An Efficient *Schistosoma japonicum* Bivalent Membrane Protein Antigen DNA Vaccine Against Schistosomiasis in Mice

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Source of support:  This work was supported by the National Natural Science Foundation of China (#30901251), Grants for Scientific Research of BSKY from Anhui Medical University (XJ201321), and the Natural Science Foundation of Anhui Province of China (KJ2012A174 and KJ2017A691)

**Background:** Schistosomiasis is one of the most important infectious parasitic diseases in the world. The most important was to control schistosomiasis is through a combination of medical therapy and immunization. The membrane antigens Tsp2 and 29 from *Schistosoma* are promising anti-schistosomiasis vaccine candidates.

**Material/Methods:** In this study, the pcDNA3.1(+)-SjTsp2, pcDNA3.1(+)-Sj29, and pcDNA3.1 (+)-SjTsp2-29 eukaryotic expression vectors were successfully constructed as DNA vaccines, and the protective abilities of these vaccines were evaluated in mice.

**Results:** The results showed that vaccination with SjTsp2, Sj29, and SjTsp2-29 reduced parasite burden and hepatic pathology compared to the control group, and the protective effect of the bivalent SjTsp2-29 DNA vaccine was better than that of the univalent SjTsp2 or Sj29 DNA vaccines. We also found high levels of IgG, IgG1, and IgG2a against SjTsp2, Sj29, and SjTsp2-29 DNA vaccines, with high expression of IFN-γ and no IL-4 in the mice.

**Conclusions:** The double-membrane antigen DNA vaccine SjTsp2-29 elicited protection against *Schistosoma* infection and might serve as a vaccine candidate.

**MeSH Keywords:** DNA • *Schistosoma japonicum* • Vaccine Potency

Full-text PDF:  [https://www.medscimonit.com/abstract/index/idArt/919195](https://www.medscimonit.com/abstract/index/idArt/919195)
**Background**

Schistosomiasis, one of the most important infectious diseases in humans, is caused by infection with 3 important Schistosome species: *Schistosoma haematobium*, *Schistosoma mansoni*, and *Schistosoma japonicum* [1,2]. There are estimated to be about 77 000 schistosomiasis patients in China and about 30 800 advanced schistosomiasis cases were diagnosed in 2015 [3]. Although praziquantel is a widely-used, high-efficiency, broad-spectrum oral antiparasitic drug, chemotherapy does not prevent reinfection or transmission of schistosomiasis in highly endemic areas [4].

Additionally, the repeated and large-scale use of praziquantel in epidemic areas has led to drug resistance and a reduced cure rate [4–6]. The best long-term strategies are intermediate host management, environmental and water sanitation, good personal hygiene maintenance, and vaccines [7–9]. Although the development of vaccines against schistosomiasis has had more setbacks than successes, some inspiring results have emerged recently based on the use of membrane protein antigens of Schistosoma [10–14].

The *Schistosome* tegument is involved in nutrition, immune evasion and modulation, excretion, osmoregulation, sensory reception, and signal transduction, and membrane proteins of parasite are vulnerable to attack by the host immune system [15–17]. Thus, the tegument is crucial to the parasite and is a target for vaccines and drugs. The membrane antigens of *S. mansoni*, the extracellular loop 2 of a tetraspanin (TSP-2) and 29, are promising anti-schistosomiasis vaccine candidates that have approximately 50% protection rate in mice infected with *S. mansoni* cercariae [1112]. The membrane antigen of *S. japonicum*, Sj-TSP-2, has been confirmed as an effective anti-schistosomiasis vaccine in mice [18,19]. The Sj29 protein also has been identified and confirmed to be a membrane protein by our group [20]. The use of multivalent vaccines is a promising method that induces greater protection than with the use of a single antigen [21,22].

