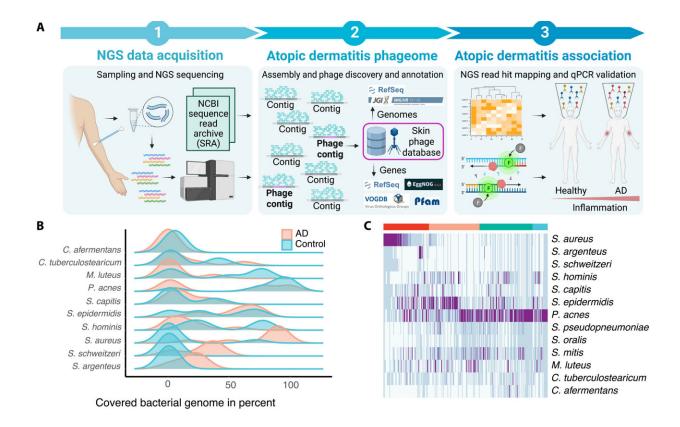


# Atopic dermatitis: Viruses discovered as new therapy option

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Study workflow and bacterial communities in healthy skin and AD. (A) Study workflow. (B) Ridge plot displaying the distribution of samples depending on the fraction of covered bacterial genome. x axis is the fraction bacterial genome covered (0% indicates that the bacterial genome was not present in the sample, while 100% indicates that the entire bacterial genome was covered). Each subpanel represents a bacterial genome. y axis is the proportion of patient samples that showed the indicated genome coverage rate. Pink are AD samples, and turquoise are healthy controls. (C) Heatmap of all reads mapped against the top abundant skin bacteria and Staphylococcus strains frequently found in AD



Samples are ordered according to an arbitrary clinical inflammation score, where red indicates high inflammation and blue indicates no inflammation in patients with AD or healthy controls. Credit: *Science Advances* (2023). DOI: 10.1126/sciadv.adg4015

Up to 15% of children and 5% of adults are affected by the chronic inflammatory skin disease atopic dermatitis. Despite advanced therapy measures, the severe itching and eczema, especially on the elbows or knees, cause great distress to the patients.

In the course of a study conducted at MedUni Wien a research team led by Wolfgang Weninger, Head of the Department of Dermatology, has discovered a new approach: bacteriophages, which colonize the skin as viral components of the <u>microbiome</u> and can drive the development of innovative atopic dermatitis therapies. The research results were recently published in the journal *Science Advances*.

Until now, the importance of bacteriophages ("bacteria eaters," also called phages) in the <a href="https://www.human.com/huma

The microbiome is the sum of all microorganisms on the skin and has been primarily investigated for its bacterial constituents. It has been unknown whether viruses also contribute to the nature of the bacterial microbiome in healthy and diseased skin. Phages are viruses of different types and functions whose sole aim is to infect bacteria, thereby either destroying them—or stimulating them to multiply.



## New phages identified

"In our study, we discovered previously unknown phages in the microbiome of the skin samples of AD patients, which help certain bacteria to grow faster in different ways," note first authors Karin Pfisterer and Matthias Wielscher from the Department of Dermatology at MedUni Vienna.

The resulting shift in the balance between phages and bacteria was not detected in the comparative samples from healthy individuals and may be one explanation for the overpopulation of the skin microbiome with bacteria called Staphylococcus aureus found in AD.

These findings contribute significantly to a better understanding of the skin bioflora in AD patients and pave the way for the development of new targeted therapeutic interventions: By identifying and culturing phages specialized for Staphylococcus aureus, a promising new option is available.

## **Specialists for targeted therapy**

Bacteriophages are found not only in the body, but in every habitat populated by bacteria. There are 10<sup>31</sup> different phage species. One of their characteristics is that they prove to be extremely specific when it comes to choosing their target of infection: Most <u>phages</u> specialize in a particular genus, and in many cases in only a single species of bacteria.

While that makes it a challenge for scientists to identify the type of phage needed for a particular purpose, it also enables them to use them in a targeted manner. Bacterial viruses do not distinguish between antibiotic-resistant and other bacteria; thus they are being researched as a possible weapon in the fight against multi-resistant pathogens. Further



studies are now planned to confirm phage therapy for topical use in <u>atopic dermatitis</u>.

**More information:** Matthias Wielscher et al, The phageome in normal and inflamed human skin, *Science Advances* (2023). <u>DOI:</u> 10.1126/sciadv.adg4015. <u>www.science.org/doi/10.1126/sciadv.adg4015</u>

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