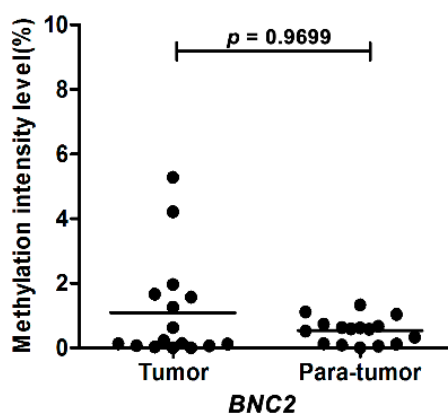
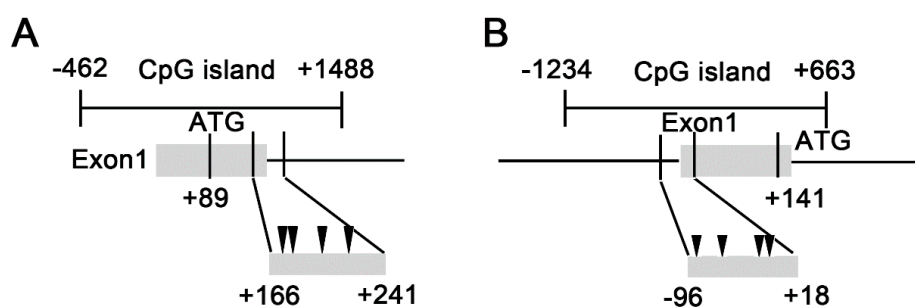


## Supplementary Materials: Decreased Expression of *BNC1* and *BNC2* Is Associated with the Genetic or Epigenetic Regulation in Hepatocellular Carcinoma

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**Figure S1.** The methylation intensity of *BNC2* in 16 paired primary HCC tissues and their corresponding adjacent non-tumor liver tissues.



**Figure S2.** Schematic representation of the detected CGI loci of the (A) *BNC1* and (B) *BNC2* genes. The transcriptional start site for gene is defined as +1. Shaded boxes are exons of *BNC1* and *BNC2*. “▼” represents the restriction enzyme cutting site.