Supplementary description

Figure S1 Relation of SLC3A2 and therapeutic responses in breast cancer. **A-C** The receiver operating characteristic curve plot of the association between SLC3A2 expression and responses to endocrine therapy, anti-HER2 therapy, and chemotherapy in breast cancer cohort considering pathological complete response. **D-F** The receiver operating characteristic curve plot of the association between SLC3A2 expression and responses to endocrine therapy, anti-HER2 therapy, and responses to endocrine therapy, anti-HER2 therapy, and responses to endocrine therapy, anti-HER2 therapy, and chemotherapy in breast cancer cohort considering pathological between SLC3A2 expression and responses to endocrine therapy, anti-HER2 therapy, and chemotherapy in breast cancer cohort considering relapse-free survival at 5 years.

Figure S2 Relation of SLC3A2 and clinicopathological characteristics in breast cancer. **A-H** Correlations between SLC3A2 mRNA expression and genders, cancer stages, nodal metastasis status, molecular subtypes, races, ages, menopause status and TP53 mutation status. **p<0.01, ***p<0.001.

Figure S3 Relation of SLC3A2 and single cell sequencing in breast cancer. **A** Correlations between SLC3A2 and diverse functional states in breast cancer from CancerSEA database. **B-I** Box diagram and scatter diagram showing the SLC3A2 expression among four single cell datasets (EXP0052, EXP0053, EXP0054, EXP0055).

Figure S4 Everolimus inhibits the proliferation of T47D cells. **A** T47D cells were treated with everolimus (10 uM) for 0, 24, 48, or 72 h, and cell viability was assayed. **B** Comparison of T47D cell proliferative capacity among control and everolimus (10 uM) groups at 72 h via Edu staining. Scale bar = 50 um. **C** T47D cells were treated with varying doses of everolimus alone or in combination with different doses of erastin for 48 h; cell viabilities were measured. *p<0.05, **p<0.01, ***p<0.001.

Figure S5 SLC3A2 influences cell viability of breast cancer cells by everolimus treatment. **A** T47D cells were treated with everolimus (10 uM) for 0, 24, 48, or 72 h with or without the overexpression of SLC3A2. Cell viability was assayed. **B** MCF7 cells were treated with everolimus (10 uM) for 0, 24, 48, or 72 h with or without the downexpression of SLC3A2. Cell viability was assayed. **C** T47D cells were treated with everolimus (10 uM) for 0, 24, 48, or 72 h with or without the downexpression of SLC3A2. Cell viability was assayed. **C** T47D cells were treated with everolimus (10 uM) for 0, 24, 48, or 72 h with or without the downexpression of SLC3A2. Cell viability was assayed. **C** T47D cells were treated with everolimus (10 uM) for 0, 24, 48, or 72 h with or without the downexpression of SLC3A2. Cell viability was assayed. **C** T47D cells were treated with everolimus (10 uM) for 0, 24, 48, or 72 h with or without the downexpression of SLC3A2. Cell viability was assayed. **C** T47D cells were treated with everolimus (10 uM) for 0, 24, 48, or 72 h with or without the downexpression of SLC3A2. Cell viability was assayed. **C** T47D cells were treated with everolimus (10 uM) for 0, 24, 48, or 72 h with or without the downexpression of SLC3A2. Cell viability was assayed. *****p<0.05.

Pathological complete response (n=1775)



Relapse-free survival at 5 years (n=1329)





A Expression of SLC3A2 in BRCA based on patient's gender





B

D





Expression of SLC3A2 in BRCA based on nodal metastasis status







TCGA samples











F







Figure S3 A

Correlations between the gene (gene list) of interest and functional states in different single- cell datasets.					orre	latio	n –0.5	5 -(.25	0	().25	0.	.5		!	₽ _
	ExpID	Name 💿	Cancer	No.cells	Angiogenesis	Apoptosis	CellCycle	Differentiation	DNArepair	EMT	Hypoxia	Inflammation	Invasion	Metastasis	Proliferation	Quiescence	Stemness
0	EXP0052	Braune EB. Stem Cell Reports. 2016 (PDX)	Breast cancer	369													
0	EXP0053	Chung W. Nat Commun. 2017 (Breast)	Breast cancer	317													
0	EXP0054	Jordan NV. Nature. 2016 (CTC)	Breast cancer	70													
0	EXP0055	Aceto N. Mol Cancer Res. 2018 (CTC)	Breast cancer	32													



B

D

 Expression of input gene(s)

 Housekeeping genes
 -2 0 2 4 6 8 10

 Input gene(s)
 -2 0 2 4 6 8 10



С

E

12





F

H Expression of input gene(s) Housekeeping genes Input gene(s) -2 0 2 4 6 8





Ι

10



