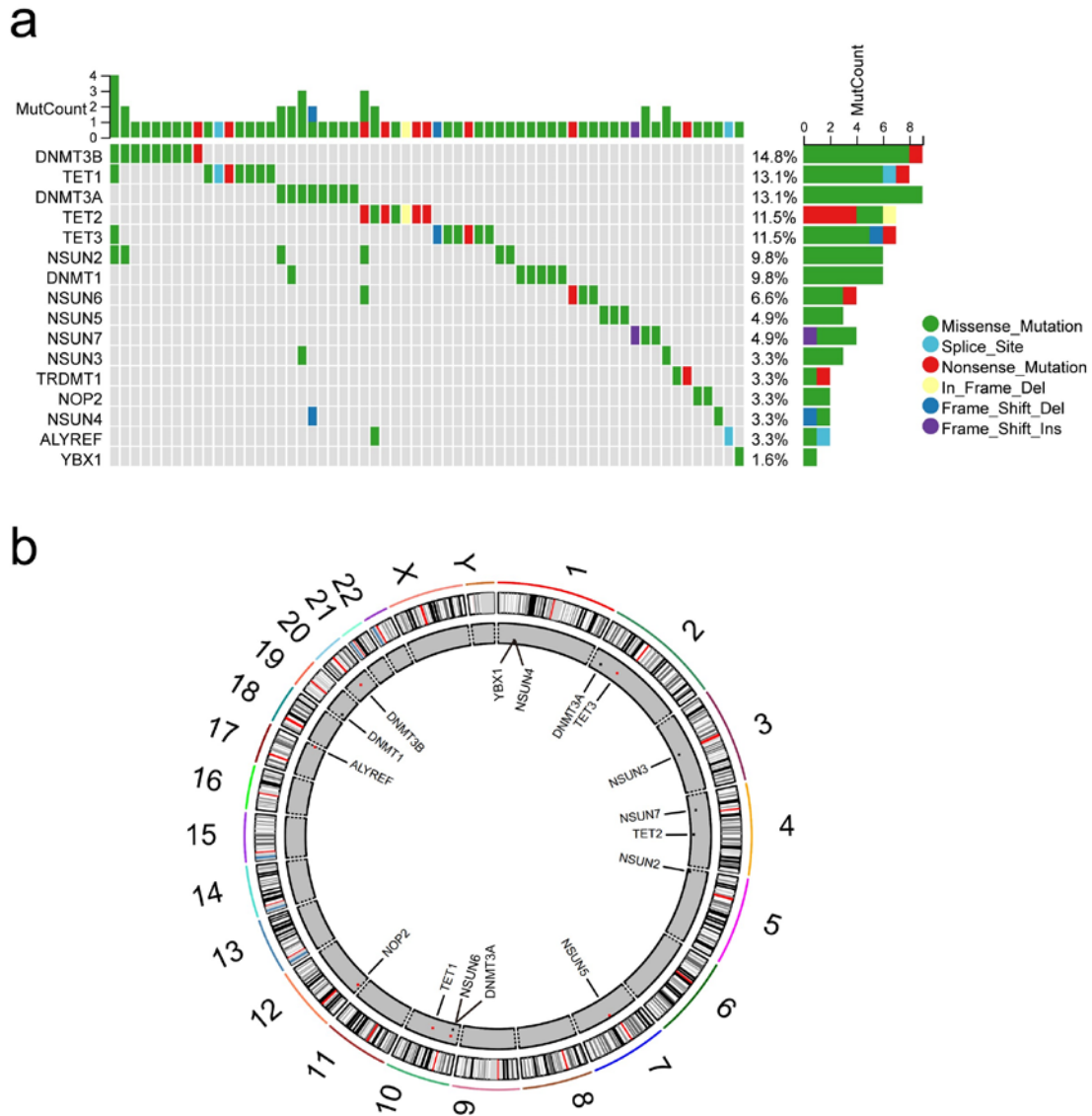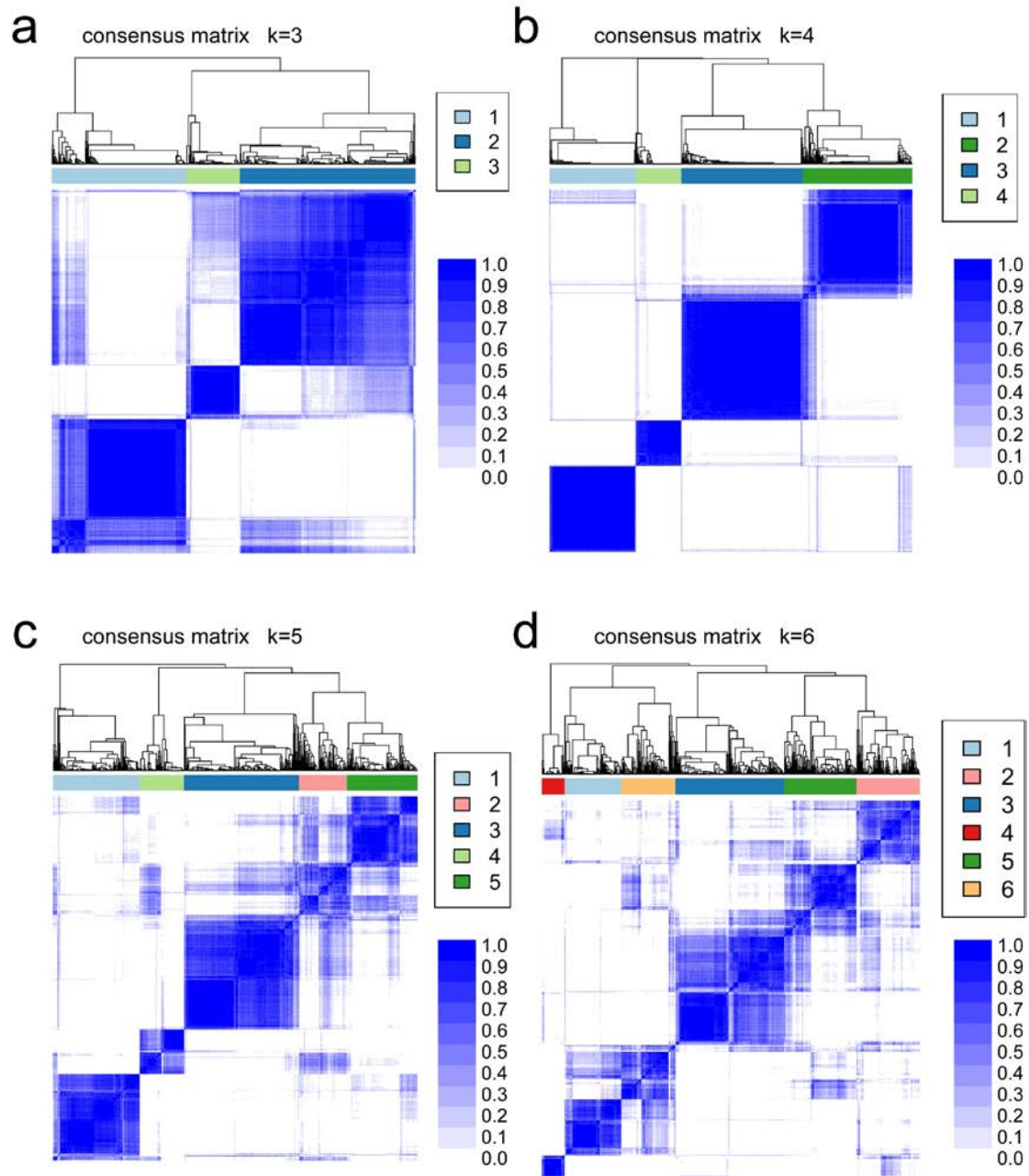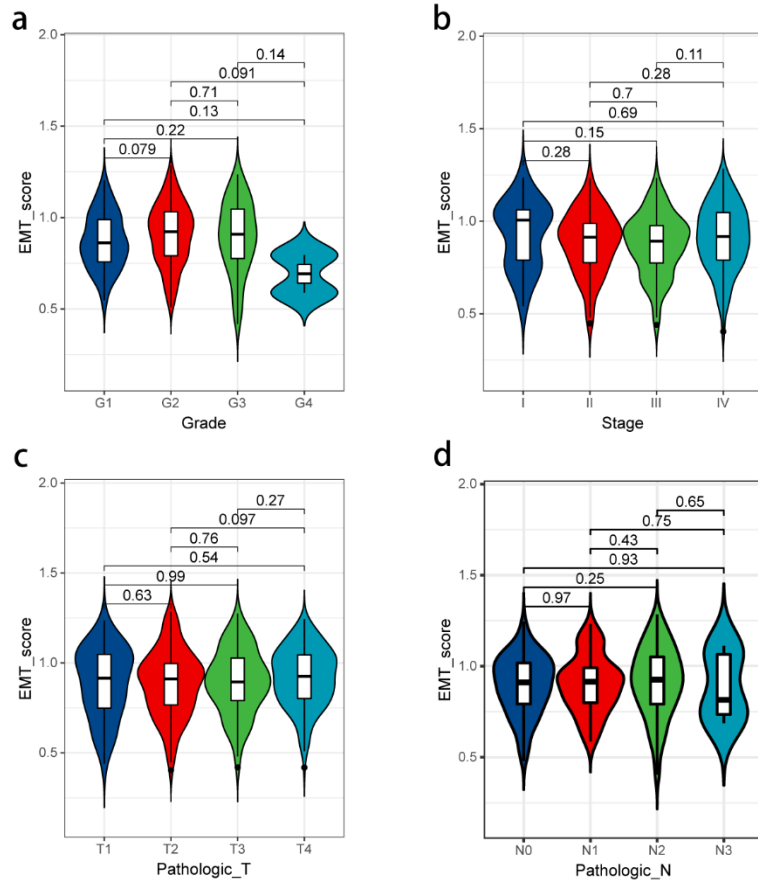# Supplementary Information

