Public relations



Press Release, October 22, 2018

dr. christiane menzfeld

phone: +49 89 8578-2824

pr@biochem.mpg.de www.biochem.mpg.de/news @MPI\_Biochem

# Disorder in the liver

Chronic excessive caloric intake leads to the deposition of fat droplets in the liver. This condition, known as fatty liver, can cause permanent damage to the organ. Researchers at the Max Planck Institute of Biochemistry (MPIB) in Martinsried have now investigated the effects of this fat overflow on liver proteins. They showed that fatty liver is associated with changes in the location and activity of numerous cellular proteins. The study, which was published in the journal *Developmental Cell*, shows the effect of lipid deposition on fundamental cellular processes in the liver.

The high-calorie and high-fat diet now common in many parts of the world, including Europe and North America, can lead to a wide range of health problems. This diet frequently results in obesity, which places a heavy strain on the cardiovascular system, but it can also damage the liver. It is estimated that around 30 percent of the population is affected by fatty liver, an initially reversible accumulation of fat droplets in liver cells. In the long run, these deposits can lead to inflammation and irreversible liver damage. The effects of fat droplets on the biology of liver cells has now been investigated in detail by researchers in Matthias Mann's Proteomics and Signal Transduction Department at the MPIB in Martinsried. For the study, the researchers collaborated with the research group headed by Ralf Jungmann at the MPIB and with scientists from the Helmholtz Center München and Harvard University, USA.

## Disorder in protein distribution

The liver is one of the most versatile organs in the human body. In addition to detoxifying the blood, it plays a key role in the metabolism of sugars, proteins and fats. These nutritional components can be temporarily stored in liver cells. Liver cells, like most cells of highly developed organisms, contain various functionally and structurally distinct areas known as organelles. The researchers used state-of-the-art mass spectrometry methods to determine the entire set of proteins, known as the proteome, of cell organelles. Mass spectrometry is a type of molecular scale used to identify proteins.





Public relations

The researchers studied the protein composition of organelles in liver samples taken from mice that were given a normal or a high-calorie diet. As in humans, increased caloric intake leads to fatty degeneration of the liver in mice. "We found that as a result of fatty liver 20 percent of the proteins occurred in different organelles than in healthy organs. The fat droplets in the cells bind hundreds of different proteins on their surface. These are drawn away from other processes and organelles," explains Natalie Krahmer, postdoc at the MPIB and lead author of the study. "The resulting disorder shows that it is not only important whether and in what amount proteins are present in cells, but also where they are." The researchers also found proteins attached to the fat droplets whose functions are still completely unknown.

### Trigger for a vicious circle

Particularly, changes in location affected proteins of the Golgi apparatus. The Golgi apparatus is an organelle that is responsible for the formation of transport vesicles in the cell. "We observed a vicious circle: the large number of fat droplets alters the structure of the Golgi apparatus and reduces its activity in liver cells. And because the Golgi apparatus is required for fat release from the cells, which counteracts the fatty degeneration of cells, the process accelerates. Restoring the structure of the Golgi apparatus could therefore be a therapeutic goal to slow the progression of fatty liver," Krahmer says.

The researchers also investigated the occurrence of molecular protein switches, a phenomenon known as phosphorylation, with the help of the EasyPhos method developed at the MPIB. Phosphorylation can alter the activity of proteins. Many of the proteins whose location was shifted in fatty liver also showed changes in their phosphorylation patterns. "However, we were not able to conclude from the findings whether this is the consequence or cause of the altered protein distribution," Krahmer explains.

MPIB Director Matthias Mann summarizes the current research findings: "At the moment, there are no treatment options for fatty liver other than a change in diet. However, until now the cellular effects of such changes were largely unknown. We hope that our research will identify therapeutic targets for slowing the progression of fatty degeneration of the liver." In any case, he says, the observed effects on the Golgi apparatus, the still unknown proteins involved in fatty deposits and the altered phosphorylation patterns open up many avenues for future research. [CW]



# max planck institute of biochemistry

**Public relations** 





### Caption:

The deposition of fat into liver cells causes fundamental changes of their biology. Via mass spectrometry, researchers at the MPIB tracked shifts in the localization of numerous proteins. © Natalie Krahmer, Max Planck Institute of Biochemistry

#### **Original Publication:**

N. Krahmer, B. Nafaji, F. Schueder, F. Quagliarini, M. Steger, S. Seitz, R. Kasper, F. Salinas, J. Cox, N.H. Uhlenhaut, T.C. Walther, R. Jungmann, A. Zeigerer, G.H.H. Borner, M. Mann: Organellar proteomics and phospho-proteomics reveal subcellular reorganization in diet-induced hepatic steatosis, *Developmental Cell*, October 2018 DOI: 10.1016/j.devcel.2018.09.017



# max planck institute of biochemistry

**Public relations** 



---

#### About Matthias Mann

Matthias Mann studied physics at the Georg August University in Göttingen and obtained his PhD from Yale University, New Haven, USA. He held group leader positions at the European Molecular Biology Laboratory (EMBL) and the University of Southern Denmark in Odense before becoming a director at the MPIB in 2005. His Department "Proteomics and Signal Transduction" uses mass spectrometry to study the proteome, the entirety of all proteins of an organism. Additionally, Mann was appointed director of the Department of Proteomics at the University of Copenhagen in 2007. Mann has received numerous awards for his research including the Louis-Jeantet Prize for Medicine, the Körber European Science Prize and the Gottfried Wilhelm Leibniz Prize.

#### 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

#### Contact:

Prof. Dr. Matthias Mann Proteomics and Signal Transduction Max Planck Institute of Biochemistry Am Klopferspitz 18 82152 Martinsried/Munich Germany

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

