

**Supplementary Material to “A comparison of mRNA and circRNA expression between squamous cell carcinoma and adenocarcinoma of the lungs.”**

**Table S4** - The differentially expressed genes related to prognosis

LUSC		LUAD	
Gene	Pvalue	Gene	Pvalue
<b>STXBP1</b>	0.00428695	IRX5	1.55E-05
EPHA2	0.00813035	MTURN	0.0002359
ADD3	0.01073184	OGT	0.0004993
SLC9A9	0.01099923	<b>PMEPA1</b>	0.0015137
EVA1A	0.01130279	IGF2BP3	0.0020889
SHANK2	0.0119727	DCUN1D1	0.0052575
FAM46A	0.02016478	DIP2B	0.0077857
<b>PMEPA1</b>	0.03070171	<b>STXBP1</b>	0.0129428
NAV2	0.04797483	TMEM243	0.0143325
		IGF2BP2	0.0204467
		WFDC2	0.0207173
		CAPN3	0.0269737
		PARM1	0.0317552
		PDCD10	0.0345579
		PARL	0.0355587
		ARRDC3	0.0365931
		CDK6	0.0396845
		DOK4	0.0408639

LUSC	LUAD
	CORO1C 0.0437517
	WSB1 0.0444235
	ERRFI1 0.0461206
	PTPRG 0.048331

The bold part is the prognostic genes associated with both LUSC and LUAD.