



Defective Interfering Viral Particle Treatment Reduces Clinical Signs and Protects Hamsters from Lethal Nipah Virus Disease

^(D)Stephen R. Welch,^a ^(D)Jessica R. Spengler,^a Jessica R. Harmon,^a JoAnn D. Coleman-McCray,^a ^(D)Florine E. M. Scholte,^a Sarah C. Genzer,^a ^(D)Michael K. Lo,^a Joel M. Montgomery,^a Stuart T. Nichol,^a Christina F. Spiropoulou^a

aViral Special Pathogens Branch, Division of High Consequence Pathogens and Pathology, Centers for Disease Control and Prevention, Atlanta, Georgia, USA

Stephen Welch and Jessica Spengler contributed equally to this work.

ABSTRACT Defective interfering particles (DIs) contain a considerably smaller genome than the parental virus but retain replication competency. As DIs can directly or indirectly alter propagation kinetics of the parental virus, they offer a novel approach to antiviral therapy, capitalizing on knowledge from natural infection. However, efforts to translate in vitro inhibition to in vivo screening models remain limited. We investigated the efficacy of virus-like particles containing DI genomes (therapeutic infectious particles [TIPs]) in the Syrian hamster model of lethal Nipah virus (NiV) disease. We found that coadministering a high dose of TIPs intraperitoneally with virus challenge improved clinical course and reduced lethality. To mimic natural exposure, we also evaluated lower-dose TIP delivery and virus challenge intranasally, finding equally efficacious reduction in disease severity and overall lethality. Eliminating TIP replicative capacity decreased efficacy, suggesting protection via direct inhibition. These data provide evidence that TIP-mediated treatment can confer protection against disease and lethal outcome in a robust animal NiV model, supporting further development of TIP treatment for NiV and other high-consequence pathogens.

IMPORTANCE Here, we demonstrate that treatment with defective interfering particles (DIs), a natural by-product of viral infection, can significantly improve the clinical course and outcome of viral disease. When present with their parental virus, DIs can directly or indirectly alter viral propagation kinetics and exert potent inhibitory properties in cell culture. We evaluated the efficacy of a selection of virus-like particles containing DI genomes (TIPs) delivered intranasally in a lethal hamster model of Nipah virus disease. We demonstrate significantly improved clinical outcomes, including reduction in both lethality and the appearance of clinical signs. This work provides key efficacy data in a robust model of Nipah virus disease to support further development of TIP-mediated treatment against high-consequence viral pathogens.

KEYWORDS Nipah virus, defective interfering particles, Syrian hamsters, animal model, henipavirus, treatment, antiviral, antiviral agents

N ipah virus (NiV) is a highly pathogenic zoonotic paramyxovirus (family *Paramyxoviridae*, genus *Henipavirus*) capable of causing severe disease in humans (1), with case fatality rates ranging from 30% to 80% (2). First recognized during an outbreak of severe febrile encephalitic and respiratory illnesses in Malaysia and Singapore from 1998 to 1999 (3–5), NiV has caused nearly annual outbreaks in India and Bangladesh since 2001 (6–8). Transmission to humans occurs predominately via direct contact with the excreta of the reservoir species, the fruit bat (genus *Pteropus*) (9, 10), with pigs also recognized as an amplifying host and source of infection during the initial outbreak (3, 5, 11). Human-to-

Editor Dimitrios Paraskevis, Medical School, National and Kapodistrian University of Athens This is a work of the U.S. Government and is not subject to copyright protection in the United States. Foreign copyrights may apply. Address correspondence to Christina F. Spiropoulou, ccs8@cdc.gov.

The authors declare no conflict of interest. Received 1 November 2021 Accepted 31 January 2022 Published 17 March 2022 human transmission is frequently reported, most commonly between close family and caregivers of NiV-infected individuals (12, 13). Currently, there are no licensed vaccines or antiviral therapies available for NiV infection (https://www.who.int/activities/prioritizing -diseases-for-research-and-development-in-emergency-contexts).

Defective interfering particles (DIs) have been shown to modulate multiple aspects of viral pathogenesis, including inhibition of viral replication and immune stimulation (14, 15), and have been characterized in numerous RNA viruses, including influenza (16), Ebola (17), canine distemper (18), and Sendai (19) viruses. DI genomes are formed by errors during nascent viral genome synthesis, in which translocation of the viral polymerase during RNA replication results in truncated and functionally inoperative versions of the genome (20). While they have lost the ability to encode the complete suite of viral proteins, DIs retain their self-replicative capacity due to intact promoter motifs at genomic termini. Shorter than the parental viral genomes, DIs are replicated faster, resulting in increased sequestration of critical viral and host-cell components, thus, in turn, outcompeting replication of full-length genomes. While direct inhibition is believed to be the main mechanism of DI-mediated inhibition, DIs can also induce anti-viral immune responses that further inhibit authentic viral replication (21–23).

