

Biophysical Society 62nd Meeting, Feb. 17-21, 2018, San Francisco, California

What Makes Circadian Clocks Tick?

Researchers discovered that how proteins wiggle, jiggle, and shape-shift by the hour is central to figuring out how circadian clocks work.

EMBARGOED for release until 8 a.m. EST on Sunday, Feb. 18, 2018

For More Information: AIP Media Line media@aip.org 301-209-3090

WASHINGTON, D.C., February 18, 2018 -- Circadian clocks are found within microbes and bacteria, plants and insects, animals and humans. These clocks arose as an adaptation to dramatic swings in daylight hours and temperature caused by the Earth's rotation. But we still don't fully understand how these tiny biological clocks work.

During the 62nd Biophysical Society Annual Meeting, held Feb. 17-21, in San Francisco, California, Andy LiWang at the University of California, Merced will present his lab's work studying the circadian clock of blue-green colored cyanobacteria. One type of cyanobacteria, called spirulina, is high in vitamins and minerals and is used as a natural food dye for candy and gum.

LiWang's group discovered that how the proteins move hour by hour is central to cyanobacteria's circadian clock function. "And now it's becoming clear that the same is true for eukaryotic [animal] clocks," LiWang said.

Cyanobacterial circadian clock proteins are unique because they can be reconstituted within a test tube in the absence of live cells. Researchers made a solution of these proteins and adenosine



triphosphate (ATP), food for the proteins, to create a circadian clock that functioned for weeks.

LiWang's structural biology lab uses nuclear magnetic resonance (NMR) spectroscopy, the parent technology for MRI, to study the protein structure and dynamics of biological molecules and then uses the structures to gain insights into their function. "We also examine how the proteins wiggle, flex, and shape-shift, because these motions ... are also critical to their biological function," LiWang said.

LiWang's lab also collaborates with X-ray crystallographers like Carrie Partch at the University of California, Santa Cruz, because X-ray crystallography is a powerful technique to capture static structures of proteins and their complexes at atomic and near-atomic resolution.

"A big surprise for us was the extent to which internal motions of circadian clock proteins dictate ... their function," LiWang said. "Static X-ray crystal structures of individual proteins, mostly solved by other labs, were invaluable to our work but told only part of the story."

Cyanobacterial clock proteins aren't exactly the same as the clock proteins of animals or human clocks, but proteins serve as the cogs, gears and springs of all circadian clockworks and the overall function of the proteins is similar.

"Because clock proteins need to keep time, there should be some basic principles of biological timekeeping shared between all clocks regardless of whether the proteins are the same or not," LiWang said. "Our structures of the complexes of the circadian clock proteins of cyanobacteria provided important mechanistic insights, but are static snapshots of a system that's continuously moving and changing hour by hour," said LiWang.

231-Pos, Board B1 "Ticking mechanism of a biological clock" is authored by Andy LiWang. It will be displayed at 1:45 p.m. PST. Sunday, Feb. 18, 2018, in the South Hall ABC of the Moscone Center South. Abstract: https://plan.coreapps.com/bpsam2018/abstract/598979c882021290aae09439cc42e19d

_____ MORE MEETING INFORMATION _____

ABOUT THE MEETING

Each year, the Biophysical Society Annual Meeting brings together more than 6,000 researchers working in the multidisciplinary fields representing biophysics. With more than 3,600 poster presentations, over 200 exhibits, and more than 20 symposia, the BPS Annual Meeting is the largest meeting of biophysicists in the world. Despite its size, the meeting retains its small-meeting flavor through its subgroup symposia, platform sessions, social activities and committee programs. The 62nd Annual Meeting will be held at the Moscone Center (South) in San Francisco, California.



PRESS REGISTRATION

The Biophysical Society invites professional journalists, freelance science writers and public information officers to attend its Annual Meeting free of charge. For press registration, contact Ellen Weiss at <u>EWeiss@biophysics.org</u> or the Media Line at the American Institute of Physics at <u>media@aip.org</u> or 301-209-3090.

NEWS RELEASES

Embargoed press releases describing in detail some of the breakthroughs to be discussed at the meeting are available on EurekAlert!, Newswise and Alpha Galileo or by contacting the Media Line at the American Institute of Physics at <u>media@aip.org</u> or 301-209-3090.

QUICK LINKS

Main Meeting Page: <u>https://www.biophysics.org/2018meeting/Home/tabid/7117/Default.aspx</u> Symposia: <u>https://www.biophysics.org/2018meeting/Program/ScientificSessions/Symposia/tabid/7192/D</u> <u>efault.aspx</u> Desktop Planner: <u>http://www.biophysics.org/2018meeting/GeneralInfo/MobileApp/tabid/7473/Default.aspx</u>

ABOUT THE SOCIETY

The Biophysical Society, founded in 1958, is a professional, scientific Society established to encourage development and dissemination of knowledge in biophysics. The Society promotes growth in this expanding field through its annual meeting, monthly journal, and committee and outreach activities. Its 9,000 members are located throughout the U.S. and the world, where they teach and conduct research in colleges, universities, laboratories, government agencies, and industry. For more information on the Society, or the 2018 Annual Meeting, visit http://www.biophysics.org.