

## **Going to extremes for enzymes**

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Extremophiles thriving in thermal springs where the water temperature can be close to boiling can be a rich source of enzymes for the deconstruction of lignocellulose.

In the age-old nature versus nurture debate, Douglas Clark, a faculty scientist with Berkeley Lab and the University of California (UC) Berkeley, is not taking sides. In the search for enzymes that can break lignocellulose down into biofuel sugars under the extreme conditions of a refinery, he has prospected for extremophilic microbes and engineered his own cellulases.

Speaking at the national meeting of the American Chemical Society

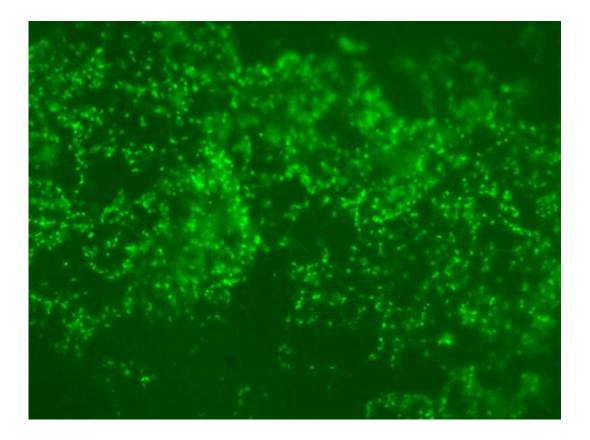


(ACS) in San Francisco, Clark discussed research for the Energy Biosciences Institute (EBI) in which he and his collaborators are investigating ways to release plant sugars from lignin for the production of liquid transportation fuels. Sugars can be fermented into fuels once the woody matter comprised of cellulose, hemicellulose, and lignin is broken down, but lignocellulose is naturally recalcitrant.

"Lignocellulose is designed by nature to stand tall and resist being broken down, and lignin in particular acts like a molecular glue to help hold it together" said Clark, who holds appointments with Berkeley Lab's Physical Biosciences Division and UC Berkeley's Chemical and Biomolecular Engineering Department where he currently serves as dean of the College of Chemistry. "Consequently, lignocellulosic biomass must undergo either chemical or enzymatic deconstruction to release the sugars that can be fermented to biofuels."

For various chemical reasons, all of which add up to costcompetitiveness, biorefineries could benefit if the production of biofuels from lignocellulosic biomass is carried out at temperatures between 65 and 70 degrees Celsius. The search by Clark and his EBI colleagues for cellulases that can tolerate these and even harsher conditions led them to thermal springs near Gerlach, Nevada, where the water temperature can be close to boiling. There they discovered a consortium of three hyperthermophilic Archaea that could grow on crystalline cellulose at 90 degrees Celsius.





Hyperthermophilic Archaea that grow on crystalline cellulose at 90 degrees Celsius yielded EBI-244, the most active high-temperature cellulase ever identified.

"This consortium represents the first instance of Archaea able to deconstruct lignocellulose optimally above 90°C," Clark said.

Following metagenomic studies on the consortium, the most active high-temperature cellulase was identified and named EBI-244.

"The EBI-244 cellulase is active at temperatures as high as 108 degrees Celsius, the most extremely heat-tolerant <u>enzyme</u> ever found in any cellulose-digesting microbe," Clark said.

The most recent expedition of Clark and his colleagues was to thermal



hot springs in Lassen Volcanic National Park, where they found an enzyme active on cellulose up to 100°C under highly acidic conditions – pH approximately 2.2.

"The Lassen enzyme is the most acidothermophilic cellulase yet discovered," Clark said. "The final products that it forms are similar to those produced by EBI244."

In addition to bioprospecting for heat tolerant enzymes, Clark and his colleagues have developed a simple and effective mutagenesis method to enhance the properties of natural enzymes. Most recently they used this technique to increase the optimal temperature and enhance the thermostability of Ce17A, a fungal cellulase that is present in high concentrations in commercial cellulase cocktails. They engineered yeast to produce this enzyme with encouraging results.

"The yeast Saccharomyces cerevisiae has often been used both in the engineering and basic study of Cel7A; however, Cel7A enzymes recombinantly expressed in yeast are often less active and less stable than their native counterparts," Clark said. "We discovered that an important post-translational modification that was sometimes absent in the yeast-expressed enzyme was the underlying cause of this disparity and successfully carried out the post-translational modification in vitro. After this treatment, the properties of Cel7A recombinantly expressed in yeast were improved to match those of the native enzyme."

## Provided by Lawrence Berkeley National Laboratory

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