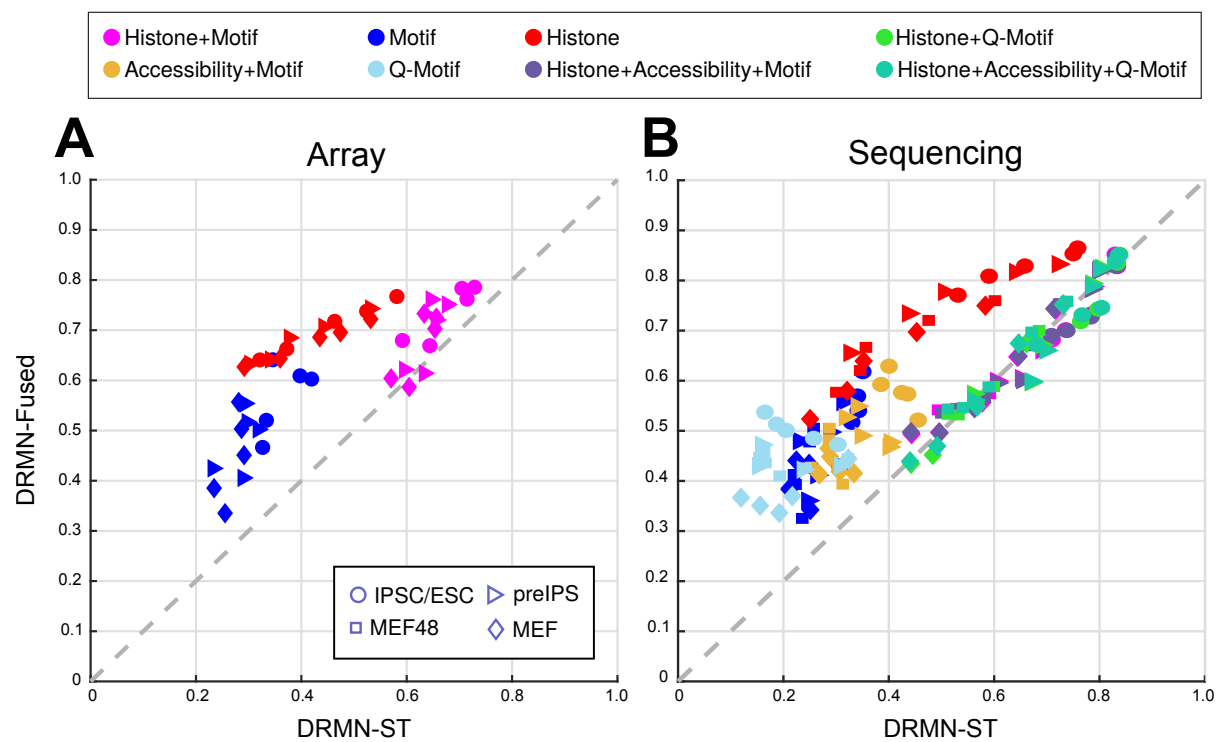




Supplemental Figure S3. Average per-module correlation for individual cell lines for DRMN-ST vs. DRMN-FUSED for (A) array dataset, and (B) sequencing dataset. Each shape correspond to a cell line and each color correspond to a different feature set. Each shape/color combination is present 5 times, corresponding to different number modules: 3, 5, 7, 9, 11. Note, for expression prediction the generative models need the information about the observed expression.