Thus, we developed a high-performance bivalent membrane DNA vaccine including the membrane proteins SjTsp2 and Sj29 of *S. japonicum*. Based on the constructions of the pcDNA3.1(+)-SjTsp2 and pcDNA3.1(+)–Sj29 groups, the SjTsp2 and Sj29 fusion gene connected with (Gly-Ser)4 was amplified by SOEing PCR and cloned into the pcDNA3.1(+) vector to construct a pcDNA3.1(+)-SjTsp2-29 DNA vaccine construct. The pcDNA3.1(+)–SjTsp2 DNA vaccine construct was modified from pET28a-SjTsp2, which contains a larger extracellular loop of TSP-2 from *S. japonicum*. Larger extracellular loop of Sj-TSP-2 gene products were digested with pET28a-SjTsp2 with BamHI/XhoI, and then subcloned into the pcDNA3.1(+) vector. Extracellular region of Sj29 gene was digested pET28a-SjTsp2 with BamHI/XhoI, and then subcloned into the pcDNA3.1(+) vector to construct pcDNA3.1(+)-Sj29 DNA vaccine. Based on the constructions of the pcDNA3.1(+)–SjTsp2 and pcDNA3.1(+)–Sj29 groups, the SjTsp2 and Sj29 fusion gene connected with (Gly-Ser)4, was amplified by SOEing PCR and cloned into the pcDNA3.1(+) with BamHI/XhoI. The fusion SjTsp2 and Sj29 genes connected by (Gly-Ser)4, were digested pET28a- SjTsp2-29 with BamHI/XhoI, and then subcloned into the pcDNA3.1(+) vector to construct the pcDNA3.1(+)-SjTsp2-29 DNA vaccine. The 3 constructs were confirmed by restriction enzyme analyses and sequencing, and a large number of endotoxin-free plasmids were then extracted in accordance with the instructions of the plasmid extraction kit (CW Biotech, Beijing, China).

**Material and Methods**

**Parasites and animals**

*Oncomelania hupensis* snails infected with a mainland Chinese strain of *S. japonicum* were provided from the Jiangsu Provincial Institute of Parasitic Diseases. Female Kunming mice (6 weeks old) were purchased from the Animal Centre of Anhui Medical University and used for the vaccine trials.

**Ethics approval**

The mice received care that complied with the Guidelines for the Care and Use of Laboratory Animals. All of the work presented here was approved by the Anhui Experimental Animal Training Base (reference number LLSC20140061).

**Preparation of DNA vaccines**

The pET28a-SjTsp2, pET28a-Sj29, and pET28a- SjTsp2-29 recombinant plasmids were constructed. The pcDNA3.1(+)–SjTsp2 DNA vaccine construct was modified from pET28a-SjTsp2, which contains a larger extracellular loop of TSP-2 from *S. japonicum*. Larger extracellular loop of Sj-TSP-2 gene products were digested with pET28a-SjTsp2 with BamHI/XhoI, and then subcloned into the pcDNA3.1(+) vector. Extracellular region of Sj29 gene was digested pET28a-SjTsp2 with BamHI/XhoI, and then subcloned into the pcDNA3.1(+) vector to construct pcDNA3.1(+)-Sj29 DNA vaccine. Based on the constructions of the pcDNA3.1(+)–SjTsp2 and pcDNA3.1(+)–Sj29 groups, the SjTsp2 and Sj29 fusion gene connected with (Gly-Ser)4, was amplified by SOEing PCR and cloned into the pcDNA3.1(+) with BamHI/XhoI. The fusion SjTsp2 and Sj29 genes connected by (Gly-Ser)4, were digested pET28a- SjTsp2-29 with BamHI/XhoI, and then subcloned into the pcDNA3.1(+) vector to construct the pcDNA3.1(+)-SjTsp2-29 DNA vaccine. The 3 constructs were confirmed by restriction enzyme analyses and sequencing, and a large number of endotoxin-free plasmids were then extracted in accordance with the instructions of the plasmid extraction kit (CW Biotech, Beijing, China).

