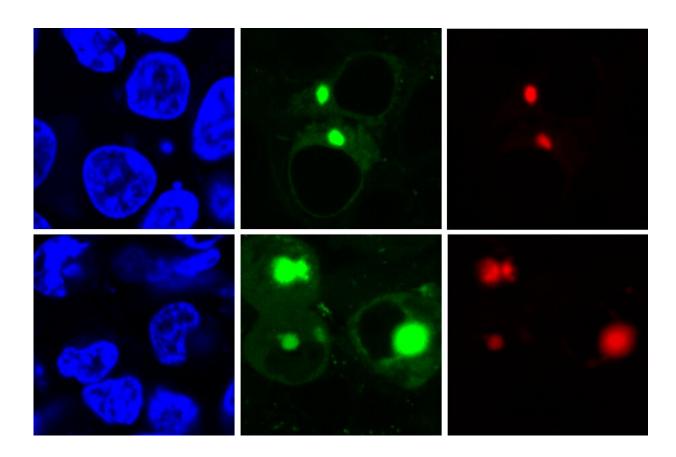


New method uses fluorescence to identify disease-causing forms of proteins

February 13 2019, by Gail Mccormick



The new AggTag method allows researchers to see the previously undetectable but potentially disease-causing intermediate forms of proteins as they misfold. The method uses fluorescence to simultaneously detect two different proteins (red, green) within the cell (blue). Credit: Zhang Lab, Penn State

A new method uses fluorescence to detect potentially disease-causing



forms of proteins as they unravel due to stress or mutations. A team of researchers from Penn State and the University of Washington reengineered a fluorescent compound and developed a method to simultaneously light up two different proteins as they misfold and aggregate inside a living cell, highlighting forms that likely play a role in several neurodegenerative diseases including Alzheimer's and Parkinson's. Two recent papers describing the method appear online in *ChemBioChem* and the *Journal of the American Chemical Society*.

"In order to function properly, proteins fold into very precise structures, but environmental stress or pathogenic mutations can cause proteins to misfold and aggregate," said Xin Zhang, assistant professor of chemistry and of biochemistry and molecular biology at Penn State and leader of the research team. "Protein aggregation is a multi-step process, and it is believed that the intermediate form, which previous imaging techniques could not detect, is responsible for a number of diseases, including Alzheimer's, Parkinson's, Type 2 Diabetes, and cystic fibrosis. We developed the Aggregation Tag method—AggTag—to see these previously undetectable intermediates—soluble oligomers—as well the final aggregates in live cells."

Previous techniques to identify protein aggregation used fluorescent compounds that were always lit up, which made it impossible to distinguish properly folded proteins from the intermediate form because both trigger low-level diffuse fluorescence. The AggTag method uses "turn-on fluorescence," so the compound only lights up when misfolding starts to occur.

"When the <u>fluorescent compound</u> has plenty of space to move, it rotates freely and remains turned off, as in the presence of a properly folded protein," said Yu Liu, assistant research professor of chemistry at Penn State and the key developer of AggTag method. "But when the protein starts to misfold and aggregate, the compound's movement becomes



restricted and it begins to light up. Diffuse fluorescence indicates that intermediate oligomers are present, while small points of brighter fluorescence indicate that the denser insoluble aggregates are present."

To allow for this distinction between forms, the research team reengineered the color-causing core of the green fluorescent protein (GFP), which is commonly used in imaging studies because it fluoresces when exposed to certain wavelengths of light. The reengineered compound binds to a tag, which in turn fuses to a protein targeted for imaging.

The research team used two different kinds of commercially available tags, Halo-tag and SNAP-tag, which when used with AggTag can induce red or green <u>fluorescence</u>, respectively. Because Halo-tags and SNAP-tags do not interact with each other, they can be used to simultaneously image two different proteins with the two colors. The team also engineered the tags so that the green and red colors can be reversed, giving researchers options for future imaging.

"We plan to continue developing this method so that we can signal the transition of oligomers into insoluble aggregates using a color change," said Zhang. "This method provides a new toolbox to study protein aggregation, which is currently a highly studied topic among scientists. Hopefully this will allow us to better understand the entire process of protein aggregation and the role of each of these forms in the progression of neurodegenerative and other diseases."

More information: Kwan Ho Jung et al, A fluorogenic AggTag method based on Halo- and SNAP-tag to simultaneously detect aggregation of two proteins in live cells, *ChemBioChem* (2019). DOI: 10.1002/cbic.201800782

Yu Liu et al. Modulation of Fluorescent Protein Chromophores To



Detect Protein Aggregation with Turn-On Fluorescence, *Journal of the American Chemical Society* (2018). DOI: 10.1021/jacs.8b02176

Provided by Pennsylvania State University

Citation: New method uses fluorescence to identify disease-causing forms of proteins (2019, February 13) retrieved 19 April 2024 from https://phys.org/news/2019-02-method-fluorescence-disease-causing-proteins.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.