

Researchers resurrect ancient viruses in hopes of improving gene therapy

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A cartoon depicting Adeno-associated viral capsid in formation by ancestral sequence reconstruction. Credit: Eric Zinn.

Researchers have recreated the evolutionary lineage of adeno-associated viruses (AAVs) to reconstruct an ancient viral particle that is highly effective at delivering gene therapies targeting the liver, muscle, and retina. This approach, published July 30 in *Cell Reports*, could be used to design a new class of genetic drugs that are safer and more potent than those currently available.

"Our novel methodology allows us to understand better the intricate structure of viruses and how different properties arose throughout evolution," says senior study author Luk H. Vandenberghe of Harvard Medical School. "We believe our findings will teach us how complex biological structures, such as AAVs, are built. From this knowledge, we hope to design next-generation viruses for use as vectors in gene therapy ."

Viruses need to efficiently transfer their genetic material into host cells in order to replicate and survive. Researchers have taken advantage of this natural property to develop viral vectors, or carriers, capable of shuttling therapeutic genes to the appropriate cells or tissues.

Early-stage clinical trials have demonstrated the safety and effectiveness of this approach for treating inherited blindness and hemophilia. But so far, AAVs used for gene therapy have been chosen from naturally circulating viral strains, which patients may already have been exposed to, which means they would have natural immunity. Because natural immunization blocks the transfer of the therapeutic gene, these individuals are often ineligible for gene therapy.



The generation of AAV vectors that reliably evade the host immune system is pivotal for the long-term success of this promising therapeutic approach. However, efforts to engineer improved AAVs have been stymied by the intricate structure of these viruses. Like pieces of a jigsaw puzzle, every protein in the shell of the virus must fit together perfectly for the virus to function normally. Altering proteins in one part of the virus to achieve a certain benefit, such as more efficient gene transfer or reduced recognition by host immune cells, could end up destroying the structural integrity of the entire shell.





Retinal targeting of retina by Anc80. Credit: Livia Carvalho

To overcome this challenge, Vandenberghe and his team at the Grousbeck Gene Therapy Center at the Massachusetts Eye and Ear Infirmary turned to evolutionary history for guidance. Under selective pressure, AAV ancestors have undergone a series of changes that altered



viral function while preserving the structural integrity and other core viral functions. "Analysis of these evolutionary intermediates provides insights into how to uncouple important vector properties to build safer and improved gene therapy vectors," says first author Eric Zinn of the Schepens Eye Research Institute.

Using computational techniques, the researchers analyzed amino acid sequences of contemporary AAVs and used this information to predict the ancestral amino acid sequence for the AAV protein shell. This analysis allowed them to reconstruct in the laboratory nine synthetic predecessors for AAVs that are currently undergoing clinical testing. The most ancient of those, named Anc80, was extensively tested for applications in gene therapy. When injected into mice, Anc80 successfully targeted the liver, muscle, and retina, with levels of gene transfer equivalent or superior to those of contemporary AAVs used in clinical trials. Moreover, the ancestral virus did not produce any toxicity and was less susceptible to the immunity induced by AAVs currently circulating in nature. "Our work is the first demonstration of a computationally designed and fully synthetic viral vector for gene therapy," Vandenberghe says.







Computational design and synthetic construction of AAV gene therapy vector is shown. Credit: Eric Zinn

According to Vandenberghe, generating novel infectious agents may raise some concerns, but several safety measures have been taken. For example, AAVs are not known to cause any diseases, and their replication machinery is crippled prior to their use in gene therapy. "With the safeguards that have been put in place, we believe our contained approach to achieve a therapeutic gene delivery vehicle with beneficial properties and a distinct immunological profile to existing technology is not less or more unethical or dangerous as compared to other efforts to discover <u>viral vectors</u>," he says.

In future studies, the researchers will characterize the interplay between the virus and host through evolution and continue to seek improved vectors for clinical applications. They will also examine the potential of Anc80 for treating liver diseases and retinal forms of blindness. "The vectors developed and characterized in this study demonstrate unique and potent biology that justify their consideration for gene therapy applications," Vandenberghe says.

More information: *Cell Reports*, Zinn et al. "In Silico Reconstruction of the Viral Evolutionary Lineage Yields a Potent Gene Therapy Vector" <u>dx.doi.org/10.1016/j.celrep.2015.07.019</u>

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