**Administration of the DNA vaccine**

Fifty 6-week-old female Kunming mice were randomly divided into 5 groups of 10 mice each: a normal saline (NS) group, a pcDNA3.1(+) plasmid group, a pcDNA3.1(+)–SjTsp2 group, a pcDNA3.1(+)–Sj29 group, and a pcDNA3.1(+)–SjTsp2–29 group. To enhance the uptake of foreign DNA vaccine, all mice were injected with 30 μl bupivacaine hydrochloride (7.5 mg/ml) (Harvest Pharmaceutical Co., Shanghai, China) in the right tibialis anterior muscle 24 h before immunization. Each group was immunized with 3 intramuscular injections of 75 μg DNA vaccine 3 times biweekly, and the control group was injected with 100 μl normal saline.
Challenge infection and worm and egg burden recovery

Two weeks after the last immunization, the 30 infected Oncomelania were exposed to sunlight for 4 h at 25–28°C and cercariae were shed in a beaker. Mice were infected with 24±2 S. japonicum cercariae by percutaneous infection. At 6 weeks after infection, the worm reduction rates and the egg reduction rates in the livers were calculated. The worms were collected and counted by perfusion from the hepatic portal system, and the eggs were also collected. The weight of liver tissues was measured before being homogenized in 50 ml PBS. We mixed 5 ml homogenate with 1 ml 10% NaOH and incubated it at 56°C for 1 h. The average of 3 counts per 100 µl mixture was considered as the number of eggs in each sample tested, and this was converted to EPG (eggs per gram). The rate of reduction in the worm and egg counts was calculated as follows: percentage reduction in worm burden = (mean worm burden of control group–mean worm burden of vaccinated group)/mean worm burden of control group×100%; percentage reduction in liver egg count = (mean EPG from control group–mean EPG from vaccinated group)/mean EPG from control group×100%.

Detection of antibodies and cytokines levels by enzyme-linked immunosorbent assay (ELISA)

Before the mice were infected with S. japonicum, the sera were collected. The anti-AWA IgG, IgG1, and IgG2 antibody levels in the serum samples were detected by ELISA. Each well of a plate was coated with 1 µg AWA (adult worm antigen). SJAWA was prepared as described previously by Abán [23]. Plates were blocked by 200 µl/well of 2.5% BSA (bovine serum albumin) in 0.02 M PBS with 0.05% Tween 20 (PBST) (pH 7.2). After the serum samples were diluted 100 times with PBST, 100 µl of the diluted sera was added to each well and incubated for 1 h. Finally, the plates were incubated with peroxidase-conjugated goat anti-mouse IgG, IgG1, and IgG2a (Sigma) diluted to 1: 10 000, 1: 5000, and 1: 5000, respectively, for 1 h at room temperature (RT). The reaction was visualized by addition of 200 µl/well of 3,3’, 5,5’-Tetramethylbenzidine substrate, and then was stopped by adding 50 µl 2 M H₂SO₄. The optical density (OD) was measured at 450 nm.

To evaluate the cytokine profile generated by the immunization of mice with DNA vaccine, the sera were collected before the mice were infected with S. japonicum. We detected the levels of a panel of cytokines (IFN-γ and IL-4) in the sera, and mouse IFN-γ and IL-4 ELISA kits (eBioscience, USA) were used according to the manufacturer’s instructions.

Histology and immunohistochemistry analysis

To detect the DNA vaccine expression in vivo, 3 days after the third immunization, 1 mouse was taken from each group, then the injected muscle tissues were removed, fixed in 10% formalin, embedded in paraffin, and sectioned for histology. Another mouse from the pcDNA3.1(+) group was chosen randomly as a control. The muscle tissue sections were incubated for 30 min with DNA vaccine immunized serum (dilution 1: 1000) for immunohistochemistry. The immunohistochemistry tests were conducted according to the instructions of the Power Vision Two-Step detection system (Zhongshan Biotechnology Co., Beijing, China), which was used to view the muscle tissues at 20× image magnification. At 6 weeks after infection, all mice were sacrificed, livers were excised and immediately fixed in 10% formalin in PBS, and embedded in paraffin. The liver tissue sections (4 µm) were stained with hematoxylin and eosin (HE) and examined for quantitative and qualitative changes. The granuloma areas were measured with computer-assisted morphometric software (Image-Pro Plus) to calculate the granuloma areas as percentages of the total areas for each side. At least 3 discontinuous slides were measured for each specimen, and the mean values of 8 mice were taken for statistical analysis [24].

Statistical analyses

The results were analyzed with one-way ANOVA with post hoc testing using the least significant difference test. All data are shown as the mean±the standard error of the mean (SEM). P values <0.05 were considered statistically significant.

