Rational Design of Zika Virus Subunit Vaccine with Enhanced Efficacy

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ABSTRACT  Zika virus (ZIKV) infection in pregnant women can lead to fetal deaths and malformations. We have previously reported that ZIKV envelope protein domain III (EDIII) is a subunit vaccine candidate with cross-neutralization activity; however, like many other subunit vaccines, its efficacy is limited. To improve the efficacy of this subunit vaccine, we identified a nonneutralizing epitope on ZIKV EDIII surrounding residue 375, which is buried in the full-length envelope protein but becomes exposed in recombinant EDIII. We then shielded this epitope with an engineered glycan probe. Compared to the wild-type EDIII, the mutant EDIII induced significantly stronger neutralizing antibodies in three mouse strains and also demonstrated significantly improved efficacy by fully protecting mice, particularly pregnant mice and their fetuses, against high-dose lethal ZIKV challenge. Moreover, the mutant EDIII immune sera significantly enhanced the passive protective efficacy by fully protecting mice against lethal ZIKV challenge; this passive protection was positively associated with neutralizing antibody titers. We further showed that the enhanced efficacy of the mutant EDIII was due to the shielding of the immunodominant nonneutralizing epitope surrounding residue 375, which led to immune refocusing on the neutralizing epitopes. Taken together, the results of this study reveal that an intrinsic limitation of subunit vaccines is their artificially exposed immunodominant nonneutralizing epitopes, which can be overcome through glycan shielding. Additionally, the mutant ZIKV protein generated in this study is a promising subunit vaccine candidate with high efficacy in preventing ZIKV infections in mice.

IMPORTANCE  Viral subunit vaccines generally show low efficacy. In this study, we revealed an intrinsic limitation of subunit vaccine designs: artificially exposed surfaces of subunit vaccines contain epitopes unfavorable for vaccine efficacy. More specifically, we identified an epitope on Zika virus (ZIKV) envelope protein domain III (EDIII) that is buried in the full-length envelope protein but becomes exposed in recombinant EDIII. We further shielded this epitope with a glycan, and the resulting mutant EDIII vaccine demonstrated significantly enhanced efficacy over the wild-type EDIII vaccine in protecting animal models from ZIKV infections. Therefore, the intrinsic limitation of subunit vaccines can be overcome through shielding these artificially exposed unfavorable epitopes. The engineered EDIII vaccine generated in this study is a promising vaccine candidate that can be further developed to battle ZIKV infections.

KEYWORDS  Zika virus, domain III, envelope protein, epitope shielding, glycan probe, intrinsic limitation of subunit vaccine designs, vaccine efficacy

Zika virus (ZIKV) is a member of the Flavivirus genus in the Flaviviridae family. Other members of the genus include dengue virus (DENV), West Nile virus (WNV), yellow fever virus (YFV), and Japanese encephalitis virus (JEV) (1, 2). ZIKV causes neurological...
diseases such as Guillain-Barré syndrome and congenital Zika syndrome (symptoms include microcephaly, brain abnormalities, and other congenital malformations) (3–6). Although several ZIKV vaccines are currently in clinical trials (7–9), no vaccine has been approved by the FDA to be used in humans.

The genome of ZIKV is a single-stranded positive-sense RNA and encodes a number of structural and nonstructural proteins (10, 11). The envelope (E) protein is a major structural protein and guides viral entry into host cells by first binding to a host receptor and then fusing viral and host membranes. It is anchored on viral envelopes and forms a dimer. Each monomer contains multiple structural domains: domain I (EDI), domain II (EDII), domain III (EDIII), the stem region, and the transmembrane domain (TM) (12, 13). The E protein is a main inducer of the host immune responses. Both the full-length E protein and its individual domains are prime targets for subunit vaccine design (14–19). Among these domains, EDI and EDII, but not EDIII, can trigger antibody-enhanced ZIKV entry (20–22). EDIII plays a critical role in viral entry by binding to host receptor(s). Because of its safety, EDIII has been our focus for subunit vaccine development. We previously showed that a ZIKV EDIII-based recombinant subunit vaccine is effective in eliciting long-term and broad-spectrum neutralizing antibodies against divergent ZIKV strains (23). However, its efficacy is not optimal. How to improve the efficacy of this subunit vaccine is key to its potential contribution to prevention of ZIKV infections.

Recombinant subunit vaccines have better safety over attenuated viral vaccines because unlike attenuated vaccines, subunit vaccines do not contain any residual infectivity. However, subunit vaccines often show insufficient efficacy. Structure-based vaccine designs have been intensively pursued to improve the efficacy of subunit vaccines, aiding battles against various viral diseases (24–26). In these vaccine designs, the envelope proteins of HIV, influenza virus, and non-ZIKV flaviviruses are modified to preserve specific neutralizing epitopes while subtracting or masking epitopes that potentially induce antibody-dependent enhancement effects or other unfavorable immune responses (24, 25, 27, 28). Trying to understand the generally low efficacy of viral subunit vaccines, we recently identified an intrinsic limitation of recombinant subunit vaccines, using Middle East respiratory syndrome (MERS) coronavirus (MERS-CoV) as a model system. That is, subunit vaccines are part of a large viral envelope protein or a virus particle, and hence much of their surface areas are buried; however, when a recombinant subunit vaccine is made, the previously buried surface areas become artificially exposed, and they contain immunodominant nonneutralizing epitopes that distract the host immune system from reacting to neutralizing epitopes (29). We also developed a neutralizing immunogenicity index (NII) to quantitatively characterize the contribution of each of these epitopes to the overall immunogenicity of the subunit vaccine (29). In spite of their initial success in MERS-CoV subunit vaccine designs, the concept of the intrinsic limitation of viral subunit vaccines and the strategy of using NII to quantitatively characterize epitopes have not been extended to subunit vaccine designs targeting another virus.

In this study, we analyzed the tertiary structure of ZIKV EDIII, identified an artificially exposed epitope on the vaccine surface, shielded it with a glycan probe, characterized its NII, and measured the efficacy of the mutant EDIII in protecting mice and their fetuses from ZIKV infections. We further investigated the molecular mechanism for the enhanced efficacy of the mutant EDIII. Overall, this study has firmly established and characterized an intrinsic limitation of viral subunit vaccines and pointed out a way to overcome this limitation. It has important implications for the design and development of novel viral subunit vaccines. The mutant ZIKV EDIII generated in this study can be a promising subunit vaccine candidate for preventing ZIKV infections in humans.

