

## 1 **Supplementary Materials and Methods**

### 2 **Acute toxicity experiment**

3 Six female Balb/c mice were randomly divided into control (3 mg PS-T, every other day) and  
4 experimental groups (30 mg PS-T, every other day), with three mice in each group. PS-T was  
5 dissolved in 100  $\mu$ L of isotonic sodium chloride solution and administered via oral gavage. Body  
6 weight and food intake were measured daily for two weeks, and weekly changes were recorded.  
7 Mortality (if any) and abnormalities in the mice were also recorded. After the mice were sacrificed,  
8 organ weights and organ coefficients were measured, and the heart, liver, spleen, lung, and kidney  
9 tissues were prepared for H&E staining.

### 10 **Cell viability assay**

11 Cell viability was assessed using the Cell Counting Kit-8 (CCK-8, CK04, Dojindo, Japan). MDA-  
12 MB-231 and 4T1 cells were seeded in 96-well plates at a density of  $5 \times 10^3$  cells per well and  
13 incubated overnight to allow attachment. Cells were then treated with PS-T at concentrations of 0, 5,  
14 and 10  $\mu$ g/mL for 24 h. Following treatment, 10  $\mu$ L of CCK-8 solution was added to each well and  
15 incubated for 2 h at 37 °C. The absorbance was measured at 450 nm using a microplate reader  
16 (Thermo Scientific). Cell viability was calculated as the percentage relative to untreated control cells.  
17 Each experiment was performed in triplicate and repeated at least three times.

### 18 **Sequencing**

19 We plated MDA-MB-231, HCC1806 and HCC1937 cells in 6-well plates and treated them with or  
20 without 50  $\mu$ g/mL PS-T for 24 h. After digestion, the cells were stored in TRIzol reagent ( $5 \times 10^6$  cells  
21 per ml of TRIzol), transferred to DNase-free cryopreservation tubes, and stored at -80 °C.  
22 Sequencing was performed by MajorBio (Shanghai, China).

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25 **Supplementary Figures**

A

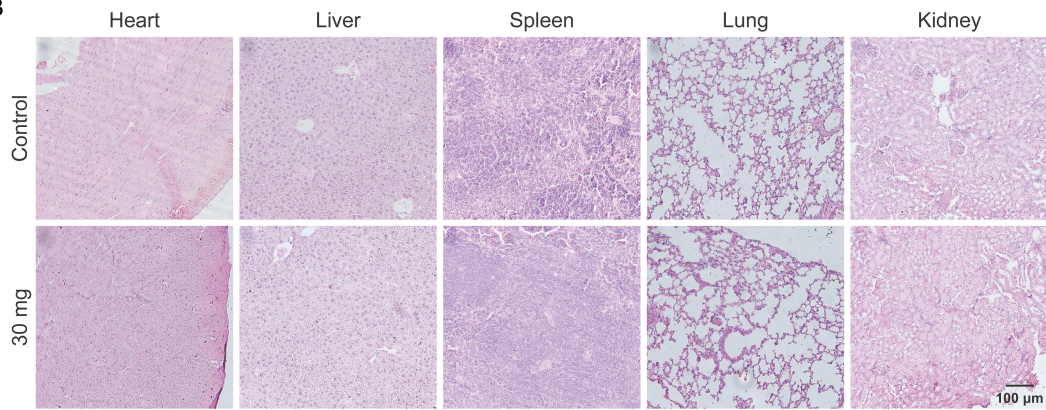
Body weight gain, food and water consumption, and organ coefficient results of mice treated with PS-T

Parameters	Acute toxicity	
	Control	30 mg
Initial weight (g)	15.867±0.507	15.820±0.286 <sup>ns</sup>
One week (g)	17.067±0.621	16.950±0.439 <sup>ns</sup>
Two weeks (g)	18.103±0.681	18.070±0.479 <sup>ns</sup>
Body weight gain (g)	2.237±0.616	2.250±0.193 <sup>ns</sup>
Food intake (g/day)	3.213±0.291	3.153±0.153 <sup>ns</sup>
Heart coefficient	0.028±0.003	0.027±0.004 <sup>ns</sup>
Liver coefficient	0.262±0.006	0.255±0.006 <sup>ns</sup>
Spleen coefficient	0.018±0.001	0.018±0.001 <sup>ns</sup>
Lung coefficient	0.034±0.001	0.036±0.002 <sup>ns</sup>
Kidney coefficient	0.042±0.001	0.045±0.004 <sup>ns</sup>

Results are expressed as mean±SD (n=6)