Results

Construction of the pcDNA3.1(+)–SjTsp2, pcDNA3.1(+)–Sj29, and pcDNA3.1(+)–SjTsp2–Sj29 DNA vaccine

pcDNA3.1(+)–SjTsp2–Sj29, pcDNA3.1(+)–SjTsp2, and pcDNA3.1(+)–Sj29 were digested with BamHI and Xhol. The sizes of digestive products were consistent with those expected: 660 bp, 225 bp, and 390 bp, respectively. According to the agarose gel results (Figure 1A) and sequencing results (Figure 1B), SjTsp2, Sj29, and SjTsp2–Sj29 were successfully incorporated into the backbone vector pcDNA3.1(+). Therefore, the pcDNA3.1(+)–SjTsp2, pcDNA3.1(+)–Sj29, and pcDNA3.1(+)–SjTsp2–Sj29 DNA vaccines were successfully constructed.

Immunohistochemistry analyses of the injected muscle tissues

Immunohistochemical analyses of the injected muscles were performed to identify the expression of the DNA vaccines in muscle following infection. The results revealed that immunostaining of the injected muscle for the DNA vaccine antibody was positive, whereas immunostaining of the pcDNA3.1(+) and NS-injected muscles was negative (Figure 2).
Antibody response to DNA vaccination

To assess antigen-specific anti-AWA antibodies production, the sera were collected before the infection of mice by *S. japonicum*. ELISA was performed to detect the levels of IgG and IgG subtypes in immunized mouse sera using AWA as antigen. High total IgG titers were recorded for all vaccinated groups relative to the infected control groups that were vaccinated with pcDNA3.1(+) or NS (P<0.05). pcDNA3.1(+)−SjTsp2-29 induced higher levels of IgG2a compared to pcDNA3.1(+)−SjTSP-2 or pcDNA3.1(+)−Sj29 alone (Figure 3).

**Figure 1.** Identification of recombinant plasmids pcDNA3.1(+)-SjTsp2, pcDNA3.1(+)-Sj29, and pcDNA3.1(+)-SjTsp2-29. (A) Identification with restriction enzyme digestion. pcDNA3.1(+)-SjTsp2-29, pcDNA3.1(+)-SjTsp2, and pcDNA3.1(+)-Sj29 were digested with BamHI and Xhol, and the sizes of digestive products were consistent with those expected (i.e., 660, 225, and 390bp, respectively). Lane M: DNA marker; lane 1: pcDNA3.1(+)-SjTsp2-29; lane 2: pcDNA3.1(+)-SjTsp2; lane 3: pcDNA3.1(+)-Sj29. (B) SjTsp2-29 nucleotide sequence alignment with Sj-TSP-2 and Sj29. a: -SjTsp2; b: (Gly4Ser)3 linker; c: Sj29

**Figure 2.** Gene expressions in the muscle tissues of mice injected with the DNA vaccines. (A) NS, (B) pcDNA3.1(+), (C) pcDNA3.1(+)-SjTsp2, (D) pcDNA3.1(+)-Sj29, (E) pcDNA3.1(+)-SjTsp2-29.

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Cytokine profile

To evaluate the cytokine profile generated by the immunization of mice with DNA vaccine, we measured the levels of cytokines (IFN-γ and IL-4) in mouse sera. Significantly higher levels of IFN-γ were detected in sera of DNA-vaccinated mice compared to the control group (Figure 4). However, lower levels of IL-4, a signature of Th2-immune response, were detected.

DNA vaccine induces immune protection against S. japonicum

To determine the protective efficacies of the pcDNA3.1(+)–SjTsp2, pcDNA3.1(+)–Sj29, and pcDNA3.1(+)–SjTsp2-29 DNA vaccines, the infected mice were sacrificed 42 days after infection to measure the adult worm and egg reduction rates in the liver. As shown in Table 1, the mice that were injected with pcDNA3.1(+)–SjTsp2 or pcDNA3.1(+)–Sj29 exhibited significantly greater reduction in worm burden and liver eggs compared to the control group (P<0.05). Most interestingly, the pcDNA3.1(+)–SjTsp2-29 group exhibited significantly greater reduction in the worm burden and liver eggs compared to...
the pcDNA3.1(+)-SjTsp2 and pcDNA3.1(+)-Sj29 groups (P<0.05). The pcDNA3.1(+)-SjTsp2-29 group exhibited a worm reduction rate of 53.2% and an egg reduction rate of 51.4%.