RESULTS
Identification and masking of a nonneutralizing epitope on ZIKV EDIII. To identify potentially immunodominant nonneutralizing epitopes on ZIKV EDIII, we analyzed the crystal structure of ZIKV E protein dimer (PDB identifier [ID] SLBV) (30). We
found a patch of surface area on EDIII that is buried in the full-length E protein dimer but becomes exposed in recombinant EDIII. Met375 is located in the center of this patch and protrudes from a bent β-strand (Fig. 1A and B). Hence, we engineered a glycan probe onto the epitope surrounding residue 375 (i.e., epitope 375) (Fig. 1C). To this end, we introduced double mutations M375N/E377T to EDIII, which changed residue 375 to an N-linked glycosylation site. We expressed and purified the mutant EDIII, along with wild-type EDIII, in mammalian 293T cells. Both proteins contain a C-terminal Fc tag and were purified to homogeneity. The results from Western blotting showed that both proteins ran as dimers without boiling (due to the Fc dimeric tag and incomplete denaturing) and as monomers after boiling (due to complete denaturing). The mutant EDIII ran slower and with a slightly larger molecular weight than wild-type EDIII (Fig. 2A), indicating that the mutant EDIII was glycosylated after addition of a glycan probe at residue 375.

To characterize the conformation of the mutant ZIKV EDIII, we investigated the binding interactions between ZIKV EDIII (wild type or mutant) and EDIII-specific monoclonal antibodies (MAbs) using enzyme-linked immunosorbent assay (ELISA). Among the MAbs used in this study, it has been shown previously that ZV-54, ZV-67, ZKA64-LALA, 2004, and SMZAb5 can potently neutralize ZIKV infections in mice and/or in vitro, whereas ZV-2 has no neutralizing activity in vitro (in vitro activities were characterized using a plaque-based neutralization assay) (Fig. 2B) (10, 23, 31, 32). Our results showed
that the mutant EDIII bound to ZV-54, ZV-67, ZKA64-LALA, Z004, and SMZAb5 but did not bind to ZV-2. In comparison, wild-type EDIII bound to all of the six MAbs (Fig. 2C).

The crystal structures of ZIKV E protein complexed with several of the MAbs have been determined previously by other groups (10, 31). Based on these previously determined structures, we analyzed the binding sites of these MAbs on EDIII. The crystal structures of ZIKV E protein complexed with ZV-67 and of ZIKV EDIII complexed with ZV-2 revealed that ZV-67 binds to an area that is exposed on the E protein dimer and a likely receptor-binding site, whereas ZV-2 binds to an area centering epitope 375 that is buried on the E protein dimer but becomes artificially exposed on the recombinant EDIII (Fig. 1B) (10). Structural analysis of EDIII complexed with Z004 identified the lateral ridge (away from epitope 375) of EDIII as its recognizing epitope (31). The structural bases for the binding of ZV-54, ZKA64-LALA, and SMZAb5 are not known, but these MAbs may also bind to functionally important regions on EDIII and away from epitope 375. Overall, these data further confirm that the mutant EDIII has successfully incorporated a glycan at residue 375, which blocked the binding of a nonneutralizing MAb ZV-2; they also reveal that despite the introduced glycan, the mutant EDIII retains its native structural conformation and antigenicity by binding to five different neutralizing MAbs.

Measurement of neutralizing immunogenicity index of EDIII epitope 375. We previously defined the neutralizing immunogenicity index (NII) of an epitope as the contribution of the epitope to the overall neutralizing immunogenicity of the vaccine (29). For epitope 375 on ZIKV EDIII, the NII can be calculated as the difference between the neutralizing immunogenicity of wild-type EDIII and that of the mutant EDIII, divided by the neutralizing immunogenicity of wild-type EDIII. To measure the NII of epitope 375 on ZIKV EDIII, we immunized mice with wild-type EDIII and mutant EDIII individually and measured the neutralizing antibody titers of the induced sera. The immunized mice included immunocompetent BALB/c and C57BL/6 mice as well as immunocompromised interferon-α/β receptor (IFNAR)-deficient (Ifnar1−/−) mice (C57BL/6 background).
The results showed that compared to the wild-type EDIII, the mutant EDIII induced significantly higher titers of anti-ZIKV neutralizing antibodies (Fig. 3A to C) but similar titers of ZIKV EDIII-specific IgG antibodies in all of the three types of immunized mice (Fig. 3D to F). These data suggest that epitope 375 makes a negative contribution to the overall neutralizing immunogenicity of EDIII; furthermore, masking epitope 375 does not affect the total IgG antibody production but instead increases the overall neutralizing immunogenicity of the vaccine. We also calculated the NII of epitope 375. The values of NII varied slightly in different mice: -0.50 in BALB/c, -0.76 in C57BL/6, and -0.34 in Ifnar1/-/- mice (Fig. 3G). The negative NII confirms the negative contribution of epitope 375 to the overall neutralizing immunogenicity of the vaccine. The values for NII suggest that masking epitope 375 increases the overall neutralizing immunogenicity of the vaccine by 50%, 76%, and 34% [calculated using formula (PRNT50-WT – PRNT50-mutant)/PRNT50-WT × 100, where PRNT50-WT and PRNT50-mutant represent PRNT50 neutralizing antibody titers for the wild-type and mutant EDIII proteins, respectively]. In a separate experiment, BALB/c mice were further immunized with different doses of above WT and mutant ZIKV EDIII proteins. Sera were collected on day 10 post-2nd immunization for detection of IgG (H) and neutralizing antibodies against ZIKV strain R103451 (I) as described above. The IgG antibody titers are expressed as the endpoint dilutions that remain positively detectable. Neutralizing antibody titers are expressed as PRNT50. The data are presented as means ± SEM of quadruplicate wells of 2-fold serially diluted sera (pooled from 5 mice in each group). The experiments were repeated three times, and similar results were obtained.

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To investigate how different antigen doses affect the neutralizing immunogenicity of the mutant ZIKV EDIII vaccine, BALB/c mice were divided into six groups and each group was immunized with one of three doses (2, 10, and 20 μg/mouse) of either the wild-type or mutant vaccine. The induced IgG and neutralizing antibodies were then compared. The results showed that for each of the doses tested, the wild-type and mutant EDIII vaccines induced the same amount of IgG antibody titers (Fig. 3H), while the mutant EDIII elicited significantly higher titers of anti-ZIKV neutralizing antibodies than wild-type EDIII did (Fig. 3I). In addition, vaccine doses were positively associated
with the induced neutralizing antibody titers (Fig. 3I). These data suggest that immunization with increased doses of epitope 375-masked EDIII potentially improves the overall neutralizing immunogenicity of the vaccine.

To identify the predominant IgG subtypes and also to investigate whether epitope masking affects subtype production, sera of BALB/c, C57BL/6, and Ifnar1−/− mice immunized with the wild-type and mutant EDIII proteins were evaluated for their IgG subtype contents. The results showed that IgG1 was the predominant subtype induced by both vaccines in all mouse strains (Fig. 4A). On the other hand, no or low titers of IgG2a were detected in Ifnar1−/− mice (Fig. 4B), a low titer of IgG2b was detected in Ifnar1−/− mice (Fig. 4C), a low titer of IgG2c was detected in BALB/c mice (Fig. 4D), and a low titer of IgG3 was detected in all mouse strains tested (Fig. 4E). Overall, compared to the wild-type EDIII, the mutant EDIII induced more IgG1, IgG2a, IgG2b, and/or IgG2c subtypes in these mouse strains. These results demonstrate that although total IgG titers were similar in mice immunized by the wild-type and mutant EDIII vaccines, the associated IgG subtypes varied depending on the mouse strains tested.

