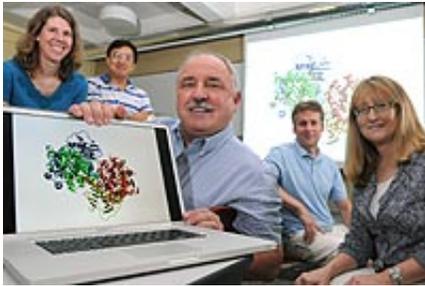


Exploring the role of gut bacteria in digestion

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Scientists at Argonne worked together to better understand bacteria and enzymes in the human gut. Pictured (l to r) are Christine Tesar, Kemin Tan, Andzej Joachimiak, Gyorgy Babnigg and Rosemarie Wilton.

They congregate in the environments that suit them best; some prefer the dry, desert-like conditions of our forearms while others thrive in the Amazon-style humidity of our feet.

Though we are born without them, bacteria inhabit our body within seconds of our birth. They live in our mouths, around our eyes, in our digestive systems, under our arms and in the shoots of our hair.

Most are helpful or at least harmless. The three or so pounds of bacteria living in our gut—mostly in the large intestine—help us digest all manner of food.

It's these tiny stowaways that interest Andrzej Joachimiak and his team the most, in part because they can have a tremendous impact on human

health.

Scientists know the bacteria inside our gut can influence our maturation, immune system development, metabolism and production of essential biocompounds.

Previous research shows that a number of diseases—including cardiovascular disease, diabetes and inflammatory bowel disease—are associated with changes in our [gut bacteria](#) or [microbiota](#). Some have been linked to obesity.

Joachimiak is the director of the Structural Biology Center and the Midwest Center for [Structural Genomics](#) at the U.S. Department of Energy's Argonne National Laboratory. He and his colleagues have recently determined the three-dimensional structure of one of these bacterial enzymes. It plays a fundamental role in human digestion by breaking down sugar polymers to simple glucose.

[Human cells](#) do the same thing, but, until recently, no one expected [gut bacteria](#) to have such a large repertoire of enzymes that break down complex sugars.

This is important because it could potentially help scientists and doctors to better understand the impact of gut organisms on human diseases like diabetes.

There are many practical applications of this type of knowledge. Joachimiak said much of the medicine prescribed today focuses on human, rather than bacterial, cells in part because we know relatively little about the hundreds of species of bacteria living in and on our bodies.

"We target human enzymes with drugs," he said. "But we don't often

target bacterial enzymes. If someone has diabetes, doctors prescribe drugs to control their production of glucose. We might want to consider whether bacterial enzymes that produce glucose should also be targeted."

Some medications—specifically, antibiotics and antimicrobials—do, in fact, target bacteria, but this arsenal could be greatly expanded if scientists knew more about exactly what these symbiotic companions do inside their hosts, Joachimiak said.

He and his group have recently published a paper on their research in the *Journal of the Federation of American Societies for Experimental Biology*, with Kemin Tan, a structural biologist as lead author.

Joachimiak said our "gut microbiome"—the colony of bacteria that live in our digestive tract—has been co-existing and evolving with us throughout human history, but only recently have scientists devoted time to understanding these tiny, unicellular organisms.

Gyorgy Babnigg, who worked with Joachimiak on this project, said that the team has determined the structure of about a dozen of these sugar-degrading enzymes, all of which play an important role in helping us with digestion.

"The human body can use starch very effectively; the degradation starts immediately when we chew our food," he said. "The di- or tri-saccharides are degraded in the small intestine and the resulting glucose is readily absorbed."

Not so for some plant material.

"The complex and branched sugars found in fruits and vegetables pose a challenge for the human body and are degraded mostly by the gut microbes," Babnigg said. "The structures of these enzymes can shed light

on their inner workings. This is a growing area of study, and numerous projects aim at sequencing the DNA of these microbes.”

Rosemarie Wilton, a molecular biologist at Argonne, played a crucial role in the study. She examined how these protein break apart sugar molecules.

"My end of the research yielded some important clues that will help us interpret some of the sequencing data coming out now," she said. "This is one little piece of the puzzle. It's a huge and complex system, and these systems are difficult to study because it's impossible to perfectly replicate the human gut where all of these different species of bacteria live together.”

And there were others who contributed; Laura Keigher cloned genes necessary for this study, and Christine Tesar purified and crystallized the proteins used in this research.

While researchers have placed increasing emphasis on this area of study in the last decade, bacterial enzymes remain a mystery, at least in part. Scientists say now is the time to focus on this type of research, especially considering that bacteria outnumber human cells by at least 10 to one.

"This is an area that has only recently been explored," Joachimiak said. "There is so much more to know."

Provided by Argonne National Laboratory

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