The N-Terminal Methionine of Cellular Proteins as a Degradation Signal

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Figure 1. Specific Binding of Ubr1 to Unacetylated N-Terminal Methionine Followed by a Hydrophobic Residue

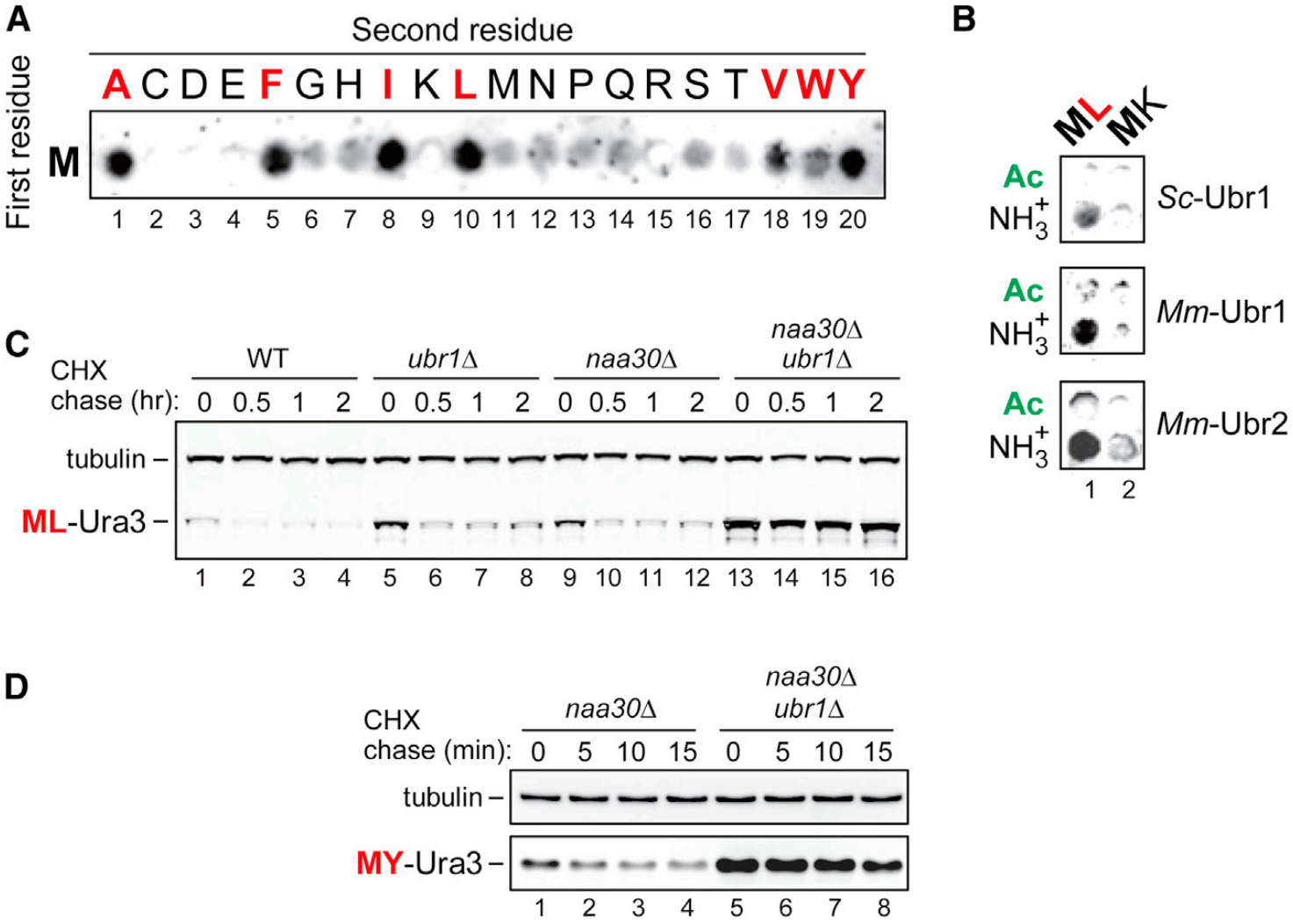


Figure 2. Unacetylated N-Terminal Methionine as an N-Degron of the Arg/N-End Rule Pathway