Enhanced efficacy of the mutant EDIII vaccine in protecting mice and their fetuses. We investigated the efficacy of the mutant ZIKV EDIII vaccine in protecting immunocompetent pregnant mice and their fetuses from ZIKV infection (immunocompetent adult mice, such as BALB/c, are nonlethal models for ZIKV infection). To this end, we immunized BALB/c mice with either the wild-type or mutant EDIII vaccine and challenged the pregnant mice with ZIKV. We then collected placenta and fetal brain 6 days postchallenge and measured the titers and RNA copies of ZIKV in these samples, respectively, using plaque and quantitative reverse transcription-PCR (qRT-PCR) assays. Embryos were also collected for comparison of fetal status. The results showed that in contrast to the wild-type-EDIII-treated group, the mutant-EDIII-immunized pregnant mice had undetectable viral titers and RNA copies in their placenta and fetal brain (Fig. 5A and B). Negative-control pregnant mice, treated with phosphate-buffered saline (PBS), contained significantly higher viral titers and RNA copies in their tissues (Fig. 5A).
and B). In addition, there were no significant morphological changes in the uteri of the wild-type- and mutant-EDIII-immunized pregnant mice, and the embryos from these mice were in good condition. In contrast, the PBS-treated mice showed severe resorption in their uteri and the embryos from these mice were invisible (Fig. 5C). Collectively, compared to the wild-type EDIII vaccine, the mutant EDIII vaccine provides enhanced efficacy in protection of immunocompetent pregnant mice from ZIKV infection as evidenced by undetectable viral titers and RNA copies, although both the wild-type and mutant EDIII vaccines protect the fetuses of immunocompetent pregnant mice from ZIKV infection.

We also evaluated the efficacy of the mutant ZIKV EDIII vaccine in protecting immunocompromised mice and their fetuses from ZIKV infection (immunocompromised mice are lethal models for ZIKV infection). To this end, we immunized female and male Ifnar1−/− mice with either the wild-type or mutant ZIKV EDIII vaccine. We then treated female and male mice differently. First, we challenged immunized female Ifnar1−/− pregnant mice with ZIKV, collected their tissues (including placenta and fetal brain) 6 days postchallenge, and measured the ZIKV titers in these tissues using the plaque assay. We also examined the embryos and uteri 6 days postchallenge. The results showed that compared to the wild-type-EDIII-treated group, the mutant-EDIII-immunized female Ifnar1−/− pregnant mice had significantly lower viral titers in their lung, liver, heart, muscle (Fig. 6A), and placenta and undetectable viral titers in fetal brain (Fig. 6B). There were no significant morphological changes in the uteri. Moreover, all embryos were in good condition in the wild-type- and mutant-EDIII-treated mice (Fig. 6C). In the negative-control group (i.e., PBS-treated female Ifnar1−/− pregnant mice), the viral titers were high in all the tested tissues and fetal brain, and there were significant morphological changes in the uteri, with severe fetal resorption or fetal death (Fig. 6). Second, we challenged adult male Ifnar1−/− mice with low-dose (10^3 PFU) and high-dose (5 × 10^4 PFU) ZIKV sequentially and observed them for survival and weight changes. The results showed that the male Ifnar1−/− mice, which were immunized with either the wild-type or mutant EDIII vaccine and then challenged with low-dose ZIKV, all survived 14 days post-1st challenge (Fig. 7A); their weight decreased slightly during days 4 to 7 postinfection but kept increasing steadily afterwards (Fig. 7B). In contrast, in the negative-control group (i.e., PBS-treated mice), all mice had
at least 25% weight loss and were humanely euthanized 10 days postinfection (Fig. 7A and B). Moreover, the mutant-EDIII-immunized mice all survived the second, high-dose ZIKV challenge (100% survival) with no obvious weight loss. In contrast, only 67% of the wild-type-EDIII-immunized mice survived the second, high-dose ZIKV challenge, with considerable weight loss during days 7 to 12 postchallenge (Fig. 7C and D). Furthermore, neutralizing antibody titers in the mutant-EDIII-immunized mice were significantly higher than in the wild-type-EDIII-immunized mice before both the 1st and 2nd challenges, particularly after the 2nd challenge (Fig. 7E). Therefore, the enhanced protective efficacy of the mutant EDIII vaccine against the high-dose ZIKV challenge was likely due to the higher serum neutralizing antibody titers that it induced. Overall, the above data reveal that the mutant EDIII vaccine protects immunocompromised female pregnant mice and their fetuses from ZIKV infection more effectively than the wild-type EDIII vaccine (as evidenced by undetectable or significantly reduced viral titers), and it also protects immunocompromised adult male mice from high-dose ZIKV infection more effectively (as evidenced by improved survival rate and steadily maintained weight).

Enhanced passive protection efficacy of the mutant-ZIKV-EDIII-induced mouse sera. To understand the association between the enhanced protective efficacy of the mutant EDIII vaccine and the increased neutralizing antibody titers that it induced, passive protection experiments were performed using the wild-type or mutant-EDIII-induced mouse sera. To this end, sera were pooled from each group and normalized for equal EDIII-specific IgG titers. Adult Ifnar1−/− mice were passively transferred with the pooled sera (either undiluted or after 1:5 dilution). The mice were then challenged with ZIKV and their viremia, survival rate, and weight changes were subsequently monitored. The results showed that viremia was significantly lower in the mice passively transferred with the mutant-EDIII-induced sera than in those transferred with the wild-type-EDIII-induced sera (Fig. 8A). In addition, while 77% of the mice transferred with the diluted mutant-EDIII-induced sera survived the ZIKV challenge, mice transferred with undiluted sera all survived the ZIKV challenge (100% survival), both of which only showed slight
In comparison, for mice transferred with diluted or undiluted wild-type-induced sera, the survival rates were significantly lower or lower than for those with diluted or undiluted mutant-EDIII-induced sera, only about 31% and 80%, respectively; the former group also had significant weight loss during days 8 to 10 postchallenge (Fig. 8B and C). Moreover, the mutant-EDIII-induced sera had significantly higher anti-ZIKV neutralizing antibody titers than the wild-type-EDIII-induced sera (Fig. 8D). Negative-control mice transferred with the PBS-induced sera had a 0% survival rate at day 10 postchallenge; they also had the highest viral titers and most weight loss (Fig. 8A).