<sup>ns</sup> Not significant

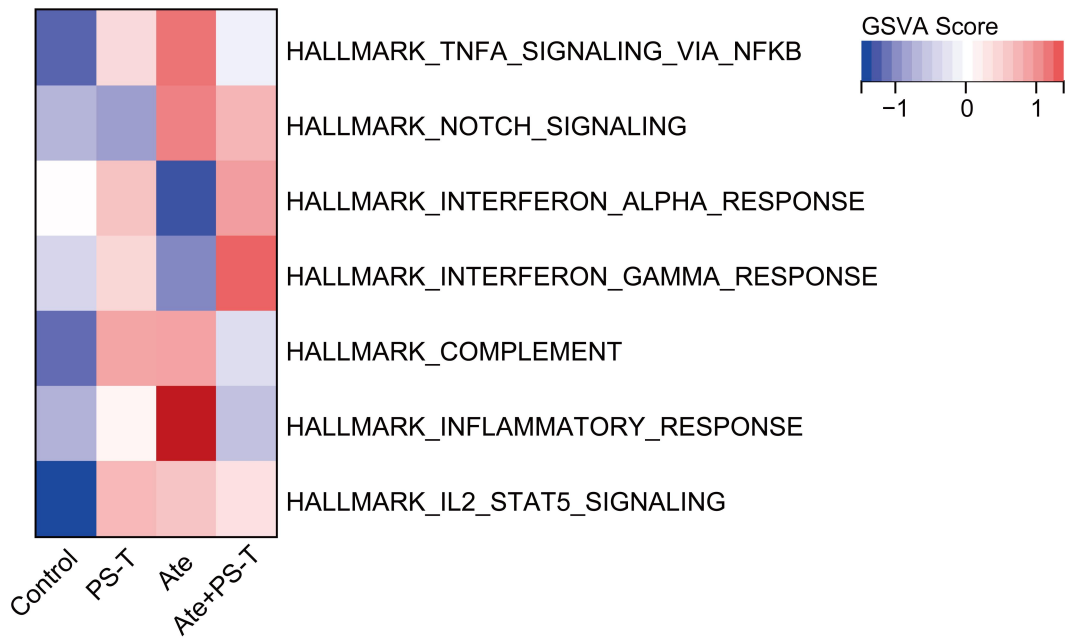
B



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27 **Fig. S1. Acute toxicity test of PS-T in mice**

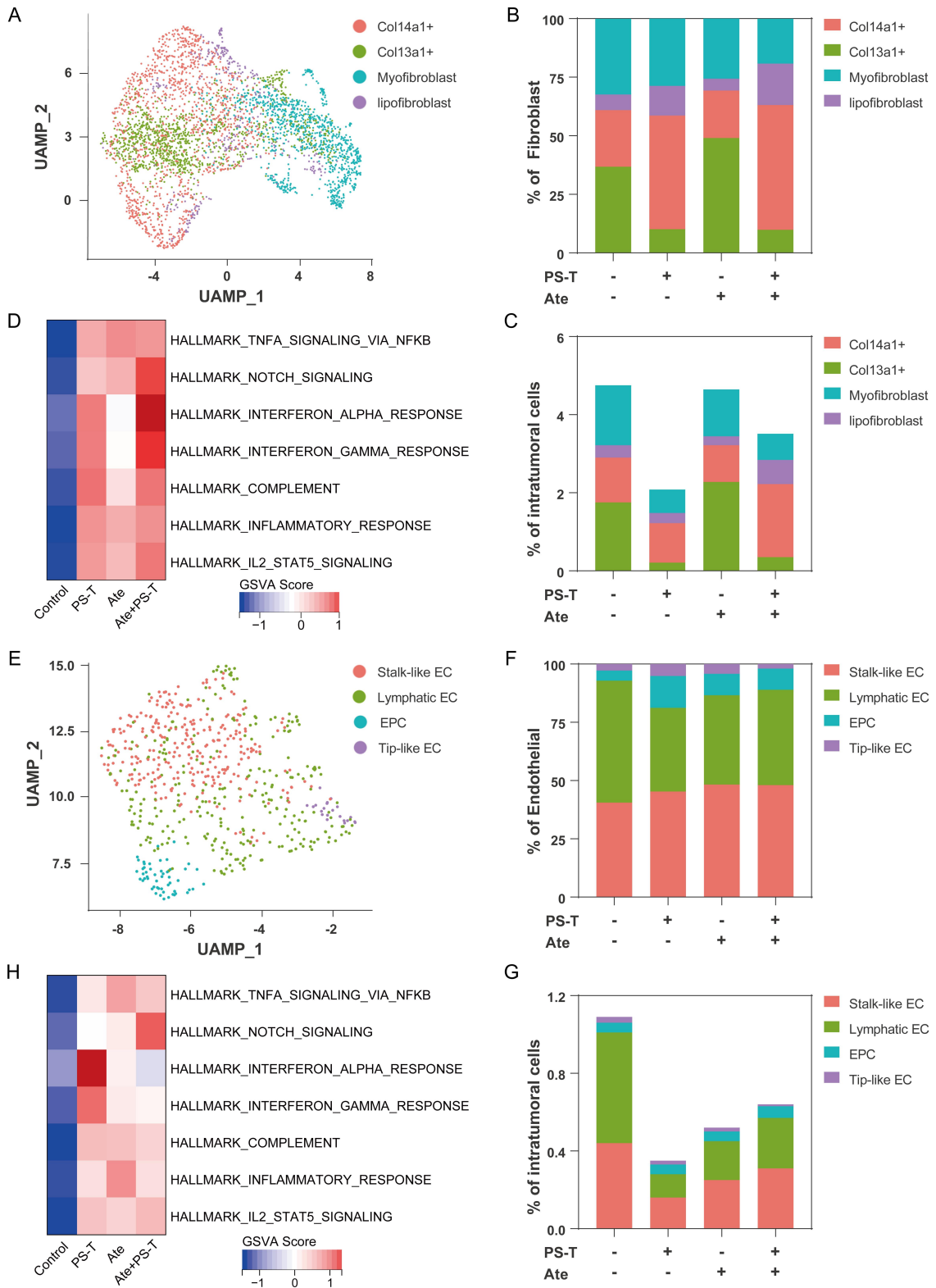
28 **(A)** Body weight gain, food and water consumption, and organ coefficient results of mice treated with  
 29 3 mg PS-T (control) and 30 mg PS-T (n=6). **(B)** Representative images of HE staining of heart, liver,  
 30 spleen, lung, and kidney tissues from the control (*up*) and 30 mg PS-T group (*bottom*) (n=6). Scale  
 31 bars = 100 µm. The data were combined based on two independent experiments. (mean ± standard  
 32 deviation; ns, not significant).



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34 **Fig. S2. GSVAscore analysis of cancer cells**

35 GSVAscore analysis of cancer cells with TNF- $\alpha$ /NF- $\kappa$ B, Notch, and IL-2/STAT5 signaling, IFN- $\alpha$ , IFN- $\beta$ , and  
 36 inflammatory response, and complement.

