP, and −28.1 mV for ds CpG-B72-PD-CaP. When RAW264 cells were stimulated with these complexes, a higher level of IFN-\(\beta\) was observed in the presence of ds CpG-B72-PD/CaP (Figure 2B). IFN-\(\beta\) level was about 1.4 times higher when induced by ds CpG-B72-PD/CaP as compared to ds CpG-B72-PD-Lipo. On the other hand, the level of IFN-\(\beta\) induction by ds CpG-B72-PD-CaP was about 30% that by ds CpG-B72-PD-Lipo (Figure 2B). This indicates that the loading method affects the adjuvant effect of ds CpG-B72-PD, and that ds CpG-B72-PD entrapment by CaP is superior to ds CpG-B72-PD adsorption onto the CaP surface for complexation. We also observed higher IFN-\(\beta\) induction with ds CpG-B72-PD/CaP than with ss CpG-B72-PD/CaP or ds CpG-free72-PD/CaP; the level of IFN-\(\beta\) induced by ds CpG-B48-PD/CaP showed a higher capacity for induction than D35/CaP (CpG-A entrapped by CaP) (Figure 2D).

We also examined the capacity of ds CpG-B72-PD/CaP to induce IFN-\(\beta\) via cytosolic DNA receptors using murine...
fibrosarcoma L929 cells and human monocyte-like THP-1 cells. Neither of these cell lines express TLR9 but have cytosolic DNA receptors. In L929 cells, IFN-β induction was higher by B-DNA complexed with Lipo (B-DNA-Lipo) than by ds CpG-B72-PD-Lipo (Figure S6A). In addition, the potential to induce IFN-β was lower for B-DNA complexed with CaP (B-DNA/CaP) and ds CpG-B72-PD/CaP as compared to B-DNA-Lipo and ds CpG-B72-PD-Lipo (Figure S6A). L929 and RAW264 cells are adherent, while THP-1 cells grow in suspension. Similar to L929 cells, in THP-1 cells IFN-β induction was higher in the presence of B-DNA-Lipo as compared to ds CpG-B72-PD-Lipo (Figure S6B). On the other hand, IFN-β was not induced by B-DNA/CaP or ds CpG-B72-PD/CaP (Figure S6B), although CaP was internalized by suspended THP-1 cells and ODN molecules were localized in the cytosol (Figure S6C). These results suggest that ds CpG-B72-PD has a lower capacity to induce IFN-β via cytosolic DNA receptors than B-DNA and that ds CpG-B72-PD delivered by CaP does not significantly activate this pathway via cytosolic DNA receptors.

Characterization of CaP

Naked CaP is an agglomerate consisting of elongated flakes with a long side ≤200 nm (Figure 3A). On the other hand, ds CpG-B72-PD/CaP is an agglomerate of petal-like flakes with a long side <150 nm (Figure 3B). The EDX spectroscopy...
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analysis detected C, O, P, and Ca in both naked CaP and ds CpG-B72-PD/CaP (Figure 3A and B). In the FTIR spectrum of naked CaP (Figure 3C), bands for the P–O vibration of the orthophosphate group were observed at 560, 570, 958, and 1,034 cm\(^{-1}\). A band attributed to the CO\(_3^{2-}\) stretching mode was observed at 862 cm\(^{-1}\). The bands for water vibration at 3,422 and 1,629 cm\(^{-1}\) provided evidence for the hydration of naked CaP. In addition to these peaks, bands attributed to NH\(^4\) were detected at 1,210, 1,400, and 1,470 cm\(^{-1}\) in ds CpG-B72-PD/CaP (Figure 3C), suggesting the presence of ds CpG-B72-PD. The XRD analysis revealed that the crystal structure of both naked CaP and ds CpG-B72-PD/CaP was identical to a low-crystalline hydroxyapatite (product number 677418; Sigma-Aldrich) (Figure 3D).

