

Bringing information into the cell

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PSI biochemist Volodymyr Korkhov (right) and Chao Qi, first author of the study, preparing a sample for cryo-electron microscopy. Credit: Paul Scherrer Institute/Mahir Dzambegovic

Researchers at the Paul Scherrer Institute (PSI) have elucidated an important part of a signal pathway that transmits information through the cell membrane into the interior of a cell.



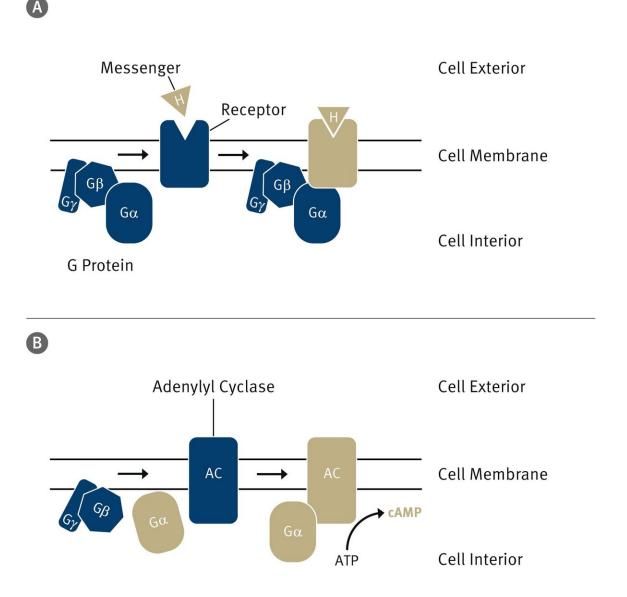
The interior of all living cells is separated from the outside world by membranes. These membranes keep the cells intact and protect them from negative influences. But they also act as a barrier for nutrients and information. For this reason, cell membranes contain mechanisms that enable selective access to desired substances or transmit information from external signals into the cell.

An important signal pathway in mammals consists of three components: The first is a receptor that recognises the signal and is activated by it. The second is a so-called G protein that binds to the activated receptor and transmits the signal to one or more effector proteins. In this case, the effector is adenylyl cyclase, the third component of the signal chain. This protein is activated by a subunit of the G protein and produces, in a biochemical reaction, a secondary messenger called cyclic AMP (cAMP).

cAMP triggers various reactions in the cell; for example, it increases the permeability of the membrane to calcium in cardiac cells, leading to an increase in the heart beat rate.

Researchers at the Paul Scherrer Institute in Villigen, Switzerland, have now examined one special type of adenylyl cyclase with the help of electron microscopy and have produced the most detailed image to date of this type of membrane protein.





The cell interior is protected against external influences by the cell membrane. To channel information into the cell, which is transmitted by a messenger (H), special signal pathways exist. A: The messenger binds to a receptor in the membrane and activates it (illustrated here by the color change). In turn, the receptor activates a G protein. B: From this, the alpha subunit splits off and binds to the last part of the signal pathway, a protein called adenylyl cyclase. This transforms ATP -- which stores energy for the cell -- into so-called cyclic



AMP (cAMP), an important secondary messenger in the cell. Credit: Paul Scherrer Institute/Markus Fischer

Helpful self-restraint

"To understand how signal pathways in the cell work, it is necessary first to know what the components involved look like in detail," says Volodymyr Korkhov, head of the research group Mechanisms of Signal Transduction in the Biology and Chemistry Division at PSI and assistant professor at the Institute for Biochemistry at ETH Zurich. "Our work is an important contribution to elucidating the exact function of adenylyl cyclase in the cAMP signal chain."

"Surprisingly, in determining the <u>structure</u> of the adenylyl cyclase bound to the alpha subunit of the G protein, we discovered that the protein appears to be able to inhibit itself," says Korkhov. One part of the protein is responsible for this self-inhibition. This part blocks the active site of the enzyme and prevents the overproduction of cAMP.

This new insight into the molecular structure of adenylyl cyclase provides a much better understanding of how external signals lead to the controlled production of the important secondary messenger cAMP. The concentration of cAMP in <u>cells</u> plays an important role in the development of cardiovascular diseases, certain tumours, and type 2 diabetes. "In the future, our new findings could make it possible to identify drugs that inhibit or activate the <u>adenylyl cyclase</u>, depending on whether overproduction or lack of cAMP is responsible for a disease," Korkhov explains.

Microscopy at low temperatures



The researchers achieved their results using cryo-electron microscopy (cryo-EM). This form of transmission electron microscopy operates at temperatures below -150 degrees Celsius. The sample to be examined is snap frozen in liquid ethane, preserving its natural structure. This method, for which the Nobel Prize for Chemistry was awarded in 2017, is increasingly used in the investigation of biological structures. "It is exciting to get a deep insight into the structure of adenylate cyclase," says Chao Qi, a doctoral candidate in Korkhov's lab and first author of the study. "The structure of this protein has been elusive for decades since its discovery, and I'm glad that I was able to elucidate this structure with cryo-EM in the course of my doctoral research."

The resolution achieved by the PSI researchers in their investigations was 3.4 angstroms. An angstrom is one ten-millionth of a millimetre. Isolated atoms have a radius of 0.3 to 3 angstroms. The researchers have now published their findings in the scientific journal *Science*.

More information: "The structure of a membrane adenylyl cyclase bound to an activated stimulatory G protein" *Science* (2019). <u>science.sciencemag.org/cgi/doi ... 1126/science.aav0778</u>

Provided by Paul Scherrer Institute

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