

Hsa-miR-19a is associated with lymph metastasis and mediates the TNF- α induced epithelial-to-mesenchymal transition in colorectal cancer

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Supplementary Materials

Table S1. Clinical and histopathological information of the 11 cases of CRC.

Case no.	Gender (M/F)	Age (years)	Location	Lymph node invasion	distant metastasis	TNM stage
1	F	70	Sigmoid colon	Yes	No	III
2	F	37	Transverse colon	No	No	II
3	F	73	Rectum	No	No	II
4	F	75	Rectum	Yes	No	III
5	M	65	Sigmoid colon	Yes	No	III
6	F	79	Right colon	No	No	II
7	M	59	Sigmoid colon	Yes	No	III
8	M	52	Rectum	Yes	No	III
9	M	85	Right colon	No	No	II
10	M	51	Left colon	Yes	No	III
11	F	85	Sigmoid colon	Yes	No	III

M, male; F, female.

Table S2. Patient demographic and clinical characteristics of 275 CRC patients.

Characteristics	of cases
Age (years)	
<60	142 (51.64%)
≥60	133 (48.36%)
Gender	
Male	152 (55.27%)
Female	123 (44.73%)
Tumor size	
<5cm	139 (50.55%)
≥5cm	136 (49.45%)
Tumor location	
Colon	132 (48%)
Rectum	142 (51.64%)
Missing data	1 (0.36%)
Differentiation	
Well	21 (7.64%)
Moderate	219 (79.64%)
Poor	31 (11.27%)
Undifferentiation	1 (0.36%)
Missing data	2 (0.73%)
Primary tumor (pT)	
T1	6 (2.18%)
T2	52 (18.91%)
T3	198 (72%)
T4	19 (6.91%)
Regional lymph node (pN)	
N0	156 (56.73%)
N1	77 (28%)
N2	35 (12.8%)
Missing data	7 (2.5%)
Distant Metastasis (pM)	
M0	250 (90.91%)
M1	25 (9.09%)
TNM stage	
I	49 (17.82%)
II	99 (36%)
III	102 (37.09%)
IV	25 (9.09%)
Relapse	
Yes	38 (13.82%)
No	233 (84.73%)
Missing data	4 (1.45%)

Table S3. Patient demographic and clinical characteristics of 218 colon cancer patients.

Characteristics	of cases
Age (years)	
<60	65 (29.8%)
≥60	153 (70.2%)
Gender	
Male	124 (56.9%)
Female	94 (43.1%)
TNM stage	
I	31 (14.2%)
II	89 (40.8%)
III	64 (29.4%)
IV	26 (11.9%)
Missing data	8 (3.7%)
Primary tumor (pT)	
TX	1 (0.5%)
T1	5 (2.3%)
T2	31 (14.2%)
T3	156 (71.6%)
T4	25 (11.5%)
Regional lymph node (pN)	
NX	1 (0.5%)
N0	128 (58.7%)
N1	53 (24.3%)
N2	36 (16.5%)
Distant Metastasis (pM)	
MX	36 (16.5%)
M0	152 (69.7%)
M1	26 (11.9%)
Missing data	4 (1.8%)

Table S4. Patient demographic and clinical characteristics of 93 rectal cancer patients.

Characteristics	of cases
Age (years)	
<60	37 (39.8%)
≥60	56 (60.2%)
Gender	
Male	50 (53.8%)
Female	43 (46.2%)
Missing data	78 (83.9%)
TNM stage	
I	13 (14.0%)
II	25 (26.9%)
III	34 (36.6%)
IV	16 (17.2%)
Missing data	5 (5.4%)
Primary tumor (pT)	
T1	5 (5.4%)
T2	14 (15.1%)
T3	63 (67.7%)
T4	10 (10.7%)
Missing data	1 (1.1%)
Regional lymph node (pN)	
NX	2 (2.2%)
N0	40 (43.0%)
N1	29 (31.2%)
N2	21 (22.6%)
Missing data	1 (1.1%)
Distant Metastasis (pM)	
MX	14 (15.1%)
M0	64 (68.8%)
M1	13 (14.0%)
Missing data	2 (2.2%)

Table S5. Correlation between miR-19a expression and clinical parameters in 218 colon cancer patients.