We recently identified numerous naturally occurring NiV DI genomes consisting of two variants, (i) trailer copybacks, which have complementary trailer nucleotide sequences at each genomic terminus, resulting in either a partial or complete hairpin structure; and (ii) deletions, which have authentic leader and trailer sequences but are missing large (>90%) sections of the internal nucleotide sequences (24). Several of these DI genomes were shown to potently inhibit NiV replication when incorporated into virus-like particles and used to treat cells. While the *in vitro* inhibitory potential of DI particles has been well documented, *in vivo* efficacy to support their use as a treatment or therapeutic candidate has not been evaluated beyond a limited number of influenza studies (25–30).

Here, we investigate the use of DI-containing particles (termed therapeutic infectious particles, or TIPs) for treating NiV infection in the Syrian hamster model of disease (31–33). Infection in Syrian hamsters accurately recapitulates the two most common phenotypic clinical presentations of human NiV disease, respiratory distress and neurological complications (34, 35). After characterizing optimal challenge parameters for this model, we applied it to assess the *in vivo* efficacy of selected TIPs to combat NiV infection. We found that high-dose intraperitoneal delivery of TIPs resulted in dramatically improved clinical outcomes and reduced NiV-associated mortality in the lethal hamster model of NiV disease. Furthermore, intranasal delivery of TIPs given at a 2,000fold lower dose also conferred protection. These data demonstrate that exploiting the natural inhibitory effects of DIs (21, 36, 37) can significantly reduce disease severity and support the continued development of this unique treatment approach to control NiV and other highly lethal viral infections.

RESULTS

NiV-infected Syrian hamsters recapitulate both respiratory and neurological disease phenotypes, providing a clinically relevant challenge model for TIP assessment. Prior to evaluating the potential of TIP treatment, we performed additional in-house characterization and clinical evaluation of the Syrian hamster model (31–33) using two inoculation routes, intranasal (i.n.) and intraperitoneal (i.p.), and a range of challenge doses. Besides monitoring body weights and temperatures, we developed a detailed clinical scoring system to better resolve NiV disease progression in hamsters. With i.n. inoculation, increased clinical severity was directly associated with increases in challenge dose (Fig. 1A); with i.p. inoculation lower doses achieved higher lethality than in i.n. inoculation (Fig. 1B).

The onset of clinical signs and overall disease progression were broadly comparable and independent of challenge dose and route. Respiratory signs (increased respiratory rate, dyspnea, and, in more severe cases, nasal and oral hemorrhage) appeared 3 to 6 days postinfection (dpi) (Fig. 1C). Disease progression appeared more acute in i.p.-



FIG 1 Optimal challenge dose determination in Nipah virus-infected Syrian hamsters. Hamsters were inoculated with Nipah virus (NiV) strain Malaysia via the intranasal (i.n.) route at 10³ (orange, n = 9), 10⁵ (blue, n = 8), or 10⁷ (red, n = 9) TCID₅₀ (A) or via the intraperitoneal (i.p.) route with 10⁵ (green, n = 10) TCID₅₀ (B). Graphs represent percent weight change from baseline (taken at -1 dpi), body temperatures, and daily clinical scores (scored from 0 to 10), with severity depicted by increasing intensity of red. Animals scoring \geq 10 were humanely euthanized; any animals that succumbed to disease prior to euthanasia were allocated a score of 10. Gray boxes indicate the end of monitoring/scoring due to euthanasia/death. Individual animals are represented, with the black line indicating the mean value each day. (C) Combined survival curves for both i.n. and i.p. challenge routes indicating typical clinical signs observed at indicated times postinfection. Independently of inoculation dose and route, the initial disease presentation is respiratory distress beginning 3 to 6 dpi followed by a primarily neurological presentation at 8 to 20 dpi.

challenged animals, transitioning from an absence of clinical signs to severe disease in 12 to 14 h (versus up to 3 days in i.n.-challenged animals). In hamsters not developing or succumbing to respiratory disease, neurological signs (mild to severe, with either a sudden or gradual onset) generally become apparent from \sim 8 dpi onwards. Mild to moderate clinical signs usually started as ataxia, limb weakness, head tilt progression

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FIG 2 Genome schematics and organization of NiV and DI genomes packaged as TIPs. (A) Virion and viral genome schematic of NiV. Viral genome is represented in the genomic polarity. 3' Le, leader sequence; N, nucleoprotein; P, phosphoprotein; M, matrix protein; F, fusion protein; G, glycoprotein; RdRp, viral RNA-dependent RNA polymerase; 5' Tr, trailer sequence. (B) Schematics of the DI genomes incorporated into TIPs.

from sporadic to persistent, and a generalized abnormality in behavior (very quiet, cautious movements, low creeping gait). These progressed to a more severe presentation that included limb paralysis, severe head tilt (>90°) associated with inability to stand on all four legs and a constant "corkscrewing" behavior, and short to constant tonicclonal seizures associated with hyperthermia. Sudden onset (<24 h after appearing healthy) of these severe neurological signs was common. Some animals exhibiting mild to moderate clinical signs maintained this presentation until study end, and some recovered. The latest observed onset of neurological signs eventually requiring euthanasia (in a previously healthy hamster) was 13 dpi in a hamster challenged i.n. with 10^6 TCID₅₀; the animal was euthanized at 16 dpi. Based on these data and previously generated data (38, 39), we chose challenge doses of 10^4 TCID₅₀ i.p. or 10^6 TCID₅₀ i.n. as optimal for evaluating efficacy *in vivo*. At these doses, we expected disease progression to be slow enough to allow manifestation of observable clinical signs and mortality rates to be sufficient to evaluate protective efficacy against lethal outcome.

