A candidate DNA vaccine elicits HCV specific humoral and cellular immune responses

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Abstract

AIM: To investigate the immunogenicity of candidate DNA vaccine against hepatitis C virus (HCV) delivered by two plasmids expressing HCV envelope protein 1 (E1) and envelope protein 2 (E2) antigens respectively and to study the effect of CpG adjuvant on this candidate vaccine.

METHODS: Recombinant plasmids expressing HCV E1 and E2 antigens respectively were used to simultaneously inoculate mice with or without CpG adjuvant. Antisera were then collected and titers of anti-HCV antibodies were analyzed by ELISA. One month after the last injection, animals were sacrificed to prepare single-cell suspension of splenocytes. These cells were subjected to HCV antigen specific proliferation assays and cytokine secretion assays to evaluate the cellular immune responses of the vaccinated animals.

RESULTS: Antibody responses to HCV E1 and E2 antigens were detected in vaccinated animals. Animals receiving CpG adjuvant had slightly lower titers of anti-HCV antibodies in the sera, while the splenocytes from these animals showed higher HCV-antigen specific proliferation. Analysis of cytokine secretion from the splenocytes was consistent with the above results. While no antigen-specific IL-4 secretion was detected for all vaccinated animals, HCV-antigen-specific INF-γ secretion was detected for the splenocytes of vaccinated animals. CpG adjuvant enhanced the secretion of INF-γ but did not change the profile of IL-4 secretion.

CONCLUSION: Vaccination of mice with plasmids encoding HCV E1 and E2 antigens induces humoral and cellular immune responses. CpG adjuvant significantly enhances the cellular immune response.

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INTRODUCTION

Hepatitis C virus (HCV) infection is a worldwide health problem[1]. Up to now, no effective medical treatment is available for the majority of HCV infected patients[2,3], while new infections are continuously emerging from blood transfusion, needle sharing, unprotected sex, close contact of HCV infected patient and other unidentified sources[4]. Thus to control the spread of HCV by vaccination becomes an urgent task, especially in developing countries including China, where there is a large infected population.

Various routes were taken to develop a vaccine against HCV infection. Recombinant HCV antigens purified from E.coli, yeast or insect cells could not protect the vaccinee from virus challenge[5], possibly due to the fact that post-translational modification of the virus antigens in mammalian cells is greatly different from that in the bacterial or yeast cells. So the study on vaccine development has been focused on mammalian systems. Although many efforts have been made in vaccine development and some encouraging results have been obtained[6-8], no effective vaccine is available.

The RNA genome of HCV has a high mutation rate, which explains the existence of many genotypes, subtypes and quasispecies[9-10]. Although there is still a possibility to develop a universal vaccine against HCV of all genotypes, the use of local HCV strain in vaccine study is preferred. Our laboratory has been focusing on DNA vaccine development[11] using an HCV strain isolated from a patient in northern China[12], since DNA vaccine is effective in eliciting cellular immune responses[13] which play key roles in viral clearance.

Envelope proteins were the first choice in the development of vaccines against virus infection. HCV envelope protein 2 has become the major target in HCV vaccine development not only because it is the putative major envelope protein[14], but also because E2 could mediate the binding of HCV particle to host cells[15] and is the ligand of the possible HCV receptor CD81 on the host cell surface[16]. It makes E2 an attractive choice for vaccine development as it has been shown to be a major target of immune response in HCV infected patients[17,18]. Previous studies also revealed multiple neutralizing epitopes in E2 proteins[19-22]. Another envelope protein of HCV, E1, is also desirable for vaccine development, since its possible role in maintaining the natural conformation of E2 protein (through E1-E2 complex)[23] and the possibility of its cooperation with E2 in mediating host cell binding and entry[24]. However, low expression levels of both E1 and E2 proteins were observed when they were expressed in a single open reading frame (data not shown). In this study, we expressed HCV E1 and E2 proteins separately and analyzed the immune responses of DNA vaccination delivered by two plasmids encoding HCV E1 and E2 proteins respectively.