DNA vaccination affects the formation of *S. japonicum* egg-induced liver granulomas

To investigate the effect of the DNA vaccines on *S. japonicum* egg-induced liver granulomas, liver tissues were fixed and stained with HE. Strikingly, compared to the control mice, the liver granulomas were attenuated in the groups that were injected with the pcDNA3.1(+)-SjTsp2, pcDNA3.1(+)-Sj29, or pcDNA3.1(+)-SjTsp2-29 DNA vaccines. pcDNA3.1(+)-SjTsp2-29 caused a significantly greater granuloma reduction in the liver compared to pcDNA3.1(+)-SjTsp2 and pcDNA3.1(+)-Sj29 (P<0.05) (Figure 5).

**Discussion**

Some schistosomiasis vaccines (e.g., Sh28GST, SmTSP2, and Sm14) are already in early human clinical trials [7,25,26], but

| Group                  | Adult worms mean±SD | Worm reduction rate (%) | Liver eggs mean±SD | Liver eggs reduction rate (%) |
|------------------------|---------------------|-------------------------|-------------------|-------------------------------|
| NS                     | 15.8±3.9            | 7.0                     | 30942±4123        |                               |
| pcDNA3.1(+)            | 14.7±4.1            | 7.0                     | 28131±3056        | 9.1                           |
| pcDNA3.1(+)–SjTsp2     | 10.2±3.1            | 35.4*                   | 19305±1978        | 37.6*                         |
| pcDNA3.1(+)–Sj29       | 11±3.9              | 30.4*                   | 21234±1927        | 31.4*                         |
| pcDNA3.1(+)–SjTsp2-29  | 7.4±2.1             | 53.2*                   | 15038±1801        | 51.4*                         |

Table 1. Parasitology data of mice vaccinated with pcDNA3.1(+)-SjTsp2, pcDNA3.1(+)-Sj29 and pcDNA3.1(+)-SjTsp2-29.

All data are expressed as the means±the SEMs (n=10 for each group). * P<0.05 versus the corresponding NS or pcDNA3.1(+) group, # P<0.05 versus the corresponding pcDNA3.1(+)–SjTsp2 or pcDNA3.1(+)–Sj29 group.
these trials are not yet finished. The failure to develop effective vaccines may be due to a variety of factors, including the fact that the current understanding of schistosomiasis immunity and immune mechanisms is largely dependent on studies in mice, but vaccines conducted in studies only in mouse models may have undesirable effects in human clinical trials. A vigorous humoral response (IgG, IgM, and IgE) is found in patients with acute and chronic schistosomiasis, which is different from that in mice [7,27,28]. Therefore, new Schistosoma vaccine molecules and vaccination are still needed for study.

Tetraspanins of Schistosoma mansoni have 4 transmembrane regions, which are connected by 2 extracellular loops, and the loops are speculated to interact with ligands [6]. The larger extracellular loop of Sm-TSP-2 has been confirmed to be valuable anti-schistosomiasis vaccine, which induces approximately 50% reduction in the worm burden [11]. Sm-TSP-2, which fuses to a thioredoxin partner, has been confirmed to be a more effective anti-schistosomiasis vaccine than Sm-TSP-2 alone [29]. Sj-TSP-2 is highly polymorphic [30,31] and therefore has limited utility as a vaccine. The Sj-TSP-2e protein is recognized by the sera of some patients who are infected with S. japonicum, but this protein does not protect mice against a S. japonicum challenge infection [30]. However, Yuan [19] used a subclass of the Sj-TSP-2 sequence to express recombinant Sj-TSP-2 protein as a vaccine, and observed significant efficacy in mice. Similarly, we found 3 subclasses of Sj-TSP and confirmed that the Sj-Tsp2-A DNA vaccine induced partial protective immunity against S. japonicum infection [18].