**FIG 7** Enhanced protective efficacy of mutant ZIKV EDIII vaccine in ZIKV-challenged lethal Ifnar1−/− mouse model. Adult male Ifnar1−/− mice were immunized with the ZIKV WT and mutant (M375N/E377T) EDIII proteins, or PBS control, and challenged with ZIKV (R103451; 10^4 PFU) 10 days post-2nd immunization to evaluate survival (A) and weight (B) for 14 days (n = 6). The surviving Ifnar1−/− mice in the WT and mutant M375N/E377T groups were further challenged with ZIKV (R103451; 5 × 10^6 PFU) and observed for survival (C) and weight (D) as described above (n = 6). The percent weight in panels B and D represents the mean percent weight of all surviving mice at the indicated days after challenge. Error bars indicate SEM. (E) Neutralizing antibodies in the sera of mice before the 1st and 2nd challenges and post-2nd challenge against ZIKV strain R103451 by plaque reduction neutralization assay. The data are expressed as mean PRNT_{50} ± SEM of quadruplicate wells of 2-fold serially diluted sera pooled from 4 to 6 mice in each group. The experiments were repeated three times, and similar results were obtained.

**FIG 8** Characterization of the protective efficacy of the passively transferred sera from mutant-ZIKV-EDIII-vaccine-immunized mice. Ifnar1−/− mice were passively transferred with mutant (M375N/E377T) EDIII-immunized mouse sera. Six hours later, they were challenged with ZIKV strain R103451 (10^3 PFU), followed by evaluation of serum viral titers on days 3 and 5 postchallenge, as well as survival and weight changes for 14 days. Mouse sera from WT-EDIII- or PBS-immunized mice were included as controls. (A) ZIKV titers in the sera of challenged mice collected on days 3 and 5 after ZIKV challenge. Error bars indicate SEM (n = 5). ** and *** $P < 0.01$ and 0.001, respectively. The detection limit was 20 PFU/ml of sera. Survival (B) and weight (C) rates of serum transferred mice after ZIKV challenge were also recorded. Error bars indicate SEM (n = 5 to 13). * in panel B, $P < 0.05$ between WT and mutant EDIII protein groups (1:5 dilution). The percent weight in panel C represents the mean percent weight of all surviving mice on days indicated after challenge. (D) Characterization of neutralizing antibodies in the above sera against ZIKV challenge using plaque reduction neutralization test. Neutralizing activity is expressed as PRNT_{50}. Error bars indicate SEM (n = 5). * and ** $P < 0.05$ and 0.01, respectively. The experiments were repeated three times, and similar results were obtained.
The above data indicate that compared to the wild-type-EDIII-induced sera, the mutant-EDIII-induced sera contained significantly higher neutralizing antibody titers as well as significantly enhanced passive protective efficacy against ZIKV challenge, which likely were positively associated with each other.

**Molecular mechanism for the enhanced efficacy of the mutant EDIII.** To explore the molecular mechanism for the enhanced efficacy of the mutant EDIII vaccine, we examined competitive binding among EDIII, EDIII-specific MAbs, and EDIII-induced sera using ELISA. More specifically, we investigated the binding interactions between the wild-type EDIII and EDIII-specific neutralizing MAbs in the presence of the mutant-EDIII-induced mouse sera. The wild-type-EDIII-induced mouse sera were used as a comparison. The result showed that compared to the wild-type-EDIII-induced mouse sera, the mutant-EDIII-induced sera from both immunocompetent BALB/c mice (Fig. 9A to D) and immunocompromised Ifnar1−/− mice (Fig. 9E to H) blocked the binding between the wild-type EDIII and EDIII-specific neutralizing MAbs more effectively. As we showed earlier that the wild-type and mutant EDIII vaccines induced similar titers of total IgG antibodies (Fig. 3), the results from the competitive binding assay suggest that the mutant EDIII vaccine induced higher neutralizing antibody titers than the wild-type EDIII vaccine. Therefore, masking the immunodominant nonneutralizing epitope surrounding residue 375 triggered the host immune system to refocus on neutralizing, but less immunodominant, epitopes, accounting for the production of more neutralizing antibodies.

**DISCUSSION**

ZIKV poses a significant threat to human health. Particularly, it has acquired the capability to cross the placenta and infect pregnant women and their fetuses, leading to severe consequences such as fetal neuropathology and fetal death (33, 34). Development of safe and effective vaccines to protect pregnant women and their fetuses is a high priority for the biomedical field. To date, a number of vaccine candidates have...
been tested in mouse and nonhuman primate models for ZIKV infection (16, 17, 35–37). Some of these vaccine candidates are currently in clinical trials (7–9, 38). In contrast to the traditional vaccines that typically focus on inactivated or attenuated virus particles, subunit vaccines contain only the antigenic components of the pathogen and hence are generally safer and more stable than traditional vaccines (http://www.who.int/vaccine_safety/initiative/tech_support/Part-2.pdf) (20, 39). The ZIKV E protein guides viral entry into cells and is a main inducer of the host immune response. It has been shown that unlike some domains of the E protein, recombinant ZIKV EDIII does not induce antibody-enhanced viral entry and hence is safe as a subunit vaccine in protecting mice from ZIKV challenge (23, 40). However, like many other recombinant subunit vaccines, ZIKV EDIII shows the following intrinsic limitation that affects its protective efficacy. As a domain of the ZIKV E protein, EDIII has large surface areas buried in the full-length dimeric E protein; when expressed individually as a recombinant EDIII, these previously buried surface areas become artificially exposed. The artificially exposed surface areas of EDIII may contain immunodominant nonneutralizing epitopes that distract the host immune system from reacting to neutralizing epitopes (e.g., epitopes in the receptor-binding regions). In this study, we aimed to overcome this intrinsic limitation of ZIKV EDIII. We first identified such an artificially exposed epitope on ZIKV EDIII around residue 375 and then introduced a glycan to cover this epitope (i.e., epitope 375). The goal was to shield epitope 375 from the host immune system, forcing the host immune system to refocus on neutralizing epitopes such as those in the receptor-binding regions.

We characterized the mutant ZIKV EDIII with epitope 375 shielded. Like wild-type EDIII, the mutant EDIII bound to EDIII-specific neutralizing MAbs through its receptor-binding regions, suggesting that the mutant EDIII retains its natural conformation. However, only the wild-type EDIII, and not the mutant EDIII, bound to an EDIII-specific nonneutralizing MAb targeting epitope 375, indicating that epitope 375 had been successfully glycosylated in the mutant EDIII. The increased molecular weight of the mutant EDIII also confirmed the successful glycosylation of epitope 375.

