



Supplementary Figure 6. Genetic conservation at primary prostate tumor chromHMM states. Boxplots showing the average phastCons conservation scores for regions defining chromHMM state in primary human tumors. Higher scores and shading indicate higher conservation scores. Kruskal-Wallis test is shown for trend across the entire dataset. Box plots indicate the median and range of scores for each state. The two most highly conserved states are prostate- lineage specific enhancers and promoters.