TIP treatment provides robust protection against NiV disease in hamsters, and efficacy is enhanced when TIPs are active and able to directly inhibit viral replication and transcription. The NiV genome is a single-stranded negative-sense RNA containing 6 genes encoding 9 proteins (Fig. 2A). Four promising TIP candidates previously characterized to result in the greatest inhibition of NiV replication in vitro were selected for in vivo evaluation (24). Three of these were copyback variants DI-07 (1,470 nucleotides [nt] in length), DI-14 (1,006 nt), and DI-35 (546 nt), and one was deletion variant DI-10 (486 nt) (Fig. 2B). In vitro experiments indicated that a high ratio of TIP to virus (>2,500:1) was required for significant virus inhibition (24). Thus, to maximize the TIP/NiV treatment ratio in hamsters for initial screening, we chose the i.p. route of inoculation because larger delivery volumes can be used than with other routes. TIPs were given simultaneously with virus challenge at a DI/NiV ratio of \sim 20,000:1 (2 mL total volume). Hamsters were challenged with a combination of NiV mixed with either: active TIPs (treated with 100 mJ/cm² UV to destroy the full-length producer NiV genomes carried over from TIP production); inactive TIPs (treated with 2,400 mJ/cm² UV to destroy both full-length NIV and TIP genomes); or Dulbecco's modified Eagle's medium (DMEM) only (control animals).

All mock-treated animals (no TIPs) exhibited weight loss between 3 to 7 dpi (Fig. 3A), as well as other clinical signs of infection (Fig. 3B); infection was lethal in 18 out of 20 (90%) hamsters. Both hamsters that survived until the end of challenge (28 dpi) exhibited persistent mild to moderate neurological signs of infection. In contrast, the majority of hamsters treated with active TIPs did not exhibit overt weight loss or temperature changes (Fig. 3A). Six of 10 (60%) of hamsters treated with active DI-07, 4 of 10 (40%) treated with active DI-10 and DI-35, and 3 of 9 (33%) treated with active



FIG 3 Intraperitoneal NiV challenge and treatment with active or inactive TIPs. Hamsters were inoculated i.p. with 2 mL DMEM containing 10^4 TCID₅₀ NiV strain Malaysia in combination with approximately 2×10^9 active TIPs (treated with 100 mJ/cm² of UV), inactive TIPs (treated with 2,400 mJ/cm² of UV), or no TIPs (mock treated). (A) Graphs represent percentage of weight change from baseline (taken at -1 dpi) and body temperatures. Individual animals are represented. DI-07, red; DI-10, green; DI-14, blue; DI-35, orange; no TIP (mock treated), black; closed circles represent hamsters treated with active TIPs, and open circles represent those treated with inactive TIPs. Lines indicate the mean value each day as follows: solid line for hamsters treated with active TIPs, and open circles represent those treated with inactive TIPs. Lines indicate the mean value each day as follows: solid line for hamsters treated with active TIPs, and open circles represent those treated with active TIPs.

closer to that of the mock-treated animals: the majority displayed weight loss between 3 to 7 dpi accompanied by other clinical signs of infection. When clinical signs were present in TIP-treated hamsters, they were more likely to appear in those treated with inactive TIPs (92.5%) than those treated with active TIPs (60.5%). Overall, TIP treatment significantly (DI-07, P = 0.0001; DI-10, P = 0.0001; DI-14,

P = 0.0009; DI-35, P = 0.0002) increased survival from 10% in mock-treated animals to 70 to 80% in active TIP-treated hamsters (Fig. 3E). Survival in hamsters treated with inactive TIPs was decreased compared to those treated with active TIPs, although these animals remained significantly protected from lethal disease (DI-07, P = 0.0367; DI-10, P = 0.0053; DI-14, P = 0.0094; DI-35, P = 0.0104) compared to mock-treated animals (30 to 50%). Corresponding to improved outcomes, levels of viral RNA in tissues were lower in survivors (both symptomatic and asymptomatic) than in animals that succumbed to infection (Fig. S2 in the supplemental material). Of the TIP-treated animals that succumbed to infection, reduction in RNA levels was more pronounced in hamsters treated with active TIPS than with inactive TIPs.

