

Supplementary Figure 3: High *CD33* expression predicts poor clinical outcomes in prostate cancer patients. A. Human prostate cancer *CD33* and *ELANE* mRNA expression data were obtained from the TCGA Provisional prostate cancer dataset (downloaded from http://cancergenome.nih.gov/) and plotted to examine correlation using two-tailed Pearson correlation analysis. B. *CD33* expression was assessed based on primary Gleason score using ordinary one-way ANOVA with Tukey's post-hoc testing (n = 491, ** p < 0.01, *** p < 0.001).