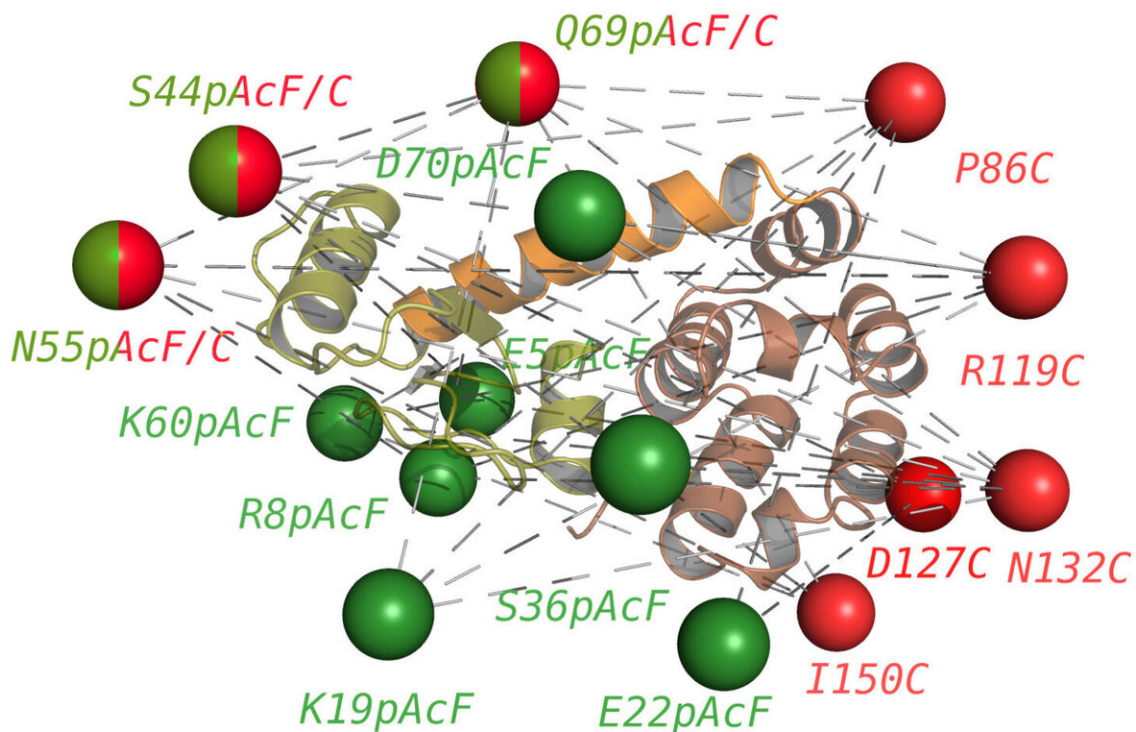


Advanced optical imaging technique may lead to structure-guided drug design

March 9 2020, by Laura Schmitt



The researchers engineered 34 different double-labeled samples for FRET experiments to measure distances across the enzyme to visualize its function. Each sphere represents one of the fluorescence markers with the corresponding engineered mutation site. Credit: Courtesy Hugo Sanabria

A Clemson University College of Science researcher, together with a

team of researchers primarily at Heinrich Heine University in Germany, developed and demonstrated new optical imaging methods to monitor a single molecule in action.

This fluorescence-based technique may accelerate the field of structural biology, helping scientists better understand how molecules are assembled, function, and interact, which in turn may aid in structure-guided drug design.

Hugo Sanabria, an associate professor of physics and astronomy at Clemson, with his colleagues used Förster resonance energy transfer (FRET) to study the lysozyme of the bacteriophage T4. They reported their findings in the paper titled "Resolving dynamics and function of transient states in single enzyme molecules," published on March 6, 2020, in *Nature Communications*.

According to co-author Claus A.M. Seidel, chair of the Institute for Molecular Physical Chemistry at Heinrich Heine University in Germany, this work underpins essential reaction steps of biomolecular machines (enzymes).

"Our FRET studies demonstrate the need of a third functional state in the famous Michaelis-Menten kinetics," said Seidel. "The Michaelis-Menten description is one of the best-known models of enzyme kinetics."

The centerpiece of this imaging tool is a FRET-based microscope, a sophisticated and powerful machine capable of visualizing biomolecules as small as a few nanometers.

To visualize biomolecules at work, Sanabria and colleagues placed two fluorescent markers on a set of [molecules](#), which created a ruler at the [molecular level](#). By using different locations of the markers, the team

collected a set of distances that describe the shape and form of the observed molecule.

In essence, this process generated a collection of data points that were computationally processed, allowing the researchers to distinguish how the molecule looks and how it moves.

"We observe changes in the structure, and because our signal is time dependent, we can also get an idea of how the molecule is moving over time," Sanabria said.

In his study, Sanabria and his team of collaborators combined the FRET-based microscope with molecular simulations to examine lysozyme, an enzyme found in tears and mucus. Lysozyme destroys the protective carbohydrate chains surrounding bacteria's cell wall. Scientists widely use lysozyme to study [protein structure](#) and function because it's such a stable enzyme.

"We can track the lysozyme of the bacteriophage T4 as it processes its substrate at near atomistic level with unprecedented spatial and temporal resolution," Sanabria said. "We've taken the imaging field to a whole new level."

Sanabria's optical method revealed that the lysozyme structure is different than previously thought. Until now, scientists have largely determined the structures of proteins like [lysozyme](#) mainly using methods such as X-ray crystallography, nuclear magnetic resonance (NMR) spectroscopy, and cryo-[electron microscopy](#).

"For the longest time, this molecule was considered a two-state molecule because of how it receives the substrate or cell wall of the target bacteria," he said. "However, we have identified a new functional state."

The team is helping to establish a database where their FRET-based structural models of similarly generated biomolecular models can be stored and accessed by other scientists. Together with the FRET community, the group is also working to establish recommendations for FRET microscopy.



Clemson University biophysics associate professor Hugo Sanabria and an international team of researchers have demonstrated new optical imaging methods that may someday aid in structure-guided drug design. Credit: Ken Scar, Clemson University

Sanabria aims to apply his imaging methodology to other biomolecules. "This optical method can be used to study protein folding and misfolding

or any structural organization of biomolecules," Sanabria said. "It can also be used for drug screening and development, which requires knowing what a biomolecule looks like in order for a drug to target it."

"This work is a milestone in structure determination using FRET to map short-lived functionally relevant enzyme states," Seidel said.

More information: Hugo Sanabria et al, Resolving dynamics and function of transient states in single enzyme molecules, *Nature Communications* (2020). [DOI: 10.1038/s41467-020-14886-w](https://doi.org/10.1038/s41467-020-14886-w)

Provided by Clemson University

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