Low-dose TIPs can protect against severe outcomes when hamsters are treated and challenged intranasally. Given the success of the simultaneous i.p. challenge and treatment experiment, we next evaluated TIP treatment using the i.n. route, which more closely mimics the natural exposure and transmission dynamics of NiV by monitoring clinical signs, outcome, and endpoint viral RNA in tissues (Fig. 4C; Fig. S3). Given the limitations regarding the maximum volume allowed for i.n. inoculation in hamsters, plus the requirement for a higher-challenge dose needed to see appreciable disease and mortality (10^6 TCID₅₀), the maximum achievable TIP/NiV ratio for this experiment was only ~100:1 (in 100 μ L) compared to ~20,000:1 achieved in the i.p. experiment (in 2 mL). Given the reduced protective effect we observed previously with inactive TIPs, hamsters were only challenged with a combination of NiV plus active TIPs.

Even with a 200-fold reduction in TIP/NiV ratio compared to i.p. delivery, i.n. TIP treatment conferred high levels of protection from NiV infection. All mock-treated hamsters (n = 20) exhibited weight loss between 3 to 6 dpi, and temperature changes associated with hypoactivity (hypothermia) or severe seizures (hyperthermia) were observed in several animals (Fig. 4A); all animals displayed clinical signs at some point during the challenge period (Fig. 4B). In contrast, only 3 of 8 (37.5%) animals treated with DI-14 and 5 of 8 (62.5%) animals treated with DI-35 displayed any clinical signs throughout the challenge period (Fig. 4C). Again, the duration of observed clinical signs was significantly reduced in some TIP-treated hamsters (DI-14 and DI-35) compared to mock-treated animals, although the majority of DI-07- and DI-10-treated animals also had reduced clinical periods of observable disease (Fig. 4D). On average, surviving active TIP-treated hamsters (n = 14) underwent 6.1 days of clinical signs, while mock-treated animals underwent 13.4 days (n = 5). Intranasally treated hamsters had a significantly higher survival rate (63 to 80%; DI-07, P = 0.0008; DI-10, P = 0.0476; DI-14, P = 0.0085; DI-35, P = 0.0203) than mock-treated animals (25%) (Fig. 4E).

FIG 3 Legend (Continued)

and dashed line for those treated with inactive TIPs. (B) Clinical signs (scored from 0 to 10), with severity depicted by increasing intensity of red. Animals scoring ≥ 10 were humanely euthanized; any animals that succumbed to disease prior to euthanasia were allocated a score of 10. Gray boxes indicate the end of monitoring/scoring due to euthanasia/death. (C) The proportion of hamsters from each group that succumbed (euthanized due to clinical signs or found dead) to NiV disease (red), those that survived to end of study at 28 dpi with having displayed at least 1 day of clinical signs (blue), or with no clinical signs (green). (D) Number of days that any hamster surviving to study end (28 dpi) exhibited clinical signs. Closed circles represent animals treated with active TIPs; open circles, those treated with inactive TIPs; and black circles, mock treated. Individual animals are represented, with lines indicating mean and standard deviation. Significance calculated by multiple *t* tests; ****, $P \leq 0.0001$. (E) Survival of hamsters in each group. In groups containing treated hamsters, solid line represents active TIPs, and dashed line represents inactive TIPs. Significance calculated by log-rank (Mantel-Cox) test as follows: ****, $P \leq 0.0001$; ***, $P \leq 0.001$; *





FIG 4 Intranasal NiV challenge and treatment with active TIPs. Hamsters were inoculated i.n. with 100 μ L containing 10⁶ TCID₅₀ NiV strain Malaysia in combination with either 1 × 10⁸ active (treated with 100 mJ/cm² of UV) TIPs or no TIPs (mock treated). (A) Graphs represent the percent weight change from baseline (taken at -1 dpi) and body temperatures. Individual animals are represented. DI-07, red; DI-10, green; DI-14, blue; DI-35, orange; no TIP (mock treated), black; line indicates the mean value each day. (B) Clinical signs (scored from 0 to 10), with severity depicted by increasing intensity of red. Animals scoring \geq 10 were humanely euthanized; any animals that succumbed to disease prior to euthanasia were allocated a score of 10. Gray boxes (Continued on next page)



FIG 5 Protection from NiV disease in hamsters treated with different TIP/NiV ratios. Hamsters were inoculated i.p. with 2 mL DMEM containing 10⁴ TCID₅₀ NiV strain Malaysia in combination with different concentrations of active TIPs DI-14 (treated with 100 mJ/cm² of UV) or with no TIPs (mock treated). Final ratios of TIP/NiV were 10,000:1 (purple), 3,333:1 (green), 1,111:1 (orange), 370:1 (blue), 123:1 (cyan), and 0:1 (black). (A) Mean daily percent weight change from baseline (taken at -1 dpi) for each group. The weights of uninfected control hamsters are shown as daily means (dashed black line) and range (gray area). (B) Survival of hamsters in each group. Also shown with a dashed gray line are findings from a previous experiment (Fig. 2) in which hamsters were treated with active TIP DI-14 at 20,000:1 ratio for comparison. Significance calculated by log-rank (Mantel-Cox) test as follows: ***, $P \le 0.001$; **, $P \le 0.01$; *, $P \le 0.05$. (C) Clinical signs (scored from 0 to 10), with severity depicted a score of 10. Gray boxes indicate the end of monitoring/scoring due to euthanasia/death.

