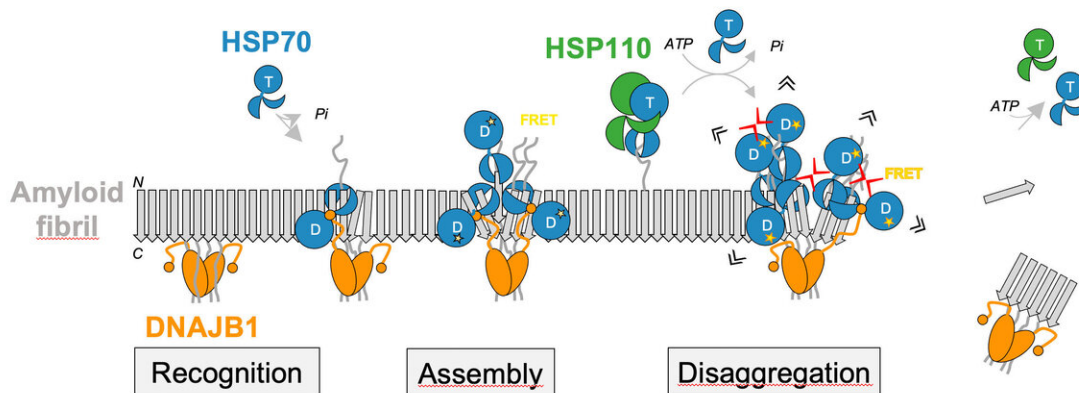


# How molecular chaperones dissolve protein aggregates linked to Parkinson's disease

November 11 2020



Disaggregation of  $\alpha$ -synuclein amyloid fibrils relies on the cooperation of the HSP70 chaperone with its co-chaperones DNAJB1 and HSP110. This cooperation ensures recognition of the fibrils, followed by the correct assembly of the Hsp70 disaggregase machinery on the fibril surface thereby allowing for productive disaggregation. Credit: Bernd Bukau / Heidelberg University

In many neurodegenerative diseases like Parkinson's, protein aggregates form in the brain and are assumed to contribute to neuronal cell death. Yet there exists a cellular defense mechanism that counteracts these

aggregates, known as amyloid fibrils, and can even dissolve fibrils already formed. This defense mechanism is based on the activity of molecular chaperones, i.e. protein folding helpers, of the heat shock protein 70 family (Hsp70).

Molecular biologists from Heidelberg University and the German Cancer Research Center investigated how the Hsp70 system disaggregates [amyloid fibrils](#) of the Parkinson-specific [protein](#)  $\alpha$ -synuclein in a test tube. The research team led by Dr. Bernd Bukau expects their research results to provide new insights into how Parkinson's disease develops and what might be done to influence it. The results were published in two articles in the journal *Nature*.

Proteins in all cells, from bacteria to human cells, need to fold to their native state. The chains of amino acid building blocks assume specific three-dimensional structures that give proteins their functionality. This state of correct folding is constantly threatened by external and internal influences that can lead to incorrectly folded and hence damaged proteins. There is a risk that damaged proteins will aggregate, or clump together into longer strands, the amyloid fibrils. This happens with  $\alpha$ -synuclein in Parkinson's disease, for example. The fibrils, in turn, are the starting point for even larger deposits.

"The formation process of such fibrillar aggregates can damage cells and even lead to cell death, as is the case in neurodegenerative diseases like Parkinson's and Alzheimer's," explains Bukau, a researcher at the Center for Molecular Biology of Heidelberg University (ZMBH) and the German Cancer Research Center (DKFZ).

Bukau's research focuses on how these [protein aggregates](#) can be dissolved. In earlier work, he and his team succeeded in identifying a cellular activity that plays a vital role in dissolving fibrillar aggregates, which is based on chaperones of the Hsp70 family. Hsp70 chaperones

help other proteins with folding and can even isolate and refold aggregated proteins. The latest research by Bukau and postdoc Dr. Anne Wentink shows the effects the Hsp70 chaperones have on the Parkinson-specific amyloid fibrils of the  $\alpha$ -synuclein protein.  $\alpha$ -synuclein is a small protein that helps in the release of messengers called neurotransmitters in the brain, although its exact function remains unclear. It became known because massive deposits of this specific protein were found in Parkinson's patients and has been causally linked to the disease.

In biochemical experiments, the Heidelberg scientists were recently able to show that the human Hsp70 chaperone relies on the assistance of two specific co-chaperone partners to dissolve the amyloid fibrils of the  $\alpha$ -synuclein protein. A precisely regulated interaction of these proteins leads to the formation of chaperone complexes on the surface of the fibrils, which then break up the aggregates.

"It is the sheer local accumulation of many chaperone proteins on the surface of the  $\alpha$ -synuclein fibrils that generates the force to break up the fibrils and detach the  $\alpha$ -synuclein molecules," explains Dr. Wentink. The close proximity between the chaperones on the restricted surface of the fibrils plays a decisive role in creating strong enough pulling forces to disrupt the fibrils.

The second study, published in *Nature*, focuses on a previously unknown regulatory mechanism, a type of molecular switch that sets in motion the overall Hsp70 chaperone activity to dissolve the amyloid fibrils. This mechanism is based on a sequence of direct interactions between the different parts of the DNAJB1 co-chaperone and the Hsp70 chaperone. This ultimately activates Hsp70 to use ATP as an energy source, making it possible to productively bind to the fibrils and effect their disintegration.

"Our latest results from the two studies give us a molecular

understanding of how amyloid fibrils are dissolved. We were able to demonstrate that the chaperones work like a machine to dissolve the fibrils," adds Bukau. According to the Heidelberg researcher, this opens up new avenues for the development of agents that specifically target the chaperone-based cellular defense mechanism against [amyloid](#) formation. A better understanding of how this [chaperone](#) activity influences the course of [neurodegenerative diseases](#) will thus be of essential importance in the therapeutic exploitation of the findings described in these studies.

**More information:** Wentink et al., Molecular dissection of amyloid disaggregation by the human HSP70 chaperone, *Nature* (2020). [DOI: 10.1038/s41586-020-2904-6](https://doi.org/10.1038/s41586-020-2904-6), [www.nature.com/articles/s41586-020-2904-6](https://www.nature.com/articles/s41586-020-2904-6)

Ofrah Faust et al. HSP40 proteins use class-specific regulation to drive HSP70 functional diversity, *Nature* (2020). DOI: [10.1038/s41586-020-2906-4](https://doi.org/10.1038/s41586-020-2906-4) , <https://www.nature.com/articles/s41586-020-2906-4>

Provided by Heidelberg University

Citation: How molecular chaperones dissolve protein aggregates linked to Parkinson's disease (2020, November 11) retrieved 25 March 2023 from <https://medicalxpress.com/news/2020-11-molecular-chaperones-dissolve-protein-aggregates.html>

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