

Decoded microbial metabolism explains biofuel yield

January 18 2017, by Blaine Friedlander



Ludmilla Aristilde, assistant professor of biological and environmental engineering. Credit: Cornell University



To unravel how intricate waste biomass converts to biofuels, a Cornell professor studied the bacterium Clostridium acetobutylicum to decipher its metabolism. Understanding the bacterium's sugar-processing complexities may lead to improved biofuel yields.

"It is recognized for over a century that Clostridium acetobutylicum [a soil bacterium] is a viable biofuel producer. Right now, though, this bacterium is not very efficient at processing complex wastes to <u>biofuel</u> <u>production</u>," said Ludmilla Aristilde, assistant professor of biological and environmental engineering.

Using nonedible plant materials for biofuel production is important to sustainable energy options, and processing these materials involves metabolizing the plant's sugars. "Now, we seek to understand the complexities of utilizing the different sugars we find in plant biomass. So instead of having to separate the different sugars, we look to save time and money by using the whole complex biomass and turn it into fuel," she said.

Think of the bacterial cell as a factory, Aristilde said.

"If we better understand the assembly line in the metabolic mill that processes all the different sugars, we will be in a better place to implement bioengineering strategies. In other words, we can use all the different plant components from biomass waste without compromising on the yield of fuel production."

Aristilde used an emerging biochemical approach, called metabolomics, to track different sugar carbon atoms inside the bacterial cells and learn why the cell factory does not efficiently convert all sugar carbons into biofuel. Specifically, she found that the bacterium incorporates pentose carbons (five-carbon sugars) in the metabolic network to make nucleotides (the building blocks of DNA), but it does not lead to making



biofuel. But some hexose carbons (six-carbon sugars) are preferentially channeled through glycolysis, a metabolic pathway that connects to biofuel production.

The United States and most other countries seek to reduce <u>greenhouse</u> <u>gas emissions</u> to stave off global warming, and most gasoline sold in the U.S. contains ethanol because such plant-based biofuels are carbonneutral. All plant materials used to produce biofuels, such as grasses and nonedible crops, capture carbon dioxide from the atmosphere through photosynthesis, offsetting car exhaust.

Significant effort is being employed to engineer bacteria to metabolize complex wastes.

"The ultimate goal is to understand the cycling of complex organic mixtures by anaerobic soil bacteria. These bacteria are important in the natural conversion of organic carbons in soils and they are exploited for industrial biofuel production," Aristilde said. "This study teaches us a little more on how the network works and what prevents the cell from routing some components to biofuel and other components not."

More information: Ludmilla Aristilde. Metabolite labelling reveals hierarchies inthat selectively channel carbons from sugar mixtures towards biofuel precursors, *Microbial Biotechnology* (2016). DOI: 10.1111/1751-7915.12459

Provided by Cornell University

Citation: Decoded microbial metabolism explains biofuel yield (2017, January 18) retrieved 26 April 2024 from https://phys.org/news/2017-01-decoded-microbial-metabolism-biofuel-yield.html



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