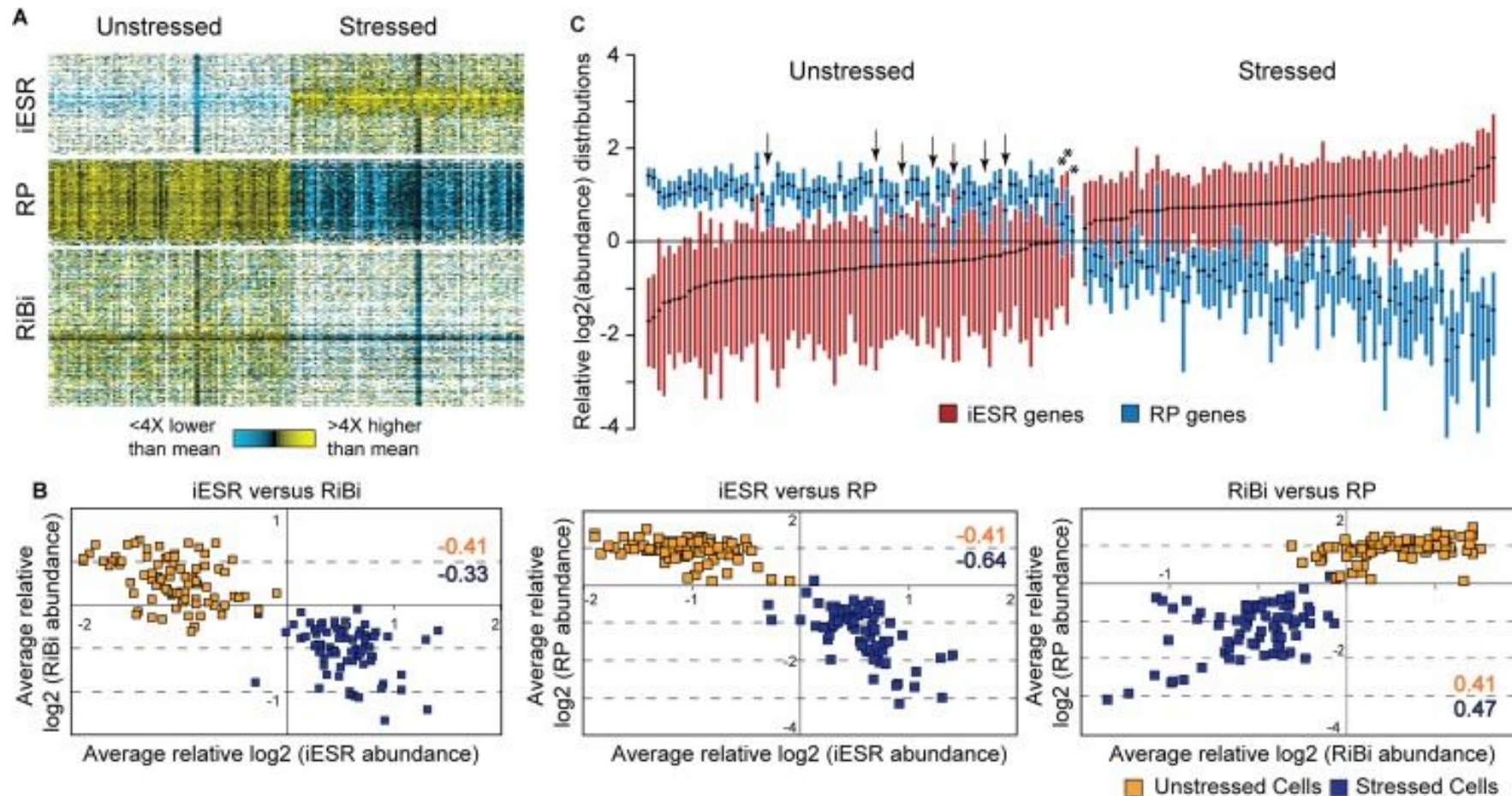
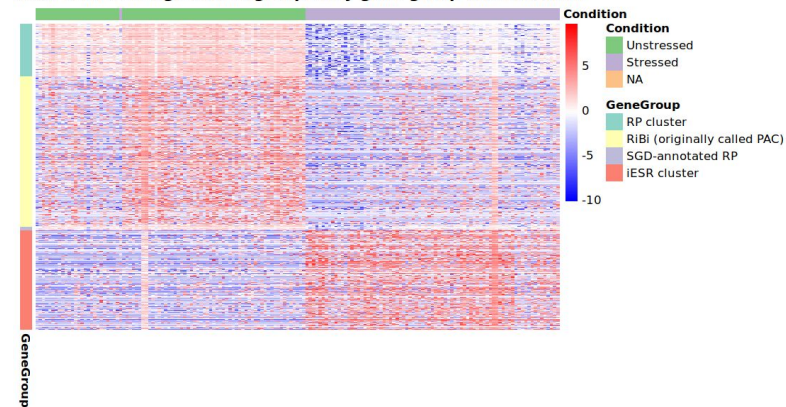


