Immune Protective Evaluation Elicited by DNA Vaccination With Neospora caninum Dense Granules Proteins in Mice

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Neospora caninum, an obligate intracellular protozoan, is the major cause for neosporosis and brings serious economic losses to cattle breeding industries worldwide. After invasion, dense granules proteins are abundantly secreted and being important components of parasitophorous vacuole and intravacuolar network where N. caninum survives and replicates. The aim of the present study was to evaluate the protective immunity induced by DNA vaccines with genes encoding dense granules proteins 1 (GRA1), GRA4, GRA9, GRA14, GRA17, and GRA23 against N. caninum tachyzoites in BALB/C mice. Eukaryotic expressing plasmids of pcNcGRAs were constructed and the mice were intramuscularly immunized with pcNcGRAs followed by challenging infection with lethal doses of N. caninum. Immune responses were evaluated through monitoring the levels of serum antibodies, measurement of lymphocyte proliferation, and secretion of cytokines. Immune protection assays were carried out through monitoring survival time, body weight, and parasite burden in the brains. Results showed that all the pcNcGRA DNA vaccines could trigger remarkably specific humoral and cellular responses, with higher levels of IgG and IgG2a antibodies as well as obviously increased secretion of Th1-type IFN-γ cytokines. The immune protective efficacy revealed that pcNcGRA4, pcNcGRA14, and pcNcGRA17 DNA vaccines could individually increase the survival rate to 50, 37.5, and 25% in comparison with 0% in the control group; prolong the survival time more than 20.88 ± 11.12, 18.88 ± 10.83, and 16.63 ± 10.66 days compared with the control group of 4 ± 1.31 days; and decrease parasite burden in the brains to 297.63 ± 83.77, 471.5 ± 110.74, and 592.13 ± 102.2 parasites/100 ng comparing with 1221.36 ± 269.59 parasites/100 ng in the control group. These findings indicated that NcGRA4, NcGRA14, and NcGRA17 are potential vaccine candidates; NcGRA4 displayed better performance in immune protective efficacy and could be further combined with other advantageous antigens applied to the development of safe and effective DNA vaccines against N. caninum.

Keywords: Neospora caninum, DNA vaccines, dense granules proteins, immune responses, protective efficacy
INTRODUCTION

*Neospora caninum*, classified as an obligate intracellular protozoan parasite of phylum *Apicomplexa*, is the major pathogen for neosporosis in cattle and dog breeding industries (1–3). Susceptible animals become infected by ingestion of oocyst-contaminated foods or vertical transmission via placenta (4). *N. caninum*-infected dogs showed nervous system dysfunction and pregnant cows faced higher risks of reproductive failure, abortion, and neonatal mortality (5). According to statistics, *N. caninum* infection in cattle industries is causing dramatic negative economic losses ranging from US $43.08 billion to US $320.98 billion each year all over the world (6, 7). As one of the cattle-breeding countries, China is also facing challenges of continuous economic losses and paying much attention to neosporosis. Nowadays, sulfonamides are still widely used to ease neosporosis and control its spread. However, large negative effects and drug residues in animals are inevitable

Vaccines based on live, attenuated or killed parasites, and recombinant antigens are available for varieties of diseases through triggering immune responses. Both protective humoral and cellular immune responses involve in attenuating the pathogenicity of neosporosis. Substantial evidence shows that T helper (Th) 1-type immune response dominated by secretion of gamma interferon (IFN-γ), interleukin (IL)-12 cytokines, and production of IgG2a antibody is not only vital for acute *N. caninum* infections but is also an effective protection for chronic infections (8, 9). Also, Th2-type immune response triggered by antibody-mediated immunity is mainly functioned in combating *N. caninum* circulating in the blood or body fluids via activating complement system and limits the development of neosporosis (10, 11).

