

Figure S1. Analysis of RNA-seq of tree shrew pancreatic cancer. (A) sample correlation of three tumor samples and normal tissue. (B) differentially expressed genes between tumor samples and normal tissue.



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Table S2. Homology between human, tree shrew, and mouse pancreatic cancer pathway genes and their modification sites.

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Table S3. The relative expression of human-Tree shrew-mouse 1:1:1 ortholog genes.

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