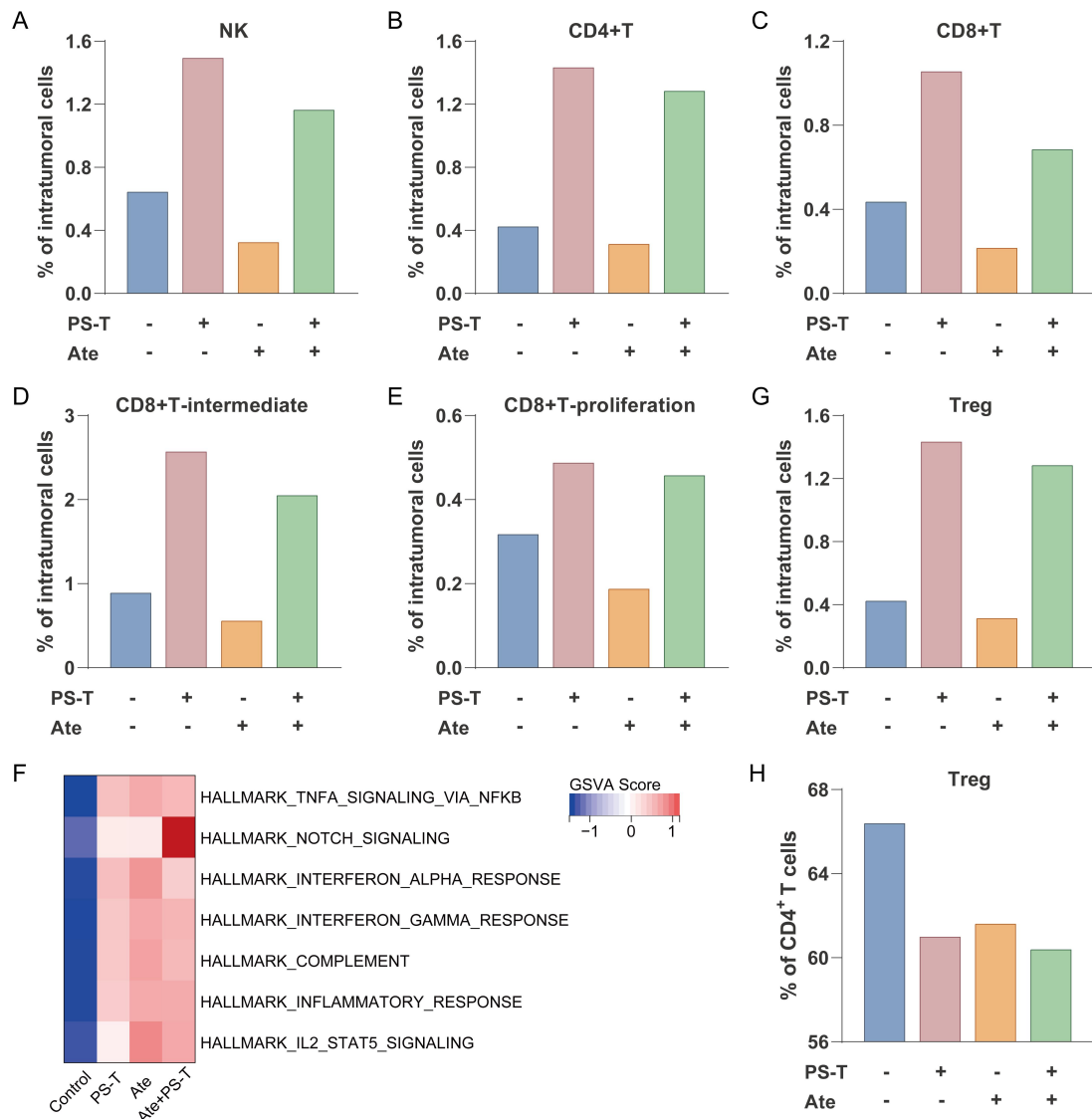


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38 **Fig. S3. Subclustering and GSVA analysis of fibroblasts and endothelial**

39 **(A)** UMAP visualization of fibroblast subsets (Col14a1+, Col13a1+, Myofibroblast, and lipofibroblast).