Recent studies showed that the immune system responded to CpG motifs by activating potent Th1-like immune responses[25]. There have also been several reports demonstrating that CpG could enhance humoral immune responses elicited by DNA vaccination[24,25], possibly due to indirect effect of enhanced cellular immune responses. The effect of CpG adjuvant on the immune response elicited by the candidate DNA vaccine was also investigated in this study.

MATERIALS AND METHODS

Plasmids, CpG oligodeoxynucleotide (ODN) and cells

For the construction of pSecTagB/sE2, the fragment of HCV
E2 (aa. 384-661) was amplified using plasmid pUC18/E[12] as template with following primers: sense, 5’GGGCTTAAGCTTAA CACCTACGTG3′ (HindIII site underlined); antisense, 5’CAG GAATTTCTACTGATCATCT3′ (EcoRI site underlined). Insertion of the PCR product into pSecTagB (Invitrogen, California) resulted in pSecTagB/E2, in which a secretion signal was provided at the N-terminal of E2 sequence. The recombinant plasmid was sequenced before further experiments.

HCV E1 encoding plasmid pSec-preS1-E1t340 (pSecTagB/sE1) was constructed as previously described[26]. CpG ODN1826 with phosphorothioate backbone was synthesized in Promega (Shanghai) according to the published sequence 5’TCCATGACGTTCCTGACGTT3′[27]. The cell line used for transient expression of the recombinant plasmids, BHK-21, was maintained in DMEM supplemented with 50 mL/L FCS and antibiotics under 50 µg/mL CO2 in a humidified 37 °C incubator.

**Transgenic expression**

Transfection of BHK-21 cells with HCV-encoding plasmid or empty vector using lipofectAMINE (Invitrogen) was done according to the manuals of the manufacturer. Cells were harvested 48 h after transfection, lysed in SDS-PAGE loading buffer, and subjected to SDS-PAGE (10%). The resolved samples were then transferred onto nitrocellulose membrane and probed with polyclonal anti-E1 (Liu et al., unpublished data) or polyclonal anti-E2 antiserum[28]. The signals were visualized with SuperSignal West Pico stable peroxide solution (Pierce, USA).

**DNA immunization and sera preparation**

Four groups of 5 female BALB/c mice (6-8 weeks old) purchased from the Shanghai Laboratory Animal Center were injected 3 times in quadriceps muscles: group 1, with 100 µg pSecTagB in 100 µL PBS; group 2, with 50 µg pSecTagB/sS1E1 and 50 µg pSecTagB/sE2 in a total volume of 100 µL PBS; group 3, with 100 µg pSecTagB together with 10 µg CpG in a total volume of 100 µL PBS; group 4, with 50 µg pSecTagB/sS1E1, 50 µg pSecTagB/sE1 and 10 µg CpG in a total volume of 100 µL PBS. Injections were performed at 0, 4 and 8 wk and blood samples were then transferred onto nitrocellulose membrane and probed with polyclonal anti-E1 (Liu et al., unpublished data) or polyclonal anti-E2 antiserum[28]. The signals were visualized with SuperSignal West Pico stable peroxide solution (Pierce, USA).

**Anti-HCV antibody analysis**

Enzyme-linked immunosorbent assay (ELISA) was used to determine the presence of antibodies against HCV E1 and E2 antigens in serum samples. E. coli-expressed HCV E1 (aa 192 to 315) or HCV E2 (aa 450 to 565[31]) was used to coat 96-well plates (MaxiSorp Surface, Nunc) at the concentration of 1 µg/mL. The antigens were suspended in PBS with 0.2 g/L sodium azide and incubated overnight at 4 °C. After the plates were washed with PBS plus 0.5 g/L Tween 20 (PBS-T) and blocked in blocking buffer (50 µL fat-free milk powder in PBS-T), twofold serial dilutions of serum samples in blocking buffer were added and incubated for 2 h at 37 °C. After three wash steps with PBS-T, horseradish peroxidase (HRP) conjugated goat-anti-mouse IgG (Dako, Denmark; 1000-fold diluted in blocking buffer) was added and incubated for 1 h at 37 °C. The plate was then developed with substrate buffer (50 mmol/L Na2HPO4, 25 mmol/L citric acid, 75 µg/mL 3,3′,5,5′-tetramethylbenzidine, 0.15 mL/L H2O2). After 30 min of incubation at room temperature, the reaction was stopped by adding 0.5 mol/L H2SO4, and absorbance was measured at 450 nm on a microplate reader (Model 450, Bio-Rad Laboratories). Antibody titers were calculated as the highest dilution which gave a positive reading. The cutoff value was set as mean absorbency (A) of sera from the control mice vaccinated with non-recombinant plasmid multiplied by 2.