Identification and Validation of Novel Prognostic Signatures Based on m5C Methylation Patterns and Tumor EMT Profiles in Head and Neck Squamous Cell Carcinoma
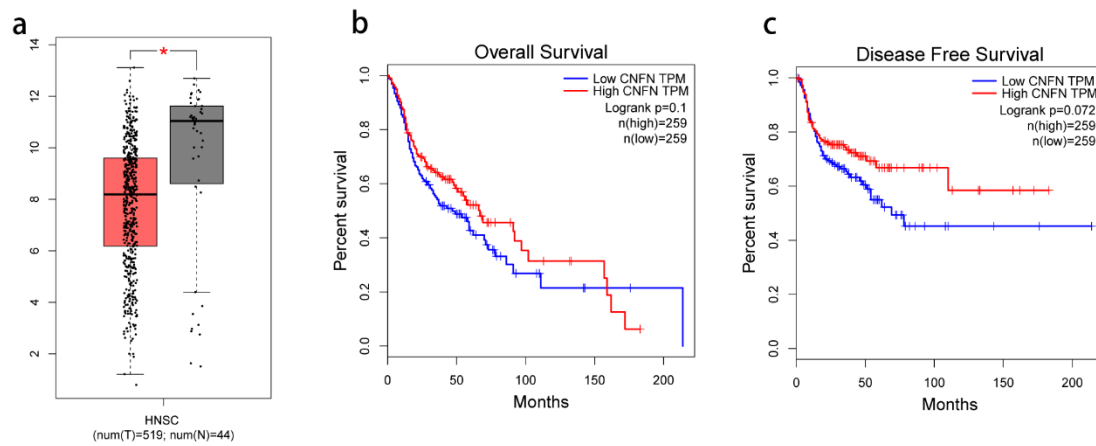

