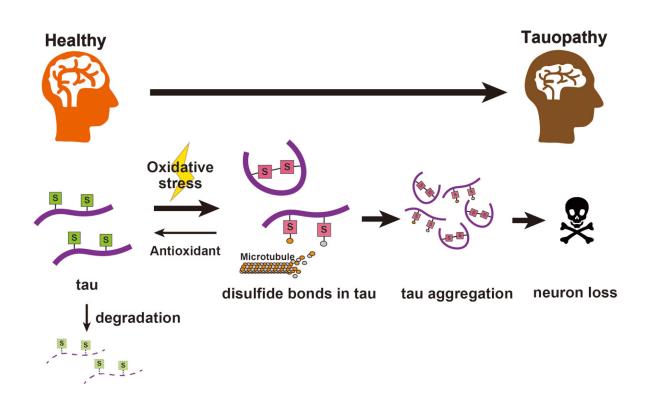


Toxicity of protein involved in Alzheimer's triggered by a chemical 'switch'

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Tau proteins with cysteine groups bearing thiol groups (S) undergo chemical changes under oxidative stress to form disulfide bonds, making a toxic mutant of the tau protein that can aggregate. These go on to cause neural degeneration. Antioxidants can help reduce these back to thiols; these normal tau proteins can then be naturally cleared away by the cell. Credit: Tokyo Metropolitan University

Researchers from Tokyo Metropolitan University have discovered that a



specific chemical feature of a key protein known as tau may cause it to accumulate in the brain and trigger illnesses like Alzheimer's. They found that disulfide bonds on certain amino acids act to stabilize tau and cause it to accumulate, an effect that got worse with increased oxidative stress. The identification of chemical targets triggering tau accumulation may lead to breakthrough treatments.

The tau <u>protein</u> is key to the healthy function of biological <u>cells</u>. It helps form and stabilize microtubules, the thin filaments that crisscross cell interiors to help keep them structurally rigid and provide 'highways' to shuttle molecules between organelles. However, when they are not formed correctly, they can accumulate and form sticky clumps. In the brain, these aggregates block the firing of neurons and cause a wide range of neurodegenerative diseases known as tauopathies, one of which is Alzheimer's disease. It is vastly important that scientists find the 'switch' that transforms tau from an indispensable part of cell function to a deadly pathology.

A team led by Associate Professor Kanae Ando of Tokyo Metropolitan University has been using model organisms like the Drosophila fruit fly to uncover how specific features of the tau protein cause it to stop working properly. Flies can be genetically altered to express the same <u>tau</u> <u>protein</u> as in humans. By systematically modifying parts of the gene encoding for tau, they have been trying to pinpoint how certain features of mutant tau proteins affect their behavior.

In their most recent work, they found that alterations to <u>amino acid</u> <u>residues</u> in the protein known as cysteines in two different locations (C291 and C322) had a drastic effect on the amount and toxicity of tau. In a further breakthrough, the team pinned down the specific chemical feature responsible for making them toxic to normal cell function, that is, <u>disulfide bonds</u> formed by these cysteine groups. The toxic accumulation of tau got worse when cells were put in an environment



with elevated levels of reactive oxygen species, as thiol groups on the cysteines were oxidized to form disulfide links. Biochemical environments with elevated oxidative stress are similar to those seen in patients with tauopathies. The co-expression of antioxidants to counter this effect helped natural processes clear away tau proteins, resulting in dramatically lower tau levels.

The team hope that knowledge of exactly which chemical groups are responsible for tau toxicity may lead to novel therapies which reduce or prevent <u>tau accumulation</u>, helping sufferers of tauopathies around the world.

More information: Taro Saito et al, Disulfide bond formation in microtubule-associated tau protein promotes tau accumulation and toxicity in vivo, *Human Molecular Genetics* (2021). <u>DOI:</u> 10.1093/hmg/ddab162

Provided by Tokyo Metropolitan University

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