**Preparation of splenocytes**

Mice were sacrificed by cervical dislocation. Spleens from these mice were ground on metal mesh to prepare single-cell suspension in grinding media (RPMI1640 supplemented with 100 mL/L FCS, 1 mmol/L Sodium pyruvate and 50 µmol/L β-ME). Red blood cells were lysed by incubating the splenocyte preparations with lysis buffer (0.15 mol/L NH4Cl, 10 mmol/L KHCO3, 0.1 mmol/L EDTA, pH 7.2-7.4) briefly and then washed with grinding media. The cells were resuspended in a small volume of grinding media and counted in the presence of Trypan blue. Splenocytes were immediately used for further experiments.

**Assay for HCV antigen specific splenocyte proliferation**

Splenocytes from vaccinated mice or naïve mice were diluted to 4×107/mL with grinding media and plated onto 24-well-plates at 2 well/well. The cells were in vitro stimulated with 500 ng E2 (aa 450 to 565[31]) or mock stimulated for 3 d. The media were collected for cytokine assay. Fresh grinding medium with 0.5 µCi/mL [3H] thymidine (Amersham Pharmacia Biotech) was added for another 24 h. Cells were then washed with PBS and 100 g/L trichloroacetic acid (TCA), incubated with 100 g/L TCA for 10 min at 37 °C. Afterwards, TCA was removed and cells were lysed with lysis buffer containing 0.33 mol/L NaOH and 10 g/L sodium dodecyl sulfate (SDS). [3H] thymidine incorporation in the cell lysates was measured by liquid scintillation counting. Antigen specific proliferation was presented by stimulation index (SI): SI = cpm of the cells stimulated by antigen/cpm of mock stimulated cells. Splenocytes from naïve mice were used to monitor the specificity of this assay.

**Analysis of HCV antigen specific cytokine secretion**

Media from the in vitro stimulated splenocytes were collected and preclarified by centrifugation at 10 000 g and 4 °C for 10 min. Concentration of INF-γ and IL-4 in the media was determined by ELISA using the mouse cytokine assay kits (Jingmei Biotech, Shenzhen) according to the protocols provided by the manufacturer. HCV E2 specific cytokine secretion was represented by stimulation index (SI): SI = cytokine secreted upon E2 stimulation/cytokine secreted upon mock stimulation. Splenocytes from naïve mice were used to monitor the specificity of this assay.

**Statistical analysis**

Statistical analysis was performed with unpaired 2-sided Student t test. Differences with P values <0.05 were considered significant.

**RESULTS**

**Transient expression of HCV antigens in mammalian cells**

Because the expression level was low when E1 and E2 were expressed in one open reading frame (data not shown), we expressed HCV E1 and E2 genes in separate plasmids in this study. To achieve high-level expression and proper post-translational modification of HCV envelope proteins, the pSecTagB vector with an efficient secretion signal of IgG molecules was chosen. Plasmid expressing HCV E2 was constructed as described in Materials and Methods. C-terminal hydrophobic sequence of E2 was truncated to facilitate the secretion of E2 and to obtain complex-type glycosylation modification, which was presented on the surface of HCV particles[29]. Secreted E2 protein was shown to have better antigenicity, possibly due to its proper modification by Golgi enzymes[30]. Plasmid pSecTagB/sE1 was taken as another component of the candidate DNA vaccine since high-level expression of E1 with this plasmid was observed previously in transiently and stably transfected NIH3T3 cells[28]. Before vaccination experiment, BHK-21 cells were transfected with pSecTagB/sE1 and pSecTagB/sE2, respectively, to check if they
would properly express the target HCV proteins in this cell line. By Western blot, both E1 and E2 were detected as glycosylated proteins with MW higher than those of polypeptide backbones, respectively (Figure 1). Secreted products were also detected for E1 and E2 (data not shown).

