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dr. christiane menzfeld

phone: +49 89 8578-2824

pr@biochem.mpg.de

www.biochem.mpg.de/news

 @MPI_Biochem

Crowd Control

Have you ever been stuck in the middle of a crowd? As people pack closer together, it becomes more difficult to move through the crowd. Sometimes it can become so tightly packed that you cannot move at all. If this sounds uncomfortable, then you probably wouldn't like to live inside a cell, which is densely packed with proteins and other molecules. This crowding is very important for the cell—it pushes the molecules together so that they can interact and perform the chemical reactions that the cell needs to live. In fact, many human diseases are likely influenced by changes in molecular crowding that cause harmful interactions between proteins. Despite its importance, it remains a mystery how the crowding inside cells is controlled. Combining biophysics, cell biology, physical modeling, and cryo-electron tomography, an international team of scientists at New York University (NYU) and the Max Planck Institute of Biochemistry (MPIB) has discovered that the mTORC1 signaling pathway controls the concentration of ribosomes inside the cell, thereby regulating crowding and the ability of proteins to interact with each other to form phase-separated compartments. This study is published in the journal *Cell*.

GEMs shed light on molecular crowding

The team led by Liam Holt at NYU School of Medicine invented a clever strategy to measure the molecular crowding within cells. They designed tiny fluorescent balls, called GEMs, which are produced inside living cells. Using light microscopy, the researchers were able to track the movement of the glowing GEMs, and thus determine how difficult it was for the GEMs to move through the crowded cellular environment under different conditions. This led to an exciting discovery—mTORC1, a protein complex that is well known to control cell growth, also controls crowding within the cell. When mTORC1 signaling was inhibited with the drug rapamycin, the GEMs traveled much quicker through the cell's cytoplasm, indicating that there was reduced crowding. This is analogous to how traffic can move faster when there are less cars on the road.





Ribosomes are the cell's natural crowding agent

To examine this change in crowding, the team led by Benjamin Engel at the MPIB imaged the native cellular environment with cryo-electron tomography. "This powerful technique allows us to see every molecule within the cell," explains Engel. "The cell's cytoplasm is full of little protein factories called ribosomes. We were able to count each ribosome in the cell with single molecule precision." When mTORC1 signaling was inhibited, the concentration of ribosomes within the cytoplasm dropped by half, dramatically reducing the molecular crowding encountered by the GEMs. The strong correlation between decreased ribosome concentration and increased GEM motility indicated that ribosomes are the main "crowding agent" within the cytoplasm.

To mix or separate?

It has recently been discovered that cells can compartmentalize their proteins by concentrating them into liquid droplets that do not freely mix with their surroundings, similar to how vinegar forms separated droplets in a dish of oil. This process, called "phase separation", allows biological reactions to occur much quicker and more efficiently by concentrating specific molecules together in a small space. When scientists examine phase separation in test tubes, they have to add an artificial crowding agent in order to get the separation to occur. Until now, it was unknown what the cell's natural crowding agent could be. Holt and colleagues discovered that changing the ribosome concentration greatly affected phase separation both in test tubes and within the cell. Thus, mTORC1 signaling controls the abundance of ribosomes, which in turn act a natural crowding agent to tune phase separation in the cytoplasm.

The effect of crowding on cellular disease

Up to a point, crowding concentrates interacting proteins together into phase-separated liquid droplets. However, if molecules in the cell become too crowded, these droplets can solidify. The current study suggests that malfunctioning mTORC1 may increase crowding, and therefore cause phase-separated droplets to become the solids found in cells with diseases of aging, such as the tau fibers that build up in the brain tissue of Alzheimer's patients. Furthermore, the mTORC1 pathway has been a common target of cancer drug development for decades because it senses whether a cell has the energy to grow and divide. These drugs have had limited success, which Holt says may be related to the crowding effect. For example, mTORC1 activation may be important to initiate cancer in some cases, but as cancer cells become too crowded with ribosomes, mTORC1 could hinder cancer growth later. Thus, the current line of work may help to set new guidelines about when to use mTORC1 inhibitors based on the stage of a patient's cancer. "The biological consequences of phase changes are an area of intense inquiry right now," says





Holt. “One of our long-term goals is to design therapies for neurodegeneration and cancer that alter molecular crowding within people’s cells.”



