

Supplementary Figure S1



Figure S1. Cloning of the full-length of porcine *YAPI* CDS. (A) The porcine *YAPI* CDS was obtained by PCR amplification and verified by Sanger sequencing. (B) Confirmed porcine *YAP* CDS sequences exhibit 96% similarity to predicted sequenced in NCBI database (XM_021062706).