We also examined the pH-dependent degradation of CaP particles. Approximately 95% of CaP was degraded in a pH 5 solution within 1 h (Figure 4A and B), implying that CaP can be easily degraded in the acidic environment of endosomes and lysosomes. RAW264 cells were incubated with FITC-labeled ODNs entrapped by CaP and examined by confocal fluorescence microscopy. A fluorescence signal corresponding to FITC was observed near the inner cell membrane and in late-endosomes/lysosomes-expressing LAMP-1 after 1 h (Figure 4C). FITC near the inner cell membrane was...
presumed to be associated with early endosomes, given that the signal was also detected in late endosomes/lysosomes, which are part of the same endocytic transport route. On the other hand, FITC was observed throughout the cell after 4 h (Figure 4C), suggesting that ds CpG-B72-PD had escaped from endosomes and/or lysosomes into the cytosol due to degradation of CaP in the acidic environment in these organelles. Although CaP degradation increases intracellular Ca\(^{2+}\) concentration, no cytotoxicity was observed (Figure 4D).

**Adjuvant effect of ds CpG-B72-PD/CaP**

We next examined whether ds CpG-B72-PD/CaP induces antigen-specific CD8\(^+\) T cells by subcutaneously injecting ds CpG-B72-PD/CaP or ds CpG-B72-PD-Lipo in conjunction with OVA (ds CpG-B72-PD/CaP + OVA or ds CpG-B72-PD-Lipo + OVA, respectively) into mice twice at a 1-week interval. One week after the last injection, the ratio of OVA-specific CD8\(^+\) T cells to total CD8\(^+\) T cells was 10-fold higher in the group vaccinated with ds CpG-B72-PD/CaP + OVA than in mice vaccinated with OVA alone (Figures 5A and S7A and B). OVA alone did not increase the ratio of OVA-specific CD8\(^+\) T cells (Figure S8A). Although we observed a higher level of IFN-\(\beta\) induction in RAW264 cells stimulated with ds CpG-B72-PD/CaP than with ds CpG-B72-PD-Lipo (Figure 2B), no significant difference was observed in the ratio of OVA-specific CD8\(^+\) T cells between mice vaccinated with ds CpG-B72-PD/CaP + OVA and ds CpG-B72-PD-Lipo + OVA (Figures 5A and S7C).

We also analyzed serum IgG1 and IgG2a antibody levels 1 week after the last injection. Administration of OVA alone significantly increased the level of OVA-specific IgG1 antibody (Figure S8B). The average level of OVA-specific IgG1 antibody in mice vaccinated with ds CpG-B72-PD/CaP + OVA and ds CpG-B72-PD-Lipo + OVA was 3-
4-fold higher than those in mice vaccinated with OVA alone, respectively, although these were not statistically significant differences (Figure 5B). On the other hand, ds CpG-B72-PD/CaP + OVA elicited 23- and 2-fold higher induction of OVA-specific IgG2a as compared to OVA alone and ds CpG-B72-PD-Lipo + OVA, respectively (Figure 5C). The average induction level of OVA-specific IgG2a in the ds CpG-B72-PD-Lipo + OVA group was 11-fold higher than that in mice injected with OVA alone, but the difference was not statistically significant. OVA alone did not trigger any detectable OVA-specific IgG2a antibody response (Figure S8C). Although no increase in anti-OVA IgG2a/IgG1 ratio was observed in the ds CpG-B72-PD-Lipo + OVA as compared to the OVA-only group, mice vaccinated with ds CpG-B72-PD/CaP + OVA showed a significantly increased ratio (Figure 5D), suggesting a shift toward a Th1-dominated response. One of the triggers for Th1-biased pathway activation is the differentiation of naïve
T cells to Th1 cells induced by IL-12 secreted by antigen-presenting cells.\textsuperscript{44–46} We therefore examined IL-12 induction in RAW264 cells stimulated with ds CpG-B72-PD/CaP and found that IL-12 was induced to a greater degree in the ds CpG-B72-PD/CaP than in the ds CpG-B72-PD-Lipo group (Figure 6A). In addition, ds CpG-B72-PD/CaP showed a higher capacity for IL-12 induction than free ds CpG-B72-PD (Figure 6B), whereas D35/CaP and ds CpG-free72-PD/CaP did not induce IL-12 (Figure 6C).