In *Toxoplasma gondii*, a closely related parasite to *N. caninum*, multiples DNA vaccines have already been developed and verified to be promising in the prevention of toxoplasmosis, such as dense granules proteins 1 (GRA1), GRA4, GRA7, and GRA14, for they could activate high levels of humoral and cellular protective immunity for relative long periods (12–14). For *N. caninum*, many studies have proposed potential antigens against neosporosis, such as surface antigens proteins 1 (SAG1) and SAG4 (15, 16); GRA1, GRA2, GRA6, and GRA7 (17–20); rhoptries proteins 2 (ROP2) and ROP40 (21–23); and micronemes proteins 1 (MIC1), MIC3, and MIC4 (24–26). However, research on DNA vaccines against neosporosis is still limited.

After invasion into host cells, *N. caninum* could secrete large amounts of GRA proteins, which are closely associated with its survival and intracellular replication (27). Many GRAs involved in the formation of parasitophorous vacuole and intravacuolar network structures, maintaining intracellular growing environment for tachyzoites. Moreover, GRAs are also found to exist in the cyst wall surrounding slowly dividing bradyzoites (7). Previous studies have verified that highly antigenic molecules of GRA2, GRA6, and GRA7 are effective vaccine candidates in *N. caninum* (20, 28, 29). In contrast, abundant GRA proteins secreted from *T. gondii* have been applied in immune protection and used as diagnostic indicators (30). Although many GRA proteins have been identified, few studies other than GRA2, GRA6, and GRA7 focus on studying its immune protection effects (31–35).

The present study aims to screen excellent DNA vaccine targets by evaluating humoral and cellular immune responses triggered by various reported *N. caninum* GRA proteins, including GRA1, GRA4, GRA9, GRA14, GRA17, and GRA23, through individual immunization with eukaryotic plasmids of pcNcGRA1, pcNcGRA4, pcNcGRA9, pcNcGRA14, pcNcGRA19, and pcNcGRA23 in BALB/c mice, and its immune protective efficacy after challenging with lethal doses of *N. caninum* tachyzoites. Our findings would lay a foundation for future research on providing new targets for DNA vaccine development.

MATERIALS AND METHODS

Ethics Statement
All the animal experiments were approved by the Animal Welfare and Research Ethics Committee of Jilin University (permission number: ppzp20190929065). Six- to eight-week-old female BALB/C mice were maintained with SPF-level food and water *ad libitum* under environment conditions of light and dark cycle every 12 h in rearing isolators. The mice were monitored every 12 h and deaths were recorded. To minimize suffering, mice were humanely euthanized by cervical dislocation after anesthetization by subcutaneous injection of atropine (0.02 mg/kg) and then inhalation of CO$_2$ (10–15 s) when they exhibited signs of illness and lost 20% body weight.

Parasites
The Nc-1 strain of *N. caninum* tachyzoites were propagated by continuous passage in monolayer bovine kidney epithelial (MDBK; ATCC, USA) cells, which were cultured in Dulbecco’s modified Eagle’s medium (Biological Industries, Israel) and supplemented with 10% fetal bovine serum (Biological Industries), 100 U/ml penicillin, and 100 μg/ml streptomycin at 37°C in 5% CO$_2$ conditions. To obtain enriched and purified tachyzoites for the following experiments, Nc-1 strain infected cells were collected at the stationary growth phase, released with a 27-gauge needle (Millipore, USA), purified with Percoll (Millipore), and enriched in the phosphate-buffered saline to a final concentration of 1 × 10$^8$ parasites/ml.

Preparation of *N. caninum* Tachyzoite Lysate Antigen
*N. caninum* tachyzoite lysate antigen (NLA) was prepared as follows: purified tachyzoite suspension was treated with Protease Inhibitor Cocktail (Sangon, Shanghai) and then lysed going through three cycles of freezing at −80°C and thawing at 4°C followed by ultrasonicaton on ice. Then, the mixture was centrifuged at 10,000 × g at 4°C for 30 min and supernatants were collected. After filtering through a 0.22-μm sterilized
membrane (Merck Millipore, USA), NLA was stored at −80°C for further use.

