

Figure S1. Appropriate cut-off value based on risk score in the training set.

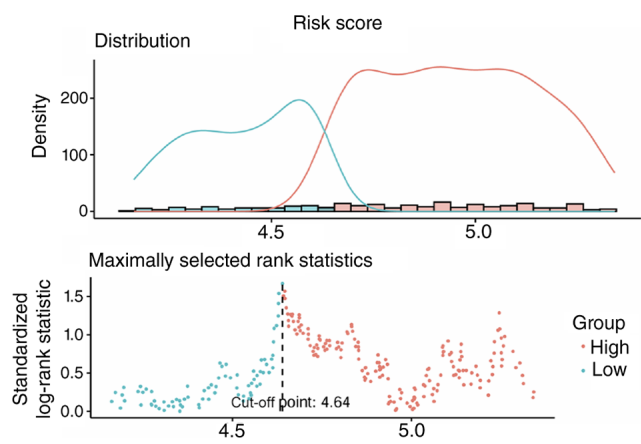


Figure S2. hnRNP C and METTL3 mRNA expression levels determined by qPCR. (A) hnRNP C mRNA expression in indicated OSCC cell lines. (B) METTL3 mRNA levels in OSCC cells after METTL3 knockdown. Data are mean $\pm$ SD from three independent experiments. *** $P < 0.001$. hnRNP C, heterogeneous nuclear ribonucleoprotein C; SCC, squamous cell carcinoma; OSCC, oral SCC; METTL3, methyltransferase-like 3; sh, short hairpin RNA; NC, negative control.