**Discussion**

Alum is widely used as a vaccine adjuvant; although it activates humoral immunity, it has poor capacity for inducing cellular immunity. Type-I IFN induced by antigen-presenting cells such as pDCs is involved in the activation of cellular immunity.\textsuperscript{22–24} It has been reported that complexation of conventional CpG-B (ss CpG-B-PT) with materials such as cationic peptides, polycations, and polysaccharides confers conventional CpG-B with the capacity to induce type-I IFN.\textsuperscript{28–32} In this study, we demonstrate that in a complex with Lipo, ds CpG-B72-PD designed based on the ss CpG-B24-PT sequence (known as the conventional CpG-B ODN2006) is a potent inducer of IFN-β as compared to ss CpG-B24-PT. ds CpG-B72-PD is a proinflammatory inducer that is similar to conventional CpG-B but has a ds native PD backbone instead of an ss artificial PT backbone.

DNA consisting of a PT backbone is more resistant to nuclease digestion than natural PD DNA, but has been reported to cause thrombocytopenia, anemia, and neutropenia in rats and arthritis in mice.\textsuperscript{33,34} Furthermore, a higher risk of adverse events was demonstrated in clinical trials using conventional CpG-B-PT.\textsuperscript{47,48} To avoid this toxicity, CpG ODNs with a native PD backbone were developed. MGN1703 consisting entirely of PD backbone, which has two 30 nt loops containing three CpG motifs and a 28 bp stem linking these loops stimulates the release of IFN-α and proinflammatory cytokines such as IFN-γ, IL-6, IL-12, and TNF-α by human peripheral blood mononuclear cells, resulting in activation of both cellular and humoral immunity.\textsuperscript{59,50} In MGN1703, CpG motifs are present in ss loops, implying that a ds CpG motif is not essential for type-I IFN induction. Additionally, MGN1703 is not required for complexation and type-I IFN induction, and its safety in humans was verified in a Phase I clinical trial.\textsuperscript{51} An X-shaped ds PD CpG ODN can be formed by self-assembly of four complementary ss PD CpG ODNs,\textsuperscript{62} while one composed of four ss PD CpG ODN molecules forms a self-ligated complex through complementarity with the sticky ACGT sequence at the 5‘ end.\textsuperscript{53} These non-self- and self-ligated X-shaped ds PD CpG ODN induced IFN-β in addition to IL-12 and TNF-α in mouse bone marrow-derived primary DCs;\textsuperscript{53} the degree of IFN-β induction was higher for the self-ligated as compared to the non-self-ligated molecule. Our uncomplexed ds CpG-B72-PD – which was incapable of inducing IFN-β – has a linear structure without branching, in contrast to the self-ligated X-shaped ds PD CpG ODN which had more branches than the non-self-ligated form. This suggests that free ds PD CpG ODN without complexation can potentially be endowed with type-I IFN induction capacity via branching. In a previous study, ds PD CpG ODN molecules with
polypod structures (tri-, tetra-, penta-, hexa-, and octapod) were synthesized via self-assembly of three to eight ss PD CpG ODNs. In RAW264 cells, the levels of proinflammatory cytokines including IL-6 and TNF-α were higher in cells stimulated with a higher branch (polypod) number. Although these authors did not demonstrate IFN-β induction in cells stimulated with polypod-like ds PD CpG ODN, CD8+ T cell activation in mice subcutaneously injected with hexapod ds PD CpG ODN has been reported.