**Fig.1 from the original study - Gasch et al. (2017)**

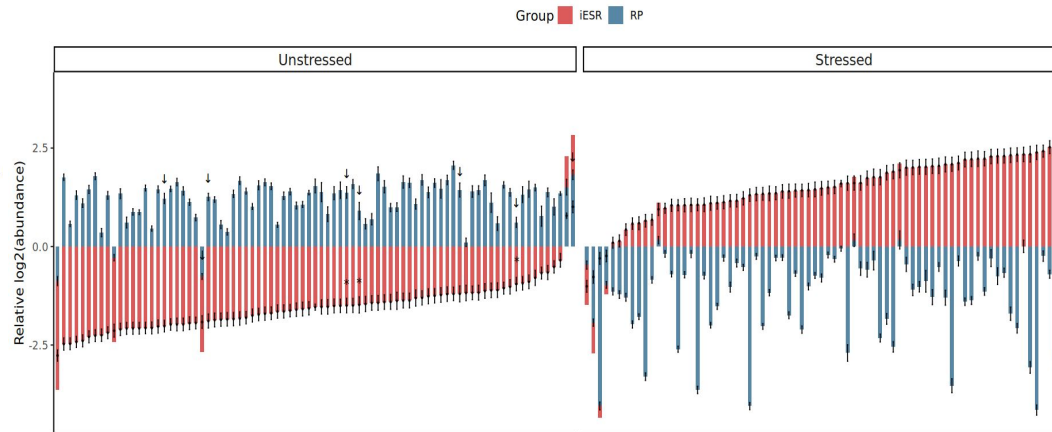


# Replication of Fig. 1: ESR-driven Transcriptomic Variation in Stressed & Unstressed Yeast Cells

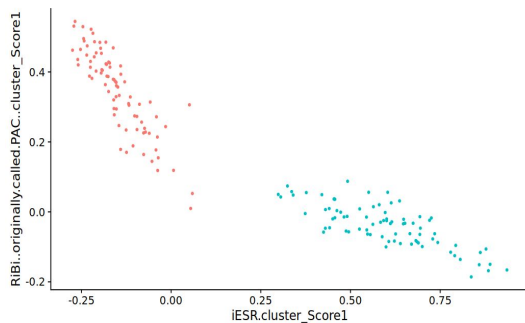
Mean-centered log2 counts grouped by gene group and condition



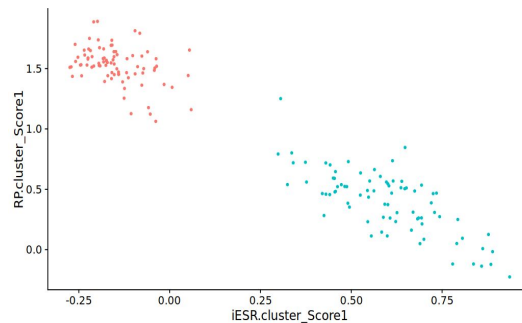
Relative log2(abundance) of RP and iESR genes per cell



