

Integrative analysis of prognostic long non-coding RNAs with copy number variation in bladder cancer

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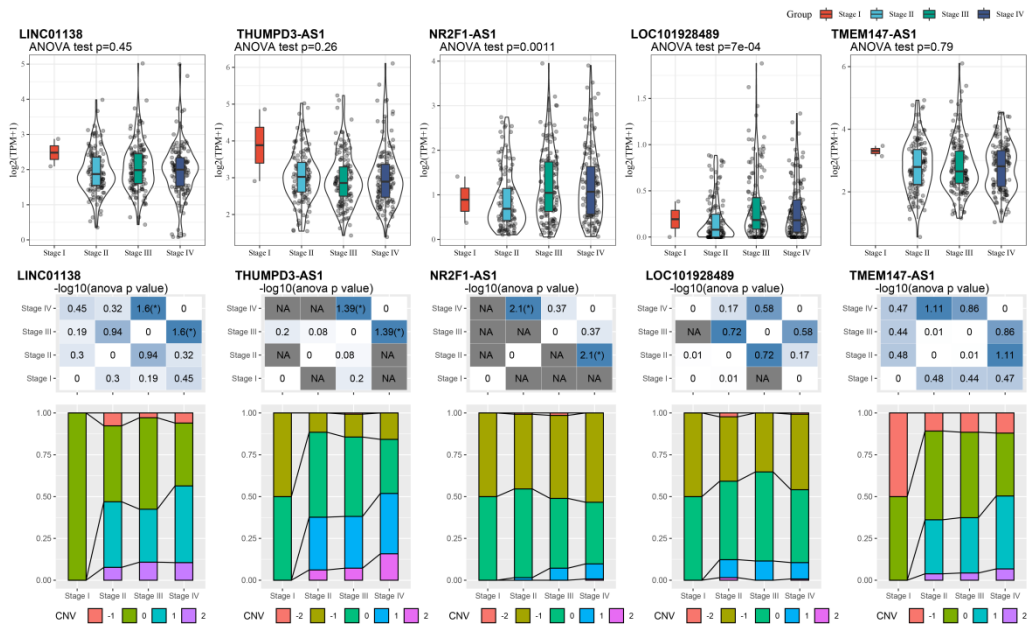


Fig. S1 Expression and copy number variation (CNV) of the five long non-coding RNAs (lncRNAs) in different stages of BLCA.

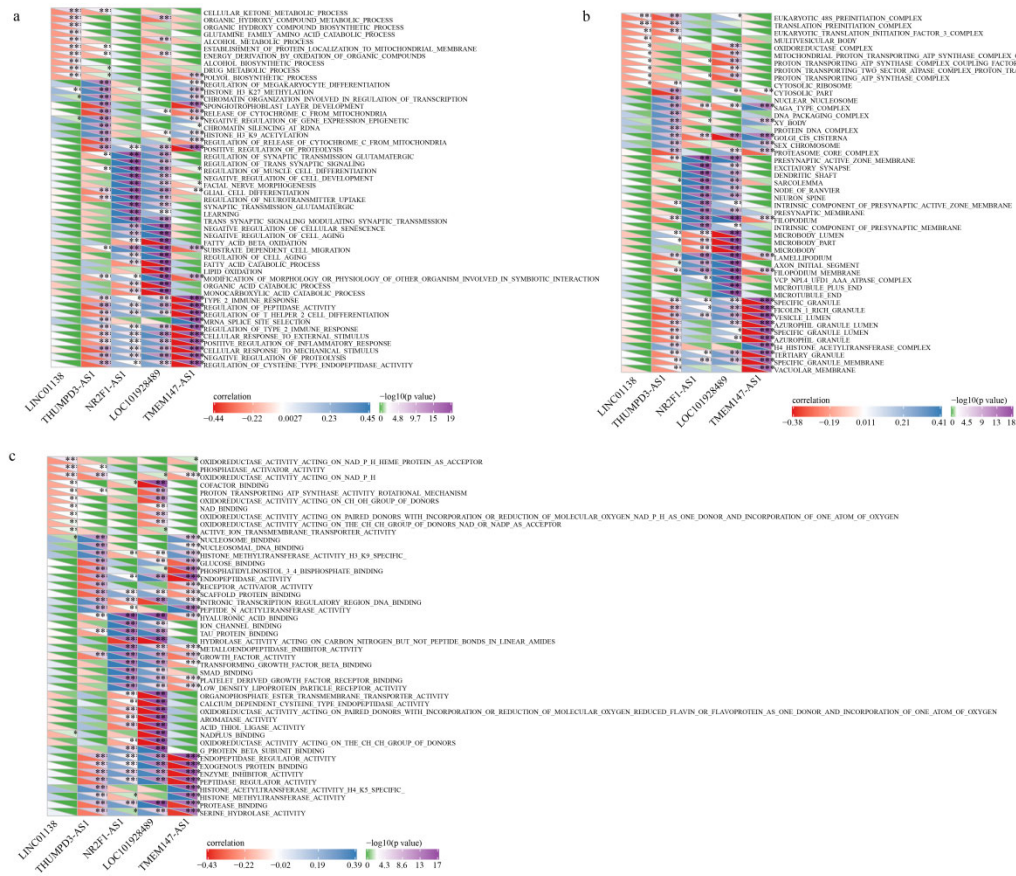


Fig. S2 Gene ontology (GO) enrichment analysis of the five lncRNAs. (a) Biological process (BP) of the five lncRNAs; (b) Cellular component (CC) of the five lncRNAs; (c) Molecular function (MF) of the five lncRNAs. * $p < 0.05$, ** $p < 0.01$, * $p < 0.001$.**