Humoral immune responses after DNA vaccination
DNA vaccination using the above characterized plasmids was carried out as described in Materials and Methods. After two injections, HCV E1 and E2 specific antibodies were detected in the sera of several mice. Without CpG adjuvant, the seroconversion rate was 2/5 for anti-E1 antibody, and 1/5 for anti-E2 antibody. When CpG was included as an adjuvant, the seroconversion rate was 3/4 for anti-E1 antibody and 2/4 for anti-E2 antibody. After the third injection, all animals became seroconverted to both anti-E1 and anti-E2 antibodies. The highest anti-E1 titer was 1:320 after the third injection for mice receiving plasmids only, while the highest titer for mice receiving plasmids with CpG was 80. The highest anti-E2 titers for both groups reached 1:280, but the average anti-E2 titer for mice receiving CpG was slightly lower than that for those receiving no CpG (Table 1).

E2 specific splenocyte proliferation
All the animals were sacrificed 30 d after the last injection to analyze cellular immune responses of the memory phase. Single-cell suspension of splenocytes was prepared for each individual animal. Splenocytes were immediately cultured in the presence of HCV E1 peptide or E2 protein. [3H] thymidine was then added to the cells to measure HCV antigen specific proliferation. In our experiments, HCV E1 specific proliferation was not observed, neither was E1 specific cytokine secretion (data not shown), possibly due to insufficient stimulation. For E2 specific splenocyte proliferation, as shown in Figure 2, animals vaccinated with pSecTagB only or pSecTagB plus CpG had stimulation indexes similar to that of naïve mice. Some of the animals vaccinated with plasmids pSecTagB/sE1+pSecTagB/sE2 showed E2 specific splenocyte proliferation while the others did not. This gave an average SI slightly higher than that of pSecTagB vaccinated mice. Animals vaccinated with HCV gene encoding plasmids together with CpG all showed E2 specific splenocyte proliferation, which resulted in a significant difference (P = 0.005) of SI between animals injected with pSecTagB/sE1+pSecTagB/sE2+CpG and animals injected with pSecTagB+CpG.

Table 1 Anti-HCV titers1 after the third injection

| Vaccine                | Anti-E1 titers          |
|------------------------|-------------------------|
|                        | 1           | 2           | 3           | 4           | 5           |
| pSecTagB/sE1+pSecTagB/sE2 | 320      | 40          | Dead animal | 320        | 320        |
| pSecTagB/sE1+pSecTagB/sE2+CpG | 80       | 80          | Dead animal | 80         | 80         |
| pSecTagB/sE1+pSecTagB/sE2 | 1        | 2           | 3           | 4           | 5           |
| pSecTagB/sE1+pSecTagB/sE2+CpG | 160      | 160         | Dead animal | 1280        | 1280       |