We measured the neutralizing immunogenicity index (NII) of epitope-375 in several mouse strains. The NIIs of epitope 375 in different mouse strains all were negative, suggesting that epitope 375 makes a negative contribution to the overall neutralizing immunogenicity of the EDIII vaccine. The NIIs of epitope 375 in different mouse strains vary from −0.34 to −0.76, suggesting that shielding of epitope 375 increases the overall neutralizing immunogenicity by 34% to 76%. Particularly, the absolute value of the NII for epitope 375 is lower in Ifnar1−/− mice (−0.34) than in BALB/c (−0.50) or C57BL/6 (−0.76) mice, suggesting that shielding of this epitope increases the overall neutralizing immunogenicity in immunocompromised Ifnar1−/− mice less than in immunocompetent BALB/c and C57BL/6 mice. Moreover, the neutralizing immunogenicity of the epitope 375-shielded mutant EDIII vaccine increases with the vaccine doses. We then examined the efficacy of the mutant EDIII in protecting mice from ZIKV challenges. Compared to the wild-type EDIII vaccine, the mutant EDIII vaccine protected female pregnant mice (both immunocompetent and immunocompromised mice) and their fetuses more effectively and also improved the survival rate of immunocompromised male mice against high-dose lethal ZIKV challenge. In addition, the mutant-EDIII-immunized mouse sera passively protected mice more effectively than the wild-type-EDIII-immunized mouse sera against lethal ZIKV challenge, and the protection was positively associated with serum neutralizing antibody titers. We further investigated the molecular mechanism behind the enhanced efficacy of the mutant EDIII vaccine. We found that shielding epitope 375 significantly increased the neutralizing antibody titers but maintained similar total IgG antibody titers in the immunized mice. Hence, shielding immunodominant nonneutralizing epitopes forces the host immune system to refocus on neutralizing epitopes of subunit vaccines. Studies on other flaviviruses, such as tick-borne encephalitis virus (TBEV), suggest that there may be variations in the specificity of antibody responses (including neutralizing antibody responses) between immunized mice and immunized (vaccinated or virus-infected) humans: while in mice...
antibody responses mainly target EDIII, in humans antibody responses mainly target EDI/EDII (41, 42). It is not clear whether this is the same case for ZIKV vaccines. Nevertheless, cautions should be taken for designing EDIII-based ZIKV subunit vaccines for human use.

It is worth noting that adjuvants play an important role in potentiating subunit vaccine-induced immune responses, including neutralizing antibodies. Aluminum hydroxide adjuvant may enhance overall immune responses and modulate specificity and functional activity of antibody responses that are induced by an EDIII-based TBEV vaccine in mice (41). Adjuvant System 04 (AS04), combining monophosphoryl lipid A (MPL) and aluminum salt, has been licensed for use in human vaccines (e.g., human papillomavirus and hepatitis B virus vaccines) with increased antibody titers (43, 44). Compared with aluminum adjuvant (hydroxide or salt) alone, the combination of MPL and aluminum salt may induce isotype-switched IgG antibody responses and significantly elevate total IgG and neutralizing antibodies as induced by subunit vaccines with an acceptable safety profile (45–47). Indeed, in this study, we found that the aluminum hydroxide and MPL adjuvant combination elicited high-titer neutralizing antibodies in conjunction with both the wild-type and mutant ZIKV EDIII vaccines. In addition, compared to the recombinant protein vaccines targeting EDIII of other flaviviruses with Freund’s adjuvants, the wild-type ZIKV EDIII vaccine with the aluminum hydroxide and MPL adjuvant combination seems to induce higher titers of neutralizing antibodies (48, 49). In general, as demonstrated in this and other studies, vaccines targeting ZIKV EDIII elicit high IgG antibody titers but relatively low neutralizing antibody titers (23, 40, 50), hence justifying the use of adjuvants. Consistent with previous data (51, 52), ZIKV EDIII vaccines elicited potent IgG1 subtype antibody responses in all mouse strains tested, suggesting that this antibody subtype might be associated with neutralizing antibody production. We also found that the aluminum hydroxide and MPL adjuvant combination helps the induction of IgG2a (for BALB/c mice), IgG2b, and IgG2c subtypes (for C57BL/6 and C57BL/6-based Ifnar1−/− mice), resulting in a more balanced immune response. Future studies will be needed to compare immunogenicity of ZIKV EDIII vaccines (wild type or mutant) in the presence of the aluminum hydroxide and MPL adjuvant combination, aluminum hydroxide alone, or MPL alone to determine whether the enhanced immunogenicity is associated with adjuvanticity of the combinational adjuvants. Such studies will also be helpful for comparing immunogenicity and efficacy of our vaccines with those of previous EDIII-based ZIKV or other flavivirus vaccines using same or different adjuvants.

To sum up, this study is the second successful application of the concept of NII in vaccine design, after the first application in MERS coronavirus subunit vaccine design (29). This study further establishes the following important principle in subunit vaccine designs. First, artificially exposed regions on viral subunit vaccines contain immunodominant nonneutralizing epitopes. Second, these epitopes limit the efficacy of viral subunit vaccines by distracting the host immune system from reacting to neutralizing epitopes. Third, shielding these epitopes can improve the efficacy of the vaccines by refocusing the host immune system on neutralizing epitopes. These principles can potentially be extended to many other viral subunit vaccines, such as vaccines against influenza and Ebola viruses. This study has also produced a safe and highly effective ZIKV subunit vaccine that holds promise to be developed into a successful vaccine to protect pregnant women and their fetuses. Therefore, the current study has important implications for both the battle against ZIKV and vaccine designs in general.

MATERIALS AND METHODS

Animals. Four- to 6-week-old female BALB/c and C57BL/6 mice and 3- to 6-week-old male and female Ifnar1−/− mice were used in the study. The Ifnar1−/− mice (breeding pairs) on the C57BL/6 background were originally purchased from the Jackson Laboratory and were maintained for self-breeding in our animal facilities. The animal studies were carried out in strict accordance with the recommendations in the Guide for the Care and Use of Laboratory Animals (53). The protocols were approved by the Institutional Animal Care and Use Committee (IACUC) of the New York Blood Center (permit numbers 344.00 and 345.01).
Expression and purification of recombinant proteins. Recombinant ZIKV EDIII proteins were prepared as previously described (23, 54). Briefly, ZIKV EDIII protein containing residues 298 to 409 (i.e., wild type) was constructed based on ZIKV E (ZikaSPH2015 strain; GenBank accession number KJU21639.1) fused with a C-terminal human IgG1 Fc tag. A mutant ZIKV EDIII protein containing a glycan probe surrounding residue 375 (i.e., M375N/E377T) was constructed by multistep mutagenesis kits (Thermo Fisher Scientific) using wild-type EDIII plasmid as the template. The recombinant proteins were transiently expressed in 293T cell culture supernatants and purified by protein A affinity chromatography (GE Healthcare).

Animal immunization. Purified ZIKV EDIII vaccines (wild type or mutant) were used to immunize mice as previously described (23, 55). Briefly, groups of six BALB/c, C57BL/6 (4- to 6-week-old), and Ifnar−/− (3-week-old) mice were intramuscularly (i.m.) immunized with the EDIII vaccine (wild-type or mutant) (10 µg/mouse) or PBS control, plus a combination of aluminum hydroxide adjuvant (500 µg/mouse, vau-alu-250; InvivoGen) and monophosphoryl lipid A (MPL) adjuvant (10 µg/mouse, vac-mpl; InvivoGen). The immunized mice were boosted once with the same immunogens at 4 weeks, and sera were collected 10 days post-last immunization to test antibody responses and neutralizing antibodies. In addition, groups of five BALB/c mice (4 to 6 weeks old) were immunized and boosted with the EDIII vaccine (wild type or mutant) at 0, 2, 10, and 20 µg/mouse, respectively, in the presence of aluminum hydroxide and MPL adjuvant combination, followed by detection of IgG and neutralizing antibodies as described above.

