

**Table S1. Correlation analysis of MTERF1 protein expression with clinicopathological parameters in 52 colorectal cancer patients** (The bold number represents the P-values with significant differences).

Patients characteristics	MTERF1 expression		$\chi^2$	P-value
	Low	High		
<i>Age (years)</i>				
<50	10	10	0.0000, 1	1.0000
≥50	16	16		
<i>Gender</i>				
Female	11	9	0.32501	0.5686
Male	15	17		
<i>Hypertension</i>				
Yes	7	5	0.4331	0.5104
NO	19	21		
<i>Diabetes</i>				
Yes	1	4	1.9911	0.1582
NO	25	22		
<i>Tumor size</i>				
<5cm	17	11	2.7861	0.0951
≥5cm	9	15		
<i>Lymph node status</i>				
0	5	15	<b>8.1251</b>	<b>0.0044</b>
1-3	21	11		
<i>ER expression</i>				
Positive	11	5	3.2501	0.0714
Negative	15	21		
<i>PR expression</i>				
Positive	12	15	0.69331	0.4050
Negative	14	11		
<i>Tumor status</i>				
T1+T2	21	10	<b>9.6651</b>	<b>0.0019</b>
T3+T4	5	16		
<i>Nodal status</i>				
N0	5	15	<b>8.1251</b>	<b>0.0044</b>
Nx	21	11		
<i>Clinical stages</i>				
1	21	11	<b>8.1251</b>	<b>0.0044</b>
2	5	15		