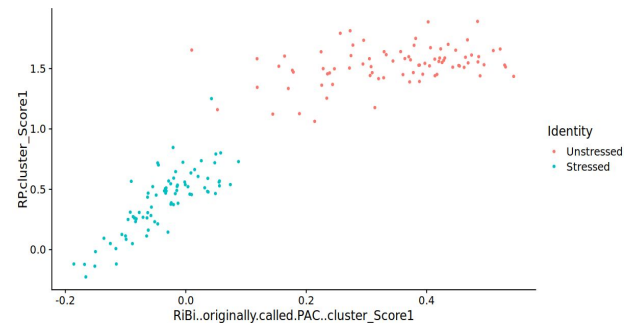
iESR vs RiBi



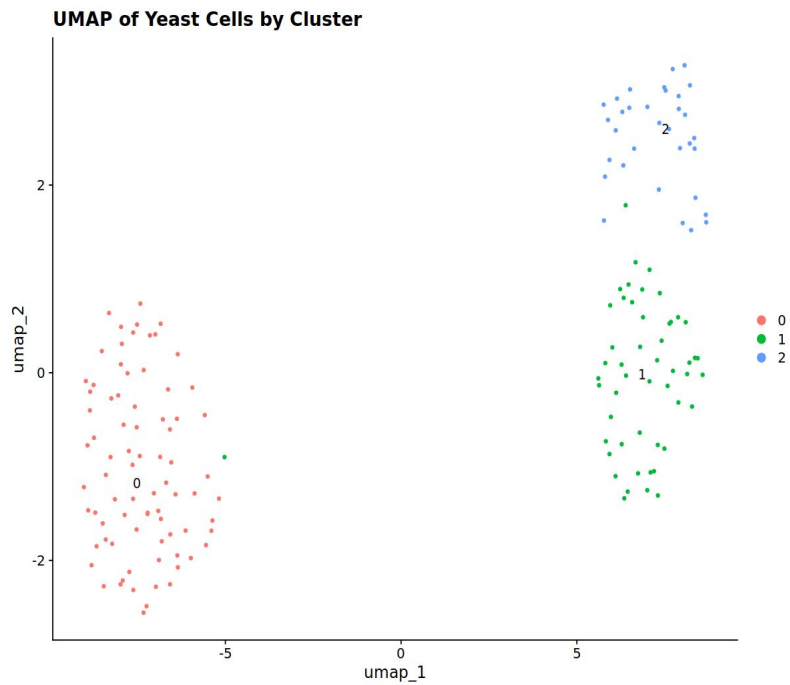
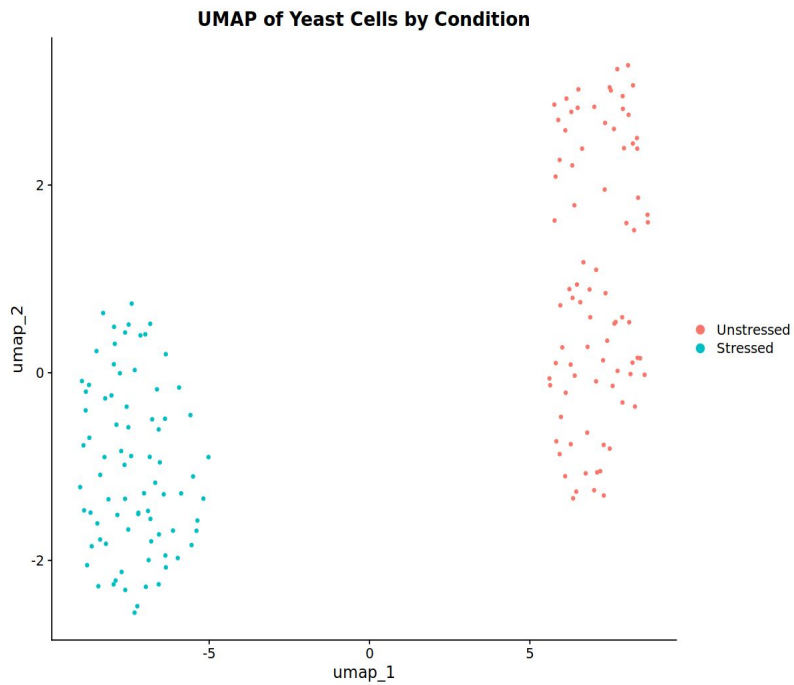
iESR vs RP



