



Supplemental Figure S21. The effect of hyper-parameters on DRMN performance for Q-Motif, Motif, and Histone+Motif features using the reprogramming sequencing dataset. Each column corresponds to type of feature sets and rows correspond to module sizes. In each panel, different colors corresponds to different values of ρ_3 , the group LASSO penalty (enforcing selection of same features for all cell lines).