Supplemental Figure 1

A) Network representation of various protein interactions, with node color indicating fitness score and edge width indicating the number of times the interaction has been published.

B) Close-up view of mitochondrial ribosomal subunits, with p-values indicating statistical significance:
- Cytosolic ribosome: p-value = 1.5348E-13
- Mitochondrial small ribosomal subunit: p-value = 6.6058E-12
- Mitochondrial large ribosomal subunit: p-value = 5.605E-13

C) OD595 growth curves for WT and rpl2bΔ strains over time (h):
- LZM100
- LZM1

Edge color:
- Genetic Interaction
- Physical Interaction
- Genetic & Physical

Edge width is an indicator of the number of times the interaction has been published.