**Construction of Recombinant pcNcGRA Plasmids and Expression in vitro**

Total RNA was extracted from purified tachyzoites using RNAiso Plus (Monad, Wuhan), genomic DNA was removed, and cDNA was synthesized with the MoScript RTII Super Mix with dsDNase (Two-Step) (Monad) according to the instructions. Specific amplification primers of NcGRAs were designed and listed in Table 1. Restriction enzyme cutting sites of BamHI and EcoRI were individually introduced into the 5’ ends of forward and reverse primers of NcGRA1. Restriction enzyme cutting sites of EcoRI and XhoI were individually added to the 5’ ends of forward and reverse primers of NcGRA4, NcGRA14, NcGRA17, and NcGRA23. Restriction enzyme cutting sites of HindIII and BamHI were individually added to the 5’ ends of forward and reverse primers of NcGRA9. Meanwhile, sequences of HA tag (TACCCATACGATGTTCCAGATTACGCT) were added into reverse primers after enzyme cutting sites. Target PCR products were obtained using PCR assays with Pfu DNA Polymerase (TIANGEN, Beijing) and then cloned into pMD18-T vector (Takara) according to the manufacturer’s manual. After sequencing, positive recombinant plasmids were cleaved with individual restriction enzymes (New England Biolabs, USA) and linked with linearized eukaryotic expression vector pcDNA 3.1(+) cleaved with same enzymes using T4 DNA ligase (Takara). After identification by restriction analysis, the positive recombinant expression plasmids of pcNcGRAs were purified using HiPure Plasmid EF Mega Kit (Magen, Guangzhou) and concentrations were measured on a Nanodrop ND-2000 apparatus (Thermo Scientific, USA). The eukaryotic recombinant plasmids of pcNcGRAs were diluted in sterile endotoxin-free PBS to a final concentration of 1 mg/ml and stored at −20°C for further use.

Plasmid of pcNcGRAs were transiently transfected into RAW264.7 (ATCC, USA) cells previously prepared on coverslips in 24-well plates using Lipofectamine 2000 Transfection Reagent (Invitrogen, USA), and empty pcDNA 3.1(+) vector was used as negative control. After 72 h, cells were washed three times with PBS and fixed in 4% paraformaldehyde at room temperature (RT) for 10 min. After permeabilization in 0.1% Triton X-100 at RT for 20 min, cells were then washed with PBS three times and blocked in 5% BSA at RT for 2 h. Next, cells were incubated with mouse anti-HA-Tag mAb (1:200; AbClonal, Wuhan) at 4°C overnight and fluorescein (FITC)-conjugated AffiniPure Goat Anti-Mouse IgG( H + L) (1:100; Proteintech, Wuhan) at 37°C for 1 h, respectively. After washing unconjugated antibodies, cells were viewed on a fluorescence microscope (Olympus, Japan).

**Mice Immunization and Infection**

As shown in Figure 1, BALB/c mice were randomly divided into 8 groups (16/group). Control groups were either injected with 100 µl of sterile PBS or 100 µg of pcDNA 3.1(+) plasmid. Experimental groups were vaccinated with 100 µg of pcNcGRA plasmids through intramuscular injection (anterior tibial muscle). This immunized protocol was conducted at weeks 0, 2, and 4. Whole blood was collected through tail-bleed at weeks 2, 4, 6, and 9, and sera were separated and used for humoral antibody assay. At weeks 6 and 9, four mice from each group were euthanized; spleens were removed and used for lymphocyte proliferation assay and cytokine measurement assay. The remaining eight mice were infected with 2 × 10⁷ N. caninum tachyzoites at weeks 9 through intraperitoneal injection while monitoring the survival time for 30 days, and brain tissues were isolated for parasite burden assays.

**Humoral Antibody Levels Assay**

The specific antibody levels against NLA were monitored at weeks 2, 4, 6, and 9 in each group. NLA (100 µg/ml) was coated in 96-well plates (100 µl per well) at 4°C overnight. The plates were then blocked with 5% skim milk at RT for 2 h. Then, 100 µl of serum (100-fold dilution in PBS) was added into wells and incubated at 37°C for 2 h. Then, unconjugated antibodies were removed from wells through washing three times with PBST buffer and the remaining antibodies were detected with 100 µl of detection secondary antibodies of HRP-conjugated goat anti-mouse IgG, IgG1, or IgG2a (1:2,000; Abcam, UK) at 37°C for 1 h. After washing three times with PBST, 100 µl of TMB substrate solution (Beyotime, Beijing) was added into each well and incubated at 37°C for 15 min. The reaction were stopped by 100 µl of 2 M H₂SO₄ per well and OD₅₄₀ nm values were measured on a microplate reader (BioTek, USA).