Caption:

This image depicts a 3D rendering of a cryo-electron tomogram of the yeast cytoplasm, with fluorescent GEM particles (yellow) lighting up a cellular environment that is crowded with ribosomes (blue) and membranes (grey). © Dimitry Tegunov and Stefan Pfeffer

Original publication

M. Delarue, G.P. Brittingham, S. Pfeffer, I.V. Surovtsev, S. Pinglay, K.J. Kennedy, M. Schaffer, J.I. Gutierrez, D. Sang, G. Poterewicz, J.K. Chung, J.M. Plitzko, J.T. Groves, C. Jacobs-Wagner, B.D. Engel & L.J. Holt. “mTORC1 controls phase separation and the biophysical properties of the cytoplasm by tuning crowding”. *Cell*, July 2018.

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About Benjamin Engel

Benjamin Engel's work focuses on characterizing the molecular architecture of organelles. Using cryo-electron tomography, he and his team are able to visualize macromolecular complexes within the native cellular environment with high spatial resolution. Engel completed his undergraduate studies in Molecular and Cell Biology at the University of California, Berkeley, in the United States. In 2011, he received his Ph.D. from the University of California, San Francisco. Since then, he has worked as a postdoctoral fellow and project leader in the "Molecular Structural Biology" department of Wolfgang Baumeister at the Max Planck Institute for Biochemistry in Martinsried near Munich.

http://www.biochem.mpg.de/5647281/12_organelle-architecture

About Liam Holt

Liam Holt's lab studies how cells process information, and how the unusual, crowded environment within the nucleus and cytoplasm is controlled. Using a combination of high-resolution, live-cell imaging and biochemical techniques, the Holt lab aims to discover fundamental principles in cell signaling. Holt completed his undergraduate studies at the University of Bath in England and his Ph.D. at the University of California, San Francisco. He then started his own group as a Bowes Fellow at the University of California, Berkeley. In 2016, he became an Assistant Professor at New York University School of Medicine in the Institute for Systems Genetics.

<https://www.liamholtlab.org/>

About the Max Planck Institute of Biochemistry

The Max Planck Institute of Biochemistry (MPIB) belongs to the Max Planck Society, an independent, non-profit research organization dedicated to top-level basic research. As one of the largest Institutes of the Max Planck Society, about 800 employees from 45 nations work here in the field of life sciences. In currently about 35 departments and research groups, the scientists contribute to the newest findings in the areas of biochemistry, cell biology, structural biology, biophysics and molecular science. The MPIB in Munich-Martinsried is part of the local life-science-campus in close proximity to the Max Planck Institute of Neurobiology, a Helmholtz Center, the Gene-Center, several bio-medical faculties of the Ludwig-Maximilians-Universität München and the Innovation and Founding Center Biotechnology (IZB).

<http://biochem.mpg.de>

About New York University

Founded in 1831, New York University is now one of the largest private universities in the United States. Of the more than 3,000 colleges and universities in America, New York University is one of





only 60 member institutions of the distinguished Association of American Universities. Enrollment has grown to more than 50,000 students at three degree-granting campuses in New York City, Abu Dhabi, and Shanghai. Today, students come from every state in the USA and from 133 foreign countries. The Institute for Systems Genetics at NYU Langone Health combines research in systems biology, genome engineering, human genetics, and computation. We take a systems approach to the wealth of information available in human biology and medicine.

<https://med.nyu.edu/institute-systems-genetics/>

Contact:

Dr. Benjamin Engel
Dept. of Molecular Structural Biology
Max Planck Institute of Biochemistry
Am Klopferspitz 18
82152 Martinsried/Munich
Germany
E-mail: engelben@biochem.mpg.de
www.biochem.mpg.de/en/rd/baumeister

Dr. Liam Holt
New York University Langone Medical Center
Science Building, 906
435 East 30th Street
New York, NY 10016
USA
E-Mail: liam.holt@nyumc.org
<https://www.liamholtlab.org/>

Dr. Christiane Menzfeld
Public Relations
Max Planck Institute of Biochemistry
Am Klopferspitz 18
82152 Martinsried/Munich
Germany
Phone: +49 89 8578-2824
E-mail: pr@biochem.mpg.de
www.biochem.mpg.de

Gregory Williams
Media Relations
NYU Langone Health
550 First Avenue
New York, NY 10016
USA
Phone: +1 212-404-3533
E-Mail: gregory.williams@nyumc.org
<https://nyulangone.org/>