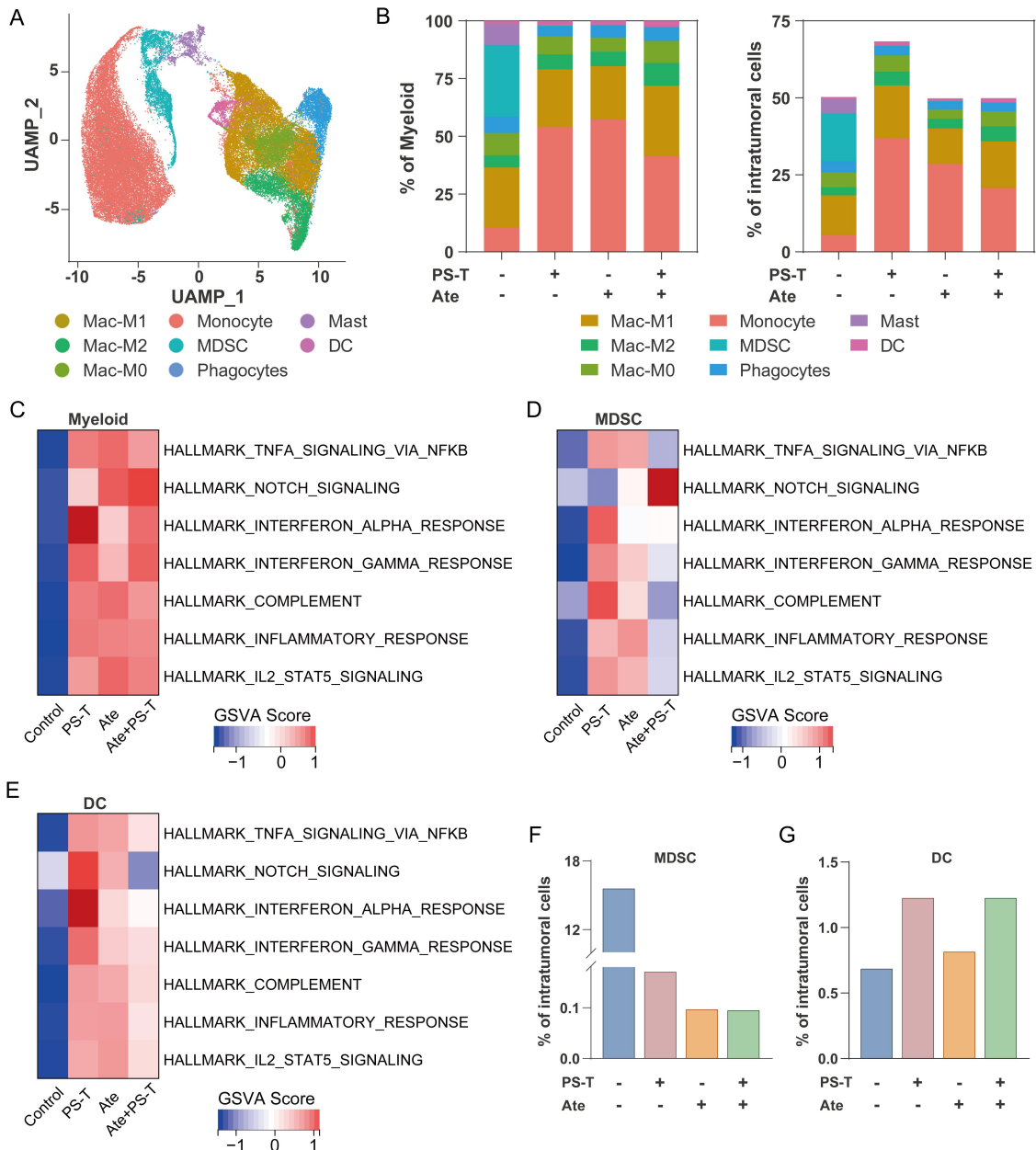
40 **(B-C)** The proportions of subsets in fibroblasts (B) and intratumoral cells (C). **(D)** GSVA analysis of  
41 fibroblasts with immune-related pathways. **(E)** UMAP visualization of endothelial cell subsets (Stalk-  
42 like EC, Lymphatic EC, EPC, and Tip-like EC). **(F-G)** The proportions of endothelial cell subsets in  
43 endothelial (F) and intratumoral cells (G). **(H)** GSVA analysis of endothelial cells with immune-related  
44 pathways.  
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47 **Fig. S4. Subclustering analysis of T cells**

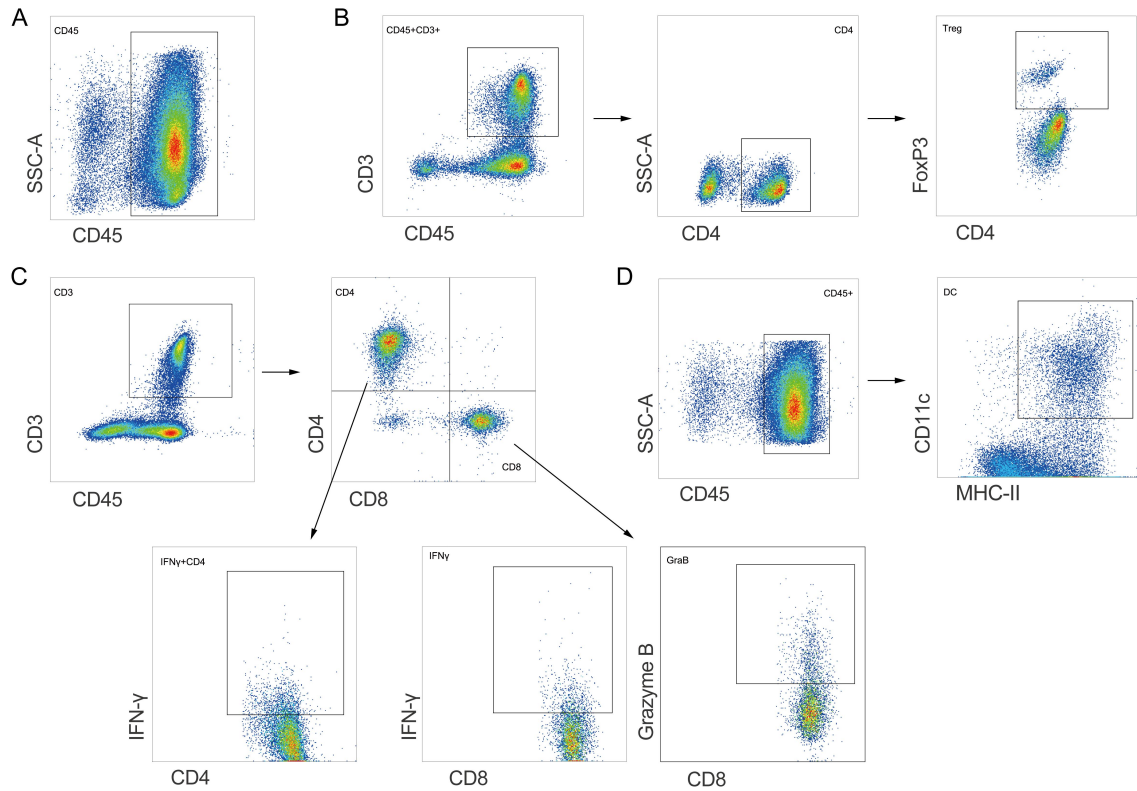
48 **(A-E)** The proportion of NK (A), CD4<sup>+</sup> T (B), CD8<sup>+</sup> T cells (C), CD8<sup>+</sup> T-  
 49 proliferation cells (E) in intratumoral cells. **(F)** GSVAs analysis of T cells with immune-related pathways.  
 50 **(G-H)** The percentage of Tregs in intratumoral cells (G) and tumor-infiltrating CD4<sup>+</sup> T cells (H).



