

Scientists discover emergency pathway to help human cells with protein damage survive

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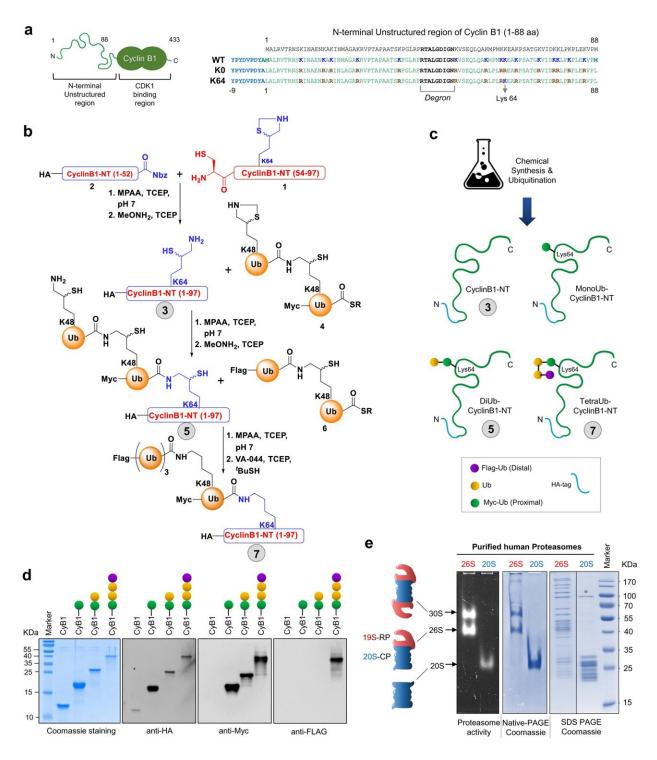


Fig. 1: Chemical synthesis and ubiquitination of ubiquitin-conjugates. a A cartoon depicting cyclin B1 with its 1–88 aa residue unstructured N-terminal region (NT), and a C-Terminal CDK1 binding region (left). The primary amino acid sequence of the N-terminal region of cyclin B1 that encompasses the



Degron and 15 Lysine residues (right). Below are the wild-type (WT) and K0-mutant (all Lysine to Arginine) Cyclin B1-NT sequences used in this study for mammalian cell expression. The K64 HA-Cyclin B1-NT sequence was used for chemical ubiquitination of substrates (bottom). b Schematic illustration of the strategy used to synthesize differentially tagged TetraUbK48-HA-Cyclin B1-NT (Nbz = N-acyl-benzimidazolinones, SR = 3-mercaptopropionic acid). The detailed chemical synthesis protocol is described in Supplementary Methods. c A cartoon representation of a set of four synthetic substrates designed to study proteasome function. Tagged (Myc/Flag) and/or untagged ubiquitin units were attached to lysine64 of the HA-Cyclin B1-NT to obtain ubiquitinated conjugates. The numbers 3, 5, and 7 correspond to the three synthetic substrates in panel (b) (see also Supplementary Methods). d All four synthetic ubiquitinated Cyclin B1-NT conjugates were resolved by SDS-PAGE and stained with Coomassie or probed with the indicated antibodies for immunoblotting (IB). e Human 30S/26S and 20S proteasomes purified from human erythrocytes were resolved by native gel for in-gel activity (left), or for Coomassie (center), or resolved by SDS-PAGE for Coomassie staining. The band marked by * near 90 kDa was identified as HSP90AA1 and HSP90AB1 in MS/MS analysis. All other bands were confirmed to contain primarily 20S (right) or 26S (to its left) subunits. Source data are provided as a Source data file. Credit: DOI: 10.1038/s41467-021-26427-0

Cell proteins damaged by oxygen radicals can be chemically "tagged" for elimination, but an "emergency pathway" bypasses strict protocol and can eliminate even without the need for prior tagging.

An international research team headed by Technion scientists has found an alternative manner for eliminating damaged proteins when the cells are impaired by "oxygen radicals," as can happen in failing human hearts where there is poor cell respiration and cells become oxygen depleted, or suffer "hypoxia," because of poor oxygen uptake.

Significantly, the researchers discovered that there can be a shift from



the tightly controlled process of eliminating proteins in the cells to a less strict mechanism when cells enter an "emergency protocol." This shift can "clear up" the toxic proteins before their toxicity levels get too high.

Their study was published on 26 October 2021 in *Nature Communications*. To carry out their study, the researchers investigated several "proteasomes," protein complexes that work by a chemical reaction to degrade unneeded or damaged cell proteins. The researchers found that elevated levels of one type of proteasome, 20S, appears to contribute to <u>cell survival</u>, even for those cells under stress from damaged proteins.

Human cells—both functional and damaged—are constantly recycled by chemically "tagging" and targeting for removal when they are under stress by the ubiquitin system (2004 Nobel Prize in chemistry). At the same time, a few proteins that are intact and functional can also be dragged into the 20S proteasome "molecular disposal unit" along with the toxic proteins that have be targeted for destruction. Nevertheless, rather than harm cells, this mode of action by 20S proteasome may aid cells in rapidly remove toxic proteins. In their conclusion, the authors raised the interesting speculation that this emergency pathway can help even damaged cells to withstand bouts of stress and allow them to "age gracefully."

To carry out the study, Technion researchers Professors Indrajit Sahu, Michael Glickman, Ashraf Brik, and Oded Kleifeld, worked with Professor Sharlene Day, from the University of Pennsylvania, and the research team of Professor Yao Cong of the Chinese Academy of Sciences in Shanghai, China.

More information: Indrajit Sahu et al, The 20S as a stand-alone proteasome in cells can degrade the ubiquitin tag, *Nature Communications* (2021). DOI: 10.1038/s41467-021-26427-0



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