TIP treatment protects hamsters from death and clinical disease in a dosedependent manner. Previous data indicated that TIP-mediated inhibition of NiV *in vitro* was dependent on the TIP/NiV ratio (24). Here, we found that DI delivery *in vivo* also confers a significant protective effect at TIP/NiV ratio of 100:1 (for i.n. inoculation), even though no *in vitro* inhibition was observed at this ratio. To investigate the dose-dependent effect of TIP-mediated protection *in vivo*, we simultaneously administered TIPs and challenge virus i.p., as this route allowed a greater range of ratios to be investigated. We used DI-14, the best-performing candidate from previous *in vivo* experiments, to assess the efficacy of different TIP:NiV ratios from a maximum of 10,000:1 to a minimum of 123:1 in hamsters, alongside mock-treated (DMEM-only) control animals.

Clinical signs and mortality were greater in animals treated with lower TIP:NiV ratios than those treated with the original 20,000:1 ratio of TIP:NiV (Fig. 5A). Compared to mock-treated hamsters, significant differences in survival were conferred with treatment ratios of both 10,000:1 (P = 0.0030) and 3,333:1 (P = 0.0114), but not at the remaining lower ratios tested (Fig. 5B). While the two lower-delivery ratios (10,000:1 and 3,333:1) improved survival outcomes (per survival and/or mean time to death) compared to no treatment, they did not prevent clinical signs. Two of the three surviving animals in the highest ratio group tested (10,000:1) exhibited mild neurological signs over 1 to 3 days between 10 to 14 dpi but were free of clinical signs by study

FIG 4 Legend (Continued)

indicate the end of monitoring/scoring due to euthanasia/death. (C) Proportion of hamsters from each group that succumbed (euthanized due to clinical signs or found dead) to NiV disease (red) or survived to end of study at 28 dpi having displayed at least 1 day of clinical signs (blue) or no clinical signs (green). (D) Number of days that any hamster surviving to study end (28 dpi) exhibited clinical signs. Individual animals are represented, with lines indicating mean and standard deviation. Significance calculated by multiple *t* test; ****, $P \le 0.0001$. (E) Survival of hamsters in each group. In groups containing treated hamsters. Significance calculated by log-rank (Mantel-Cox) test as follows: ***, $P \le 0.001$; **, $P \le 0.01$; *, $P \le 0.05$.



FIG 6 Early interferon signaling and cytokine/chemokine expression in hamsters treated with active TIPs. Hamsters inoculated i.n. with 100 μ L DMEM containing 10⁶ TCID₅₀ NiV strain Malaysia in combination with either 1 × 10⁸ active TIPs (treated with 100 mJ/cm² of UV) or no TIPs (mock treated) were euthanized at 1 or 4 dpi (n = 5 each group). Lung tissues were collected for mRNA analysis. Heatmaps represent fold changes in transcript levels (compared to mock-treated hamsters at each day) of mRNA associated with genes from interferon induction and signaling or cytokine and chemokine signaling pathways. Intensity of color corresponds to magnitude of fold change over baseline.

completion at 28 dpi (Fig. 5C). The remaining animal exhibited persistent mild to moderate neurological signs from 8 dpi until study end (20 days total). Similarly, the 1 survivor from the 3,333:1 ratio group exhibited mild to moderate neurological signs from 6 dpi until study end (19 days total). These data indicate that reduction in clinical disease and decreased mortality require high delivery ratios to allow optimal TIP-mediated protection.

TIP treatment in hamsters results in strong but transient immunostimulatory response. To investigate the immunostimulatory properties of TIP treatment in the hamster model and to assess its potential contribution to protective efficacy, groups of 5 hamsters were treated i.n. with (i) active TIP only (either DI-07 or DI-14), (ii) a combination of active TIP (either DI-07 or DI-14) and 10⁶ TCID₅₀ NiV, or (iii) 10⁶ TCID₅₀ NiV alone. Hamsters were euthanized at 1 and 4 dpi, and RNA extracted from lung tissues was examined for differentially expressed immune genes in interferon (IFN; sensing, induction, and signaling) and cytokine/chemokine expression pathways.

