



Figure S1

Figure S1 Gene Set Enrichment Analysis (GSEA) between the Cluster 1 and Cluster 2.

(A-D) Results of GSEA in TCGA-BRCA cohort.



Figure S2 Differential signaling pathways and cellular processes between the highrisk subgroup and the low-risk subgroup

(A-C) Biological processes of differentially expressed genes between the high-risk subgroup and the low-risk subgroup. (D-F) KEGG pathways in which differentially expressed genes were distributed.







(A) The CCK-8 assay was performed to measure the proliferation capacity of CAL51 cells; (B) The colony formation assay and corresponding statistical analysis of CAL51 cells; The effect of FBXL6 on the migration of CAL51 cells determined by transwell(C) and wound healing (D) assays; (E) Western blot analysis revealed that downregulation of FBXL6 inhibited the cell cycle progression in CAL51 cells.





Figure S4 upregulation of PDZRN3 inhibited the proliferation and migration of CAL51 breast cancer cells.

(A) The CCK-8 assay was performed to measure the proliferation capacity of CAL51 cells. (B) The colony formation assay, and corresponding statistical analysis of CAL51 cells. The effect of PDZRN3 on the migration of CAL51 cells determined by transwell (C) and wound healing (D) assays. (E) Western blot analysis revealed that upregulation of PDZRN3 inhibited the cell cycle progression in CAL51 cells.

	Antibodies	Sourse	Cat.No.
Western blot	β-actin	Cell Signaling Technology	3700
	P-21	Proteintech	10355-1-AP
	P-16	Proteintech	10883-1-AP
	Cyclin B1	Santa Cruz	SC-752
	Cyclin D1	Santa Cruz	SC-753
	GAPDH	Sangene Biotech	KM9002
	FBXL6	Bioss Antibodys	bs-16041R
	PDZRN3	Abcam	ab272628
	β-catenin	Cell Signaling Technology	8480
	Vimentin	Immunoway	YT4880
	Ki67	BD pharmingen	PAC047Ra01

Table S1 the Antibody information used in this research.

Name	Sequence (5' to 3')	
DCAF13 up	GAACTCCTAGCGGACACCT	
DCAF13 low	CAACTTGGTTTCGCGGACATA	
USP39 up	TTGGAAGAGGCGAGATAA	
USP39 low	AGGAGCATCAATCATCATC	
PSMD14 up	TCGGAAGCCTAACTACAGCGA	
PSMD14 low	ATTATTGAGGTCAACGGCAG	
PDZRN3 up	ATTATTGAGGTCAACGGCAG	
PDZRN3 low	AGGGCCATGATATGTTCAAAG	
TLE3 up	AACCACCATGAACTCGATCAC	
TLE3 low	TCACTGTCGTATCGGCTCAAG	
SOCS2 up	TTAAAAGAGGCACCAGAAGGAAC	
SOCS2 low	AGTCGATCAGATGAACCACACT	
SKP2 up	CTTTACTATTAGTGACAAGAGCTGG	
SKP2 low	TGGCTGGACTTGAGTTTGGA	
FBXL6 up	GGAGACCGCATTCCCTTGG	
FBXL6 low	AAAACCGATTGGGCATAAGCC	
PSMD14 up	TGTGGAGGCAGTTGATCCAG	
PSMD14 low	TCCACACCAGAAAGCCAACA	