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52 **Fig. S5. Subclustering analysis of myeloid cells**

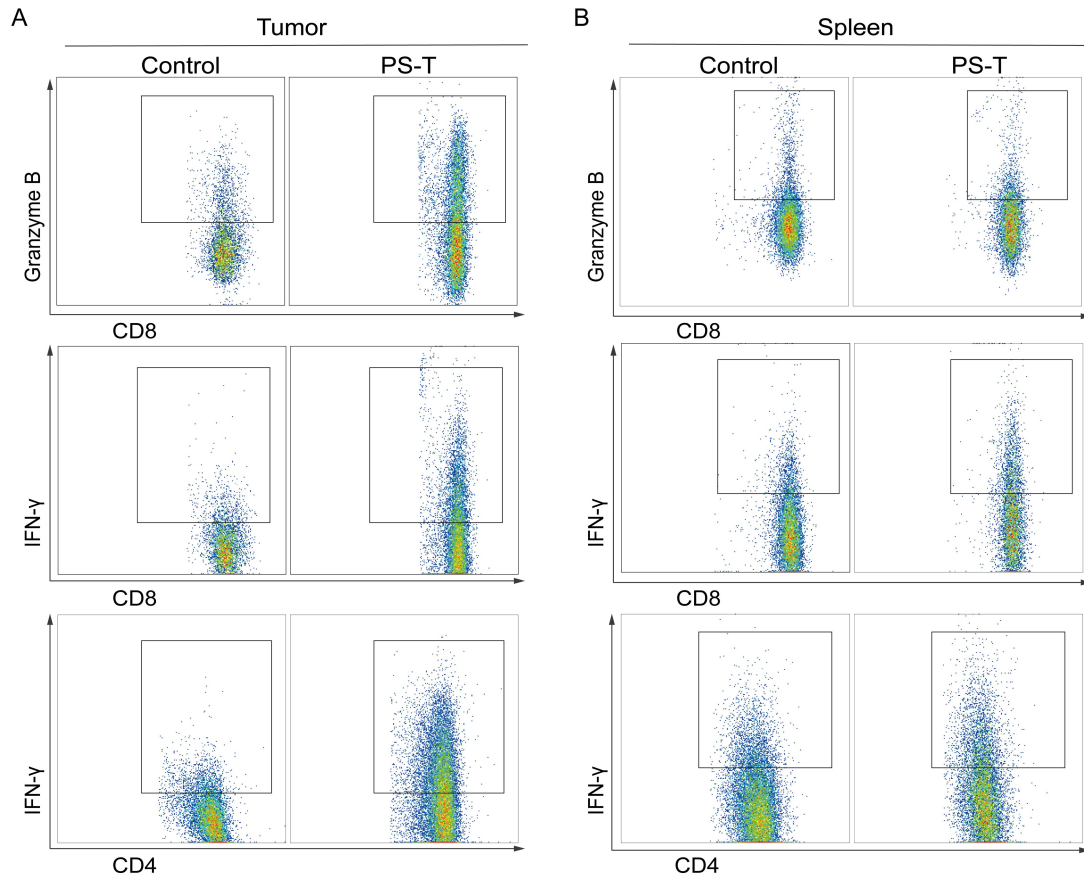
53 **(A)** UMAP visualization of myeloid cell subsets (Mac-M1, Mac-M2, Mac-M0, Monocyte, MDSC,  
 54 Phagocytes, Mast, and DC). **(B)** The proportions of myeloid cell subsets in myeloid (*left*) and  
 55 intratumoral cells (*right*). **(C-E)** GSVAscores analysis of myeloid cells (C), MDSC (D), and DCs (E) with  
 56 immune-related pathways. **(F-G)** The proportion of MDSC (F) and DCs (G) in intratumoral cells.



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58 **Fig. S6. Gating strategies of flow cytometry analysis**

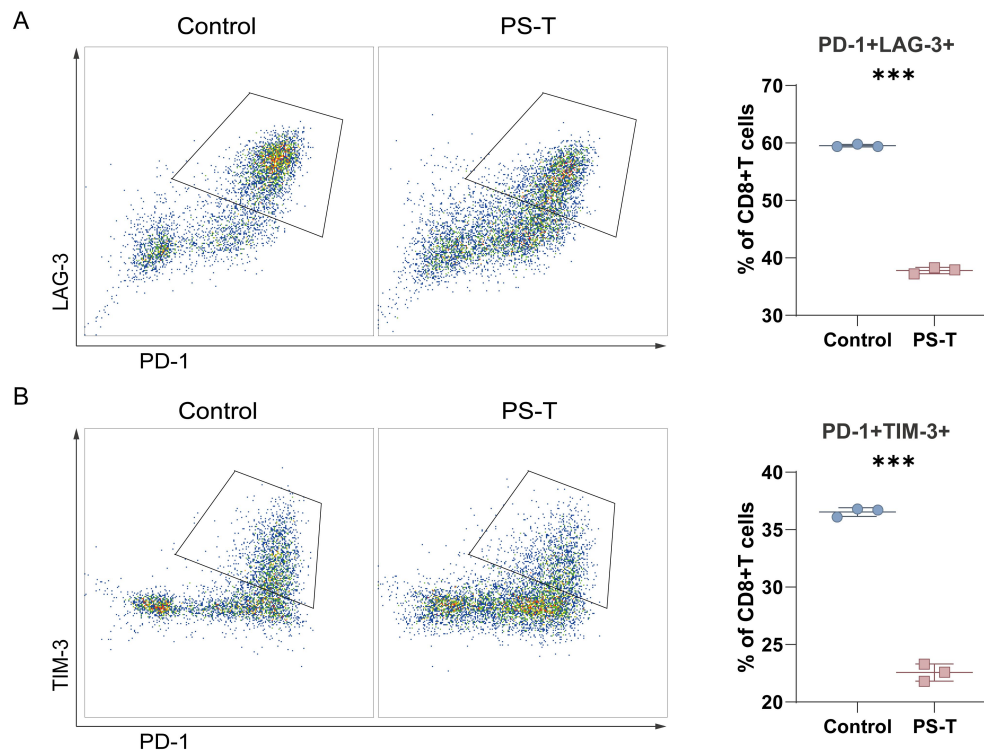
59 **(A-D)** Gating strategies of CD45<sup>+</sup> (A), regulatory T cells (Tregs) (B), CD3<sup>+</sup>, CD4<sup>+</sup>, and CD8<sup>+</sup> T cells,  
 60 cytotoxic T cell subsets (granzyme B<sup>+</sup> CD8<sup>+</sup> T, IFN- $\gamma$ <sup>+</sup> CD8<sup>+</sup> T, and IFN- $\gamma$ <sup>+</sup> CD4<sup>+</sup> T) (C), and dendritic  
 61 cells (DCs) (D).