As expected, due to well-characterized IFN antagonism by NiV (40, 41), upregulation of IFN or interferon-stimulated genes (ISGs) was not observed in infected untreated hamsters (NiV alone). In TIP-treated hamsters, type I and type II interferon (IFNA-23-like, IFNA-13-like, IFNA-9-like, IFNE, IFNG, and IFNK) were strongly upregulated at 1 dpi (Fig. 6). However, this was not seen in TIP-treated infected hamsters (TIP plus NiV). Since TIP propagation initially relies on viral replication, some level of cell-to-cell spread of virus is likely to occur early in infection. This spread may confer immunosuppressive activity, counteracting the TIPinduced upregulation seen in tissues from animals receiving TIP alone. Pattern recognition receptor (PRR) genes involved in the antiviral response (DDX58, DHX58, and IFIH1) and those associated with IFN-mediated effector pathways (MX1 and OAS2) were slightly upregulated (2- to 3-fold) in DI-14 treated, but not DI-10 treated, hamsters. This may be due to the structural differences between the two DI variants (copyback versus deletion) and may explain the small difference in treatment efficacy noted between the two. However, these effects were transient, with changes observed at 1 dpi largely absent at 4 dpi. In hamsters treated with TIP plus NiV, any differentially expressed (DE) genes were largely similar to those observed in the NiV-alone control animals. A similar pattern was true for genes involved in cytokine and chemokine expression. At 1 dpi, in hamsters treated only with active TIP, several genes involved in inflammatory pathways and response to infection (CCL1, CSF3, IL-10, IL-12A, IL-17A, and IL-2) were strongly upregulated, which was not seen in TIP plus NIV and NiV-only groups (Fig. 6). Again, these effects were transient and absent by 4 dpi, and at both time points, the DE gene pattern was largely similar in both TIP plus NiV and the NiV-alone control animals.

DISCUSSION

Here, we demonstrate that TIPs can provide robust protection in a lethal hamster model of NiV disease. A single-dose TIP treatment reduced mortality, and TIP-treated hamsters were less likely to exhibit clinical signs of infection. If present, clinical signs in TIP-treated animals were shorter in duration than signs in mock-treated animals. These results are encouraging compared to other reported NiV treatment options. While some treatments, including the antiviral compounds T-705 (42) and remdesivir (43) and the neutralizing antibodies m102.4 (44, 45) and h5b3.1 (46), have been shown to protect against disease and decrease mortality when given pre- and/or postexposure, they all require multiple doses to achieve protection. Here, we assess single-dose concurrent delivery of TIPs with virus challenge. Additional characterization of TIPs given prophylactically and postexposure will be key to determining optimal timing for TIP treatment alone or in conjunction with other interventions and will provide data to help develop putative treatment plans for patient care.

To date, few reports have investigated inhibitory effects of DIs in vitro and in vivo. Inoculation with a high dose of highly purified, uncharacterized vesicular stomatitis virus (VSV) DI particles protected against lethal VSV challenge in mice (47, 48). Similarly, Semliki Forest virus (SFV) preparations high in uncharacterized DI genomes protected against lethal SFV challenge in mice (49), and further studies using a single cloned characterized DI candidate also protected mice against SFV challenge (50). The majority of work investigating DI treatment in vivo centers on influenza A virus (IAV). IAV DI244 (deletion in genome segment PB2), given either prophylactically or postexposure, increased survival and reduced clinical signs in IAV-infected mice and ferrets (25, 27-29), and DIG-3 (deletions in genome segments PB2, PB1, and polymerase acidic [PA]) protected against IAV challenge in mice (26). In addition, DI244 protected against heterologous influenza B virus challenge in mice (30). Due to differences in DI genome structures, DI quantification methods, experimental processes, challenge doses, and viral pathogenesis, comparisons between these data and ours are difficult. However, these promising findings for influenza and our findings regarding NiV support broader development and application of DIs for other diverse pathogens.

Hamsters were protected from NiV challenge using both a high-TIP/NiV-ratio i.p. dose (~20,000:1) and a low-TIP/NiV-ratio i.n. dose (~100:1). Dimmock et al. achieved protection using a ratio of DI244:IAV between 100,000:1 to 1,000,000:1 (51), but their studies did not evaluate lower ratios. Our data demonstrate that lower doses of DIs can provide protective effects and that efficacious ratios determined *in vitro* may underestimate the treatment potential of lower doses. However, our data also show that protective ratios are associated with route of delivery and infection. For example, in hamsters, NiV disease progresses more rapidly, and time to death is shorter with i.p. challenge than i.n. challenge (31, 52). Intranasal challenge may therefore allow TIPs more time to exert their protective effects. Alternatively, i.n. administration may allow more targeted delivery of TIPs. Respiratory disease in NiV-infected hamsters is characterized by extensive viral replication in the respiratory tract and lungs (32, 33, 53). Direct delivery of TIPs to the cells where initial NIV infection occurs may thus facilitate protection at lower-ratio doses.

