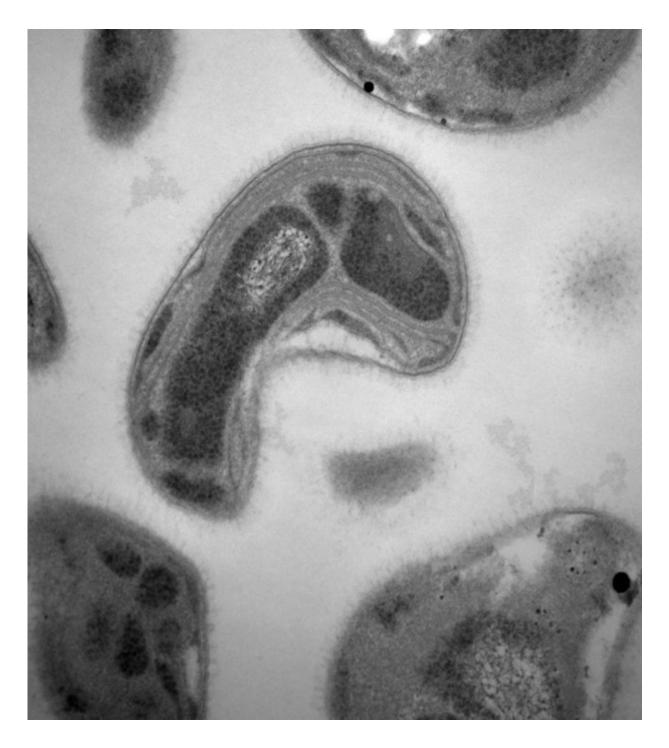


Newly identified microbial process could reduce toxic methylmercury levels

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An ORNL-led team along with University of Michigan and Iowa State University investigated many different methanotrophs to single out how and which ones take up and break down methylmercury. Credit: Jeremy Semrau, University of Michigan



A team led by the Department of Energy's Oak Ridge National Laboratory has identified a novel microbial process that can break down toxic methylmercury in the environment, a fundamental scientific discovery that could potentially reduce mercury toxicity levels and support health and risk assessments.

Methylmercury is a neurotoxin that forms in nature when mercury interacts with certain microbes living in soil and waterways. It accumulates at varying levels in all fish—particularly large predatory fish such as tuna and swordfish—and, when consumed in large quantities, can potentially cause neurological damage and developmental disorders, especially in children.

A previous ORNL-led study, <u>published in *Science* in 2013</u>, unlocked the genetic code that led scientists to accurately identify microbes responsible for methylmercury production in the environment. Following this finding, the ORNL team has now discovered which bacteria perform the reverse process, called demethylation. Details are published in *Science Advances*.

"Much attention has focused on mercury methylation or how methylmercury forms, but few studies to date have examined microbial demethylation, or the breakdown of methylmercury at environmentally relevant conditions," said Baohua Gu, co-author and a team lead in ORNL's Mercury Science Focus Area.

Bacteria called methanotrophs feed off methane gas and can either take up or break down methylmercury, or both. Methanotrophs are widespread in nature and exist near methane and air interfaces, and both methane and methylmercury are usually formed in similar anoxic, or oxygen-deficient, environments.

To single out how and which methanotrophs perform demethylation, the



ORNL-led team—along with methanotroph experts from the University of Michigan and Iowa State University—investigated the behavior of many different methanotrophs and used sophisticated mass spectrometry to analyze methylmercury uptake and decomposition by these bacteria. They discovered that methanotrophs such as *Methlyosinus trichosporium* OB3b can take up and break down methylmercury, while others such as *Methylococcus capsulatus* Bath only take up methylmercury.

In either case, the bacteria's interactions can lower mercury toxicity levels in water.

"If proven environmentally significant through future studies, our discovery of methanotrophs' behavior could be a new biological pathway for degrading methylmercury in nature," Gu said. This approach differs greatly from a previously recognized enzymatic pathway, which is only effective at very high mercury concentrations.

The methanotrophs identified in this study "open new opportunities to explore how nature detoxifies <u>methylmercury</u> and could improve our prediction of mercury toxicity levels and support better <u>risk assessments</u> and remediation efforts at mercury contamination sites," Gu added.

More information: X. Lu el al., "Methylmercury uptake and degradation by methanotrophs," *Science Advances* (2017). DOI: 10.1126/sciadv.1700041, advances.sciencemag.org/content/3/5/e1700041

Provided by Oak Ridge National Laboratory

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