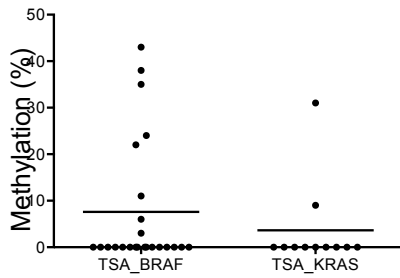
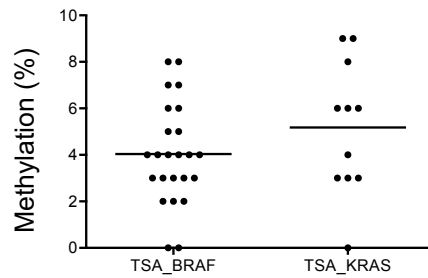


S4 Figure. Levels of methylation of the indicated genes and LINE-1 in TSAs with *BRAF* or *KRAS* mutation.

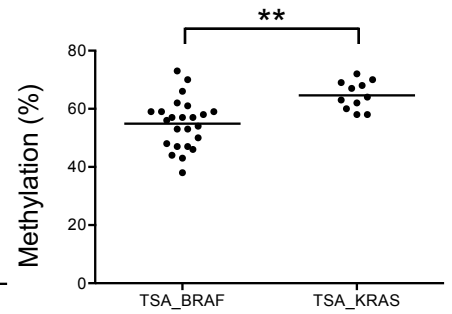
(A) *CDKN2A*



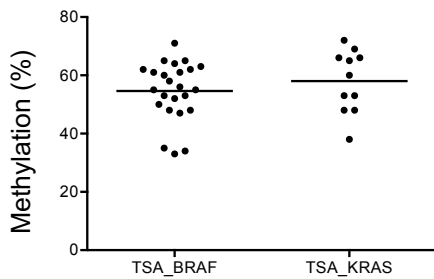
(B) *MLH1*



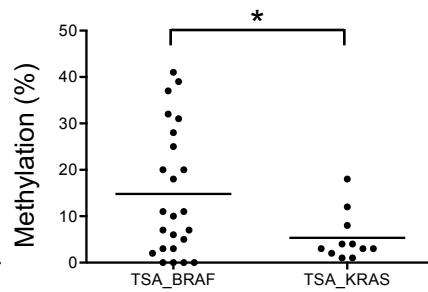
(C) *SFRP1*



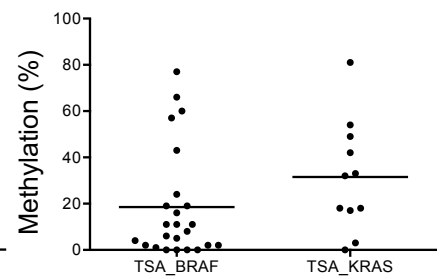
(D) *SFRP2*



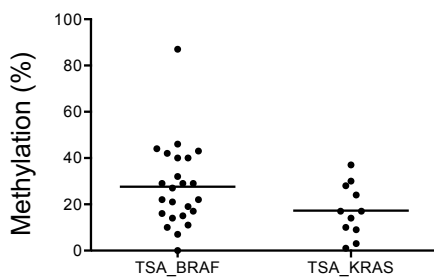
(E) *IGFBP7*



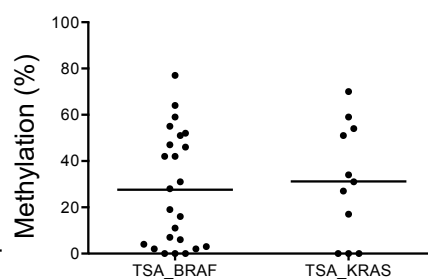
(F) *SMOC1*



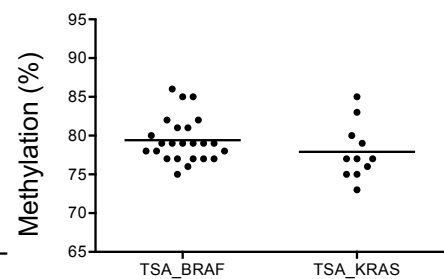
(G) *GALNT14*



(H) *SOX5*



(I) LINE-1



* $P < 0.05$, ** $P < 0.01$.