

Figure S1. Reverse transcription-quantitative PCR results for circ_0004365 in TE11 5-FU, TE11 DTX and TE11 cells.

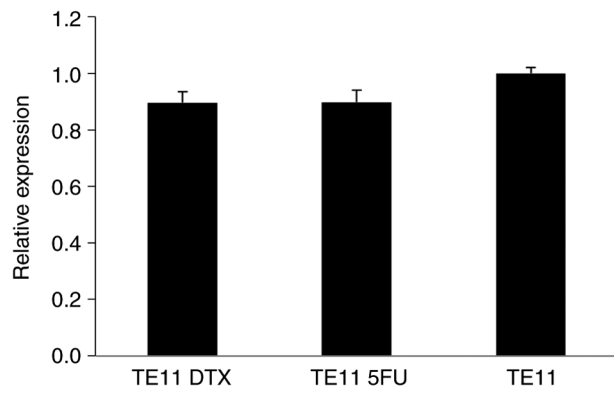


Figure S2. Relative expression of circ_0004365 in ESCC tissues measured via reverse transcription-quantitative PCR. (A) Relative expression of circ_0004365 from ESCC tumor and adjacent non-tumor tissues before NAC. (B) Relative expression of circ_0004365 from ESCC tumor tissues and adjacent non-tumor tissues after NAC. (C) Relative expression of circ_0004365 from matched ESCC tumor tissues before and after NAC compared using the paired-samples t-test. N, nontumor tissues; T, tumor tissues; ESCC, esophageal squamous cell carcinoma; NAC, neoadjuvant chemotherapy.

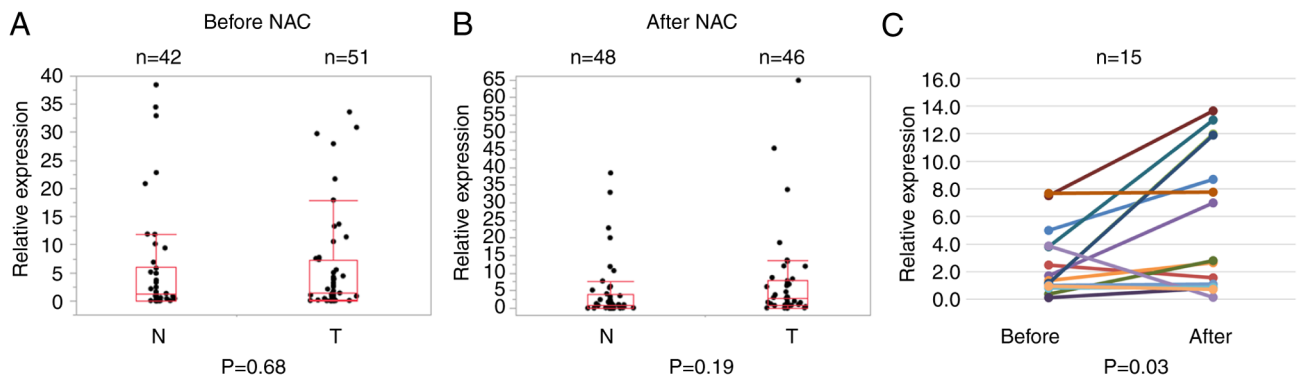
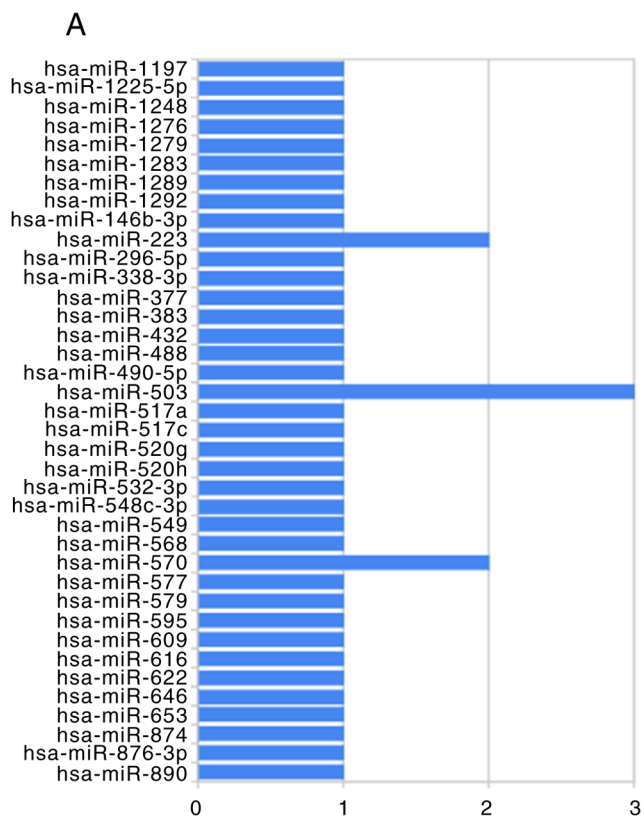


Figure S3. (A) CircInteractome predicting the binding sites between hsa_circ_0004365 and miR. (B) Sequences of the binding sites between hsa_circ_0004365 and miR-503. circRNA, circular RNA; miRNA, microRNA.



B

hsa_circ_0004365 (5'-3')	
hsa-miR-503 (3'-5')	
1	UUCAUGGGGACAGAUGCUGCUAU GACGUCUUGACAAGGGCGACGAU
2	UGUGGAUCGAGUGAACGCUGCUG GACGUCUUGACAAGGGCGACGAU
3	UUCAUGGGGACAGAUGCUGCUAU GACGUCUUGACAAGGGCGACGAU