Explore Prostate Cancer Datasets Data Input Gene Profile Gene Correlation Copy Number Construcing a heatmap from the gene list you uploaded in the Analysis Parameters tab. If you haven't uploaded a gene list, an example gene list of three genes will be used Method to calulate distances Euclidean Color Key Correlation and Histogram Method of hierachical clustering ward single complete -2 Row Z-Score average mcquitty MELK median AURKA centroid Re-order Rows? AURKB Yes No ETV1 CDC20 Scaling? row UBE<sub>2</sub>C column none IDH1 Cut the dendrogram into k groups, or h(eight) TMPRSS2 h FOXA<sub>1</sub> SPOP Select k groups from the dendrogram