1Titers were determined as described in Materials and Methods. Serial dilution of anti-sera started from 1:40, thus titers lower than 40 were not determined and considered 0. 2The cutoff value was established as mean A of sera from the control mice vaccinated with pSecTagB. 3The cutoff value was established as mean A of sera from the control mice vaccinated with pSecTagB+CpG.
genomes. Recent studies have shown that the immune DNAs but are heavily methylated and suppressed in vertebrate. These CpG motifs are prevalent in bacterial and many viral. It was first found to be immunoactive in activating B cells[38]. Th1-like responses might be more important in evaluating a candidate against HCV infection was positively correlated with cellular expression. Studies of HCV patients showed that protection which could activate CTL to remove cells with HCV antigen. Further analysis revealed that the cellular immune response is Th1 type, antigen specific splenocyte proliferation indicated that our candidate DNA vaccine was able to induce cellular immune responses makes it a favorable choice in the development of prophylactic and therapeutic vaccine against HCV. CpG motif, that is, CpG dinucleotide in particular base context (XCGY, where X is any base but C, and Y is any base but G), was first found to be immunooactive in activating B cells[38]. These CpG motifs are prevalent in bacterial and many viral DNAs but are heavily methylated and suppressed in vertebrate genomes[38,39]. Recent studies have shown that the immune system could respond to CpG motifs by activating potent Th1-like immune responses[23]. Several reports demonstrated that CpG could enhance humoral immune responses elicited by DNA vaccination[24,25], possibly due to indirect effect of enhanced cellular immune responses. In our vaccination experiment, we observed that after the second injection, the seroconversion rate for mice receiving CpG adjuvant was higher than that for those not receiving CpG. This enhanced humoral immune response at the early stage was likely due to the indirect positive effect of CpG on antigen presentation (via activation of antigen presenting cells by Th1). However, after the third injection, when all animals became seroconverted, those who received CpG as an adjuvant showed lower titers of both anti-E1 and anti-E2. This was consistent with the notion that when Th1-like response is the major immune response, Th2-like response is usually inhibited, followed by decreased antibody production. Further analysis of antigen specific cytokine secretion was consistent with this explanation. E2 specific INF-γ secretion was enhanced in splenocytes from those animals receiving CpG in vaccination, while this adjuvant did not change the E2 specific IL-4 secretion profile. These results indicated that CpG enhanced the Th1-like cellular immune response.

Apart from DNA vaccine, live virus vaccine and peptide vaccine are also efficient in inducing cellular immune response. Live virus vaccine using several different viral vectors has been under investigation[22,28,40], with continuous promising progresses. However, the interest in peptide vaccine is diminishing. The problem lies in the polymorphism of MHC molecules in the population. Certain individuals may have the MHC molecules necessary to bind a specific antigenic peptide, while others may not. Thus, the universality of a vaccine based on a specific peptide becomes a concern. Recently, combinational vaccination strategy with different species of vaccine components has been successful in laboratory in protecting against some elusive infectious reagents[41,42]. Combinational vaccination could reduce host immune responses against the viral vector itself, thus enhancing the efficiency of delivery. More importantly, combinational vaccination with different components could activate different components of the host immune surveillance system and result in enhanced immune responses. Our next effort will be focused on combinational vaccination with DNA vaccine described above as one component.

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**DISCUSSION**

In this study, target HCV antigens encoded by the DNA vaccine candidate, E1 and E2, were delivered by two separate plasmids that achieved higher-level expression and appropriate glycosylation of both antigens. DNA immunization results demonstrated that this DNA vaccine candidate could induce HCV specific humoral and cellular immune responses in mice. Anti-E2 titer obtained in this study was similar to our previous[31-35]. We also detected comparable levels of anti-E1 antibodies in the sera of vaccinated mice, which together with anti-E2 antibodies, might be important in neutralizing the circulating HCV virus and preventing its spread in patients. HCV antigen specific splenocyte proliferation indicated that our candidate DNA vaccine was able to induce cellular immune response, which was critical in viral clearance. Cytokine secretion analysis revealed that the cellular immune response is Th1 type, which could activate CTL to remove cells with HCV antigen expression. Studies of HCV patients showed that protection against HCV infection was positively correlated with cellular immune responses[36,37], indicating that cellular immune responses might be more important in evaluating a candidate vaccine. The ability of our candidate vaccine to induce HCV specific cellular immune responses makes it a favorable choice in the development of prophylactic and therapeutic vaccine against HCV.

**Figure 3** E2-specific cytokine secretion. Error bars represent the standard errors. Asterisk indicates the very significant difference of the average SI between animals injected with (pSecTagB/sE1+pSecTagB/sE2+CpG) and animals injected with (pSecTagB+B+CpG) in terms of INF-γ secretion (P = 0.0003).
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