

The virus detectives—The secret of brown trout dying uncovered

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Dr. Jens-Eike Taubert examining a trout infected by a previously unknown virus, which is related to a virus that infects salmon in the North Atlantic and the Pacific. Credit: Chair of Zoology / Technical University of Munich

Every summer in Southern Germany, Austria and Switzerland, tons of brown trout perish. An interdisciplinary team from the Technical University of Munich (TUM) has now discovered the culprit of the mysterious dying afflicting brown trout. It is triggered by a previously unknown virus, which is related to a virus that infects salmon in the North Atlantic and the Pacific.

The setting is always in the same river sections. And the victims are always brown trout. Within a matter of days, their skin turns dark and the animals perish. Researchers and authorities have been hunting down the cause of this mysterious mass mortality for decades—unsuccessfully, until now.

Using cutting-edge analysis methods, Prof. Ralph Kühn and a [research team](#) from the Technical University of Munich have now identified the pathogen that triggers the proliferative darkening syndrome (PDS) in brown trout.

From the river to the lab

"The biggest challenge was identifying a pathogen nobody knew. At the outset, it was not even clear whether it was a bacterium, a [virus](#), a parasite or an environmental toxin," recalls the biologist.

Ten years of scientific detective work followed. The researchers set up two experimental stations at the Iller river—one in the headwaters, near Obersdorf, where the [fish](#) disease had never been observed; a second one in the lower course near Kempten, where brown trout die every summer.

At both stations, the researchers set up aquariums fed with river water. From May to September they observed the fish in the aquariums and took [tissue samples](#) that were frozen and sent to the TUM laboratory for analysis.

Searching for clues with next generation technology

The observations indicated that PDS advances in three phases. Initially, the fish look healthy. Then changes in internal organs such as the liver and kidneys appear. In the third phase, the brown trout skin turns dark and the animals die shortly thereafter. "The progression of the disease quickly led us to suspect that PDS was a viral disease," says Kühn.

To detect the virus in the samples, the researchers deployed a set of modern molecular genetic procedures, known as next generation technology. Genomes can be analyzed rigorously using these procedures.

First, the progression of the disease in the fish was characterized at the gene response level. Subsequently, the researchers looked at animals with a similar immune response to determine the [nucleotide sequence](#) of all inherited [genetic material](#)—of the fish and its possible pathogens.

The needle in a genetic haystack

The researchers deployed computer programs to pinpoint the pathogenic virus in this mountain of genetic data. Using deep bioinformatic processing, they successfully identified parts of the pathogen's genetic profile and compared them to the profiles of known viruses.

Extended, virus-specific sequencing led to the determination of the viral genome. The result: The culprit behind the demise of the brown [trout](#) is a piscine reovirus. It is related to the virus that infects salmon in the North Atlantic and the Pacific, where it causes significant economic damage.

In the next step, the [researchers](#) hope to grow the virus in the laboratory and investigate why it only occurs in certain sections of the Alpine rivers

and to what extent global fish trade facilitates its spread.

More information: Ralph Kuehn et al, Identification of a piscine reovirus-related pathogen in proliferative darkening syndrome (PDS) infected brown trout (*Salmo trutta fario*) using a next-generation technology detection pipeline, *PLOS ONE* (2018). [DOI: 10.1371/journal.pone.0206164](https://doi.org/10.1371/journal.pone.0206164)

Provided by Technical University Munich

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