**Lymphocyte Proliferation Assay**

Lymphocyte proliferation assay was carried out by isolation of spleens at weeks 6 and 9 (four mice per time-point and
FIGURE 1 | Immunization schedule and evaluation of protective efficacy. One hundred and twenty-eight BALB/c mice were randomly divided into 8 groups (16/group), including PBS control group, pcDNA control group, and pcNcGRA1, pcNcGRA4, pcNcGRA9, pcNcGRA14, pcNcGRA17, and pcNcGRA23 experimental groups. Mice were immunized three times with 2-week intervals. Four blood samples were collected at each time points of weeks 1, 4, 6, and 9 for humoral immune evaluation. Four spleen tissues were isolated at weeks 6 and 9 for cell immune evaluation. Eight BALB/c mice from each group inoculated with N. caninum tachyzoites at weeks 9 for immune protection evaluation.

per group). Single-cell suspensions were prepared as previously described (20). Splenocytes were seeded into a 96-well plate at a density of $3 \times 10^5$ cells per well and cultured in RPMI 1640 medium with 10% fetal bovine serum (Biological Industries). Cells were then stimulated with NLA at a final concentration of 10 µg/ml, concanavalin A (ConA, 5 µg/ml; Sigma, USA) as positive control, or non-treatment as negative control at 37°C/5% CO₂ for 72 h. The supernatants were discarded and replaced with fresh medium containing 10 µl of Cell Counting Kit-8 solution (CCK8; Beyotime). After incubation at 37°C/5% CO₂ for 3 h, the OD₄₅₀nm values were measured on a microplate reader (BioTek).

Cytokine Measurement Assay
Cytokine levels of IL-4 and IFN-γ in the splenocyte culturing supernatants were measured using IFN gamma or IL-4 Mouse ELISA Kit (Invitrogen, USA) according to the manufacturer's instructions. The OD₄₅₀nm values were read and converted to picograms per milliliter. Concentrations were calculated based on the standard curves generated in each plate.

Determination of Parasite Burden
Parasite burden was measured in brain tissues from immunized and infection mice at weeks 9 (eight mice per group) using real-time quantitative qPCR method. Genomic DNA was extracted using TIANamp Genomic DNA Kit (TIANGEN, China) according to instructions and standard curve was established with serially diluted N. caninum tachyzoite Nc-1 strain DNA samples ($1.0 \times 10^2 - 1.0 \times 10^7$ parasites/ml). The parasite burden was measured by amplifying the Nc5 gene (36) and parasite burden in 100 ng of tissue samples was calculated.

Statistical Analysis
Statistical analyses were conducted using SPSS (version 21.0; SPSS, Chicago, IL). All data represented mean ± SDs. A one-way ANOVA followed by Levene's test was used to analyze the difference between immunized groups and control groups or among different immunized groups. Correlation analysis was evaluated using non-parametric Spearman correlation with two-tailed tests. Kaplan–Meier survival curves were analyzed using log-rank (Mantel–Cox) tests. Differences were regarded as significant at *p < 0.05, **p < 0.01, or ***p < 0.001. Graphs were generated in GraphPad Prism 7.00 (GraphPad Software, La Jolla, CA, USA). All the experiments were performed three times with three technical replicates.