For the crystal structure of CaP prepared for ds CpG-B72-PD complexation, we detected a band attributed to the CO$_3^{2-}$ group at 862 cm$^{-1}$ in the FTIR spectrum. The band for the CO$_3^{2-}$ group of hydroxyapatite generally appears around 1,400–1,600 cm$^{-1}$ (Figure S9), but we did not observe this band in our CaP. Therefore, the crystal structure of our CaP was deduced to be identical to that of low-crystalline hydroxyapatite, but its chemical composition is different from that of hydroxyapatite, where the molar ratio of Ca to P is 1.67. We loaded ds CpG-B72-PD by using adsorption and entrapment methods. Although the detailed mechanism by which ds CpG-B72-PD binds to the CaP surface in the adsorption method remains unclear, it has been reported that both C and P sites are involved in the electrostatic interaction, and play a critical role in the binding of DNA to the hydroxyapatite surface. In the complexation of ds CpG-B72-PD with CaP, the loading method influenced IFN-β induction. In ds CpG-B72-PD-Lipo, ds CpG-B72-PD binds to the Lipo surface via electrostatic interactions (Figure S4C). Although ds CpG-B72-PD was also located on the surface of CaP in ds CpG-B72-PD-CaP (Figure S4B), it showed a lower capacity for inducing IFN-β than ds CpG-B72-PD-Lipo. ds CpG-B72-PD bound to the Lipo surface is thought to interact with endosomal TLR9. In contrast, in ds CpG-B72-PD-CaP, most of the ds CpG-B72-PD molecules may be immediately released upon dissolution of CaP in the acidic environment of endosomes/lysosomes (Figure 7A) and thus lack the capacity to induce IFN-β. In ds CpG-B72-PD/CaP, which induced IFN-β to a greater extent than ds CpG-B72-PD-CaP, the CaP surface had randomly multimerized ds CpG-B72-PD exposed by the dissolution of the original surface (Figure 7B), which were presumed to interact with TLR9 to induce IFN-β and IL-12.

![Proposed model of CaP degradation and ODN release.](https://www.dovepress.com/)

**Figure 7** Proposed model of CaP degradation and ODN release.

**Notes:** (A) ds CpG-B72-PD molecules adsorbed onto the surface of CaP (cross-sectional view of ds CpG-B72-PD-CaP). Most ds CpG-B72-PD molecules are immediately released by the degradation of CaP. (B) ds CpG-B72-PD molecules entrapped by CaP (cross-sectional view of ds CpG-B72-PD/CaP). Surface degradation of CaP releases ds CpG-B72-PD molecules entrapped near the surface, but ds CpG-B72-PD molecules inside CaP are exposed on the new surface.

**Abbreviations:** CaP, calcium phosphate; ODN, oligodeoxynucleotide; ds, double stranded; CpG-B, class B cytosine-guanine; PD, phosphodiester; TLR, Toll-like receptor; IFN, interferon; IL, interleukin.
RAW264 cells express cytosolic DNA receptors in addition to TLR9, including IFN-γ-inducible protein 16, dead-box helicase 41, and cyclic GMP-AMP synthase, which recognize ds PD DNA in a sequence-independent manner and induce type-I IFN via stimulator of IFN genes (STING). It is possible that IFN-β induction in RAW264 cells stimulated with ds CpG-B72-PD-CaP was due to cytosolic DNA receptors, since ds CpG-B72-PD molecules released from CaP were localized throughout cells including in the cytosol after 4 h of stimulation with ds CpG-B72-PD-CaP. However, the degree of IFN-β induction by ds CpG-B72-PD was much higher than that by ds CpG-free72-PD, implying that IFN-β induction by ds CpG-B72-PD/CaP is heavily dependent on TLR9 but not cytosolic DNA receptors. In L929 and THP-1 cells expressing only cytosolic DNA receptors, IFN-β induction by ds CpG-B72-PD-Lipo was lower than that by B-DNA-Lipo. In addition, both ds CpG-B72-PD/CaP and B-DNA/CaP barely induced IFN-β in L929 and THP-1 cells, although ds CpG-B72-PD and B-DNA induced IFN-β when complexed with Lipo. Free ds CpG-B72-PD and B-DNA molecules released from CaP may be degraded before or during escape from endosomes/lysosomes, while these molecules released from Lipo could still be protected by cationic lipids. However, it is unclear why ds CpG-B72-PD/CaP and B-DNA/CaP barely induced IFN-β, and there is no information on whether MGN1703 and polypod-like ds PD CpG ODN induce type-I IFN via interaction with cytosolic DNA receptors.