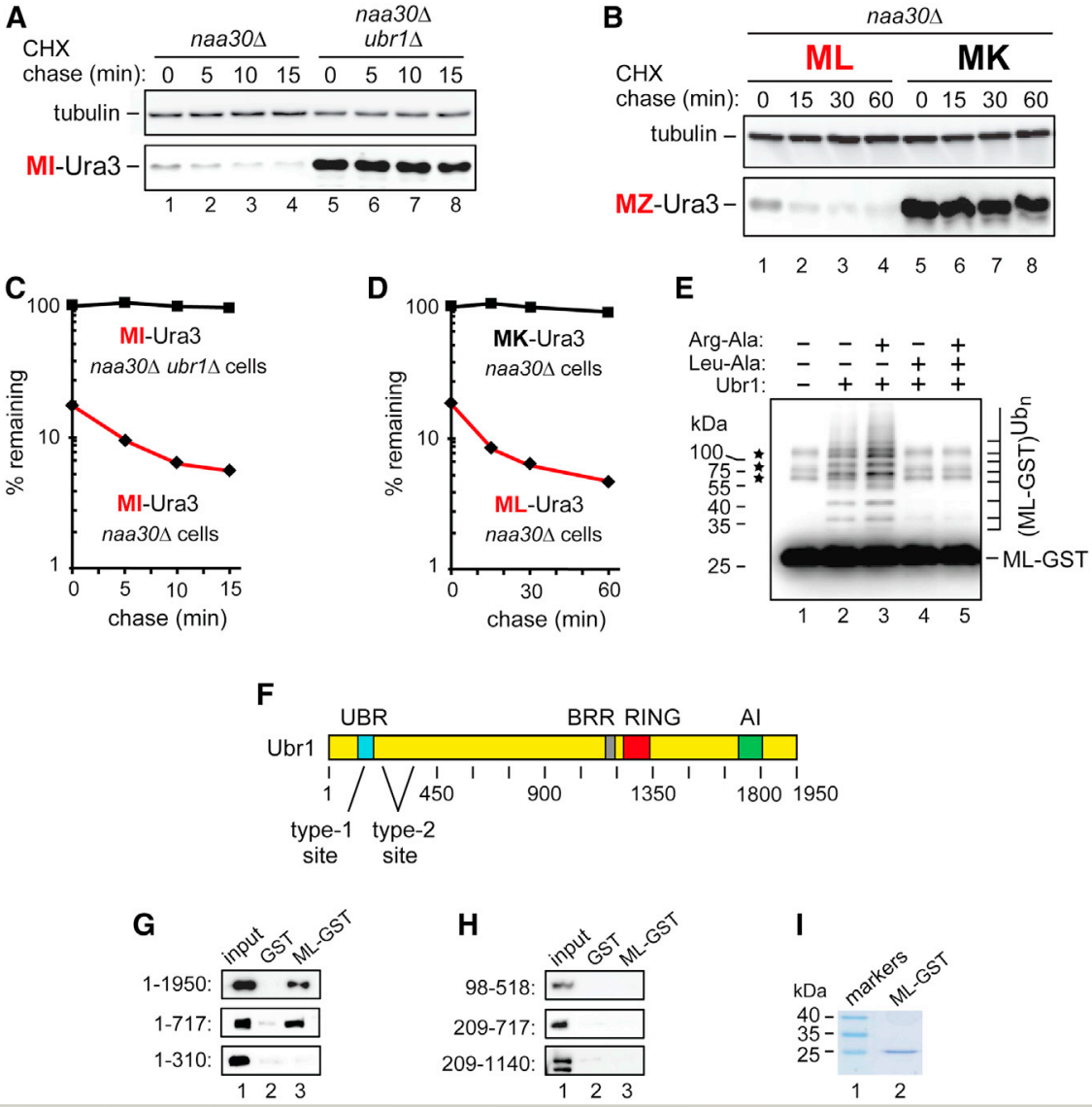


Figure 3. Misfolded Proteins Containing Met-Based N-Degrons

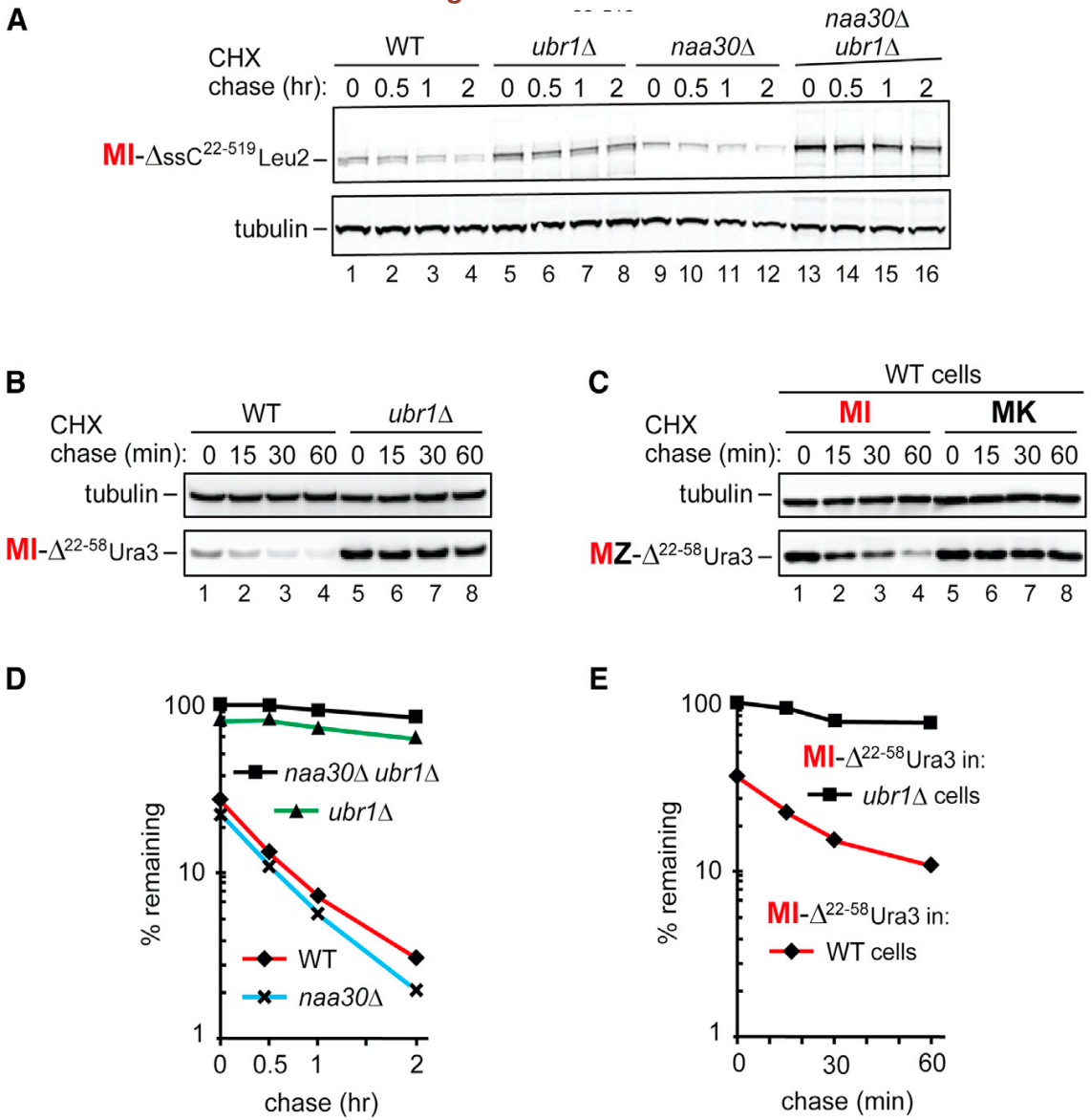


