

Table S1 The risk coefficients of 14 lncRNAs

Gene	Coef
<i>AP006621.2</i>	0.080625191
<i>AL590369.1</i>	0.943370712
<i>ITGB1-DT</i>	0.674312339
<i>AC013652.1</i>	0.132492235
<i>AC016394.3</i>	0.099422829
<i>AC092944.1</i>	1.558696937
<i>AC099850.4</i>	-0.036462077
<i>AL513550.1</i>	0.296482356
<i>AC073896.3</i>	-0.655491862
<i>TNFRSF10A-AS1</i>	-0.079711559
<i>AC007128.1</i>	0.03146985
<i>AC245041.1</i>	0.074254937
<i>LRP4-AS1</i>	0.150206033
<i>AL137782.1</i>	-0.335651208

Table S2 The DEGs in the high- and low-risk groups

Gene	Low mean	High mean	logFC	P value	FDR
<i>REG1A</i>	520.62766	208.711771	-1.31873998976008	0.00868142	0.02818453
<i>RNA5SP383</i>	0.81742774	2.5766383	1.65632693	0.00315067	0.01301231
<i>EEF1A2</i>	0.84311722	2.50878565	1.57318408	0.00407908	0.01590618
<i>CDKN2A</i>	1.56511682	3.23301847	1.04661141	0.00398524	0.01562694
<i>REG1B</i>	72.7147996	25.3915608	-1.5178999451226	0.01007801	0.03173523
<i>DKK1</i>	0.84165342	1.73726478	1.04551948	0.00030671	0.00214479
<i>EPHB6</i>	1.04424242	2.34710695	1.16842691	1.14E-05	0.00016869
<i>CST6</i>	0.74731565	2.00010459	1.42028581	0.00314691	0.01300129
<i>KRT5</i>	1.22918855	4.93199962	2.00446645	0.00772583	0.02586971
<i>VGF</i>	1.40602569	3.89894732	1.47146171	0.00174431	0.00829052
<i>C7</i>	1.32699805	2.66760723	1.00738002	0.00703338	0.02398819
<i>PLIN4</i>	0.66800041	3.11847949	2.22292187	0.00519277	0.01910147
<i>COMP</i>	4.7195529	11.2456984	1.25265116	0.00085103	0.00480146
<i>CXCL8</i>	80.0270785	36.4405966	-1.13494166171531	3.65021419605745E-06	7.03E-05
<i>CCL19</i>	3.17685667	7.02636538	1.1451786	0.01596173	0.04482656
<i>S100A2</i>	4.7648246	9.7841678	1.03802604	0.00040813	0.00270214
<i>KRT14</i>	0.08776144	4.14600262	5.56199001	0.00059089	0.00360451
<i>H3P16</i>	1.91661241	0.93487364	-1.03571533109378	0.00395678	0.01553065