Western blotting. The purified proteins were analyzed using Western blotting as previously described (23, 56). Briefly, proteins were first run on 10% Tris-glycine SDS-PAGE gels and were then transferred to nitrocellulose membranes for Western blot analysis. The transferred blot was blocked with 5% fat-free milk in PBS-Tween 20 (PBST) overnight at 4°C, followed by sequential incubation with ZIKV EDIII-specific antibodies or mouse IgG1, IgG2a, IgG2b, IgG2c, or IgG3 (1:2,000), or anti-human IgG-Fab antibody (1:5,000) or anti-mouse IgG-Fab antibody (1:5,000; Thermo Fisher Scientific) or anti-human IgG-Fab (for ZV-67, ZKA64-LALA, Z004, and SMZAb5, 1:5,000; Abcam) antibody for 1 h at 37°C. The reaction was detected using the substrate 3,3′,5,5′-tetramethylbenzidine (Sigma) and stopped with 1 N H2SO4. Absorbance at 450 nm was measured using an ELISA plate reader (Tecan).

ZIKV EDIII-specific antibodies in mouse sera were tested using ELISA as described above except that the plates were coated with the wild-type ZIKV EDIII protein, followed by sequential incubation with serially diluted mouse sera and HRP-conjugated anti-mouse IgG Fab (for ZV-2 and ZV-54, 1:5,000; Thermo Fisher Scientific) or anti-human IgG-Fab (for ZV-67, ZKA64-LALA, Z004, and SMZAb5, 1:5,000; Abcam) antibody for 1 h at 37°C. The signal was visualized with ECL Western blot substrate buffer and Amersham Hyperfilm (GE Healthcare).

ELISA. The binding interactions between ZIKV EDIII proteins and EDIII-specific MAbs were characterized using ELISA as previously described (57). These MAbs include mouse MAbs ZV-2 (BEI Resources) and ZV-54, as well as human MAbs ZV-67, ZKA64-LALA, Z004, and SMZAb5 (Absolute Antibody) (10, 21, 31, 32). Briefly, ELISA plates were coated with the wild-type or mutant ZIKV EDIII protein overnight at 4°C and blocked with 2% fat-free milk in PBST for 2 h at 37°C. After three washes, the plates were sequentially incubated with serially diluted MAbs and HRP-conjugated anti-mouse IgG-Fab (for ZV-2 and ZV-54, 1:5,000; Thermo Fisher Scientific) or anti-human IgG-Fab (for ZV-67, ZKA64-LALA, Z004, and SMZAb5, 1:5,000; Abcam) antibody for 1 h at 37°C. The reaction was detected using the substrate 3,3′,5,5′-tetramethylbenzidine (Sigma) and stopped with 1 N H2SO4. Absorbance at 450 nm was measured using an ELISA plate reader (Tecan).

The competition between ZIKV EDIII-specific human MAbs (including ZV-67, ZKA64-LALA, Z004, or SMZAb5) and mutant EDIII-induced mouse sera for the binding of wild-type ZIKV EDIII was performed using ELISA as described above, except that the binding between wild-type EDIII and the MAbs (0.5 µg/ml) was tested in the presence of serially diluted mouse sera induced by the ZIKV EDIII vaccine (wild-type or mutant) or PBS. The EDIII- MAb binding was measured by a EDIII-MAB antibody (1:5,000) and subsequent enzymatic reaction.

ZIKV plaque-forming assay and plaque reduction neutralization test. ZIKV human strain R103451 (2015/Honduras) was amplified in Vero E6 cells for determination of viral titers using a standard plaque-forming assay. Viral titers in the tissues and/or sera from challenged mice were determined as described above. The ZIKV titers were detected from around 40 or 50 mg of tissue samples, or 50 µl of sera, so the detection limit was about 25 or 20 PFU/g of tissues, or 20 PFU/ml of sera. A plaque reduction neutralization test was carried out to measure neutralizing antibody titers in immunized mouse sera or activities of EDIII-specific MAbs as previously described (23). Briefly, ZIKV (100 PFU) was incubated with pooled mouse sera (at 2-fold serial dilutions from 1:10 to 1:2,560, quadruplicates of each dilution) or MAbs (4-fold serial dilutions from 1 µg/ml to 0.0005 µg/ml, quadruplicates of each dilution) for 1.5 h at 37°C, followed by addition of antibody-virus mixtures to Vero E6 cells and incubation for 1 h at 37°C. The cells were overlaid with medium (1% carboxymethyl cellulose in Dulbecco modified Eagle medium [DMEM] containing 2% fetal bovine serum [FBS]) and cultured at 37°C for 4 to 5 days before being stained with 0.5% crystal violet. PRNT50 and 50% neutralization dose (ND50) were calculated through dilutions and neutralizations of sera or MAbs at 50% plaque reduction using the CalcuSyn computer program (which is based on the median effect principle) (58–62). Specific PRNT50 and ND50 of sera or MAbs were obtained after inputting individual dilutions and neutralization values for each serum or MAB to the program.

ZIKV challenge and protection evaluation. ZIKV challenge experiments were carried out as previously described (23, 63). Briefly, 10 days post-last immunization, female Ifnar−/− and BALB/c mice immunized with ZIKV EDIII (wild type or mutant) were mated with respective naive male mice within the same strains. Mice injected with PBS were included as controls. Pregnant female mice (E10 to E13) were intraperitoneally (i.p.) challenged with ZIKV strain R103451 (10³ PFU for Ifnar−/− mice and 2 × 10³ PFU for BALB/c mice; 200 µl/mouse). ZIKV titers and RNA copies in tissues of adult mice as well as placenta and fetal brain (collected at 6 days postinfection) were detected using plaque-forming assay (described...

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above) and qRT-PCR (described below), respectively. In a separate experiment, male Ifnar1−/− mice were (i.p.) challenged with ZIKV (R103451; 10^4 PFU) 10 days post-last immunization and evaluated for survival and weight for another 14 days. Ten days after completion of the 1st challenge experiment, the surviving Ifnar1−/− mice in the wild-type or mutant ZIKV EDIII groups were further challenged with ZIKV (R103451; 5 × 10^4 PFU) and observed for survival and weight as described above. Mice with ≥25% body weight loss were humanely euthanized.