RESULTS
Evaluation of Humoral Immune Response in Immunized Mice
After sequencing, positive recombinant plasmids of pcNcGRAs were individually transfected into RAW264.7 cells and proteins were successfully expressed in vitro (Supplementary Figure 1).
To explore the humoral immune levels triggered by pcNcGRA DNA vaccines, serum was collected from mice tail at weeks 2, 4, 6, and 9. The IgG antibody levels against N. caninum were measured using indirect ELISA. As shown in Figure 2A, no significant differences were observed in the total IgG antibody levels when comparing serum from empty pcDNA3.1(+) vector (pcDNA)-treated group and PBS-treated group throughout the monitoring periods from 2 to 9 weeks ($p > 0.05$), whereas each pcNcGRA-immunized group triggered remarkably increased levels comparing with the control group. The pcNcGRAs significantly triggered high IgG antibodies at week 2 ($p < 0.05$ or $**p < 0.01$), the levels were increased at week 4 ($**p < 0.01$ or $***p < 0.001$), and the OD$_{450nm}$ values peaked at week 6 ($***p < 0.001$). Moreover, the high antibody levels lasted for 5 weeks after the last immunization ($***p < 0.001$).

To identify the Type 1 T helper (Th1) and Type 2 T helper (Th2) cells stimulated by pcNcGRA DNA vaccines, the IgG1 and IgG2a antibody levels in serum were determined at week 6. As shown in Figure 2B, there was no obvious difference between the pcDNA-immunized groups and negative control group when measuring the OD$_{450nm}$ values of IgG1 and IgG2a ($p > 0.05$). The IgG1 levels in pcNcGRA-immunized groups slightly increased ($p < 0.05$), whereas the IgG2a levels remarkably upregulated ($**p < 0.01$ or $***p < 0.001$) in comparison with that in the control groups. These data indicated that immunization with pcNcGRA DNA vaccines mainly induced Th1 immune responses in mice.

### Evaluation of Cell Immune Response in Immunized Mice

#### Lymphocyte Proliferation

To assess the cell-mediated immune responses, spleens were isolated from mice after 2 and 5 weeks since the last immunization. Lymphocyte proliferation assays were conducted using ConA as positive control, NLA as stimulators, and medium as negative control. As shown in Figures 3A,D, mice immunized with PBS or pcDNA both triggered no statistically significant proliferative response ($p > 0.05$), by contrast, the pcNcGRA-immunized mice caused obviously high levels of lymphocyte proliferation both at week 6 and 9 after stimulation with 10 µg/ml of pcNcGRAs ($**p < 0.01$, $***p < 0.001$) or 5 µg/ml of ConA ($***p < 0.001$). Overall, these results indicated that mice immunized with pcNcGRAs induced cell immune response.

#### Cytokine Production

The culturing supernatants were harvested from NLA-stimulated splenocytes for cytokine measurement. The secretion levels of Th1 (IFN-γ) and Th2 (IL-4) were determined and results are displayed in Figures 3B,C,E,F. All the splenocytes from pcNcGRA-immunized mice could induce high levels of IFN-γ ($**p < 0.01$), on the contrary, insignificant changes of IL-4 levels ($p > 0.05$). Furthermore, correlation analysis was performed to explore the relationship between humoral immune response and cellular immune response. As shown in Figure 4, significantly positive correlation not only occurred between IFN-γ secretion and IgG antibody levels ($p = 0.0012$, $r^2 = 0.3390$) but also appeared between IFN-γ secretion and IgG2a antibody levels ($p = 0.0034$, $r^2 = 0.2855$). There was a negative correlation between IL-4 and IgG1, but was not significant ($r = -0.02586$, $p > 0.05$). Overall, these data illustrated that all the pcNcGRAs triggered Th1 response.

