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A lung-targeted therapeutic siRNA against highly conserved viral M1 mRNA effectively limits highly pathogenic influenza A infection in mice

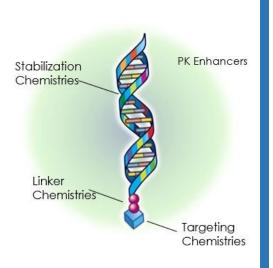
Keivan Zandi¹, Casi Schienebeck¹, Zhao Xu¹, Alireza Saeidi¹, Nicholas Sheets¹, Tingting Yuan¹, David Kasahara¹, Pinghan Huang², Cassio Pontes Octaviani², Jason Hsu³, Bi-Hung Peng⁴, Chien-Te Kent Tseng⁵, James Hamilton¹, Tao Pei¹, Erik W. Bush¹ . Arrowhead Pharmaceuticals, 2. Department of Microbiology and Immunology, The University of Texas Medical Branch, Galveston, Texas, 3. Department of Biochemistry and Molecular Biology, The University of Texas Medical Branch, Galveston, Texas, 4. Department of Neuroscience/Cell Biology/Anatomy, The University of Texas Medical Branch, Galveston, Texas, 5. Departments of Microbiology and Immunology, Neuroscience/Cell Biology/Anatomy, and Pathology, Centers for Biodefense and Emerging Diseases, The University of Texas Medical Branch, Galveston, Texas, 5. Departments of Microbiology and Immunology, Neuroscience/Cell Biology/Anatomy, and Pathology, Centers for Biodefense and Emerging Diseases, The University of Texas Med Branch, Galveston, Texas.

RATIONALE

- Human infections of highly pathogenic avian influenza A (bird flu) are rare but represent a significant global health concern due to the continued widespread circulation of the H5N1 virus in wild bird, poultry and more recently different mammalian populations.
- Therapeutic small interfering RNAs (siRNAs) offer an alternative approach by using the RNAi mechanism to silence viral transcripts encoding essential gene products.
- Here, an evaluation of siRNAs directed against highly conserved influenza A viruses (IAV) including highly pathogenic avian influenza (H5N1) genomic regions demonstrated a unique viral vulnerability to M1 silencing.

TRIMTM PLATFORM

Optimized therapeutic siRNAs (RNAi triggers) for specific target gene silencing and limited potential offtarget interactions.



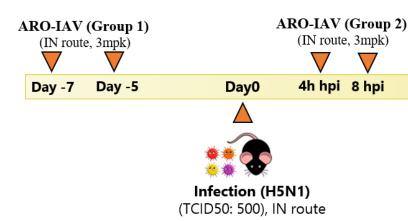
- Rational use of modifying chemistries maximize innate stability and potency.
- Integrin $\alpha v\beta 6$ targeting moiety facilitates delivery to pulmonary epithelium.

METHODS

- Nine-week-old BALB/C mouse model of H5N1 (A/whooper swan/Mongolia/244/2005 strain) infection was used to assess in vivo efficacy of influenza M1-silencing siRNA conjugates.
- Prophylactic (pre-infection) and therapeutic (post-infection) efficacy was evaluated in two groups of mice (Groups 1 and 2)
- G1 mice received two intranasal doses (3 mg/kg each) of siRNA conjugate or vehicle (saline) on day 7 and 5 prior to infection.
- H5N1 infection was initiated by intranasal inoculation of virus.
- G2 infected mice received two intranasal doses (3 mg/kg each) of siRNA conjugate or vehicle (saline) at 4- and 8-hours postinfection.
- Lung tissue was collected 6 days post-infection for viral load analyses (TCID50 and qRT-PCR assays) and histopathological assessments.
- A subset of mice from each treatment group was reserved for clinical scoring of morbidity and mortality.
- Histological analysis of pulmonary inflammation was performed using hematoxylin and eosin (H&E) staining.