Factors	n	Low miR-19a expression (%)	High miR-19a expression (%)	P - value
Age				0.104
<60	65	47 (72.3)	18 (27.7)	
≥60	153	93 (60.8)	60 (39.2)	
Gender				0.108
Male	124	74 (59.7)	50 (40.3)	
Female	94	66 (70.2)	28 (29.8)	
TNM stage				0.299
I	31	21 (67.7)	10 (32.3)	
II	89	54 (60.7)	35 (39.3)	
III	64	42 (65.6)	22 (34.4)	
IV	26	21 (80.8)	5 (19.2)	
pT				0.433
T1+T2	36	21 (58.3)	15 (41.7)	
T3+T4	181	118 (65.3)	63 (34.8)	
pN				0.302
N0	128	79 (61.7)	49 (38.3)	
N1+N2	89	61 (68.5)	28 (31.5)	
pM				0.062
M0	152	94 (61.8)	58 (38.2)	
M1	26	21 (80.8)	5 (19.2)	

Table S6. Correlation between miR-19a expression and clinical parameters in 93 rectal cancer patients.

Factors	n	Low miR-19a expression (%)	High miR-19a expression (%)	P - value
Age				0.638
<60	37	16 (43.2)	21 (56.8)	
≥60	56	27 (48.2)	29 (51.8)	
Gender				0.086
Male	50	19 (38.0)	31 (62.0)	
Female	43	24 (55.8)	19 (44.2)	
TNM stage				0.104
I	13	7 (53.8)	6 (46.2)	
II	25	15 (60.0)	10 (40.0)	
III	34	10 (29.4)	24 (70.6)	
IV	16	8 (50.0)	8 (50.0)	
pT				0.728
T1+T2	19	8 (42.1)	11 (57.9)	
T3+T4	73	34 (46.6)	39 (53.4)	
pN				0.004
N0	40	25 (62.5)	15 (37.5)	
N1+N2	50	16 (32.0)	34 (68.0)	
pM				0.579
M0	64	30 (46.9)	34 (53.1)	
M1	13	5 (38.5)	8 (61.5)	

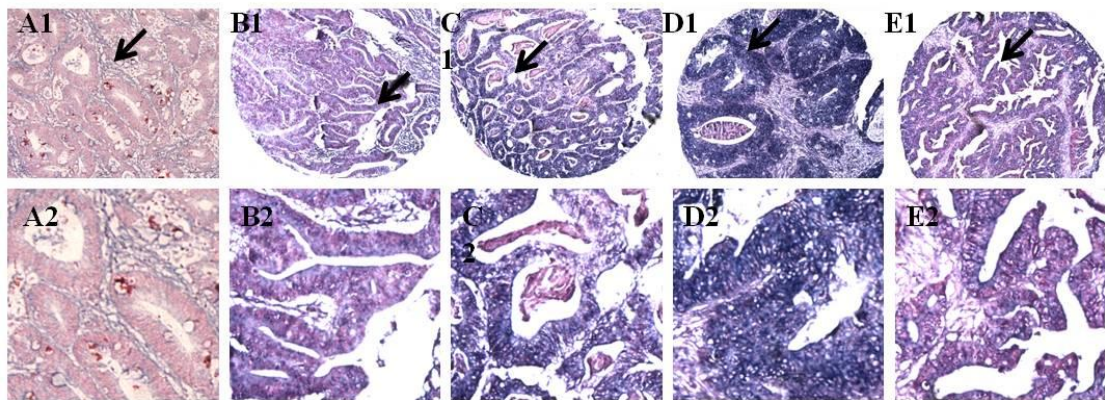


Fig. S1 Expression of miR-19a in CRC using *in situ* hybridisation analysis.

Tissue sections were incubated with DIG-labelled LNA probe to miR-19a. The positive staining of tissue was expressed as blue-violet. No *in situ* hybridization signal was obtained in the absence of DIG-labeled probe (A1, A2). Weak staining of miR-19a was detected in stage I (B1, B2) and IV (E1, E2), stage II moderately positive expression of miR-19a (C1, C2), stage III strongly positive expression of miR-19a (D1, D2).