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63 **Fig. S7. PS-T enhances T cell function in TNBC**

64 **(A-B)** Representative plots of IFN- $\gamma$  and granzyme B in tumor-infiltrating CD4<sup>+</sup> T and CD8<sup>+</sup> T cells in  
 65 the breast tumor (A) and spleen (B) (n=10).



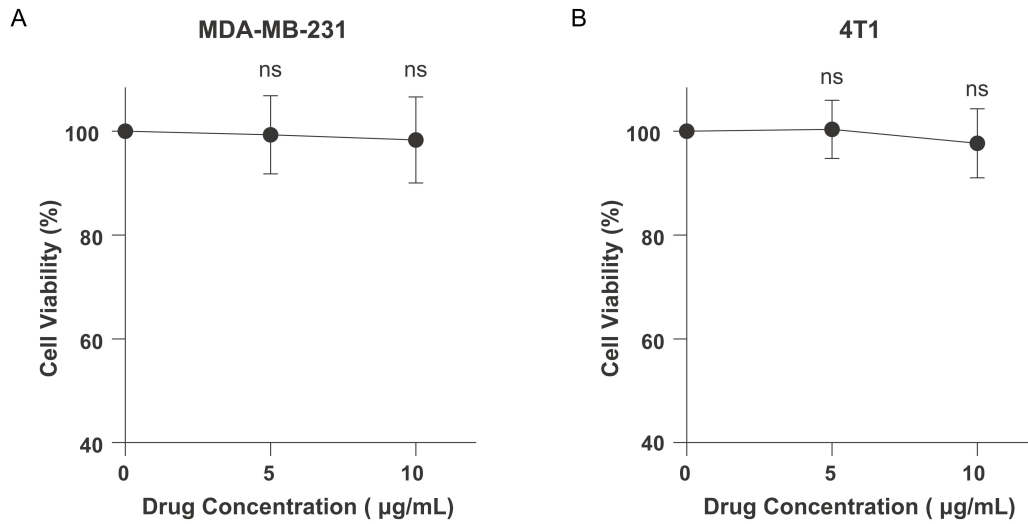
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67 **Fig. S8. PS-T alleviates T cell exhaustion in tumor-infiltrating CD8+ T cells**

68 **(A-B)** Representative flow cytometry plots showing PD-1<sup>+</sup> LAG-3<sup>+</sup> (A) and PD-1<sup>+</sup> TIM-3<sup>+</sup> CD8<sup>+</sup> T cells

69 (B) from 4T1 tumor-bearing mice treated with PS-T (n = 3). (mean ± standard deviation; \*\*\*P < 0.001).

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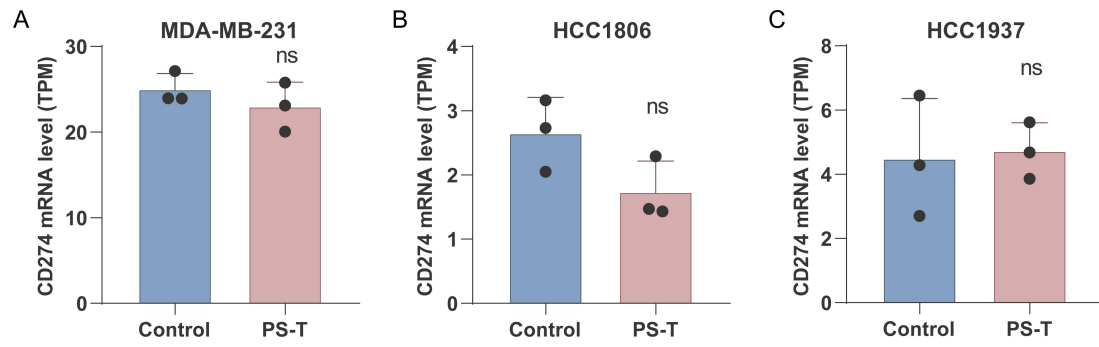


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72 **Fig. S9. Effect of PS-T on cell viability in TNBC cell lines**

73 **(A-B)** MDA-MB-231 (A) and 4T1 (B) cells were treated with PS-T at indicated concentrations (0, 5,  
 74 and 10 µg/mL) for 24 h, and cell viability was assessed by CCK-8 assay (n = 9). (mean ± standard  
 75 deviation; ns, not significant).

