

Supplementary Table S1. GO terms and KEGG pathway enrichment of BZW2 coexpressed genes.

Supplementary Table S2. KEGG pathway enrichment of eIF5 coexpressed genes.

Supplementary Figure S1. Expression of BZW2-related regulators and association with survival outcome. **(A)** Top 5 kinase regulators of BZW2 coexpressed genes. All of these kinase genes were significantly highly expressed in tumor tissues. All these genes have a significant association with OS in HCC. **(B)** Transcription factors of BZW2 coexpressed genes.

Supplementary Figure S2. The expression of BZW2 is highly correlated with that of eIF5, eIF5A, eIF5B, eIF3a, eIF3b, and eIF3e.