Mechanisms of TIP-mediated viral inhibition can be broadly divided into two categories, direct and indirect. Direct inhibition is a result of DI genomes interfering with replication of viral genomes, whereas indirect inhibition may be due to nonspecific immunostimulatory properties of the TIPs (54–58). Several inherent factors in the DI structure have been shown to influence their direct inhibitory potential, including DI length (17, 24, 59, 60), relative strengths of the replication promoter elements incorporated into the DI (61–63), and efficiency of DI genome packaging (64, 65). However, these structural features can also influence mechanisms related to indirect interference. For example, double-stranded hairpin loops formed by nonencapsidated copyback genomes are readily recognized by cytosolic innate immune pattern recognition receptors like RIG-I, MDA5, and LGP2 (66–69). Our data suggest that while both direct and indirect mechanisms play a role, direct interference is the predominant mechanism for protection by NiV TIPs. When the RNA component of TIPs was destroyed by UV irradiation, eliminating direct interference capacity (inactive TIPs), we saw a significant reduction in the protective effect of TIP treatment. Furthermore, of the candidates evaluated, copyback Dls conferred the highest levels of protection. Copyback Dls have two copies of the 5'-trailer sequence, which contains the stronger of the two *cis*-acting replication promoter elements found in the NiV genome. Therefore, copyback Dl genomes are generated at a higher rate than deletion Dis. They may thus exert greater direct inhibitory pressure on full-length viral genome replication and may also result in comparably higher, albeit still low, nonspecific immunostimulatory effects. Interestingly, the shortest copyback (DI-35) performed more poorly than the slightly longer copybacks (DI-07 and DI-14). The shorter length of DI-35 may have resulted in the removal of one or more elements involved in efficient packaging, suggesting that the optimal length of DI genomes is 1.0 to 1.5 kb.

Destruction of the RNA component of TIPs did not completely abolish their inhibitory potential. Inactive TIPs are structurally similar to classical VLPs, and protection from respiratory diseases via VLP vaccination is reported for respiratory syncytial virus (RSV) (70, 71), influenza (72), and SARS1 (73). The low-level protection we observed by TIPs in this "inactive" state was likely due to indirect stimulation of the innate immune system. "Inactive" TIPs are likely not completely inert and may still produce pathogenassociated molecular patterns (PAMPs) that can activate some level of innate immune response; this is supported by the observed upregulation of several pathways for IFN production and cytokine signaling in animals treated with TIPs alone. While fast-acting and strong for many genes, the immunostimulatory effects of the TIPs in our study were predominantly transient in effect, with a return to basal levels by 4 dpi. TIPs are presumed to be inert in the absence of parental virus, however, low-level replication activity may occur due to co-packaging of viral polymerase with DI genome. This activity may have led to the transient nature of the immunostimulation. As nucleotide sequences and motifs can be manipulated to enhance and prolong immunostimulatory effects (74, 75), DI genomes may be further optimized to increase the contribution of indirect inhibition, thereby improving the overall protective efficacy of TIPs.

Here, we provide support for continued studies of the clinical utility of TIP treatment. NiV is a highly lethal disease circulating in areas with exceptionally dense human populations. Developing a therapeutic candidate that is easily administered and achieves high levels of protection is both indicated and required to prevent human disease and to provide intervention options in outbreak response. Here, we provide the first evidence that TIPs can confer protection against disease and lethal outcome in a robust animal model of NiV, supporting further investigations of both treatment and therapeutic use of DI for NiV and other high-consequence pathogens.

MATERIALS AND METHODS

Biosafety. All work with infectious virus or infected animals was conducted in a biosafety level 4 (BSL-4) laboratory at the Centers for Disease Control and Prevention (CDC) following established BSL-4 standard operating procedures approved by the Institutional Biosafety Committee. All recombinant virus work was approved by the Centers for Disease Control and Prevention Institutional Biosafety Committee.

Cells and viruses. Vero-E6 and BSR-T7/5 cells were cultured in Dulbecco's modified Eagle's medium (DMEM) supplemented with 5% (vol/vol) fetal calf serum, nonessential amino acids, 1 mM sodium pyruvate, 2 mM L-glutamine, 100 U/mL penicillin, and 100 μ g/mL streptomycin. HSAEC-KT1 cells (ATCC, USA) were cultured in airway epithelial cell basal medium paired with the bronchial epithelial cell growth kit medium (both ATCC, USA). NiV strain Malaysia (NiV-M; GenBank accession no. AF212302) was originally obtained from a clinical isolate, passaged once on Vero-E6 cells for isolation from clinical sample, and further amplified on Vero-E6 cells. Viral titers were calculated as 50% tissue culture infective dose (TCID₅₀) in Vero-E6 cells (76). All viral stocks were verified by next-generation sequencing and confirmed to be mycoplasma free.

TIP production. TIPs were produced as previously described (Fig. S1 in the supplemental material) (24). To generate these virus-like particles containing the DI genomes, a modified NiV reverse genetics rescue system (40) was developed in which a plasmid transcribing the DI genome plus support plasmids expressing the proteins required for transcription and genome replication were transfected into BSR-T7/

5 cells. After 2 days, the transfected cells were overlaid with Vero-E6 cells and infected with a recombinant NiV-M-expressing ZsG fluorescent protein (rNiV-M/ZsG) in order to supply the necessary proteins for assembly and release of the virus-like particles. To remove the infectious rNiV-M/ZsG from the final product, a precise dose of UV radiation was used (100 mJ/cm² of UV radiation [CX-2000 Crosslinker; UVP] in a 6-well plate [1 mL per well]), which has been previously shown to result in breakage of full-length RNA genomes (uncoupling the leader and trailer *cis*-acting elements and therefore removing the ability to replicate full-length genome) while leaving the shorter DI genomes intact (24). The resulting products were termed active TIPs. The TIPs were quantified using a DI genome-specific digital droplet quantitative reverse transcription-PCR (RT-qPCR) assay; the above method produced an average of approximately 1 \times 10° DI genome copies/mL. To create stocks depleted of both the producer virus and DI genome (termed inactive TIPs), stocks were treated with 10 consecutive treatments of 240 mJ/cm² of UV radiation (2,400 mJ/cm² total).

