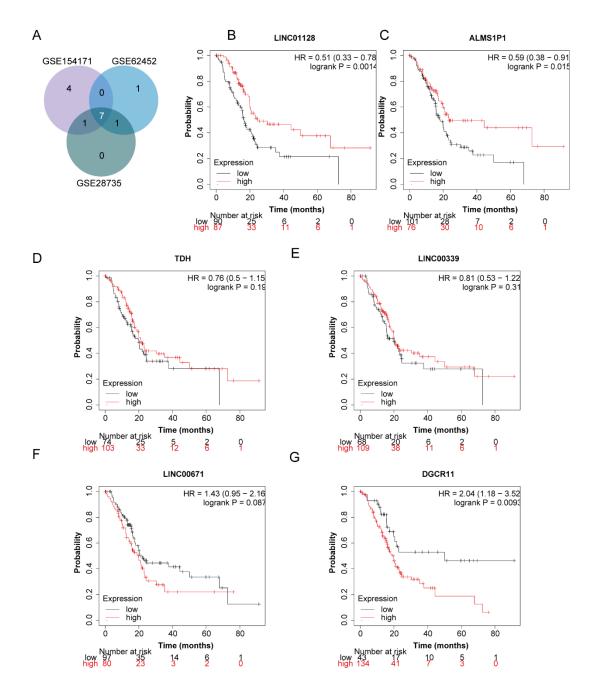


**Supplementary Figure S1. Bioinformatics analysis of pancreatic cancer chips from the Gene Expression Omnibus (GEO)** (A,B) GSE15471. (C,D) GSE62452. (E,F) GSE28735.



Supplementary Figure S2. The association between the expression of lncRNA and pancreatic carcinoma patients' survival rate in Kmplot

No.	Age	Gender	Tumor location	TNM	Stage
1	54	F	Head	T2N0M0	IB
2	68	F	Head	T1N0M0	IA
3	70	F	Head	T3N0M0	IIA
4	53	М	Head	T1N0M0	IA
5	64	М	Head	T2N1M0	IIB
6	50	М	Head	T1N1M0	IIB
7	66	F	Head	T1N0M0	IA
8	57	F	Head	T1N1M0	IIB
9	38	F	Head	T1N0M0	IA
10	67	М	Head	T2N0M0	IB
11	70	F	Head	T2N0M0	IB
12	65	Μ	Head	T1N1M0	IIB
13	75	F	Head	T1N0M0	IA
14	56	М	Head	T1N0M0	IA
15	53	Μ	Head	T2N0M0	IB

Supplementary Table S1. The clinical character of patients