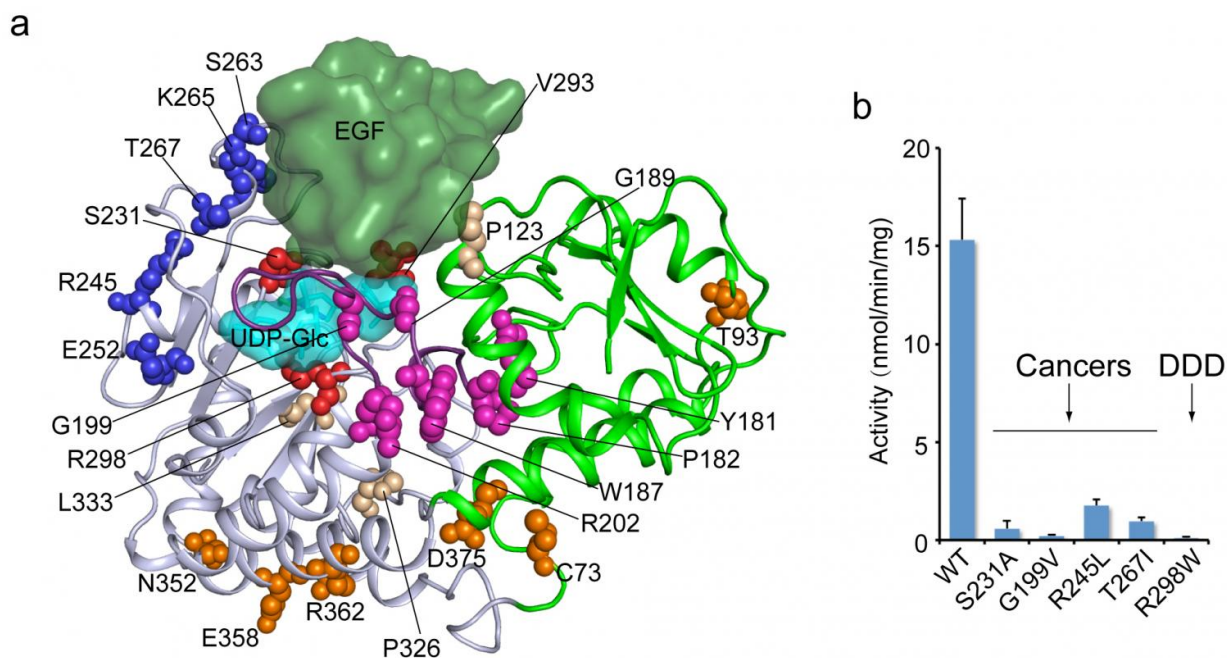


Scientists determine structure of enzyme linked with key cell-signaling protein

July 18 2016



Panel a: This is a cartoon "ribbon" diagram of the structure of Rumi in complex with the donor substrate UDP-Glucose (UDP-Glc, shown in cyan) and acceptor substrate hFA9 EGF (dark green). The spheres represent mutations to Rumi that have been found in human diseases including cancers and Dowling-Degos disease (DDD). Panel b: Disease-related Rumi mutations abolish the enzyme's activity relative to the "wild type" (WT) unmutated enzyme, likely leading to compromised Notch signaling and pathogenesis. Credit: Brookhaven National Laboratory

Scientists have captured atomic level snapshots showing how one key enzyme modifies a protein involved in turning genes on or off inside cells. Understanding this process-which is particularly important when cells are first taking on specialized identities such as nerve cells, muscle, skin, and so on-helps explain how complex organisms can arise from a finite number of genes. The research, published in *Nature Chemical Biology*, also identifies links between defects in this particular enzyme and certain cancers, potentially pointing to new targets for anticancer drugs.

"Our work describes the structure and function of an important enzyme called Rumi, which adds a [glucose molecule](#) to several signaling proteins to modify their activities," said the study's lead author, Huilin Li, a biologist at the U.S. Department of Energy's Brookhaven National Laboratory and Stony Brook University. "One of these target proteins, called Notch, is the major receptor that controls the development of multicellular organisms. It plays an important role in how cells sense their neighbors and, by controlling gene expression, determines which cells should develop into different types and how much they should grow-like a master controller."