IFN-β level in RAW264 cells was higher in the presence of ds CpG-B72-PD/CaP than of ds CpG-B72-PD-Lipo, but there was no difference in the ratio of OVA-specific CD8+ T cells to total CD8+ T cells in the peripheral blood of mice administered these complexes in conjunction with OVA. This suggests that IFN-β induction by ds CpG-B72-PD-Lipo may be sufficient for activation of cellular immunity. In addition, ds CpG-B72-PD/CaP stimulated the production of OVA-specific IgG2a antibody to a greater extent than ds CpG-B72-Lipo. This may be attributed to the high level of IL-12 induction in the presence of ds CpG-B72-PD/CaP as compared to ds CpG-B72-PD-Lipo. Although ds CpG-B72-PD-Lipo did not increase the ratio of IgG2a to IgG1, an increase was observed for ds CpG-B72-PD/CaP, implying that the latter can potentially shift the immune response to one that is Th1 predominant. Enhancement of Th1/Th2 balance is a therapeutic strategy for allergies caused by pollen and house dust. However, we obtained variable results when ds CpG-B72-PD/CaP was used as an adjuvant in mice, which could be due to a problem with ds CpG-B72-PD/CaP itself such as aggregation and not to individual differences among mice.

In this study, we demonstrated that ds CpG-B72-PD/CaP can activate both cellular and Th1-type humoral immunity. ds CpG-B72-PD/CaP preparation is simple and easily scalable, making it a promising adjuvant for high-coverage vaccines such as influenza vaccine.

Acknowledgments
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Disclosure
The authors report no conflicts of interest in this work.

References


Supplementary materials

Table S1 Primer sequences for the analysis of transcript expression in RAW264.7 and L929 cells

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<th>Gene</th>
<th>Forward sequence</th>
<th>Reverse sequence</th>
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<td>GAPDH</td>
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<td>5′-TGTGAGGGAGATG</td>
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<td>GATCTCCA-3′</td>
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Abbreviations: IFN, interferon; IL, interleukin.

Table S2 Primer sequence for the analysis of transcript expression in THP-1 cells

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<td>5′-CAATGTCAGTCCA</td>
</tr>
<tr>
<td></td>
<td>TACC-3′</td>
<td>GAGG-3′</td>
</tr>
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Abbreviation: IFN, interferon.

Figure S1 Polyacrylamide gel electrophoresis.
Notes: Lane 1, ds CpG-B72-PD before purification. Lane 2, ds CpG-B72-PD after purification; ds CpG-B72-PD recovered from the gel was concentrated with EtOH. Lane 3, ds CpG-B72-PD after purification; ds CpG-B72-PD recovered from the gel was concentrated with an ultracentrifugal filter unit.
Abbreviations: ds, double stranded; CpG-B, class B CpG; PD, phosphodiester; CpG, cytosine-guanine; bp, base pairs.

Figure S2 The level of IFN-β secreted into the culture medium.
Notes: IFN-β level in the culture medium was determined by ELISA. ds CpG-B-PD ODNs and ds PD CpG-free ODNs were complexed with Lipofectamine 2000. Data represent mean ± SD (n=3). ***p<0.001.
Abbreviations: IFN, interferon; ds, double stranded; PD, phosphodiester; CpG-B, class B CpG; ODNs, oligodeoxynucleotides; CpG, cytosine-guanine.

Figure S3 Free ds CpG-B72-PD is a proinflammatory cytokine inducer.
Notes: RAW264.7 cells were stimulated with free ss CpG-B72-PD, ds CpG-B72-PD, and ds CpG-free72-PD. After 6 h, total RNA was recovered, and relative IL-12 and IFN-β transcript levels were evaluated by quantitative real-time PCR. ODN concentration was 50 µg/ml. IFN-β transcript was not detected in cells stimulated with any ODN. *p<0.05.
Abbreviations: ds, double stranded; CpG-B, class B CpG; PD, phosphodiester; ss, single stranded; CpG, cytosine-guanine; IL, interleukin; IFN, interferon; ODN, oligodeoxynucleotide; ND, not detected.
A  ds CpG-B72-PD  
\( \text{Ca}^{2+} + \text{PO}_{4}^{2-} \rightarrow \text{ds CpG-B72-PD/CaP} \)

B  ds CpG-B72-PD  
\( \text{Ca}^{2+} \rightarrow \text{CaP} \rightarrow \text{ds CpG-B72-PD-CaP} \)

C  ds CpG-B72-PD  
\( \text{Lipo} \rightarrow \text{ds CpG-B72-PD-Lipo} \)

**Figure S4** Methods for loading ds CpG-B72-PD onto CaP.

**Notes:** (A) Entrapment method. ds CpG-B72-PD molecules were entrapped by CaP. (B) Adsorption method. ds CpG-B72-PD molecules were adsorbed onto the surface of CaP. (C) ds CpG-B72-PD complexed with Lipo via electrostatic interactions.

