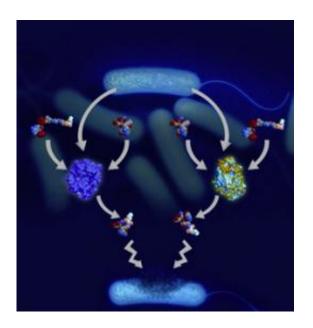


Outsmarting bacteria: Researchers develop faster method to generate new antibiotics

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This illustration shows the concept of using enzymes that evolved in bacteria to confer resistance to antibiotics as a chemical tool to generate novel antibacterials with broad-spectrum activity. M. Fridman and S. Garneau-Tsodikova et al. show how two aminoglycoside acetyltransferases (in purple on the left and multi-color on the right) isolated from resistant bacterial strains catalyze the region-specific acyl transfer from an acyl-coenzyme A (the two molecules on the left and the right sides of the picture, the acyl atoms are shown in brighter colors) to aminoglycosides to create novel N-acylated antibiotics. Illustration courtesy of Sylvie Garneau-Tsodikova.

(PhysOrg.com) -- Researchers at the University of Michigan's Life Sciences Institute have developed a new method to rapidly generate and



test novel antibiotic-drug candidates. The technique could provide scientists with a new tool in the fight against antibiotic-resistant bacteria.

LSI research assistant professor Sylvie Garneau-Tsodikova and her colleagues essentially outsmarted <u>bacteria</u> by using the microbes' own defenses against them. The research team reports its findings in the January edition of the journal *ChemBioChem*.

"We've taken what the bacteria used to kill the antibiotic as a tool to make new <u>antibiotics</u>," said Garneau-Tsodikova, who is also the John G. Searle assistant professor of medicinal chemistry.

Decades of clinical use—and overuse—have made bacteria smarter and trickier to subvert. As a result, many antibiotics are losing their efficacy against bacteria that have built up resistance to them.

"There's currently a huge problem associated with bacterial resistance to antibiotics as bacteria come up with tricks to become resistant to all the drugs that scientists discover," said Garneau-Tsodikova.

Resistance is a problem in the aminoglycoside family of antibiotics, which are commonly used to fight serious bacterial infections and <u>genetic diseases</u>, and as anti-HIV drugs.

Because they have complex structures containing numerous chemically identical amine and hydroxyl functionalities, attempts to create new aminoglycoside antibiotics have been limited. They are hard to synthesize, and one compound takes multiple chemical steps to build, which can translate into years of work. And results are not guaranteed: The compound could turn out to be inactive, and the search would start all over again.

Bacterial resistance to this type of antibiotic happens because the



substrate promiscuity is strong, which means the bacteria are able to take a lot of aminoglycosides, modify them, and kill them by using, among others, enzymes called aminoglycoside acetyltransferases. The enzymes modify and eventually kill the drug by adding a small part of the cosubstrate, co-enzyme A, to the drug. "If the enzyme acetylates the drugs, the drugs become inactive," said Garneau-Tsodikova.

While much is known about the substrate promiscuity of these enzymes, little was known about the co-substrate promiscuity itself—the enzyme created by the bacteria to kill the antibiotics and render them ineffective.

Garneau-Tsodikova's lab, along with collaborator Micha Fridman at the University of Tel Aviv, sought to determine whether these enzymes would be co-substrate promiscuous and found that they are. Because they are now using an enzyme—which is easier to target and control—they are able to make new compounds and test them in one step, taking a day or two at the most.

"We've taken what the bacteria used to kill the antibiotic as a tool to make new antibiotics by using the co-substrate promiscuity to generate novel <u>drug candidates</u>. The method rapidly tells us which drugs are worth further pursuing," said Garneau-Tsodikova.

Their method also requires a much smaller amount of the drugs generated for testing—as little as a couple of micrograms. Garneau-Tsodikova has also used this method on other acetyltransferase systems with success. Her lab continues to build on this concept.

More information: ChemBioChem paper: <u>www3.interscience.wiley.com/jo ... e/cover/current.html</u>



Provided by University of Michigan

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