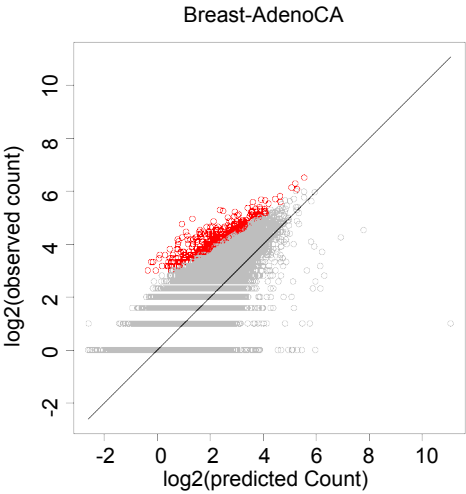
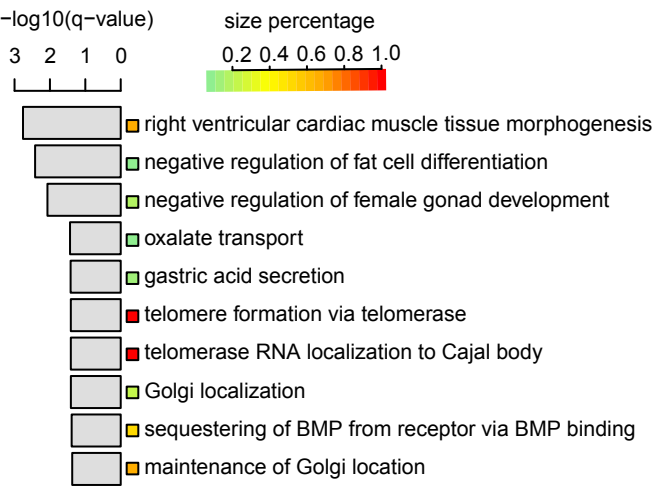


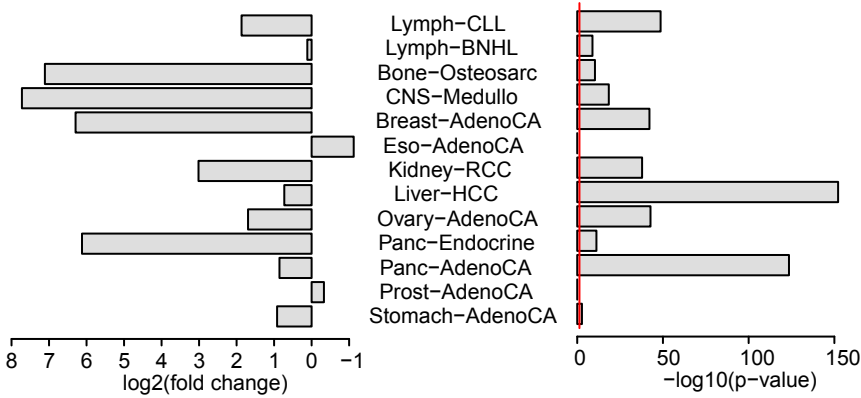
a The cancer-specific regions



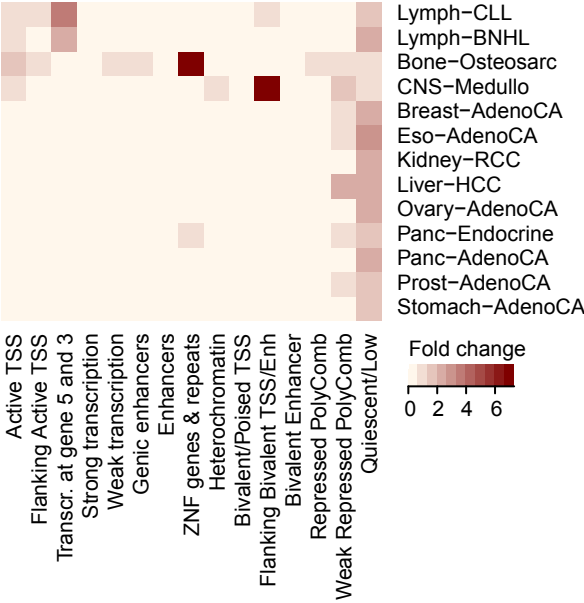
b The enriched GO terms for Breast-AdenoCA specific regions



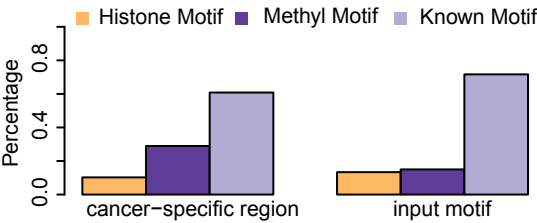
c The motif disruption rates in the 13 tumors



d The fold change of chromHMM for cancer-specific regions



e The percentages of motif types in cancer-specific regions



f Classification model performance