Digital droplet quantitative reverse transcription-PCR. A $20 \times \text{mix}$ of DI-specific or NiV-specific primers and probes (18 μ M each primer and 5 μ M probe) were used in conjunction with 1-Step RT-digital droplet PCR (ddPCR) advanced kit for probes and droplet generation oil for probes (both from Bio-Rad). Duplex RT-qPCR assays were set up and run per manufacturers' conditions, with droplets generated using QX200 droplet generator (Bio-Rad). Results were analyzed using QX200 droplet digital PCR system, and quantification data were generated using QuantaSoft software (both from Bio-Rad).

Hamster studies. All animal experiments were approved by the CDC Institutional Animal Care and Use Committee and performed in an AAALAC International-approved facility. Data are based on four independent hamster studies using 6-week-old HsdHan:Aura Syrian hamsters (Envigo; catalog no. 8903F or 8903M). The DI dose-effect study used experimental groups of 8 to 10 hamsters (half female and half male). All other hamster studies used experimental groups of 5 to 10 female hamsters. Hamsters were inoculated either intranasally (i.n.; 100 µL divided bilaterally) or intraperitoneally (i.p.; 2 mL) with Dulbecco's modified Eagle's medium (DMEM) alone, NiV strain Malaysia (NiV-M) in DMEM, or a mixture of NiV-M and TIP in DMEM. Hamsters were kept in a climate-controlled laboratory with a 12-h day/night cycle, provided Teklad global 18% protein rodent diet (Envigo) and water ad libitum, and group housed on corn cob bedding (Bed-o'Cobs, ¼ in.; Anderson Lab Bedding, Maumee, OH, USA) with cotton nestlets and crinkle paper in an isolator caging system (Thoren Caging, Inc., Hazleton, PA, USA) with a HEPA-filtered inlet and exhaust air supply. Microchip transponders (BMDS; IPTT-300) were placed subcutaneously in the interscapular region at -4 dpi for individual identification and to assess body temperature. Baseline weights were taken -1 dpi, and hamsters' weight change, body temperatures, and clinical signs were assessed daily. Animals were scored by the following criteria: quiet, dull responsive, hunched back/ruffled coat, hypoactivity, mild neurological signs, each 2 points; abnormal breathing (i.e., increased respiratory rate, dyspnea), hypothermia (<34°C), moderate neurological signs, each 5 points; and paralysis, frank hemorrhage, moribund, weight loss >25% of baseline (measured at -1 dpi), severe neurological signs, each 10 points. Neurological signs were classified as (i) mild, with abnormal gait or movement and/or mild head tilt (\sim 0 to 30° from vertical or sporadic); (ii) moderate, with tremors, ataxia, circling, absence seizures, and/or moderate head tilt (\sim 30 to 90° from vertical and persistent) while retaining the ability to walk, feed, and drink; or (iii) severe, with limb paralysis, tonic clonal seizure, inability to return to normal posture, and/or severe head tilt (>90° from vertical). Hamsters were euthanized with isoflurane vapor when they met euthanasia criteria (score \geq 10) or at the completion of the study (28 dpi). Survival data from the 10 hamsters challenged with 10^5 TCID_{50} i.p. in Fig. 1B were previously reported unvaccinated control animals (38).

Quantitative RT-PCR. Viral RNA was extracted from homogenized tissue samples and whole blood using the MagMAX-96 total RNA isolation kit (Thermo Fisher Scientific) on a 96-well ABI MagMAX extraction platform with a DNase I treatment step according to the manufacturer's instructions. RNA was detected using a RT-qPCR assay targeting the NiV NP gene sequence (NiV forward, 5'-CTGGTCTCTGCAGTTATCACCATCGA-3'; NiV reverse, 5'-ACGTACTTAGCCCATCTTCTAGTTTCA-3'; and NiV probe, 5'-FAM-CAGCTCCCGACACTGCCGAGGAT-BHQ1-3'; all from IDT), with levels normalized to 18S RNA values using a commercial endogenous control assay (Thermo Fisher). Genome copy numbers were determined using standards prepared from *in vitro*-transcribed NP RNA.

Statistics and graphing. Survival statistics were calculated using the log-rank (Mantel-Cox) test. Significance in duration of observable clinical signs was calculated using multiple *t* tests. All graphs and statistical analyses were created in GraphPad Prism (v9.3.0).

SUPPLEMENTAL MATERIAL

Supplemental material is available online only. FIG S1, TIF file, 0.6 MB. FIG S2, TIF file, 1.4 MB. FIG S3, TIF file, 1.2 MB.

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We have no reported conflicts of interest.

The findings and conclusions in this report are those of the authors and do not necessarily represent the official position of the Centers for Disease Control and Prevention.

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