RESULTS

- Previous work evaluated over 200 candidate siRNA sequences targeting the most highly Conference, May 2023).
- influenza.



viral load, M1 mRNA expression in H5N1-infected mouse lung

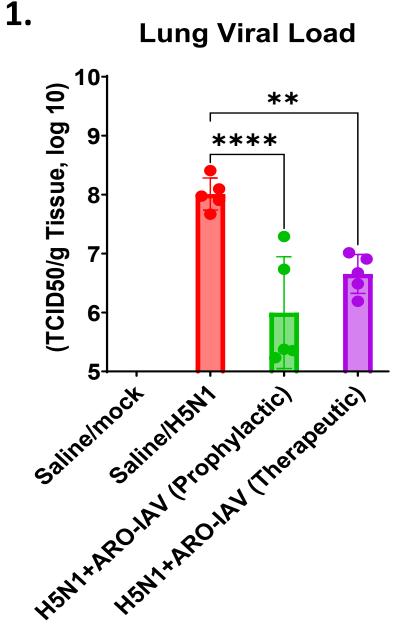


Figure 1. ARO-IAV treatment demonstrated prophylactic efficacy (reducing H5N1 viral load in the lung by >2 log of 10 vs untreated infected controls) and therapeutic efficacy (reducing H5N1 viral load in the lung by \approx 1.5 log of 10 vs untreated infected controls). Figure 2. ARO-IAV treatment reduced lung H5N1 genomic RNA and mRNA expression by approximately 1 log of 10 or greater compared to untreated infected controls.

H5N1 infection conserved regions of approximately 10,000 IAV genomes of different subtypes (H1N1, H3N2, Figure 3. Survival index showed that ARO-3. H5N1, H5N6, H5N8, H7N2, H7N3, H7N4, H7N7, H7N9 and H9N2) (Presented at 7th ISIRV IAV treatment significantly improved the survival rate among the infected-treated Saline/mock ARO-IAV, an siRNA conjugate targeting a conserved region of the influenza A M1 gene, has 90-Saline/H5N1 mice while by day 10 post-infection, 100% 80been previously shown to have potent antiviral activity against H1N1 and H3N2 viruses. H5N1+ARO-IAV mortality was observed in untreated Survival (%) 70-Here, we extend observations of ARO-IAV antiviral activity to highly pathogenic **H5N1** avian (Prophylactic) 60infected mice. At 14 days post-infection, H5N1+ARO-IAV 50-(Therapeutic) 80% of mice receiving therapeutic ARO-IAV 40treatment survived and 60% of mice 30-**Daily Observation** 20receiving prophylactic ARO-IAV treatment 10 Day14 poi Day6 poi survived. 10 15 **Days post-infection Tissue Collection** 4. (Viral Load and Histopathology) Saline/mock Silencing M1 expression with intranasally delivered ARO-IAV significantly reduces **Figure 4.** Clinical score index showed that Saline/H5N1 Score 3 treatment with ARO-IAV could improve the H5N1+ARO-IAV (Prophylactic) clinical score in mice. However, clinical H5N1+ARO-IAV Clinical 2. score improvement was more significant (Therapeutic) withing the mice group which received the Lung viral RNA copies ARO-IAV in prophylactic mode as all remain copies/μg RNA active and considered recovered from the disease. 0 1 2 3 4 5 6 7 8 9 101112131416 Days post infection 9. 5. 1101 ral Saline/mock Log10 vii Saline/H5N1 Figure 5. Body weight index showed that, H5N1+ARO-IAV Weight (%) infected mice receiving therapeutic or (Prophylactic) prophylactic ARO-IAV treatment lost less H5N1+ARO-IAV (Therapeutic) weight than untreated infected controls. H5N1+AROIAN (Prophylactic) H5N1+ARO-IAV [Therapeutic] 90 However, weight loss prevention was more pronounced with ARO-IAV prophylaxis. 0 1 2 3 4 5 6 7 8 9 101112131416 Days post infection

CONCLUSION

We demonstrate a vulnerability of H5N1 avian influenza to silencing of M1, an essential viral gene product that has been challenging to drug with traditional small molecule antivirals. ARO-IAV, an inhaled M1-silencing therapeutic siRNA, offers a novel first-in-class approach for the

potential treatment of highly pathogenic influenza A infections.

Intranasal ARO-IAV significantly improves survival and clinical score in a model of severe