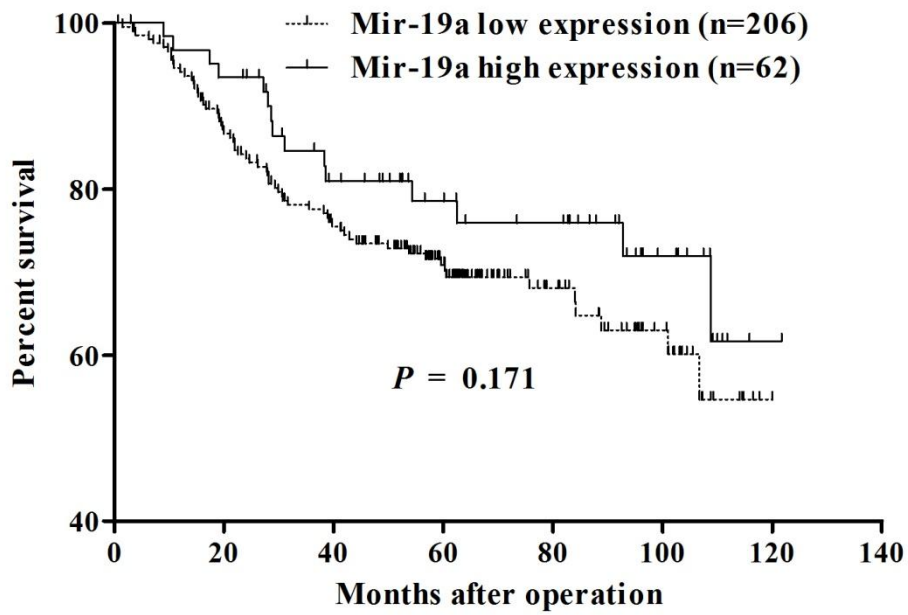


Fig. S2 Association between miR-19a expression in colorectal cancer and overall survival in 268 CRC patients.

In situ hybridization from paraffin blocks of CRC specimens followed by analysis of miR-19a levels. The Kaplan-Meier curve and log-rank test showed that miR-19a expression wasn't associated with survival ($P = 0.171$).

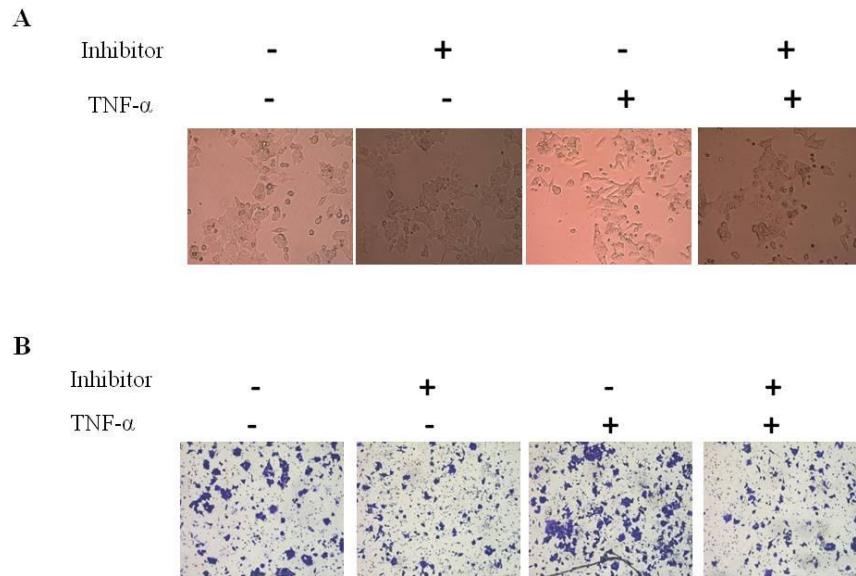


Fig. S3 MiR-19 inhibitor reversed the TNF- α induced EMT transition.

(A) The HCT 116 cells transfected either miR-19a inhibitor or negative control were treated with or without TNF- α . Cell morphology was examined and photographed.