### Protective Efficacy Evaluation in pcNcGRA-Immunized Mice

To assess the protective function of pcNcGRA vaccines, eight mice per group were infected with lethal doses of 2 $\times 10^7$ N. caninum tachyzoites and monitored the survival time for 30 days. As shown in Figure 5A, mice in control groups immunized with PBS or pcDNA died off within 5–6 days after infection. All the other immunized mice (pcNcGRA1: 9.5 ± 5.5 days, $p = 0.0074$; pcNcGRA4: 20.88 ± 11.12 days, $p = 0.0001$; pcNcGRA9: 7.75 ± 4.65 days, $p = 0.0056$; pcNcGRA14: 18.88 ± 10.83 days, $p < 0.0001$; pcNcGRA17: 16.63 ± 10.66 days, $p = 0.0003$; pcNcGRA23: 10.25 ± 7.03 days, $p = 0.0074$) displayed significantly longer survival time than the PBS-treated control groups (4 ± 1.31 days). During the monitoring periods, pcNcGRA4 and pcNcGRA14-immunized mice prolonged 10 more days of life and increased survival rate to 50 and 37.5%, respectively. In contrast, the death rates in pcNcGRA1 and pcNcGRA9-immunized groups were both 100%. For the pcNcGRA17 group, the death rate decreased to 75%. In addition, further analysis was made between different immunized groups. The pcNcGRA4-immunized mice showed significantly higher survival percentage than that in the pcNcGRA1 or pcNcGRA23 ($**p < 0.05$) and pcNcGRA9 ($***p < 0.01$). Moreover, pcNcGRA4-immunized mice showed significantly increased survival percentage than pcNcGRA1, pcNcGRA9, or pcNcGRA23 ($p < 0.05$). Comparisons between other immunized groups manifested no obvious significance ($p > 0.05$).

The parasite burden in infected mice brain tissues was determined in Figure 5B. Comparing with the PBS-treated group (1221 ± 269.59 parasites/100 ng), numbers of parasites in pcNcGRA-immunized mice brain tissues were significantly reduced (pcNcGRA4: 297.63 ± 83.77 parasites/100 ng, $p < 0.001$; pcNcGRA9: 738.36 ± 159.42 parasites/100 ng, $p = 0.029$; pcNcGRA14: 471.5 ± 110.74 parasites/100 ng, $p = 0.001$; pcNcGRA17: 592.13 ± 102.2 parasites/100 ng, $p = 0.005$; pcNcGRA23: 656 ± 144.3 parasites/100 ng, $p = 0.009$) except for pcNcGRA1-immunized group (818.38 ± 150.18 parasites/100 ng, $p = 0.096$). Comparisons between immunized groups with low parasite burden presented that pcNcGRA9, pcNcGRA17, or pcNcGRA23 shared similar levels with pcNcGRA1 group ($p > 0.05$). It is worth mentioning that the pcNcGRA4-immunized mice shared the lowest levels of parasite burden, which were remarkably decreased than other immunized groups ($**p < 0.01$ or $***p < 0.001$) except for pcNcGRA14 ($p = 0.096$). Meanwhile, body weight was also recorded throughout the whole survival period. As shown in Figure 5C, minimal body weight decrease in the survived mice occurred in the pcNcGRA4- or pcNcGRA14-immunized groups compared with other groups. After comprehensive evaluation, these results illustrated that pcNcGRA4, pcNcGRA17, and pcNcGRA23 DNA vaccines could be used for protection against N. caninum infection, and pcNcGRA4 DNA vaccine elicited the best protection effects on
FIGURE 2 | Time course assessment of humoral immune levels against pcNcGRAAs in mice. (A) IgG total levels of serum from immunized and control mice at weeks 2, 4, 6, and 9. (B) IgG1 and IgG2a levels of serum from immunized and control mice at week 6. Each bar represented the mean ± SD from four mice per group. Significance was analyzed between experimental groups and the PBS control group and presented as *p < 0.05, **p < 0.01, or ***p < 0.001.

infected mice and could improve the survival rate to 50% as well as decrease the parasite burden in brain tissues to 24.38%.

**DISCUSSION**

Neosporosis have threatened the cattle industries for a long time; however, no effective drugs or vaccines are available to use (5, 37). Up to now, there was only one licensed Neospora vaccine of Bovilis Neoguard composed of tachyzoite lysate available in partial countries (38). However, it has been stopped to be further used now for its uncompleted efficacy and risks of early embryonic death (39, 40). Along with the increasing demands for control and treatment with neosporosis, especially for countries with large numbers of cattle breeding industries, safe, and effective vaccines against neosporosis are urgently needed.