Figure 4. The Natural ML-Msn4 and MF-Arl3 Proteins Contain Met-Based N-Degrons

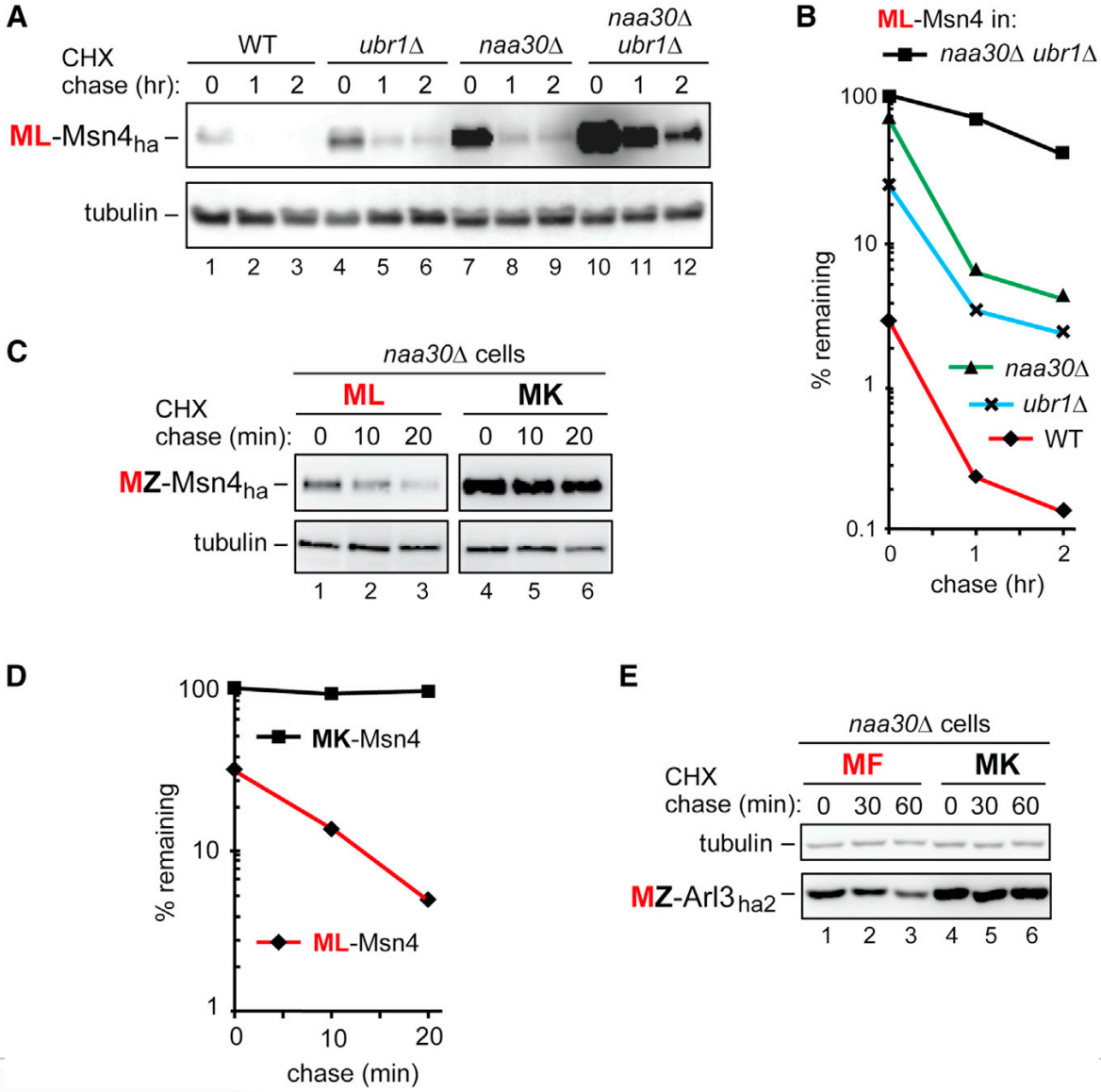


Figure 5. The Natural MF-Pre5 and MI-Sry1 Proteins Contain Met-Based N-Degrons

