



Supplemental Figure S22. The effect of hyper-parameters on DRMN performance for Histone, Accessibility, and Histone+Q-Motif features using the reprogramming sequencing dataset. Each column corresponds to a type of feature set 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).