

★ NEWS FEATURE

Can microbes keep time for forensic investigators?

Forensic scientists are building a “clock” from the bacteria and other microscopic scavengers that make up the postmortem microbiome. But how reliably will it tick?

Carolyn Beans, *Science Writer*

On a small hill outside of Grand Junction, CO, the sun beats on dry earth, and human bodies lie face up amid low sagebrush. Inside each corpse, on it, and beneath it, a rich ecosystem of bacteria, nematodes, and other microbes flourishes. As these tiny forces of nature—constituents of the postmortem microbiome—steadily decompose the remains, scientists document their every move.

At Colorado Mesa University’s Forensic Investigation Research Station (FIRS), researchers are studying these bodies—now up to 11—to learn about the waves of microbes that bloom and fade at each stage of human decomposition.

With advances in microbiology and genomics and the declining costs of DNA sequencing, microbes are

poised to play a big role in forensics. Researchers have shown that microbes have the potential to alter toxicology results, point to causes of death, and even place a suspect at the scene of a crime (1). If researchers can identify collections of microbes that predictably turn up at specific time points as a body decomposes, then this “microbial clock” could become a powerful tool for crime scene investigators looking to estimate the time and even the circumstances of death.

Recent studies suggest that a microbial clock may well exist. But before this forensic tool can be deployed at crime scenes and in courtrooms, researchers first must prove that the clock ticks reliably whether a body falls in lush forest or dry field, whatever the season.



In a fenced-in outdoor research area just beyond the laboratory at the Forensic Investigation Research Station near Grand Junction, CO, human remains are allowed to decompose so researchers can study the ebb and flow of resident microbes. Close-up photographs are not allowed due to privacy concerns. Image courtesy of Melissa Connor (Colorado Mesa University, Grand Junction, CO).

Published under the [PNAS license](#).

that allows them to study entire pools of microbes at once. For example, they use the DNA sequences of a bacterial gene known as 16S rRNA to identify and gauge the relative abundance of species in each sample and a gene known as 18S rRNA to do the same for eukaryotic microbes such as nematodes and amoebas. Pieter Dorrestein of the University of California, San Diego, is now gearing up to conduct a metabolomics analysis on the samples to pick up the chemical signatures of microbes in the process of decomposing remains. "We think that it is very likely that as these microbial communities are going through a succession, the byproducts they're producing are also going to be changing," says Metcalf.

In all this data, Metcalf believes she'll pick up at least one reliable signal of microorganisms keeping time. It probably won't come, she says, in the form of a single metronomic microbe or even a series of individual species that pop up at different time points. She expects the clock will be made from groupings of microbial species that rise and fall in a complex, yet predictable fashion over time. "The power here is that we are not looking at individual species," Metcalf says.

Steady Ticking

The first bodies were placed at the three sites in the spring of 2016, and Metcalf expects to publish results from that season soon. Based on pilot studies and animal work, she has reason to be hopeful.

In 2011, Metcalf followed the microbiomes of decomposing mice over 48 days and found dramatic shifts in the microbial community that occurred at about the same time points for each mouse (4). The results were so consistent that Metcalf could use the microbiome to accurately estimate the time since death to within approximately 3 days, even at 48 days out. "It was a surprise to me even at that point how impressively repeatable the microbial succession during decomposition is across different mice with all of the variables controlled," she says.

In 2012, Metcalf placed mice on different soil types (5). Most of the decomposing microbes came from the soil, she notes, but changing the soil type did not affect the pace of the clock. "That was a real surprise," says Metcalf. "Certainly you could see that the biggest driver of change was time."

Metcalf had also studied four human cadavers at the Texas facility, one pair placed in winter and one pair in spring of 2013 (5). After she controlled for seasonal temperature and how it would affect microbial activity, Metcalf found the microbial successions were remarkably consistent across different human bodies and different seasons. Although humans and mice don't share the exact same postmortem microbial communities, microbes from the same families emerged at similar times in both species. The human microbial clock, it seemed, ticks much like that of the mouse.

In the case of Metcalf's current work in humans, the microbial species that make up the postmortem microbiome appear to vary at least in part by geographic area. But she's betting enough microbial signatures will remain constant. "Not every microbe in a

community is equally important in a model," Metcalf says, noting that she employs machine learning to find the steady signals in the noise.

Flies in the Ointment?

To fine-tune models, researchers may also have to account for the diversity of the postmortem microbiome, tracking its multitude of contributors. At least some of these microbes are opportunistic members of the living-human microbiome (5). "Most likely there are some microbes that are just waiting for us to keel over," Metcalf says. Soil microbe populations are geographically diverse, too, potentially affecting which organisms are available to colonize a new corpse (6).

In a previous collaboration with Bucheli, Metcalf found that at least a small percentage of the postmortem microbiome may also be deposited by flies colonizing a body (5). In one study Bucheli designed, she and her team found that the community of microbes on flies visiting recently placed bodies varied in diversity and abundance across different seasons and even across the same seasons in different years.*

Still, Metcalf and Bucheli don't think these many potential sources of variability spell trouble for a microbial clock. First, the clock may keep time without

"It would be really, really cool to find microbiome patterns that vary according to geographic location."

—Carl Schmidt

relying on any of the microbes that vary by host, season, geography, or blowfly visitor. And even if these variations do prove to be important, the team expects they could be used to calibrate the clock to specific circumstances, rather than disabling it altogether.

That, says Bucheli, is the approach forensic entomologists have long been honing. Instead of throwing off estimates, variation in insects often adds clues to a case; for example, insect activity might suggest that a body found in one climate actually came from a different climate. "I think we're going to see that with the microbes," says Bucheli. "If we do find changes or variation, it might be able to tell us fine level pieces of information."

True Crime Stories

Despite promising results in the FIRS project and others so far, the odds of finding a forensically useful clock may depend on the time window of interest. The first 48 hours after death are particularly tricky. "There have been no significant reproducible changes in the microbiome within 48 hours in the cases we've seen," says Schmidt.

He has been working with Jen Pechal of Michigan State University to see if she can identify a clock based

*King K, Berry R, Dane D, Lynne A, Bucheli S. Microbiome of forensically significant flies (Diptera) associated with human decomposition. *ASM Microbe*, June 1–5, 2017, New Orleans, LA.