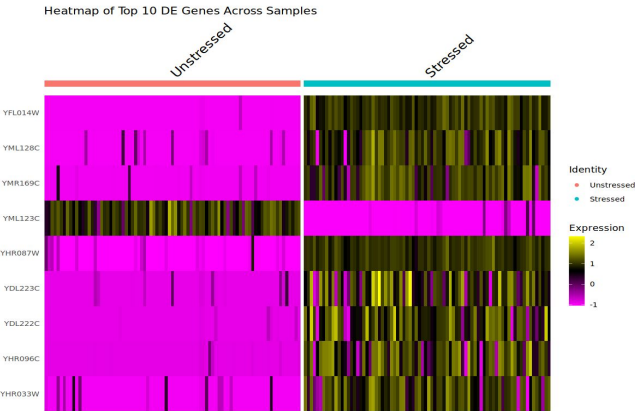
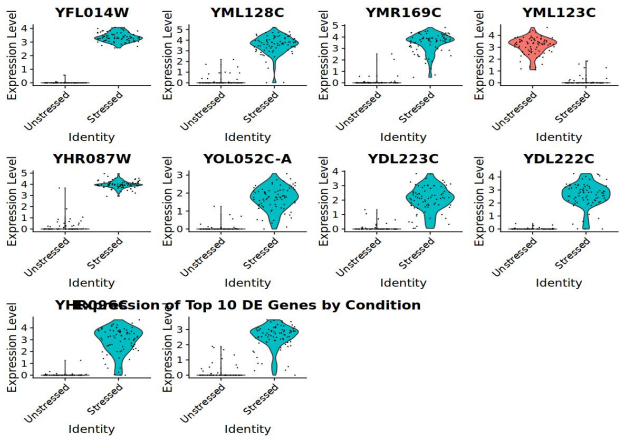
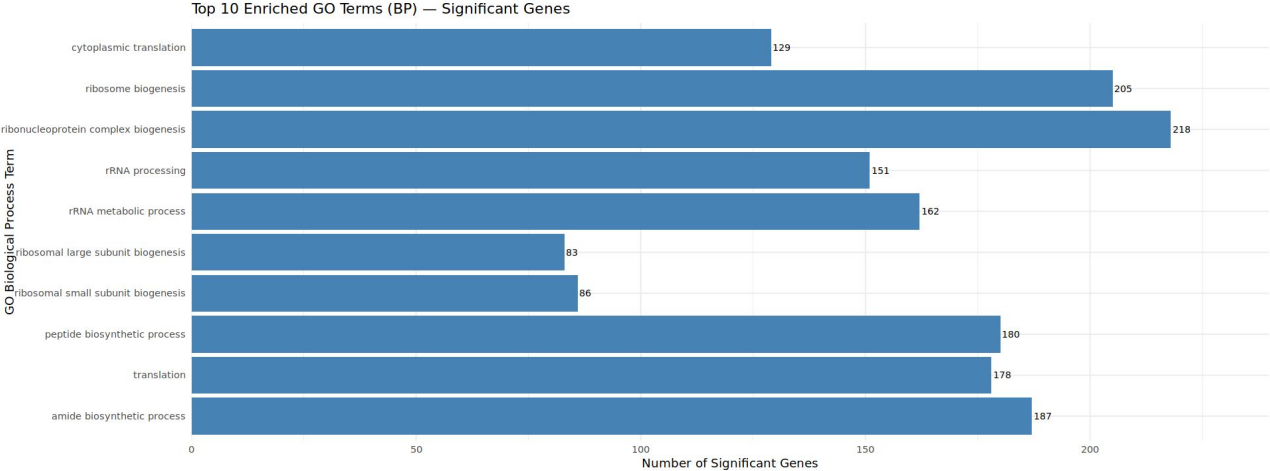
RiBi vs RP



# PCA and UMAP Reveal Distinct Transcriptomic Clusters by Condition in Yeast Cells



# Functional Enrichment & Expression Signatures Define Condition-Specific Responses in Yeast



# Gene Module Score Distributions by Condition

Gene Module Scores by Condition

