

Environmental DNA in rivers offers new tool for detecting wildlife communities

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Ecologists in England, Scotland and the University of Massachusetts Amherst, report on a new method of identifying an 'entire community of mammals,' including elusive and endangered species such as red deer that are otherwise difficult to monitor, by collecting DNA from river water. Credit: UMass Amherst/Joseph Drake



Ecologists in England and Scotland, collaborating with ecologists Christopher Sutherland and Joseph Drake at the University of Massachusetts Amherst, report this week on a new method of identifying an "entire community of mammals"—including elusive and endangered species that are otherwise difficult to monitor—by collecting DNA from river water.

"Some <u>mammal species</u> are notoriously difficult to monitor," says environmental conservation Ph.D. student Drake. He adds that traditional survey methods are often tailored to a specific species, and therefore don't guarantee the detection of many other important species that are also present. Camera traps have improved the way conservation scientists study wildlife, but environmental DNA (eDNA) methods may offer a monitoring tool that could revolutionize conservation and ecology research, Sutherland adds, but the method required testing.

He adds, "We knew the potential of eDNA was massive, but when it comes to conservation, it is extremely important that we validate new approaches, and that's what we set out to do in this study." Details of their international collaborative work are in the *Journal of Applied Ecology*.

Naiara Sales of the University of Salford, U.K. and UMass's Drake led this study, in collaboration with researchers from the University of Tromsø, Norway, the universities of Aberdeen, Hull and Sheffield, and Liverpool John Moores University in the U.K. The research takes advantage of the fact that DNA shed from animals, either directly in the water or washed into the river, provides a snapshot of the local mammal community.

Mammologist Allan McDevitt of the University of Salford points out, "We currently use many ways of detecting and monitoring mammals, from looking for signs such as footprints or feces, to using <u>camera traps</u>



to take photos of them over several weeks. Now, we may just simply need to collect a few bottles of water and take it to the laboratory and look at the DNA we find."

To test this, the researchers collected water and sediment from streams and rivers in Scotland and England. They found DNA from over 20 wild British mammals and compared the results to historical records, field signs such as fecal samples and cameras. They report that eDNA "provided a similar or better performance in detecting water voles, for example, when compared to looking for water voles using field signs or cameras."

They add that accurately assessing the conservation status and distribution of mammals is increasingly important as many species' populations decline worldwide. Further, surveys using traps, trail cameras and fields signs are time-consuming and costly.

Collaborators for several studies, McDevitt's group, collaborating with Sutherland's group at UMass Amherst, as well as at the universities of Aberdeen and Hull, now believe that using water bodies is an effective way of capturing all the mammals present within a watershed. McDevitt says, "We are always looking for ways to improve biodiversity assessments and monitoring, and we need to find methods which can be applied universally and cost-effectively."

He adds, "We have demonstrated that environmental DNA collected from water bodies is effective for providing us with information about the presence or absence of mammals of <u>conservation</u> concern. This could be used at national levels for monitoring declining or recovering populations, or the early detection of harmful invasive species."

More information: Naiara Guimarães Sales et al, Fishing for mammals: Landscape-level monitoring of terrestrial and semi-aquatic



communities using eDNA from riverine systems, *Journal of Applied Ecology* (2020). DOI: 10.1111/1365-2664.13592

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