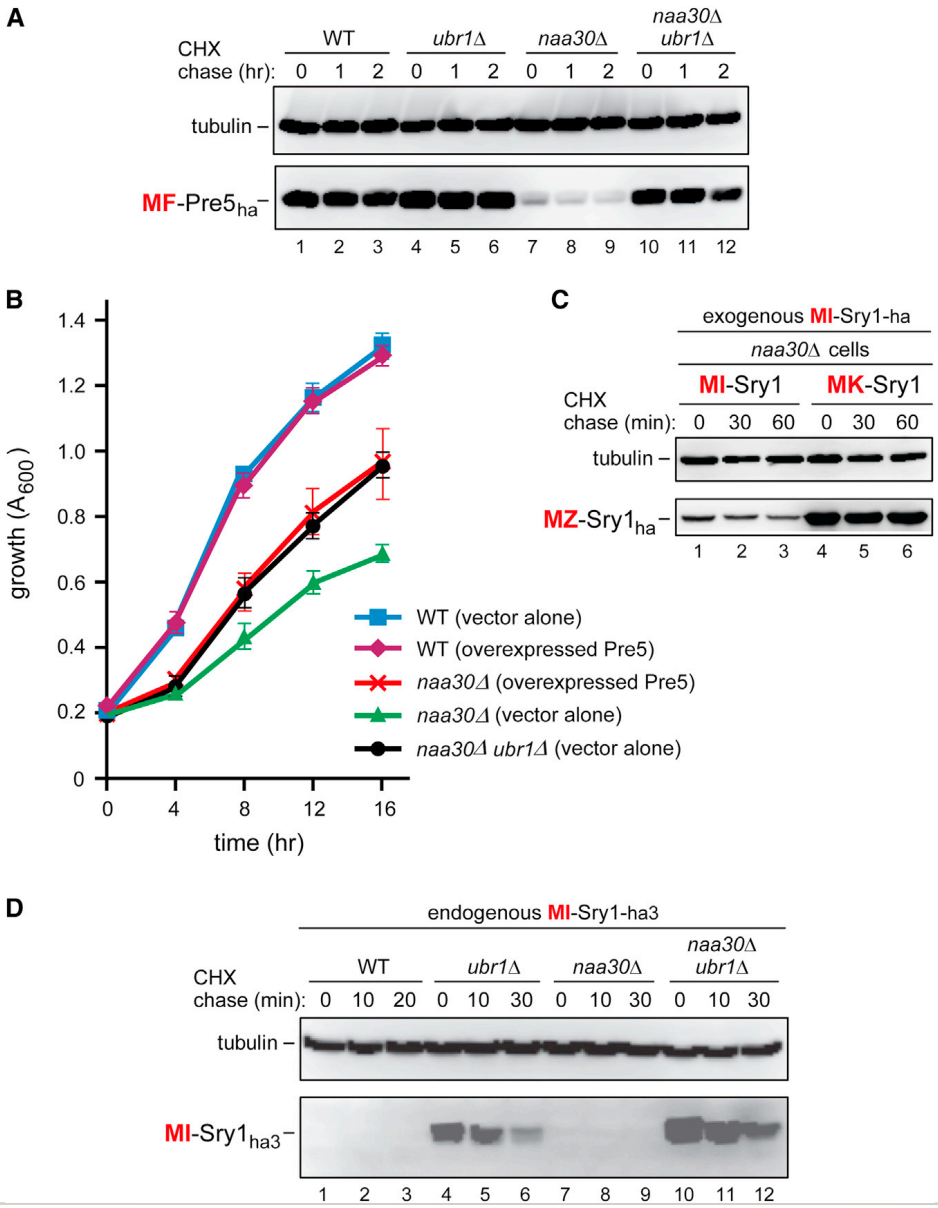


Figure 6. Complementary Specificities of the Arg/N-End Rule Pathway and the Ac/N-End Rule Pathway