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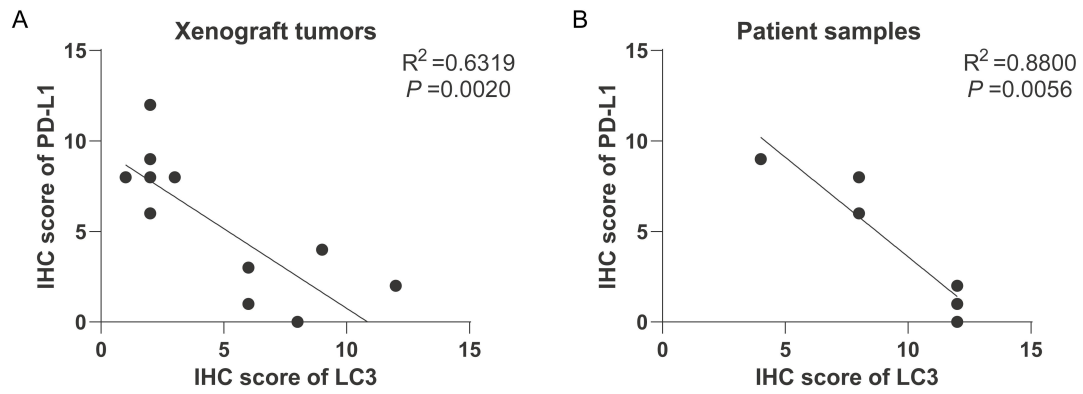


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78 **Fig. S10. PS-T modulates PD-L1 but not at the transcriptional level**

79 **(A-C)** Transcriptome sequencing was performed to measure the TPM of *CD274* mRNA levels in the  
 80 control and PS-T-treated MDA-MB-231 (A), HCC1806 (B), and HCC1937 cells (C). (mean  $\pm$  standard  
 81 deviation; ns, not significant).

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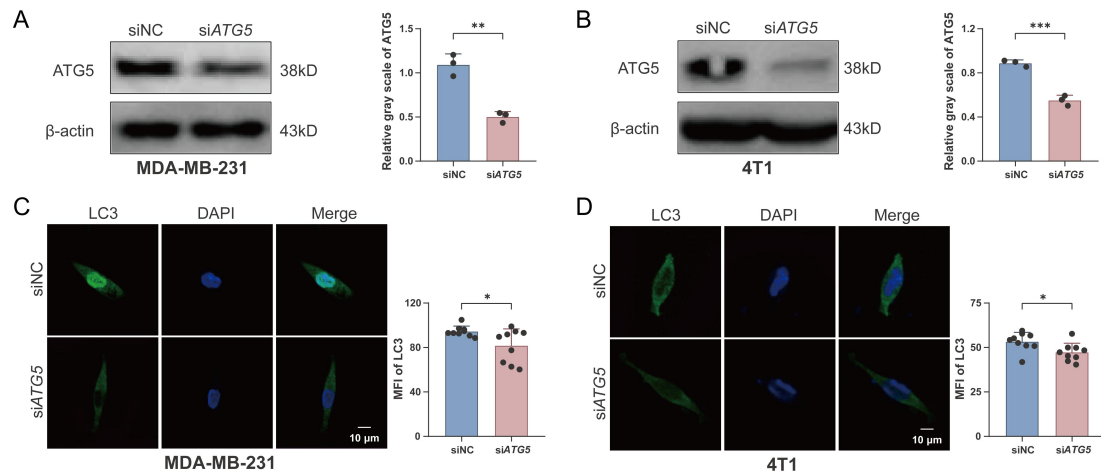


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84 **Fig. S11. Correlation analysis between LC3 and PD-L1 expression**

85 **(A)** Correlation analysis of LC3 and PD-L1 expression in 4T1 tumor-bearing mice ( $R^2 = 0.6319$ ,  $P =$   
 86  $0.0020$ ). **(B)** Correlation analysis of the IHC score of LC3 and PD-L1 in TNBC patients ( $R^2 = 0.8800$ ,  $P =$   
 87  $0.0056$ ). The Pearson correlation test was used for the statistical analysis. The data were combined  
 88 based on two independent experiments.

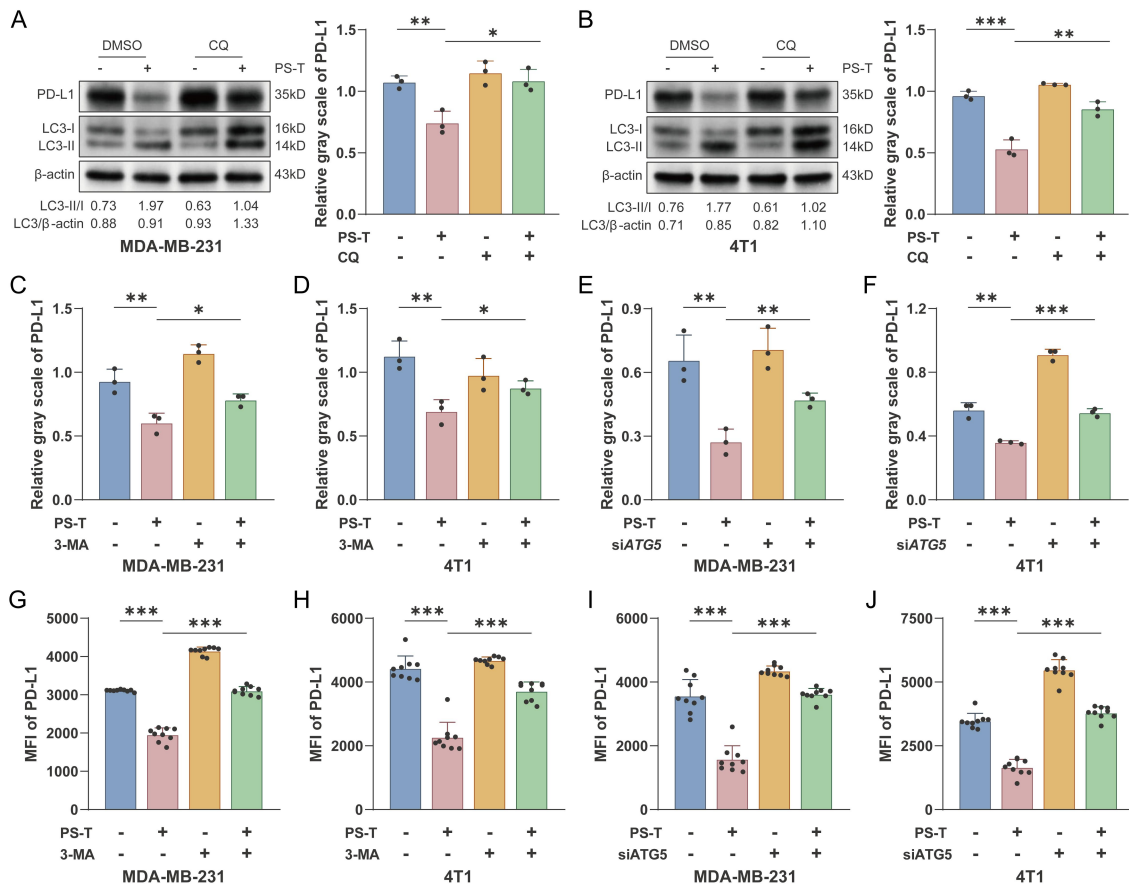
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91 **Fig. S12. Verification of ATG5 knockdown effect in TNBC cell lines**