Supplementary Figure 1. Mutation mapping of m5C regulatory genes in HNSCC cohort. **a**: Mutational status of the m5C regulatory genes; **b**: CNV profile of the m5C regulatory genes on chromosomes
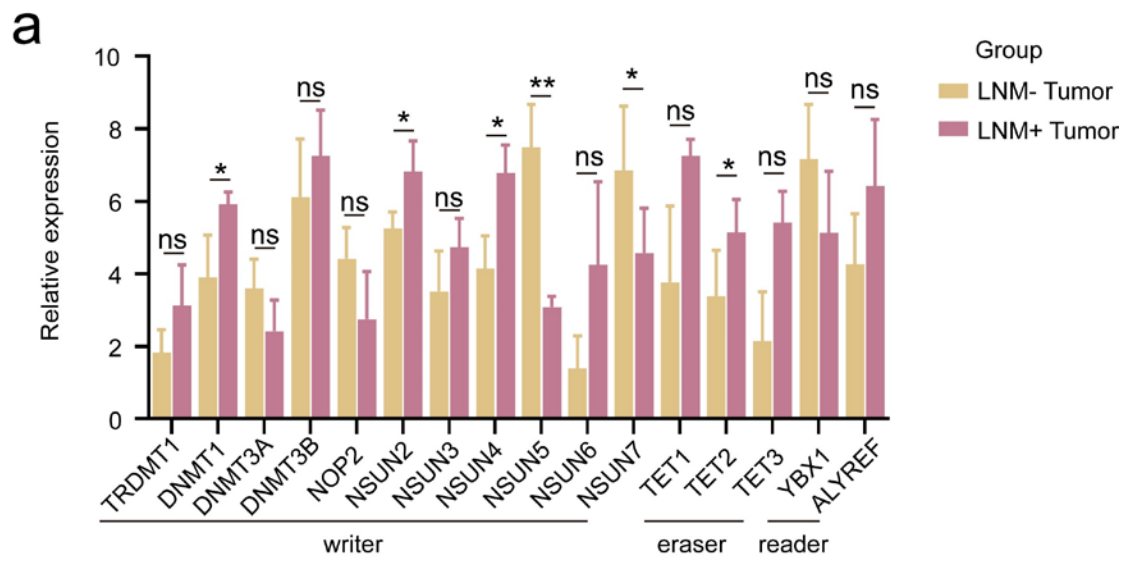
Supplementary Figure 2. Consensus clustering matrix for k = 3, 4, 5, 6.

Supplementary Figure 3. EMT score of different grades, stages, T stages, and N stages.

Supplementary Figure 4. expression levels of CNFN in HNSCC. a. expression levels of CNFN in normal tissues compared with HNSCC tissues; b and c. OS and DFS of different CNFN expression levels.
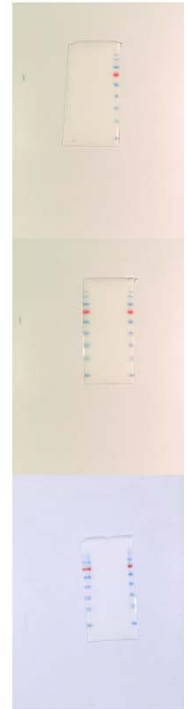
Supplementary Figure 5. Expression levels of 16 m5C regulators in clinical samples about LNM+ vs LNM-.

Uncropped western blots

ZO-1

E-Cadherin

N-Cadherin

SLUG

Vimentin

GAPDH