There have been several studies focused on live vaccines and subunit vaccines. In contrast, live vaccines shared excellent advantages of efficacy via triggering strong humoral and cellular immune responses; however, they were always regarded as alternatives and restricted its applications in the pharmaceutical industry for its high production costs, limitation in transportation and storage, and high risks for virulence return (41). Thus, vaccines of recombinant antigens are better choices for effective protection. In-depth knowledge of the underlying pathogenic mechanisms of *N. caninum* would help to combat with neosporosis. For apicomplexan parasites, GRA proteins are advantage weapons during invasion and construct the core contents of parasitophorous vacuole and...
intravacuolar network structure where parasites survive and replicate within host cells (32, 42). For instance, deletion of TgGRA6 gene would replace intravacuolar network with small vesicles and limit the replication of *T. gondii* (43). TgGRA2 gene disruption would inhibit Rab11A vesicle formation, disrupt intravacuolar network structure, and weaken the pathogenicity of *T. gondii* (44, 45). Moreover, many studies also proved to have protective effects in mice immunized with recombinant NcGRA antigens (20, 46).

Recently, DNA vaccines prepared with eukaryotic expression vectors could protect hosts from intracellular parasite invasion. Meanwhile, it shared advantages of low costs, easy transportation, and excellent immune protection through inducing lasted humoral and cellular responses (47). Many DNA vaccines have...
been evaluated in *T. gondii* by constructing with different antigens and verified to have the ability to trigger humoral and cellular immune responses lasting for long periods (48–50). For instance, TgGAR14 shares functions of intervacuolar transportation and mice immunized with TgGAR14 DNA vaccines would trigger specific immune responses, prolong the survival time, and reduce the parasite burden in target tissues (14). Abundant GRA proteins are secreted to make *N. caninum* survival and replication during invasion into hosts. Although many NcGRAs have been proved to be dominant antigens, there are few studies that focus on the evaluation of immune protection other than NcGRA1, NcGRA2, NcGRA6, and NcGRA7 up to now (17, 20, 28, 29). Ellis et al. (17) expressed GRA1 recombinant protein and evaluated the immunization effects using mice model. Results showed that GRA1 protein vaccine could produce protection in vertical propagation in mouse models (17). Considering that GRA1 protein was secreted after *N. caninum* invasion into host cells, the GRA1 DNA vaccine would stimulate a stronger Th1 immune response than GRA1 protein vaccine. Thus, GRA1 DNA vaccine was used as a positive control in our research. The present study evaluated the protective immunity of DNA vaccines encoding GRA1, GRA4, GRA9, GRA14, GRA17, and GRA23 of *N. caninum* in mice for the first time. Results demonstrated that mice immunized with these NcGRA DNA vaccines could both lead to humoral responses and cellular responses. Moreover, these vaccines exhibited different degrees of protective efficacy in survival time, body weight, and parasites in the brain tissues.

Humoral response plays vital roles in killing intracellular parasites (51). In the present study, the IgG total levels in serum isolated from pcNcGRA DNA–immunized mice were significantly upregulated than that in the PBS control group. In addition, the IgG secretion amount increases as time goes on and peaks at 2 weeks since the last immunization, and the antibody levels lasted for 5 weeks. Also, the IgG1 and IgG2a antibody levels were also measured after 2 weeks since the last immunization, and results demonstrated that pcNcGRA DNA–immunized mice could both upregulate the IgG1 levels ($\ast p < 0.05$) and dramatically increase the IgG2a antibody levels ($\ast\ast p < 0.01$ or $\ast\ast\ast p < 0.001$) in comparison with the control group.
These data were in accordance with other DNA immunizations that trigger mixed Th1/Th2 immune response, and Th1 immune response occupies the predominant function with a higher level of IgG2a (51, 52).

The Th1-type immune response was closely related with the secretion of IFN-γ cytokine (9, 53). To further explore the relationship between IgG2a response and IFN-γ cytokine, splenocytes were individually isolated from pcNcGRA DNA-immunized mice and control mice and then stimulated with NLA in vitro. An obviously increased level of IFN-γ cytokine was generated followed by different pcNcGRA DNA immunization (\( ^{**}p < 0.01 \)). Correlation analysis was also made and a significant difference occurred when comparing IgG total with IFN-γ or IgG2a with IFN-γ. The high levels of IFN-γ would contribute to enhance the macrophage function, disrupt parasitophorous vacuole through GTPases and p65 guanylate-binding proteins, and further eliminate the intracellular parasites (54, 55). Meanwhile, the Th2-type cytokine of IL-4 levels was also measured in stimulated splenocytes, however, there was no significance between all the pcNcGRA DNA-immunized groups and control group (\( p > 0.05 \)). As expected, the \( p \) values calculated from correlation analysis showed that no significant difference was observed between IgG and IL-4. These data indicated that immunization with pcNcGRA DNA vaccines could not cause sufficient B-cell proliferation and mast cell responses. This may be the main reason for the incomplete protection of mice from challenging with lethal dose of Neospora caninum.