The Sm29 protein is strongly recognized by IgG1 and IgG3 antibodies of naturally resistant individuals, as well as by patients who are resistant to reinfection and live in areas that are endemic for schistosomiasis [12,32]. Sm29 is a promising candidate as a vaccine against schistosomiasis, and is confirmed to have about a 50% protection level against experimental S. mansoni infection [12]. Pinheiro et al. [33] confirmed that vaccination using Sm-TSP-2 linked with the terminus of Sm29 induced reductions in parasite burden and hepatic pathology relative to control; the resultant protection levels ranged from 27.84% to 34.83%. The combination of Sm29 and Sm14 induces significant protective immunity in mice infected with S. mansoni [34]. A vigorous humoral immune response can be induced by a vaccine formulation containing rSm29 adsorbed to alum, which successfully protects against S. mansoni reinfection in mice, indicating a potentially effective vaccine formulation that could be used in humans [35]. Sj29 is homologous to Sm29 of S. mansoni, which was characterized by our group as a membrane-bound antigen in S. japonicum [20].

S. japonicum is a multi-cellular parasite with complex life cycle stages and can evade the immune recognition system via mechanisms of antigenic variation. Furthermore, not all individuals have immunogenic responses to the identical antigens that occur in natural infections. The bivalent vaccine SjTsp2-29, which is composed of the Sj-TSP-2 and Sj29 schistosomal genes, can induce more broad-range protection against Schistosoma than can the univalent Sj-TSP-2 and Sj29 DNA vaccine. The method of using 2 promising vaccine molecules in combination merits further investigations for improving protection efficacy. In the present study, we used a flexible peptide (Gly5Ser), as a linker to connect the Sj-TSP-2 and Sj29 schistosomal genes to construct the DNA vaccine SjTsp2-29. Immunohistochemistry analyses revealed the expressions of specific antigens in the quadriiceps muscles of the Sj-TSP-2, Sj29, and SjTsp2-29 groups. The present study shows that it is possible to induce protection in a murine schistosomiasis model by DNA vaccines Sj-TSP-2, Sj29, and SjTsp2-29. Kunming mice immunized with DNA vaccines Sj-TSP-2, Sj29, and SjTsp2-29 revealed high levels of IgG1, IgG1, and IgG2a after the last immunization. However, the highest levels of IgG2a were induced after immunization with SjTsp2-29 compared to Sj-TSP-2 or Sj29 alone. The Th1-type of immune response was confirmed to be induced by DNA vaccines Sj-TSP-2, Sj29, and SjTsp2-29, which was shown by the production of IFN-γ and no significant IL-4 by cytokine analysis. Some researchers [31,36–38] have found that the Th1 type of immune response plays a more important role in providing protection against Schistosoma infection than does Th2. Moreover, these results were involved in increases in the expression of IFN-γ. The protective response mechanism of IFN-γ is macrophage activation to impede schistosomula migration and kill parasites in a nitric oxide-dependent approach [39,40]. Sm-TSP-2 and Sm29 are promising anti-schistosomiasis vaccine candidates that have been shown to elicit approximately 50% protection in mice [11,12]. However, S. japonicum differs from S. mansoni, and we found that the Sj-TSP-2 and Sj29 DNA vaccines achieved more than 30% protection and reduced the pathological inflammation in the mouse livers. The reduction rate of the granuloma areas elicited in the Sj-TSP-2 and Sj29 groups was approximately 30%. We found that the SjTsp2-29 DNA vaccine achieved greater than 50% protection and also reduced the pathological inflammation in mice. The reduction rate of the granuloma area in the SjTsp2-29 group was 50%. Thus, the protective effect of the bivalent DNA vaccine was better than that of the univalent DNA vaccines. We found that the bivalent DNA vaccine not only reduced worm and egg burden, but also reduced the pathological lesions caused by Schistosoma infection.

Conclusions

The findings of the present study suggest that the bivalent membrane SjTsp2-29 DNA vaccine might be a promising strategy for immunoprophylaxis against S. japonicum and might also provide a basis for the development of an effective vaccine against Schistosoma in the future.
Availability of data and materials

The datasets used and analyzed are available from the corresponding author on reasonable request. The results of analysis

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of the datasets are presented as tables, figures, and supplementary information files.