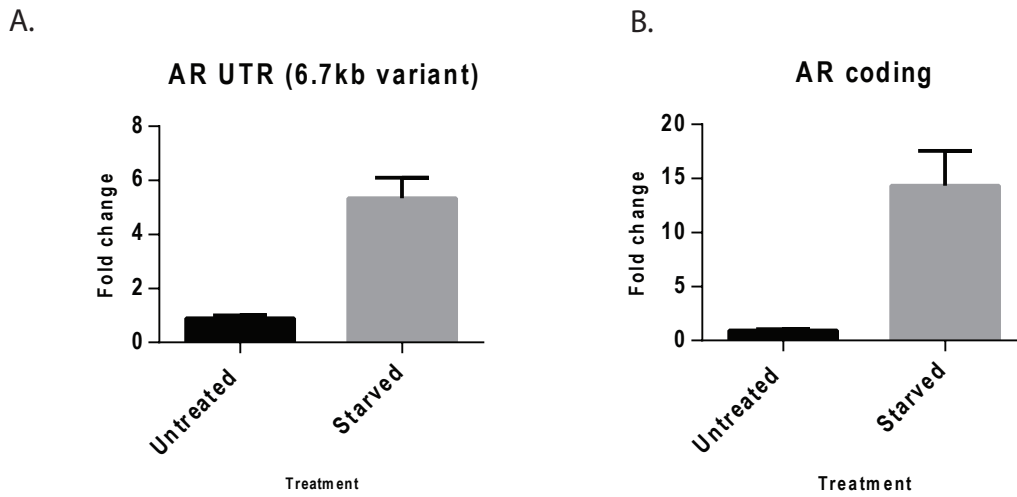


Supplemental figure 1.



Q-PCR analysis of A, AR UTR and B, AR coding region in LNCaP cells grown in starvation medium for 72 hours. Data represents the mean and standard error from three independent replicates. Data is normalised to housekeeping genes GAPDH, RPL19 and b-actin.