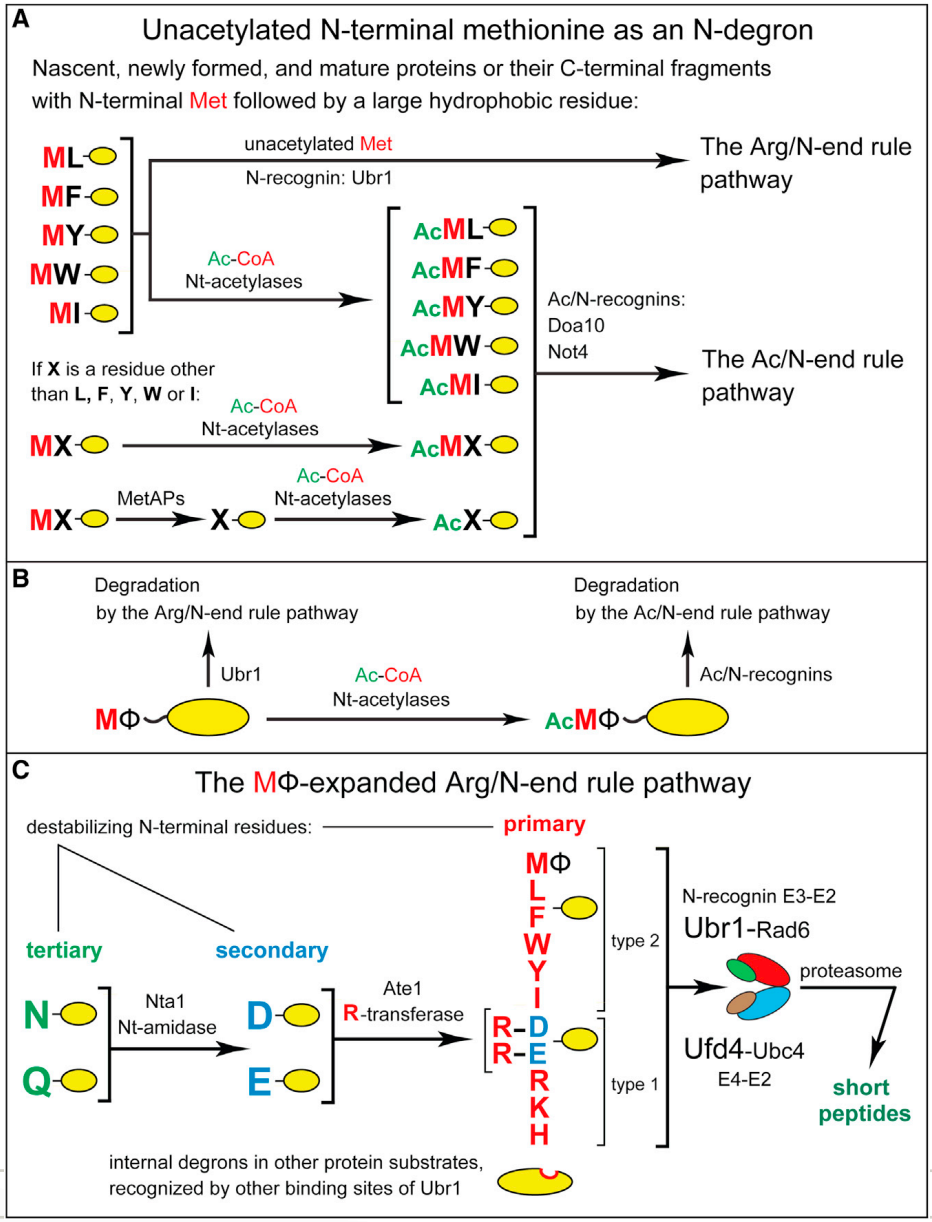


Figure 7. Conditionality of Ac/N-Degrons and Protein Remodeling by the N-End Rule Pathway