(B) The HCT 116 cells transfected either miR-19a inhibitor or negative control with or without TNF- α were subjected to invasion assays. The HCT 116 cells transfected with miR-19a inhibitor reversed TNF- α effect on cell invasion.

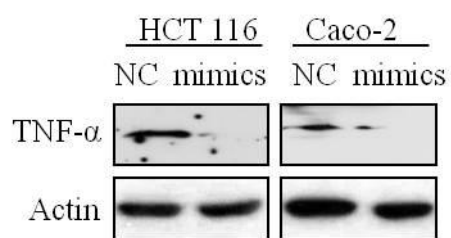


Fig. S4 miR-19a down-regulated the expression of TNF- α .

HCT 116 and Caco-2 cells were transfected with miR-19a mimics or negative control. Western blot analysis showed that miR-19a down-regulated the expression of TNF- α .

Supplementary information

Full-length blots:

Figure S4 A

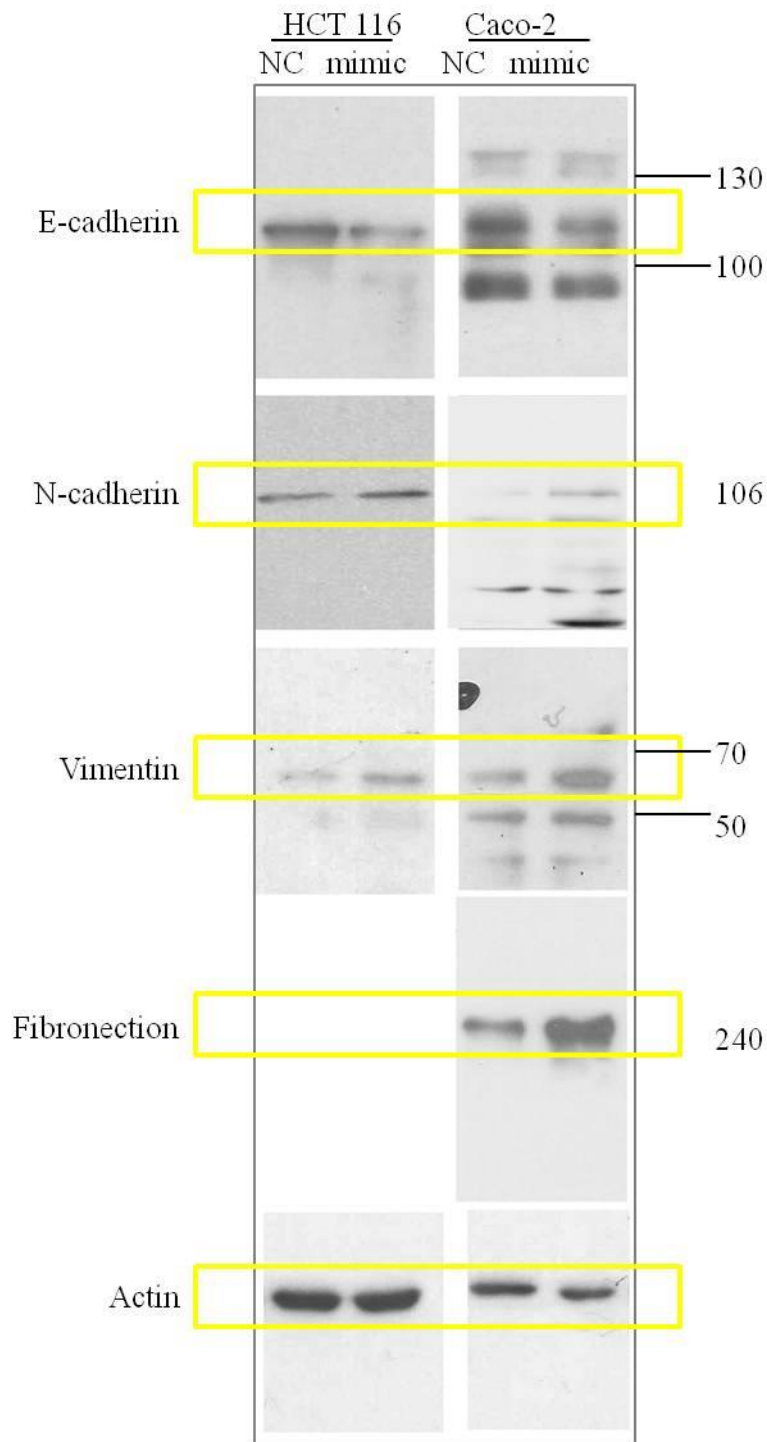


Figure S4 C

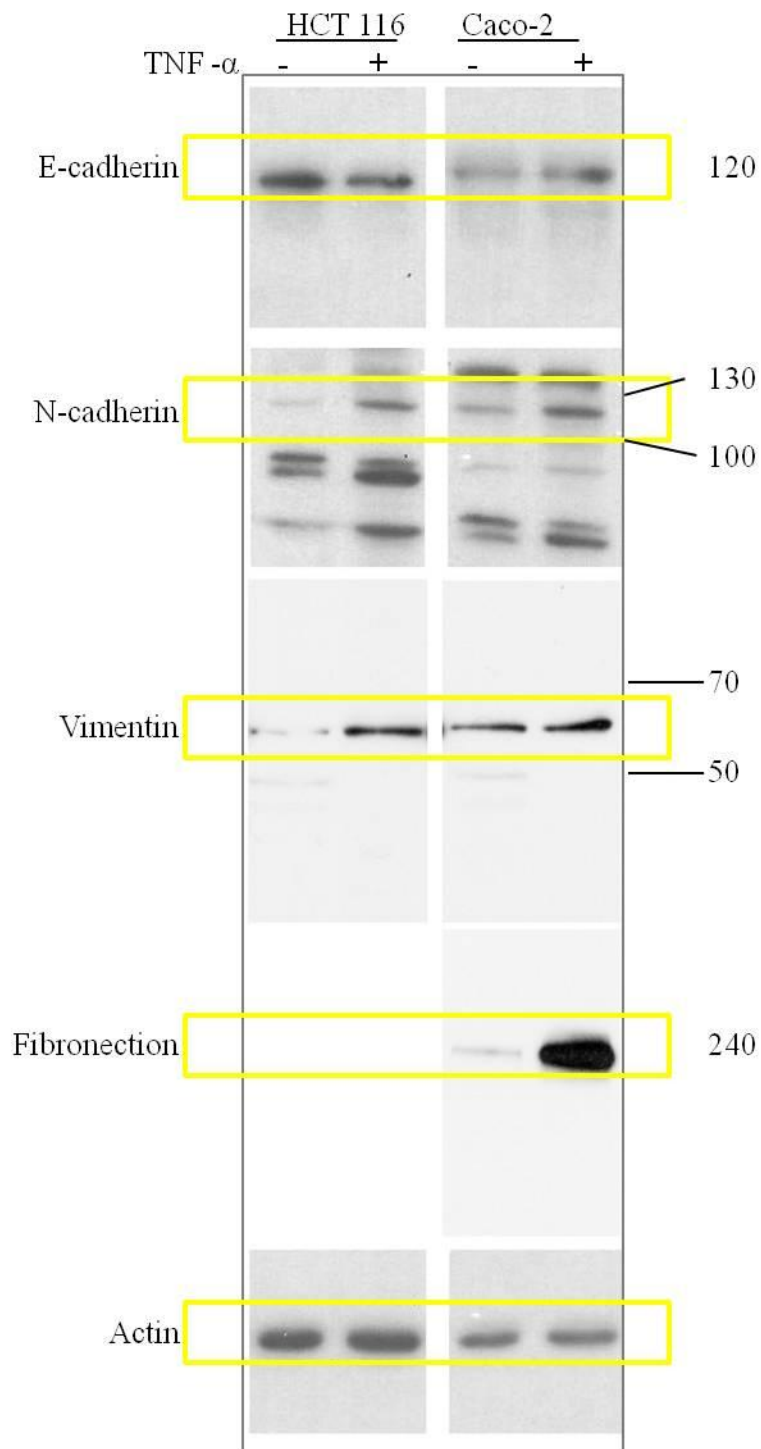


Figure S5

