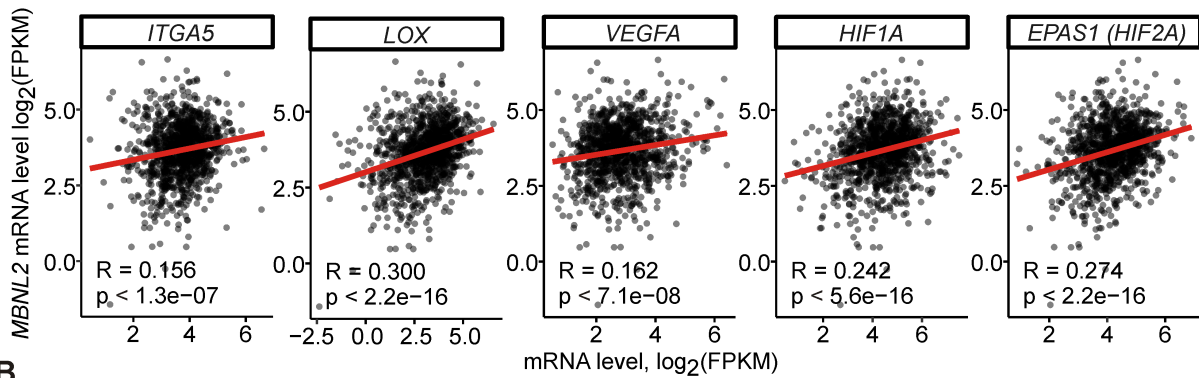
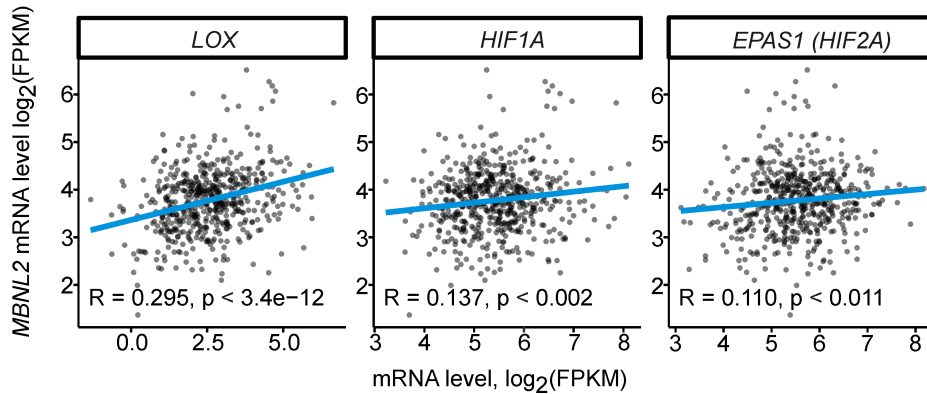


A**B**

Supplementary Figure 12. *MBNL2* expression levels correlate with hypoxia response genes in breast and lung cancer patients. (A) *MBNL2* mRNA levels positively correlate with *ITGA5*, *LOX*, *VEGFA*, *HIF1A* and *EPAS1 (HIF2A)* in breast cancer patients. Scatter plot of fragments per kilobase transcript per million mapped reads (FPKM) for 1,088 patients with breast invasive carcinoma (BRCA) from The Cancer Genome Atlas (TCGA) PanCancer Atlas. Pearson correlation coefficient and associated *P* value are given. Colored lines correspond to linear regression. (B) In lung cancer patients, *MBNL2* mRNA levels positively correlate with *LOX*, *HIF1A* and *EPAS1 (HIF2A)*. Visualization as in (A) for 533 patients with lung adenocarcinoma (LUAD).