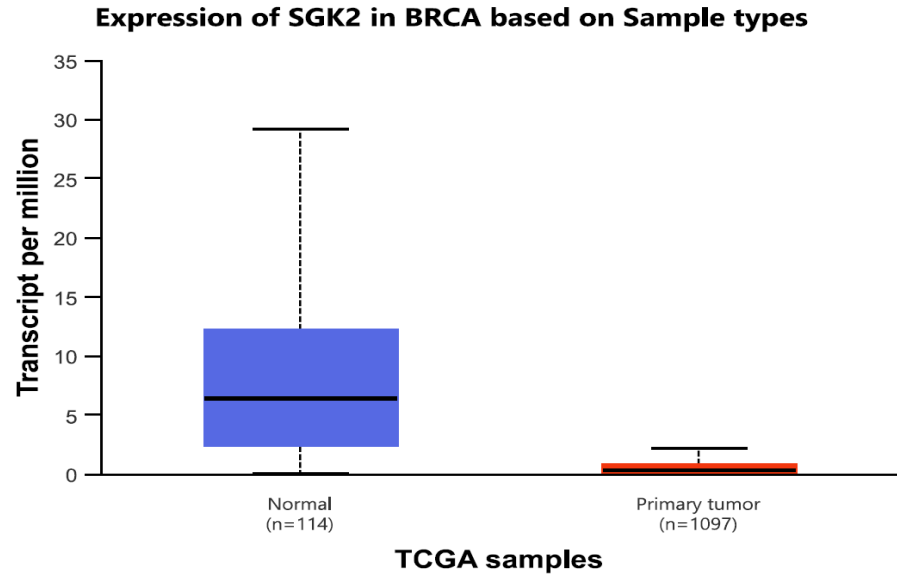
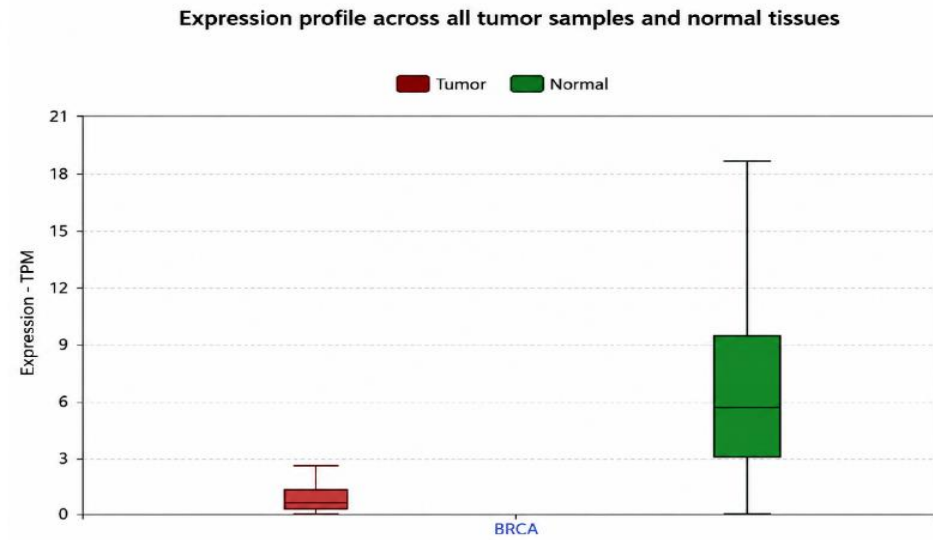
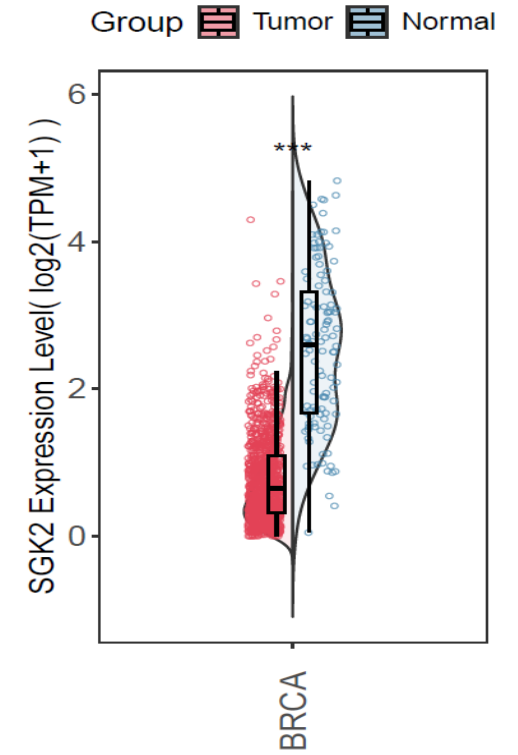


a.**b.****c.**

Supplementary Figure S1. TCGA validation of SGK2 expression. Evaluation of SGK2 expression profile using TCGA-based databases **(a)** UALCAN, **(b)** GEPIA and **(c)** TIMER3.0 revealed significant downregulation of SGK2 in breast cancer tissues compare with normal tissues.