Supplementary Information

Supplementary Figures and Figure Legends



Supplemental Figure S1. Twist promotes the expression of ROR1. (A)T47D cells expressing the vector or the Twist were examined for morphological changes indicative of EMT by phase microscopy and the expression of ROR1, E-cadherin, and Twist (green) by immunofluorescence staining (40X).

Figure S1





Supplemental Figure S2. Both Twist and ROR1 are highly expressed in BLBC cells.

(A)The indicated proteins were analyzed by Western blotting in indicated cells.



Supplemental Figure S3. The promoter activity of ROR1 is marginally affected by mutating E-box in fragment 3. (A)Schematic illustration of wild-type ROR1 promoter and its mutant. (B) ROR1 promoters (wild-type or mutant) linked to luciferase were transfected into MDA-MB-231 cells, and after 48h luciferase activity was assayed.





Supplemental Figure S4. Silencing ROR1 inhibits the growth of BLBC cells. (A) The viability of indicated cells stably expressing NC or shROR1 was measured with an MTT. The dots represent the means, and the bars indicate the SD. The results are

expressed as the mean \pm SD of three independent experiments. **(B)** Images of colony formation of indicated cells. **(C)** colony-forming efficiencies were calculated after 12-day conventional culture. Measurements were carried out in triplicate, and experiments were repeated three times. Data are presented as mean \pm SD. P values were calculated using Student's t test. **P < 0.01.







Supplemental Figure S5. The knockdown of ROR1 enhances migration and invasion of BLBC cells. (A) Western blot analyses of ROR1 expression in the Hs 578T cells with control or ROR1 shRNA. (B-C) Cell migration and invasion were assessed in Hs 578T (B) and MDA-MB-231 (C) cells stably expressing control or shRNA-ROR1, as described in the Materials and Methods (40X).



Supplementary Figure S6. The promotion of cell migration and invasion after Twist overexpression primarily depends on ROR1. Hs 578T(A) and MDA-MB-231(B) cells stably expressing control, shRNA-ROR1, Twist or both, as indicated, were subjected to cell migration and invasion assays as described in the Materials and Methods (40X).

	Twistexpression		T - 4 - 1
	Low	High	Iotal
ROR1 expressio	n	•	
Low	60	22	82
High	29	23	52
Total	89	45	134

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Supplementary Figure S7. The combination of Twist and ROR1 correlates with clinical prognosis in breast cancer. (A) A positive correlation was observed between Twist and ROR1 expression in 134 paraffin-embedded breast cancer tissues (p = 0.038, χ^2 tests. R: Spearman correlation coefficient). Overall survival curve was generated on the basis of the protein level of Twist (**B**) or ROR1(**C**) in 134 paraffin-embedded breast cancer tissues.

Supplementary Materials and Methods: Primers used in qRT-PCR, the luciferase reporter assay, and the chromatin immunoprecipitation (ChIP) assay.

RNA extraction and qRT-PCR

The primers used to amplify the indicated genes are as follows.

	qRT-PCR Primers		
Gene	Forward Sequence (5'-3')	Reverse sequence (5'-3')	
Twist	GTCCGCAGTCTTACGAGGAG	GCTTGAGGGTCTGAATCTTGCT	
ROR1	CAGTCAGTGCTGAATTAGTGCC	TCATCGAGG GTCAGGTAAGAAT	
Vimentin	GACGCCATCAACACCGAGTT	CTTTGTCGTTGGTTAGCTGGT	
ZO-1	GTATCCGATTGTTGTGTGTTCC	TCACTTGTAGCACCATCCGC	
GAPDH	ACAGTCAGCCGCATCTTCTT	GACAAGCTTCCCGTTCTCAG	

The luciferase reporter assay

The primers used for cloning the indicated promoters are as follows.

	Primers for promoter clone		
Gene	Forward Sequence (5'-3')	Reverse sequence (5'-3')	
ROR1-	CCCAGGGCGACTCACGCCCACT	CGTGAGTCGCCCTGGGTCGGA	
Mut-1	GGTGCGACCCGGACAGCCTGGG	CTCCGAGAACAGCGAAAATTT	
	ACTGACCCGCC	CGCCCAGGGGCTG	
ROR1-	CCCAGGGCGACTCACGCCCACT	CGTGAGTCGCCCTGGGTCGGA	
Mut-2	GGTGCGACCCGGAAAATTTTGG	CTCCGAGAACAGCGCACCCTG	
	ACTGACCCGCC	CGCCCAGGGGCTG	
ROR1-	CCCAGGGCGACTCACGCCCACT	CGTGAGTCGCCCTGGGTCGGA	
D-Mut	GGTGCGACCCGGAAAATTTTGG	CTCCGAGAACAGCGAAAATTT	
	ACTGACCCGCC	CGCCCAGGGGCTG	
ROR1-	TAAATCTTGCCAGGAGGCAGTA	CTCCTGGCAAGATTTATTGAGG	
Mut-3	AATTTACATTGA	ATTA	

The chromatin immunoprecipitation (ChIP) assay

The primers for the indicated promoters are as follows.

	ChIP-qPCR Primers		
Gene	Forward Sequence (5'-3')	Reverse sequence (5'-3')	
ROR1-1	GGTGCCAAGGAGTATGGA	TCGGGAGTTGTTGCTGTT	
<i>ROR1-2</i>	CTGAGTGCTGGCTTGGCTAC	CTGGAGGAACAGGAAGAAGC	
ROR1-3	AAAGTTAAACCAGGAGAAATGG	CTGGCAAGATTTATTGAGGATT	
<i>ROR1-4</i>	CCTCGGTTTCCCCTTCTG	CTCCTCAAACTCGCCACC	
ROR1-5	GACCCAGGGCGACTCACG	CAGAACATCCACGGGCTCTTC	
Vimentin	CAGCCTATCACAGCCCAGAG	ATGTCTCCTGGAATGGGCAC	
GAPDH	CCCAAAGTCCTCCTGTTTCA	GTCTTGAGGCCTGAGCTACG	