



Figure S2.
Mutations identified in CRPC, Related to Figures 4 and 5.

Schematics of protein changes in PIK3CA (A), AKT1 (B), FOXA1 (C), and SPOP (D). The red triangles represent Indels, including insertions and deletions. The green circles represent missense mutations. ABD=adaptor-binding domain that binds to p85, RBD=Ras-binding domain, C2=putative membrane-binding domain, PH=Pleckstrin homology domain, S_TKc=serine/threonine kinase domain, TAD=transactivation domain, FH=Forkhead Winged Helix DNA binding domain, MATH=meprin and TRAF homology domain, BTB=BTB-POZ domain.