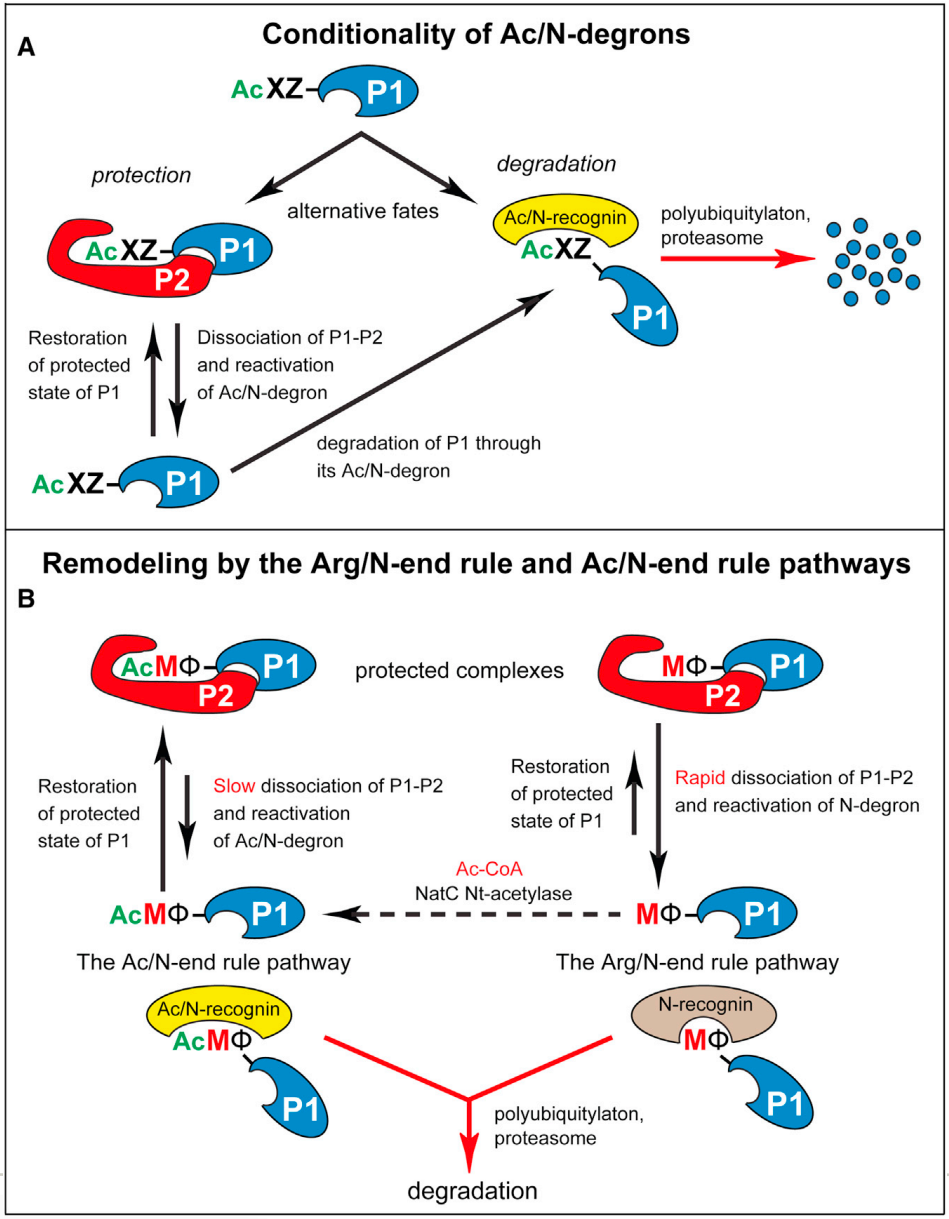


Figure S1. The Ac/N-End Rule Pathway and the Arg/N-End Rule Pathway, Refers to Figures 1, 2, 3, 4, 5, 6, and 7

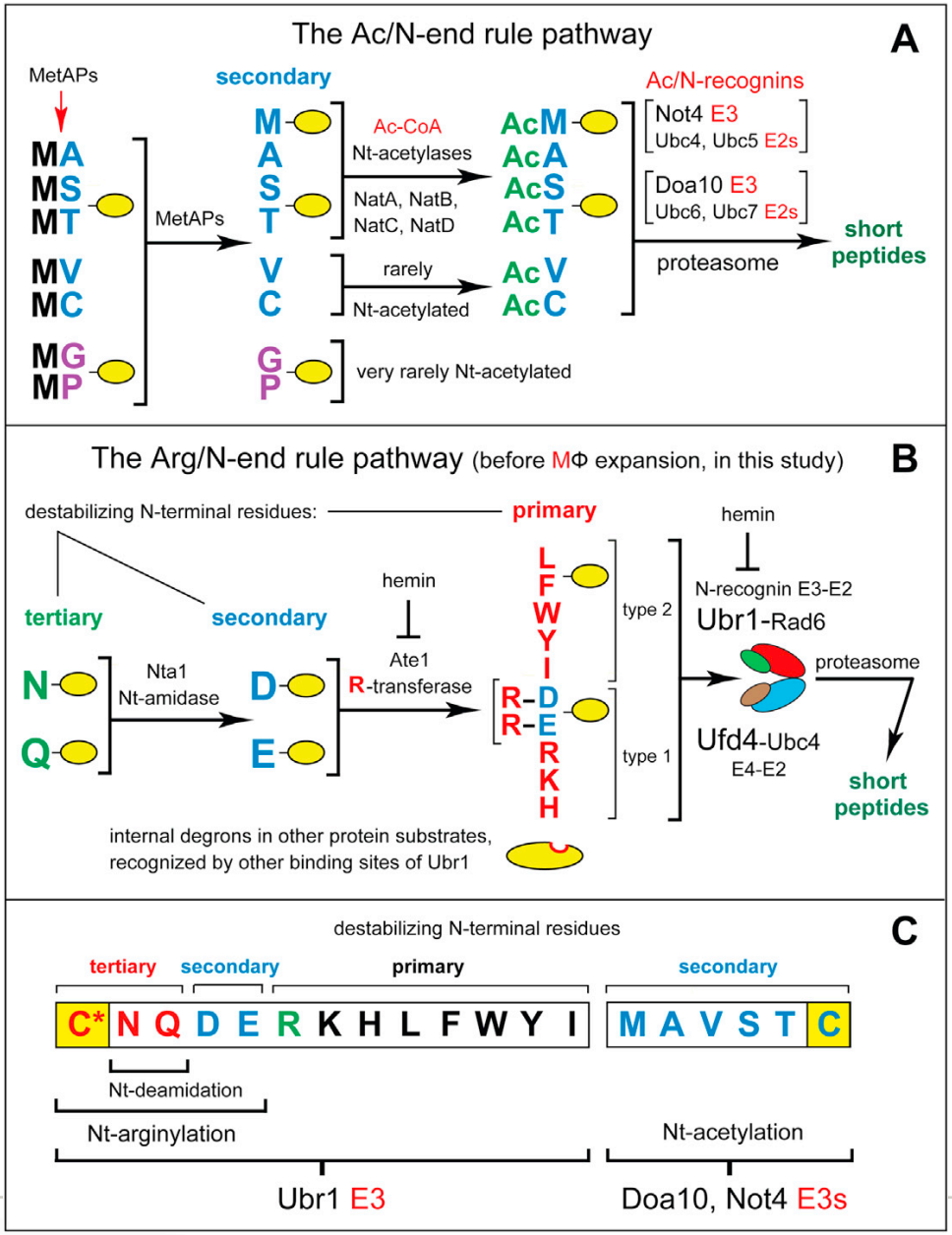


Figure S2. Substrate Specificities and Subunit Compositions of *S. cerevisiae* N$^{\alpha}$-Terminal Acetylases (Nt-acetylases), Refers to Figures 1, 2, 3, 4, 5, 6, and 7