**Abbreviations:** ds, double stranded; CpG-B, class B CpG; CpG, cytosine-guanine; PD, phosphodiester; CaP, calcium phosphate; Lipo, Lipofectamine 2000.

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**Figure S5** Hydrodynamic size distribution of naked CaP and ds CpG-B72-PD complexed with CaP.

**Notes:** Naked CaP, ds CpG-B72-PD/CaP, and ds CpG-B72-PD-CaP were collected by centrifugation at 15,000 \( \times \) g for 15 min at 4\(^\circ\)C, and then resuspended in pure water at a concentration of 850 \( \mu \)g CaP/ml. Size distribution was measured at 25\(^\circ\)C after sonication for 1 min.

**Abbreviations:** CaP, calcium phosphate; ds, double stranded; CpG-B, class B CpG; CpG, cytosine-guanine; PD, phosphodiester.
Figure S6 IFN-β induction via cytosolic DNA receptors.
Notes: (A) IFN-β induction in murine fibrosarcoma L929 cells stimulated for 24 h. (B) IFN-β induction in human monocyte-like THP-1 cells stimulated for 6 h. **p<0.025. (C) Internalization of FITC-labeled ODNs complexed with CaP by THP-1 cells, as observed by confocal fluorescence microscopy. The three images show cross sections of the same cell at different heights.
Abbreviations: IFN, interferon; FITC, fluorescein isothiocyanate; ODNs, oligodeoxynucleotides; CaP, calcium phosphate; ds, double stranded; CpG-B, class B CpG; CpG, cytosine-guanine; PD, phosphodiester; Lipo, Lipofectamine 2000.

Figure S7 (Continued)
Figure S7 Responses of CD8+ T cells in peripheral blood analyzed by cell sorting.

Notes: OVA tetramer-vs-CD8 density plot in mouse vaccinated with (A) OVA alone, (B) OVA + ds CpG-B72-PD/CaP, and (C) OVA + ds CpG-B72-PD-Lipo. Ratio of OVA-specific CD8+ T cells to total CD8+ T cells in each mouse was calculated from these density plots. UR (C-Q2) and LR (C-Q4) regions in quadrant region of each density plot indicate the ratios of OVA-specific and OVA-nonspecific CD8+ T cells in total CD3+ T cells. Ratio of OVA-specific CD8+ T cells was calculated from the equation: 100 × (% in UR)/{(% in UR) + (% in LR)}.

Abbreviations: OVA, ovalbumin; ds, double stranded; CpG-B, class B cytosine-guanine; PD, phosphodiester; CaP, calcium phosphate; Lipo, Lipofectamine 2000; UR, upper right; LR, lower right; PE, phycoerythrin; FITC, fluorescein isothiocyanate; SSC, side scatter; FSC, forward scatter; Cy5, cyanine 5.
Figure S8 OVA-specific immune responses in mice.
Notes: (A) Ratio of OVA-specific CD8⁺ T cells in total CD8⁺ T cells in peripheral blood. Each circle corresponds to the ratio of OVA-specific CD8⁺ T cells in an individual mouse, and horizontal lines indicate mean ± SD. (B) OVA-specific IgG1 antibody level in serum. Each circle corresponds to OVA-specific IgG1 antibody level in an individual mouse. (C) OVA-specific IgG2a antibody level in serum. Each circle corresponds to OVA-specific IgG2a antibody level in an individual mouse. *p<0.05 (analysis of variance followed by Tukey’s honestly significant difference post hoc test and Ryan’s method).
Abbreviations: NS, not significant; OVA, ovalbumin.

Figure S9 FTIR spectrum of hydroxyapatite.
Abbreviation: FTIR, Fourier transform infrared.