

Supplementary Figure 1. Correlation analysis of HMGA2 expression with EGFR signaling components and EMT-related markers in TCGA OSCC samples.

Pearson correlation analyses were performed to assess the relationships between HMGA2 and EGFR, MAPK1, AKT1, STAT3, and representative epithelial markers (CDH1 and EPCAM) using transcriptomic data from The Cancer Genome Atlas (TCGA). Gene expression levels are presented as $\log_2(\text{TPM} + 1)$.