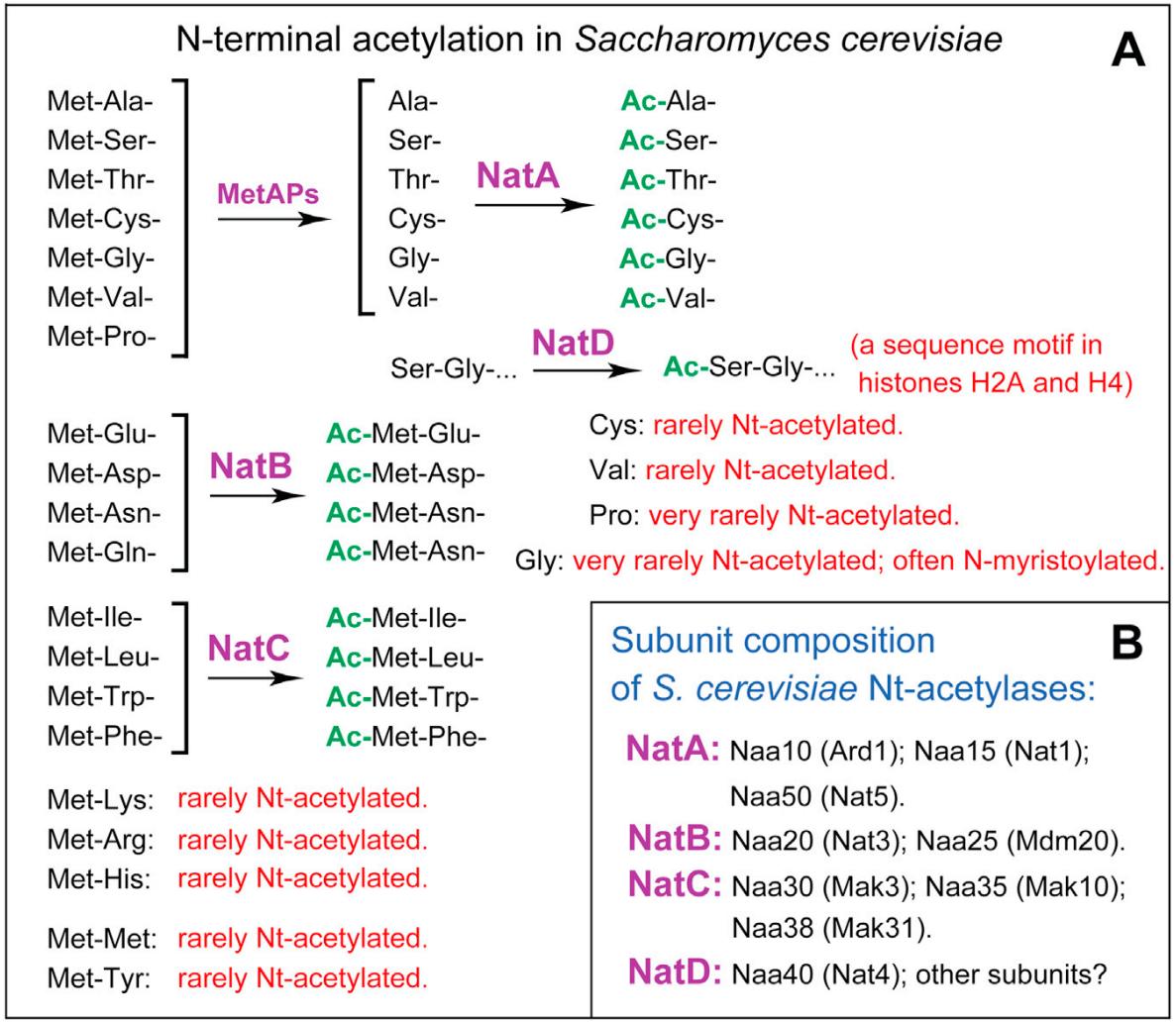


Figure S3. ^{35}S-Pulse-Chase and Cycloheximide-Chase Assays with ML-Ura3 and MI-Sry1, Refers to Figures 1 and 5

