## **Supplementary materials**

**Figure S1 (A)** Screening of miR-30a expression in breast cancer cell lines and normal breast cell line by qRT-PCR. \*\*\* P < 0.001. (B) qRT-PCR results of miR-30a in breast cancer cells stably transfected with miR-30a. \*\*\* P < 0.001. (C) Relative expression of miR-30a in HBL-100 stably transfected with miR-30a inhibitor. (D) Western blot results of expression for EMT related markers in HBL-100 cells transfected with miR-30a sponge. (E) The expression of miR-30a was downregulated in breast cancer in TCGA database. \* P < 0.05.

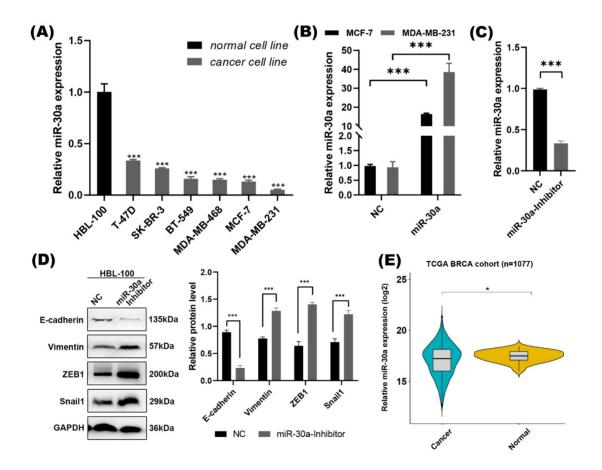
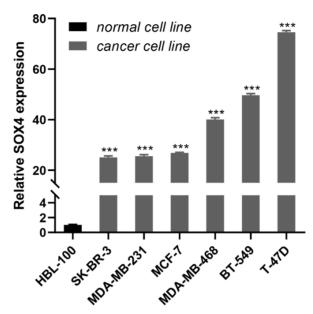
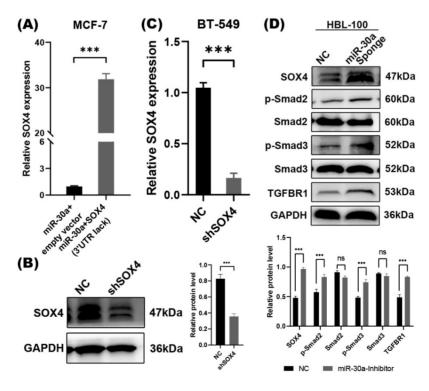


Figure S2 Screening of SOX4 expression in breast cancer cell lines and normal breast cell line. \*\*\* P < 0.001.



**Figure S3 (A)** qRT-PCR results of SOX4 expression in cells transfected with miR-30a or co- transfected with miR-30a and SOX4 lacking 3'UTR. **(B)** Western blot results of SOX4 expression in BT549 cells transfected with shSOX4. **(C)** Relative expression of SOX4 in BT549 stably transfected with shSOX4. **(D)** Western blot results of SOX4 and TGF- $\beta$  related markers in HBL-100 cells transfected with miR- 30a sponge.



Antibodies	Working concentration			Sources
	WB	IF	IHC	Sources
E-cadherin	1:1000	1:100	1:400	CST
Vimentin	1:1000	1:200	1:100	CST
Snail1	1:1000	-	-	CST
ZEB1	1:1000	-	-	CST
SMAD2	1:1000	-	-	CST
p-SMAD2	1:1000	-	1:200	CST
SMAD3	1:1000	-	-	CST
p-SMAD3	1:1000	-	1:200	CST
TGFBR1	1:500	-	1:100	R&D
SOX4	1:500	-	1:200	Abcam
BMI1	-	-	1:200	Proteintech
OCT4	-	-	1:200	Proteintech
GAPDH	1:3000	-	-	Servicebio

Supplementary table 1. Details for primary antibodies.

Abbreviations: WB: Western blot; IF: Immunofluorescence; IHC: immunohistochemistry; CST: Cell Signaling Technology (Beverly, MA, USA); R&D: R&D Systems GmbH, Germany.

Genes	Туре	Primer sequences	
miR-30a-5p	Forward	5'-CAGTGCAGGGTCCGAGGT-3'	
	Reverse	5'-AACACGCTGTAAACATCCTCGAC-3'	
U6	Forward	5'-CTCGCTTCGGCAGCACA-3'	
	Reverse	5'-AACGCTTCACGAATTTGCGT-3'	
SOX4	Forward	5'-CTGCGCCTCAAGCACATG-3'	
	Reverse	5'-TTCTTCCTGGGCCGGTACT-3'	
GAPDH	Forward	5'-GGGAGCCAAAAGGGTCATCA-3'	
	Reverse	5'-TGATGGCATGGACTGTGGTC-3'	

Supplementary table 2. The information for specific primer sequences of genes.

	Tissue specimens	TMA
	(n=42)	(n=133)
Age, years (range)	59 (26, 74)	55 (29, 87)
ER Status		
positive	23 (54.8%)	90 (69.2%)
negative	19 (45.2%)	40 (30.8%)
PR Status		
Positive	19 (45.2%)	71 (53.8%)
negative	23 (54.8%)	61 (46.2%)
HER2 Status		
Positive	22 (52.4%)	23 (17.7%)
negative	20 (47.6%)	107 (82.3%)
Subtype		
TNBC	10 (23.8%)	27 (20.8%)
Non-TNBC	32 (76.2%)	103 (79.2%)
Tumor Size		
≤2cm	14 (32.6%)	52 (39.1%)
>2cm	29 (67.4%)	81 (60.9%)
lymph node involved		
0	22 (51.2%)	71 (53.4%)
1-3	16 (37.2%)	19 (14.3%)
>3	5 (11.6%)	43 (32.3%)

Supplementary table 3. Clinical characteristics of breast cancer patients.

Abbreviations: TMA: tissue microarray; ER: estrogen receptor; PR: progesterone receptor; HER2: human epidermal growth factor receptor-2; TNBC: triple-negative breast cancer.