



Figure S3: DDR and Mitochondrial strains of interest. A, B) 3-way, area-proportional Venn diagrams demonstrating overlap between CF and LVS-nominated gene sets with the DDR and mitochondrion-annotated gene sets, respectively. C) LVS-specific subnetwork of YeastNet V3 with edge weight thresholded to ≥ 1.5 . Black node border denotes predicted human ortholog, green node border denotes established human complementation, yellow nodes denote CCR4-NOT complex components, blue nodes denote tRNA wobble modification components, and orange nodes denote autophagy components.