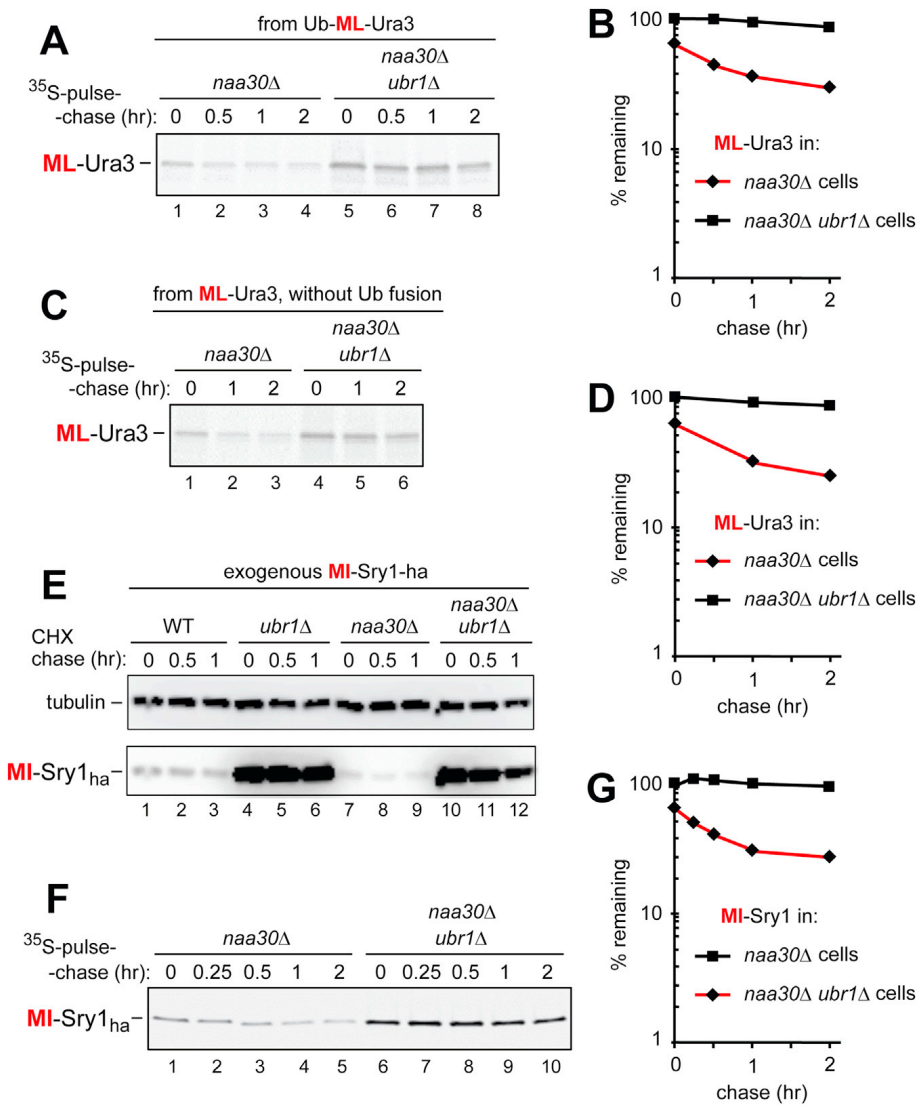
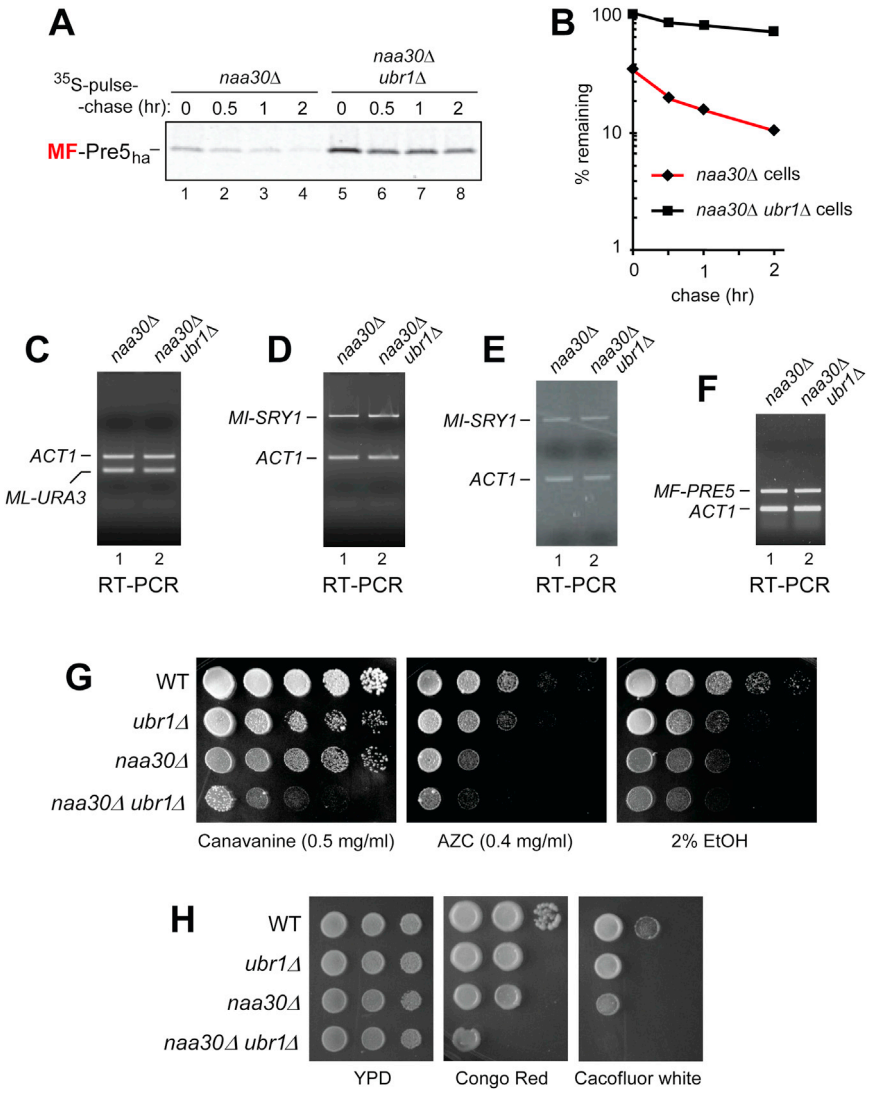


Figure S4. Instability of the MF-Pre5 Proteasomal Subunit and Hypersensitivity of *naa30Δ* and *naa30Δ ubr1Δ* Cells to Multiple Stresses, Refers to Figures 2 and 5



That's all Folks!