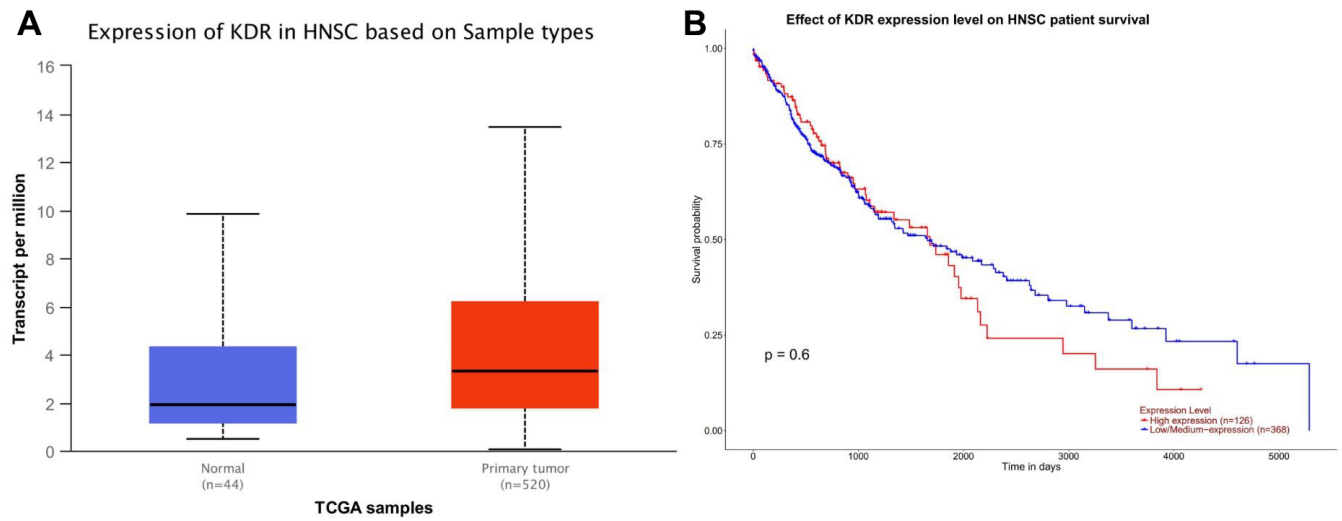


SUPPLEMENTARY FIGURE



Supplementary Figure 1. (A) Compared with the normal group (blue), the expression of VEGFR2 was high in tumor samples (red); (B) Survival curves of VEGFR2 expression for prognosis in the TCGA-HNSC dataset. Red represents the high expression group and blue represents the low expression group.