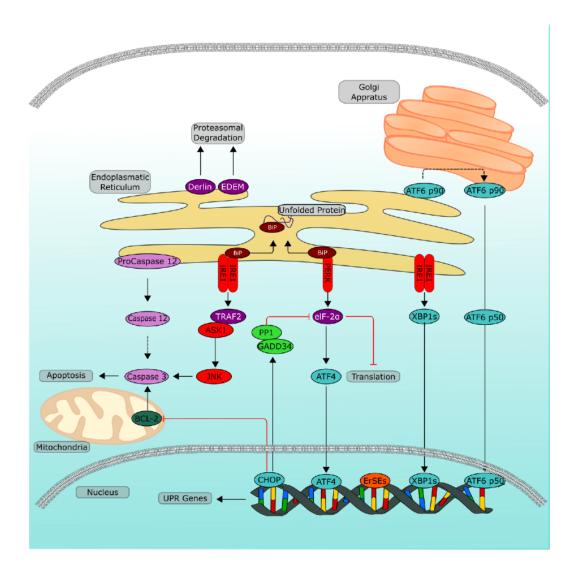
Unfolded Protein Response



The unfolded protein response (UPR) is a cellular stress response activated in eukaryotic cells in response to endoplasmic reticulum (ER) stress, an accumulation of unfolded proteins in the ER lumen. ER Stress can be caused by different conditions such as high protein demand, viral infection, energy deprivation or excessive oxidative stress. UPR signalling is highly regulated and dynamic and integrates information about the type, intensity, and duration of the stress stimuli, thereby determining cell fate1.

The regulation of UPR is mediated by three major proteins, namely inositol-requiring enzyme 1 (IRE1), activating transcription factor 6 (ATF6) and protein kinase RNA-like ER kinase (PERK)2. The unfolded Protein Response Regulator BiP/GRP78 has the abiliity to bind and inactivate them3. Misfolded proteins bind and sequester BiP which releases the UPR effector proteins and therefore reactivate them. Activation of PERK leads to phosphorylation of eukaryotic initiation factor (eIF). Phosphorylated elf-2α inhibits protein translation in order to restore homeostasis and enables ATF4 as well. In response to ER stress, ATF6p90 transits to the Golgi where it is cleaved by site-1 protease (S1P) and site-2 protease (S2P), yielding the active transcription factor, ATF6p504. IRE1 oligomerises and activates its ribonuclease domain through auto phosphorylation. Activated IRE1 catalyses the excision of a 26 nucleotide intron from XBP1u mRNA, in a similar manner to pretRNA splicings. Removal of this intron causes a frame shift in the XBP1 coding sequence resulting in the translation of a 376 amino acid, 40 kDa, XBP-1s isoform.

Active ATF6p50 and XBP1 subsequently bind to the ER stress response element (ERSE) and the UPR element (UPRE), leading to expression of target genes encoding ER chaperones and ER-associated degradation (ERAD) factors involved in degradation of unfolded proteins6. The outcome of UPR activation increases protein folding, transport and ER-associated protein degradation (ERAD), while attenuating protein synthesis to restore ER homeostasis. If these adaptive mechanisms cannot resolve the protein-folding defect, cells enter apoptosis.

UPR inhibitor drugs have been developed to tackle some types of cancers as well as neurodegenerative disorders. UPR inhibitors are also tested as antivirals against SARS-CoV-27.

References:

- 1. Hetz C, (2012): "The unfolded protein response: Controlling cell fate decisions under ER stress and beyond." *Mol Cell Biol.*;13:89–102.
- 2. Ron D, (2007): "Signal integration in the endoplasmic reticulum unfolded protein response." *Nat Rev Mol Cell Biol.*, Jul;8(7):519-29.
- 3. Bertolotti A, Zhang Y, (2000): "Dynamic interaction of BiP and ER stress transducers in the unfolded-protein response." *Nat Cell Biol*, Jun;2(6):326-32.
- 4. Jin Ye (2000): "ER Stress Induces Cleavage of Membrane-Bound ATF6 by the Same Proteases that Process SREBPs" *Molecular Cell*, Volume 6, Issue 6, Pages 1355-1364
- Yanagitani K (2009): "Cotranslational targeting of XBP1 protein to the membrane promotes cytoplasmic splicing of its own mRNA." Mol Cell, Apr 24;34(2):191-200
- 6. Zhang K, Kaufman RJ. (2006): "The unfolded protein response: a stress signaling pathway critical for health and disease." *Neurology*, 66:S102-9
- 7. Echavarría-Consuegra L, Cook GM et al (2021): "Manipulation of the unfolded protein response: A pharmacological strategy against coronavirus infection." *PLOS Pathogens*, 10.1371/journal.ppat.1009644