Passive protection. Groups of 5 to 13 male and female Ifnar1−/− mice (6- to 8-week old) were injected (i.p.) with pooled sera, (200 μL/mouse; normalized for equal ZIKV EDIII-specific IgG titers at 10^6 or a 1:5 dilution in PBS) of wild-type- or mutant-EDIII-vaccine-immunized Ifnar1−/− mice collected before the 1st and 2nd ZIKV challenges. Six hours later, mice were i.p. challenged with ZIKV strain R103451 (10^4 PFU). Mice injected with PBS-induced Ifnar1−/− mouse sera were included as controls. Neutralizing activity of passively transferred mouse sera was detected using the plaque reduction neutralization test. ZIKV titers were detected from sera collected at 3 days and 5 days postchallenge using the ZIKV plaque-forming assay. Mouse survival and weight were recorded for 14 days postchallenge. Mice with ≥25% body weight loss were humanely euthanized.

qRT-PCR. ZIKV RNA copies in sera and tissues of challenged mice were detected using qRT-PCR as previously described (23). Briefly, RNAs were extracted using QiAmp MinElute virus spin kit (for sera) (Qiagen) and RNeasy minikit (for tissues) (Qiagen) and quantified through one-step qRT-PCR using Power SYBR Green PCR master mix, MultiScribe reverse transcriptase, and Ambion RNase inhibitor (Thermo Fisher Scientific) in a Viia 7 Master Cycler PCR system (Thermo Fisher Scientific). The forward and reverse primers for the amplification were 5′-TGGTGCTAGATGCTGATTGC-3′ and 5′-CCTCCACAAGTGCCCTATTGC-3′,

A standard curve of qRT-PCR was established by 10-fold serial dilutions of a recombinant plasmid containing ZIKV membrane (M) and E genes. A linear standard curve at 10^3 to 10^12 RNA copies (correlation coefficient [R²] value > 0.98; detection limit = 10^1 RNA copies) was selected for calculation of ZIKV RNA in the samples. The ZIKV RNA was purified from around 40 mg of tissue samples, so the detection limit was about 2.5 × 10^2 RNA copies/g of tissue (64).

Statistical analysis. The values were presented as means with standard errors of the means (SEM). Statistical significance of antibody titers among different groups was calculated using the Mann-Whitney test. Statistical significance of viral titers among different groups, as well as serum inhibition, was calculated through Student’s two-tailed t test. Statistical significance of survival curves among different groups was calculated through the log rank test using GraphPad Prism statistical software.

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W.T., Y.Z., F.L., and L.D. designed the study. W.T., J.C., G.Z., Q.G., L.H., and Y.C. conducted the experiments and analyzed the data. Q.G. and F.L. performed the structural analysis. W.T., Y.Z., F.L., and L.D. wrote, revised, and proofread the paper.

We declare no competing interests.

REFERENCES

1. Lazear HM, Diamond MS. 2016. Zika virus: new clinical syndromes and its emergence in the Western Hemisphere. J Virol 90:4846–4875. https://doi.org/10.1128/JVI.00252-16.

2. Gatherer D, Kohl A. 2016. Zika virus: a previously slow pandemic spreads rapidly through the Americas. J Gen Virol 97:269–273. https://doi.org/10.1099/jgv.0.003381.

3. Kassavetis P, Joseph JB, Francois R, Perloff MD, Berkowitz AL. 2016. Zika virus: new clinical syndromes and its emergence in the Western Hemisphere. J Virol 90:4846–4875. https://doi.org/10.1128/JVI.00252-16.

4. Salinas JL, Walters DM, Styczynski A, Garzon F, Quijada H, Bravo E, Chaparro P, Madero J, Acosta-Reyes J, Ledermann J, Arteta Z, Borland E, Burns P, Gonzalez M, Powers AM, Mercado M, Solano A, Sejvar JJ, Ospina ML. 2017. Zika virus–associated Guillain-Barre syndrome variant in Haiti. Neurology 87: 336–337. https://doi.org/10.1212/WNL.0000000000002759.

5. Salinas JL, Walters DM, Styczynski A, Garzon F, Quijada H, Bravo E, Chaparro P, Madero J, Acosta-Reyes J, Ledermann J, Arteta Z, Borland E, Burns P, Gonzalez M, Powers AM, Mercado M, Solano A, Sejvar JJ, Ospina ML. 2017. Zika virus–associated Guillain-Barre syndrome variant in Haiti. Neurology 87: 336–337. https://doi.org/10.1212/WNL.0000000000002759.

6. Modjarrad K, Lin L, George SL, Stephenson KE, De La Barrera RA, Jarman RG, Sondergaard E, Tennant J, Ansel JL, Mills K, Koren M, Robb ML, Barnett J, Thompson J, Kosel AE, Dawson P, Hale A, Tan CS, Walsh SR, Meyer KE, Crowell TA, Blazevic A, Mosby K, Larocca RA, Abbink P, Boyd M, Bricault CA, Seaman MS, Basil A, Walsh M, Tonwe V, Hoft DF, Thomas SJ, Barouch DH, Michael NL. 2018. Preliminary aggregate safety and immunogenicity results from three trials of a purified inactivated Zika virus vaccine candidate: phase 1, randomised, double-
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38. Cohen J. 2018. Deep drop in Zika cases undermines vaccine trial. Science 361:1055–1056. https://doi.org/10.1126/science.361.6407.1055.

39. Zhang N, Jiang S, Du L. 2014. Current advancements and potential strategies in the development of MERS-Cov vaccines. Expert Rev Vaccines 13:761–774. https://doi.org/10.1586/14760584.2014.912134.

40. Yang M, Dent M, Lai H, Sun H, Chen Q. 2017. Immunization of Zika virus envelope protein domain III induces specific and neutralizing immune responses against Zika virus. Vaccine 35:4287–4294. https://doi.org/10.1016/j.vaccine.2017.04.052.

41. Zlatkovic J, Tsouchnikas G, Jamrer J, Koessl C, Stiasny K, Heinz FX. 2013. Aluminum hydroxide influences not only the extent but also the fine specificity and functional activity of antibody responses to tick-borne encephalitis virus in mice. J Virol 87:12187–12195. https://doi.org/10.1128/JVI.01690-13.

42. Jamrer J, Zlatkovic J, Tsouchnikas G, Vratsikhi O, Strauss J, Aberle JH, Chmeliak V, Kundl M, Stiasny K, Heinz FX. 2014. Variation of the human antibody responses after tick-borne encephalitis virus infection and vaccination. J Virol 88:13845–13857. https://doi.org/10.1128/JVI.02086-14.

43. Didierlaurent AM, Morel S, Lockman L, Giannini SL, Bisteau M, Carlsen H, Kielland A, Vosters O, Vanderheyde N, Schiavetti F, Larocque D, Van MM, Moon SH, Shin EC, Noh YW, Lim YT. 2015. Evaluation of hyaluronic acid-based combination adjuvant containing monophosphoryl lipid A and aluminum salt for hepatitis B vaccine. Vaccine 33:4762–4769. https://doi.org/10.1016/j.vaccine.2015.08.006.

44. Apostolico JS, Boscardin SB, Yamamoto MM, De Oliveira-Filho JN, Kalil J, Cunha-Neto E, Rosa DS. 2016. HIV envelope trimer specific immune response is influenced by different adjuvant formulations and heterologous prime-boost. PLoS One 11:e0145637. https://doi.org/10.1371/journal.pone.0145637.

45. Ko EJ, Lee YT, Kim KH, Lee Y, Jung YJ, Kim MC, Lee YN, Kang T, Kang SM. 2017. Roles of aluminum hydroxide and monophosphoryl lipid A adjuvants in overcoming CD4+ T cell deficiency to induce isotype-switched IgG antibody responses and protection by T-dependent influenza vaccine. J Immunol 198:279–291. https://doi.org/10.4049/jimmunol.1600173.