92 **(A-B)** The expression levels of ATG5 in MDA-MB-231 (A) and 4T1 cells (B) were analyzed using  
 93 western blotting (n=3). **(C-D)** The expression levels of LC3 in MDA-MB-231 (C) and 4T1 cells (D)  
 94 were evaluated by immunocytochemistry (n=9). The data were combined based on three independent  
 95 experiments. (mean  $\pm$  standard deviation; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ ).

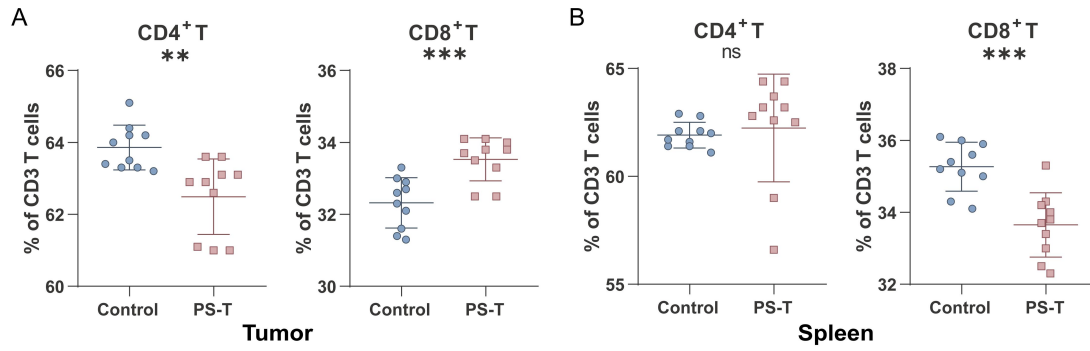


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97 **Fig. S13. Autophagy inhibition reverses the PS-T–induced degradation of PD-L1**

98 **(A-B)** The relative gray density of PD-L1 in MDA-MB-231 (A) and 4T1 cells (B) that autophagy was  
 99 inhibited by CQ (n=3). **(C-F)** The relative gray density of PD-L1 in MDA-MB-231 and 4T1 cells that  
 100 autophagy was inhibited by 3-MA (C-D) and siATG5 (E-F) (n=3). **(G-J)** The mean fluorescence  
 101 intensity (MFI) of PD-L1 protein in MDA-MB-231 and 4T1 cells that autophagy was inhibited by 3-MA  
 102 (G-H) and siATG5 (I-J) (n=9).

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105 **Fig. S14. Percentage of CD4<sup>+</sup> T and CD8<sup>+</sup> T cells in T lymphocytes**

106 **(A-B)** The percentage of CD4<sup>+</sup> T (*left*) and CD8<sup>+</sup> T cells (*right*) in tumor (A) and splenic (B) infiltrating T  
 107 lymphocytes (n=10). The data were combined based on two independent experiments. (mean ±  
 108 standard deviation; ns, not significant; \*\**P* < 0.01; \*\*\**P* < 0.001).

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110 **Supplementary Tables**

111 **Table. S1. Clinicopathological characteristics of TNBC patients**

No.	Age (years)	Tumor Size (cm)	Lymph Node Status	Histological Grade	Ki67 Index (%)	LC3 Expression Level	PD-L1 Expression Level
1	57	1.0	Positive	I	10	High	Low
2	75	5.0	Positive	II	40	High	Low
3	50	3.0	Negative	II	90	High	Low
4	68	3.0	Positive	II	80	High	Low
5	39	3.0	Positive	I	20	High	Low
6	41	2.0	Negative	II	40	High	Low
7	39	3.0	Negative	II	8	High	Low
8	50	3.0	Positive	II	30	High	Low
9	43	4.0	Positive	II	30	High	Low
10	55	5.0	Negative	II	30	High	Low
11	28	4.0	Positive	II	80	Low	High
12	71	2.5	Negative	II	30	Low	High
13	78	4.5	Negative	II	20	Low	High
14	68	3.0	Negative	II	50	Low	High
15	51	3.0	Positive	II	40	Low	High
16	57	4.0	Negative	II	40	Low	High
17	61	2.0	Negative	I	30	Low	High
18	66	2.5	Positive	II	15	Low	High
19	47	3.0	Positive	II	15	Low	High
20	39	2.8	Negative	II	50	Low	High

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113 **Table. S2. The TPM of CD274 mRNA levels in the tumor infiltrating immune cells**

	Control	PS-T	P value
<b>Myeloid</b>	0.24	0.64	0.08
<b>MDSC</b>	0.11	0.23	0.41
<b>DC</b>	0.30	0.60	0.15

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