

Table S1 Correlation between FoxM1 expression and clinicopathologic characteristics

	Cases	FoxM1 protein expression(n)			P-value
		Negative	Weak	Strong	
Age					
<60	90	26	37	27	0.663
≥60	65	15	27	23	
Gender					
male	92	19	42	31	0.131
female	63	22	22	19	
Tumor Size					
<4(cm)	105	32	44	29	0.123
≥4(4cm)	50	9	20	21	
Stage					
I	51	29	18	4	0.001
II	41	8	25	8	
III	63	4	21	38	
T stage					
T1	44	19	21	4	0.001
T2	95	22	37	36	
T3	15	0	6	9	
T4	1	0	0	1	
N stage					
N0	65	30	28	7	0.001
N1	29	7	15	7	
N2	58	4	19	35	
N3	3	0	2	1	
Metastasis					
M0	71	34	30	7	0.001
M1	84	7	34	43	
Vessel invasion					
No	115	38	47	30	0.002
Yes	40	3	17	20	
Differentiation					
Well	88	32	39	17	0.001
Poor	67	9	25	33	

Fig. s1

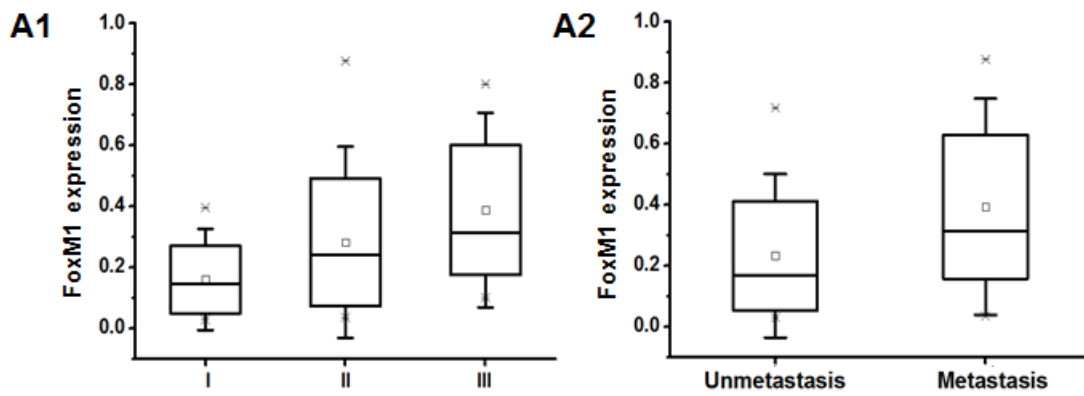


Figure S1. The FOXM1 expression levels were detected in 78 frozen lung adenocarcinoma specimens using Real-time PCR analysis. FOXM1 expression was elevated in stage II and III groups (I vs II, $P < 0.05$, I vs III, $P < 0.05$, A1), as well as in metastasis group ($P < 0.05$, A2)