Notch is also known to be a tumor suppressor. But its action is far more complicated than a simple on and off switch. It's a protein chain that spans the cell membrane with complicated 3D folding-including 36 repeating "beads on a necklace" folds that can each be modified in different ways to affect the protein's overall function.

"We wanted to understand the Notch structure upon different types of modification," said Li, who collaborated with Robert Haltiwanger and Hideyuki Takeuchi, biochemists at Stony Brook University at the time of the experiment who are now at the University of Georgia.

Picking up on recent discoveries that the Notch-modifying enzyme

known as Rumi is necessary for animal development and that various mutants of Rumi are linked with a form of skin hyperpigmentation known as Dowling-Degos disease (made by scientists at the Baylor College of Medicine and the University of Bonn, Germany), Li's group became focused on this particular enzyme. Studying the Rumi-Notch interaction at the molecular level might help scientists understand the disease mechanisms, and could even lead to the design of drugs to combat these diseases.

Background knowledge directs precision study

Prior to the current study, the scientists already knew that Rumi modifies Notch by adding glucose molecules to the protein. There are other examples of proteins modified by other simple sugars, but Rumi is the first enzyme known to specifically add glucose.

Previous research had also revealed that the first glucoses added by Rumi are essential for Notch signaling. "Without the glucose molecules, Notch cannot take on the right 3D folding to mature into a functional receptor, and therefore cannot function," said Li.

"Also, because Notch is known to be a tumor suppressor, the essential role of Rumi in Notch activation suggests that Rumi could be a [tumor-suppressor](#) as well. If a chemical biologist wants to design drugs that affect this process, he or she would need to know the mechanism of the complex enzyme reactions," said Li.

"We set out to do just that-to find out what features in the Notch protein Rumi is looking for and how it puts a sugar (glucose) on it. "

Details of the enzyme-protein interaction

Li's team studied samples of a fruit-fly derived version of Rumi in

complex with a Notch "surrogate"-a human protein with a Notch-like fold-that were provided by Haltiwanger.

"It was very hard to get sufficient samples," said Hongjun Yu, a postdoctoral scientist working with Li and the first author of the paper, noting that the proteins make very small crystals with a hard to solve structure. But eventually, using intensely bright beams of x-rays at the National Synchrotron Light Source (NSLS) at Brookhaven and the Advanced Photon Source (APS) at Argonne National Laboratory-both DOE Office of Science User Facilities-they gathered sufficient data on the arrangement of the proteins' atoms. They averaged data from multiple crystals to generate stronger signals, and used computers to reconstruct 3D images of how the Rumi and Notch pieces fit together.

The scientists found that Rumi recognizes a six-amino-acid sequence on many of the 36 repeat folds of Notch. A water-repelling patch with a separate recognition sequence was also essential to the interaction. When Rumi binds, the six-amino-acid "signature" sequence that accepts the glucose molecule fits very precisely into the active site on Rumi in a classic "lock-and-key" manner. "Adding to or taking away even one amino acid from this signature sequence would make it impossible for the proteins to join in such a manner, which would prevent the glucose from being added," Yu said.

These studies revealed how the cancer-linked mutations to Rumi alter key structural components that render Rumi inactive. "Our study suggests that mutated forms of Rumi could be the cause of cancers because these mutant enzymes are unable to modify Notch. This, for the first time, links Rumi defects with cancers," Li said. The scientists emphasize, however, that other complex modifications to Notch likely also play a role.

"We are now looking at the details of the study to determine what the

biomedical impact of our new understanding might be," said Li. This includes thinking about whether drugs might be developed to affect the behavior of Notch by targeting specific sites on Rumi.

More information: Structural analysis of Notch-regulating Rumi reveals basis for pathogenic mutations, *Nature Chemical Biology*, [nature.com/articles/doi:10.1038/nchembio.2135](https://doi.org/10.1038/nchembio.2135)

Provided by Brookhaven National Laboratory

Citation: Scientists determine structure of enzyme linked with key cell-signaling protein (2016, July 18) retrieved 29 March 2023 from <https://phys.org/news/2016-07-scientists-enzyme-linked-key-cell-signaling.html>

This document is subject to copyright. Apart from any fair dealing for the purpose of private study or research, no part may be reproduced without the written permission. The content is provided for information purposes only.