46. Moon SH, Shin EC, Noh YW, Lim YT. 2015. Evaluation of hyaluronic acid-based combination adjuvant containing monophosphoryl lipid A and aluminum salt for hepatitis B vaccine. Vaccine 33:4762–4769. https://doi.org/10.1016/j.vaccine.2015.08.006.

47. Apostolico JS, Boscardin SB, Yamamoto MM, De Oliveira-Filho JN, Kalil J, Cunha-Neto E, Rosa DS. 2016. HIV envelope trimer specific immune response is influenced by different adjuvant formulations and heterologous prime-boost. PLoS One 11:e0145637. https://doi.org/10.1371/journal.pone.0145637.

48. Block OK, Rodrigo WW, Quinn M, Jin X, Rose RC, Schlesinger JJ. 2010. A tetravalent recombinant dengue domain III protein vaccine stimulates neutralizing and enhancing antibodies in mice. Vaccine 28:8085–8094. https://doi.org/10.1016/j.vaccine.2010.10.004.

49. Zhao H, Jiang T, Zhou XZ, Deng YQ, Li XF, Chen SP, Zhu SY, Zhou X, Qin ED, Qin CF. 2014. Induction of neutralizing antibodies against four serotypes of dengue viruses by MixBiDEIII, a tetravalent dengue vaccine. PLoS One 9:e86573. https://doi.org/10.1371/journal.pone.0086573.

50. Yang M, Lai H, Sun H, Chen Q. 2017. Virus-like particles that display Zika virus envelope protein domain III induce potent neutralizing immune responses in mice. Sci Rep 7:7679. https://doi.org/10.1038/s41598-017-08247-9.

51. Wu Y, Li S, Du L, Wang C, Zou P, Hong B, Yuan M, Ren X, Tai W, Kong Y, Zhou C, Lu L, Zhou X, Jiang S, Ying T. 2017. Neutralization of Zika virus by germline-like human monoclonal antibodies targeting cryptic epitopes on envelope domain III. Emerg Microbes Infect 6:E89. https://doi.org/10.1038/emi.2017.79.

52. Keeffe JR, Van Rampay KK, Olsen PC, Wang Q, Gazumyan A, Azzopardi SA, Schaefer-Babajev D, Lee YE, Stuart JB, Singapari A, Watanabe J, Usachenko J, Ardesir A, Saeed M, Agudelo M, Eisenreich T, Bournazos S, Oliveira TY, Rice CM, Coffey LL, Macdonald MR, Bjorkman PJ, Nussenzweig MC, Robbiani DF. 2018. A combination of two human monoclonal antibodies prevents Zika virus escape mutations in non-human primates. Cell Rep 25:1385–1394. https://doi.org/10.1016/j.celrep.2018.10.031.

53. National Research Council. 2011. Guide for the care and use of laboratory animals, 8th ed. National Academies Press, Washington, DC.

54. Tai W, Wang Y, Fett CA, Zhao G, Li F, Perlman S, Jiang S, Zhou Y, Du L. 2017. Recombinant receptor-binding domains of multiple Middle East respiratory syndrome coronaviruses (MERS-CoVs) induce cross-neutralizing antibodies against divergent human and camel MERS-CoVs and antibody escape mutants. J Virol 91:e01651-16. https://doi.org/10.1128/JVI.01651-16.

55. Zhang N, Channappanavar R, Ma C, Wang L, Tang J, Garron T, Tao X, Tasneem S, Lu L, Tseng CT, Zhou Y, Perlman S, Jiang S, Du L. 2016. Identification of an ideal adjuvant for receptor-binding domain-based subunit vaccines against Middle East respiratory syndrome coronavirus. Cell Mol Immunol 13:180–190. https://doi.org/10.1038/cmi.2015.03.

56. Ma C, Wang L, Tao X, Zhang N, Yang Y, Tseng CT, Li F, Zhou Y, Jiang S, Du L. 2014. Searching for an ideal vaccine candidate among different MERS coronavirus receptor-binding fragments—the importance of immunofocusing in subunit vaccine design. Vaccine 32:6170–6176. https://doi.org/10.1016/j.vaccine.2014.08.086.

57. Du L, Zhao G, Yang Y, Qiu H, Wang L, Kou Z, Tao X, Yu H, Sun S, Tseng CT, Jiang S, Li F, Zhou Y. 2014. A conformation-dependent neutralizing monoclonal antibody specifically targeting receptor-binding domain in Middle East respiratory syndrome coronavirus spike protein. J Virol 88:7045–7053. https://doi.org/10.1128/JVI.00433-14.

58. Koide F, Goebel S, Snyder B, Walters KB, Gast A, Hagelin K, Kalkeri R, Rayner J. 2016. Development of a Zika virus infection model in cynomolgus macaques. Front Microbiol 7:2028. https://doi.org/10.3389/fmicb.2016.02028.

59. Xu S, Wu W, Wang Q, Wang C, Hua C, Li W, Lu L, Jiang S. 2018. Peptide-based membrane fusion inhibitors targeting HCoV-229E spike protein HR1 and HR2 domains. UMS 19:487. https://doi.org/10.3399/jimms/19020487.

60. Chou TC. 2006. Theoretical basis, experimental design, and computerized simulation of synergism and antagonism in drug combination studies. Pharmacol Rev 58:621–681. https://doi.org/10.1124/pr.58.3.10.

61. Hachiya A, Reeve AB, Marchand B, Michailidis E, Ong YT, Kirby KA, Leslie MD, Oka S, Kodama EN, Rohan LC, Mitsuya H, Parniak MA, Sarafianos SG. 2013. Evaluation of combinations of 4-ethyl-2-fluoro-2-deoxyadenosine with clinically used antiretroviral drugs. Antimicrob Agents Chemother 57:4534–4538. https://doi.org/10.1128/AAC.00283-13.

62. Lin B, He S, Yam HJ, Jiang TJ, Hu Z. 2016. Evaluation of antiviral drug synergy in an infectious HCV system. Antivir Ther 21:595–603. https://doi.org/10.3851/IMP3044.

63. Yu Y, Deng YQ, Zou P, Wang Q, Dai Y, Yu F, Du L, Zhang NN, Tian M, Hao JN, Meng Y, Woo JFW, Yuen KY, Qian CF, Jiang S, Lu L. 2017. A peptide-based viral inactivator inhibits Zika virus infection in pregnant mice and fetuses. Nat Commun 8:15672. https://doi.org/10.1038/ncomms15672.

64. Tai W, Voronin D, Chen J, Bao W, Kessler DA, Shaz B, Jiang S, Yazdanbaksh K, Du L. 2019. Transfusion-transmitted Zika virus infection in pregnant mice leads to broad tissue tropism with severe placental damage and fetal demise. Front Microbiol 10:7209. https://doi.org/10.3389/fmicb.2019.00209.