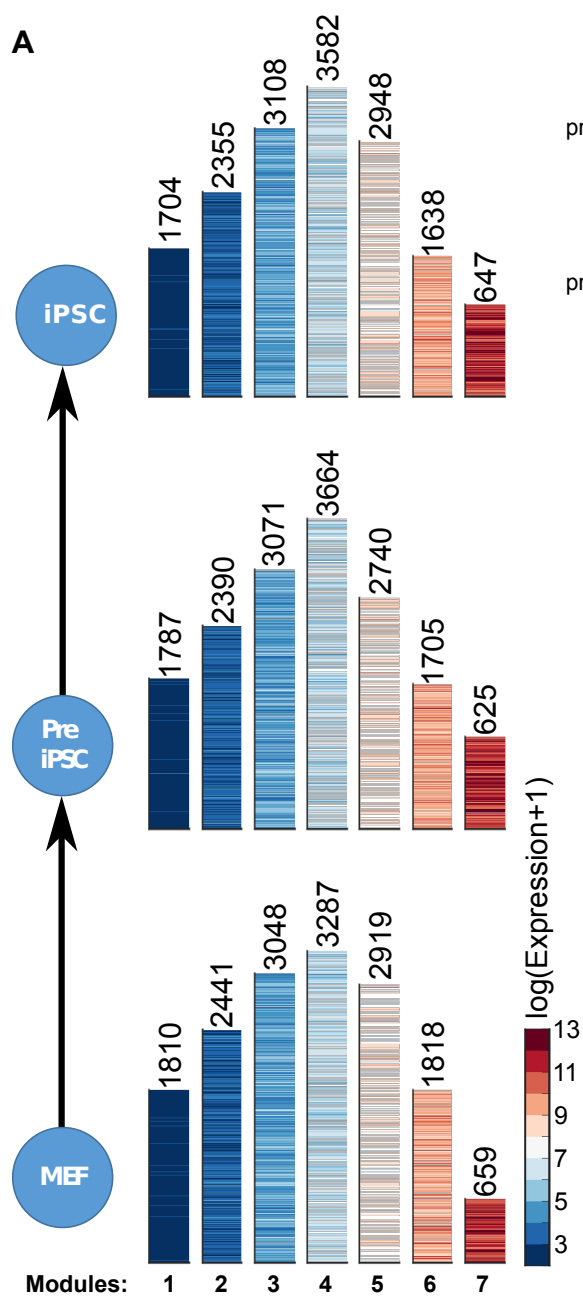
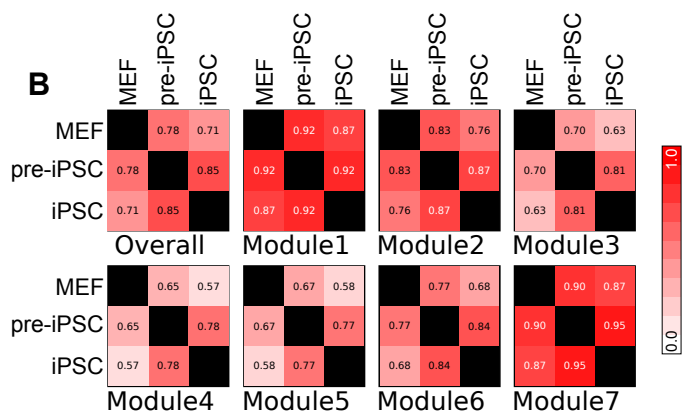


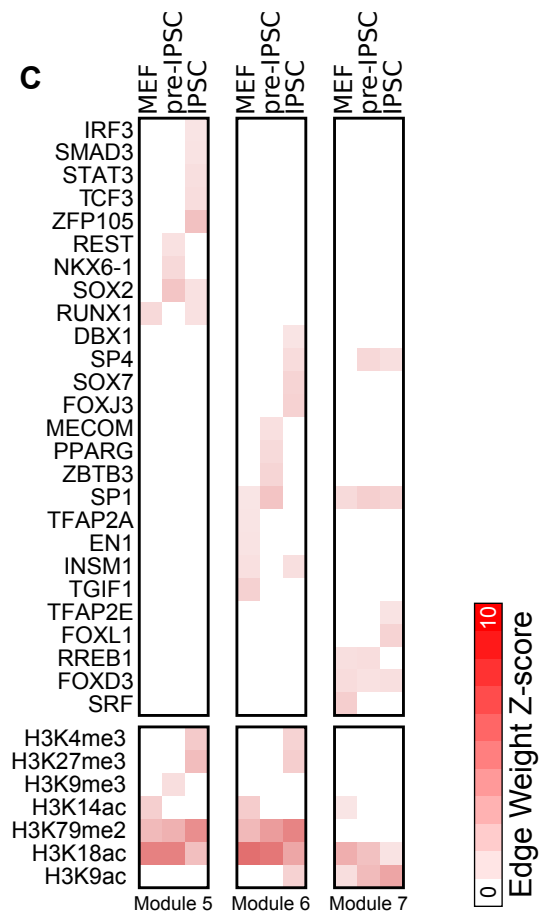
A



B



C



Supplemental Figure S20. Application of DRMNs to the cellular reprogramming array dataset using histone marks and motifs for the feature set. **A.** Shown are gene expression patterns of the $k = 7$ modules. The number above the heatmap is the number of genes in that module. **B.** Similarity of modules across cellular stages as measured by Jaccard index. The color intensity is proportional to the match. **C.** Inferred regulators for the three most highly expressed modules across cell stages.