T cell–mediated immunity mainly controls intracellular parasite infection. Splenocyte proliferation assays were carried out to determine the difference between immunized groups and the control group. A slightly increased level occurred in the pcNcGRA1 group (\( ^{*}p < 0.05 \)); obviously upregulated levels existed in the pcNcGRA9, pcNcGRA14, and pcNcGRA23 groups (\( ^{**}p < 0.01 \)); and an extremely significant increase occurred in the pcNcGRA4 and pcNcGRA17 groups (\( ^{***}p < 0.001 \)).

The aforementioned results could only illustrate those pcNcGRAs that trigger humoral and cellular responses and each protective immune response still needed to be determined in vivo. In the present study, the immunized mice were further challenged with lethal doses of N. caninum after 5 weeks since the last immunization. Results showed that the longest survival time occurred in the pcNcGRA4-, pcNcGRA14-, and pcNcGRA17-immunized groups and the highest survival rate occurred in the pcNcGRA4-immunized group, which were consistent with the measurement of parasite burden in brain tissues. Surprisingly, pcNcGRA1-, pcNcGRA9-, and pcNcGRA23-immunized mice triggered high levels of humoral and cellular responses; however, their performance relatively weakened protective immunity with 100% death rates even though the survival time was prolonged to some extent. These data indicated that pcNcGRA4, pcNcGRA14, and pcNcGRA17 DNA have the priority to be chosen as candidate vaccines and pcNcGRA4 DNA vaccine performed the best. All these data were evaluated on the non-pregnant mice. Although evaluation of vaccines using this immunocompetent mice model is regarded as the gold standard to explore protective antigens against N. caninum, some effective antigens display poor immune protection abilities in vertical transmission (5). Also, vaccines that triggered a strong immune response in cattle may not be available to protect against abortion (56). Thus, the protective efficacy of these DNA vaccines on pregnant mice to simulate vertical transmission and susceptible animals of cattle or other livestock still need to be determined in the future. The protective efficacy may differ in different animal models, physiological status, adjuvant types, immune methods, and antigen candidates from different strains; thus, promotion of safe, and effective vaccines should take comprehensive factors into consideration.

CONCLUSION

The present study evaluated the immunogenicity and potency of NcGRA1, NcGRA4, NcGRA9, NcGRA14, NcGRA17, and NcGRA23 using BALB/c mice as models. These eukaryotic expression vector constructed pcNcGRA DNA vaccines could trigger a strong Th1-type immune response accompanying with significantly increased levels of IgG2a antibodies and IFN-γ cytokine production in vitro. The immune protective efficacy revealed that pcNcGRA4, pcNcGRA14, and pcNcGRA17 DNA vaccines had relatively strong effects on prolonging the survival time and decreasing the parasite burden in brains. Although none of them completely resist the challenge of Neospora caninum infection, they would probably be useful for further vaccine development against Neospora caninum through combining multiple protective antigens.

DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/Supplementary Material, further inquiries can be directed to the corresponding author/s.

ETHICS STATEMENT

All animal experiments have been reviewed and approved by research ethics from the Animal Welfare and Research Ethics Committee of Jilin University (Permission No. ppzx20190929065).

AUTHOR CONTRIBUTIONS

JD and WL designed the research. GY, WL, QY, and JW conducted the research. GY, YW, TZ, XZ, and HF analyzed the data. PZ, QY, and WL wrote the article. JD and LC directed the project. All authors have read and approved the article.

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SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fvets.2021.638067/full#supplementary-material
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Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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