

**Bioeliminable Pt@Bi<sub>2</sub>Se<sub>3</sub>-RGD Nanoassembly for Enhancing  
Photoacoustic Imaging-Guided Tumor Immuno-Radiotherapy by  
Inducing Apoptosis via Areg Pathway**

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## Materials

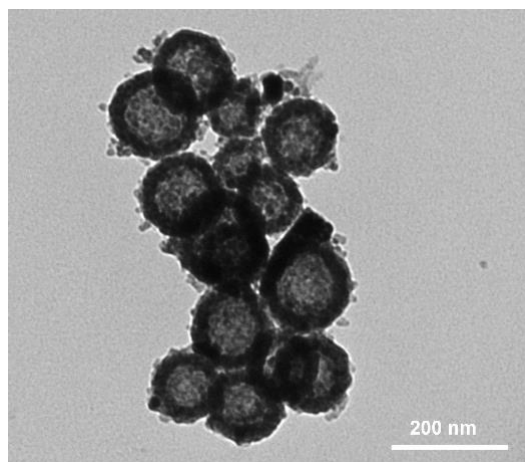
Bismuth nitrate pentahydrate ( $\text{Bi}(\text{NO}_3)_3 \cdot 5\text{H}_2\text{O}$ ), sodium hydroxide (NaOH), polyvinylpyrrolidone (PVP), ethylene glycol (EG), silk sericin (SS), ascorbic acid (AA), sodium selenite ( $\text{Na}_2\text{SeO}_3$ ), chloroplatinic acid ( $\text{H}_2\text{PtCl}_3$ ), sodium borohydride ( $\text{NaBH}_4$ ), 1-(3-dimethylaminopropyl)-3-ethylcarbodiimide (EDC), N-hydroxysuccinimide (NHS), Crgd-PEG-amine (cRGD-PEG-NH<sub>2</sub>), dimethyl sulfoxide (DMSO), terephthalic acid (TPA) and hydrogen peroxide ( $\text{H}_2\text{O}_2$ , 30%) were bought from Shanghai KeShi Technology Co. Ltd. (China). Nystatin and Me- $\beta$ -CD were bought from MACKLIN (China). Amiloride was bought from Aladdin (China). CPZ was bought from Sigma. Dulbecco's modified Eagle's medium (DMEM), a penicillin ( $10000 \text{ U} \cdot \text{mL}^{-1}$ )/streptomycin ( $10000 \mu\text{g} \cdot \text{mL}^{-1}$ ) mixture, fetal bovine serum (FBS), and TrypLE™ express enzyme were acquired from Thermo Fisher Scientific (Waltham, MA, USA). FITC and IR783 were obtained from Sigma-Aldrich (St. Louis, MO, USA). Calcein/PI Cell Activity and Cytotoxicity Assay Kit, Lyso-Tracker Red (Lysosomal Red Fluorescent Probe), 2',7'-dichlorofluorescein diacetate (DCFH-DA), TUNEL apoptosis assay kit, phosphate-buffered saline (1×PBS), enhanced cell counting kit-8 (CCK-8), DAB horseradish peroxidase color development kit, Ki67 rabbit polyclonal antibody, phospho-EGFR (Ser695) rabbit monoclonal antibody, HRP-labeled goat anti-rat IgG(H+L), HRP-labeled goat anti-rabbit IgG(H+L), Alexa Fluor 488-labeled goat anti-rabbit IgG(H+L), and Alexa Fluor 647-labeled goat anti-mouse IgG(H+L) were purchased from Beyotime Biotechnology (China). All the cell lines were obtained from the Type Culture Collection of the Chinese Academy of Sciences (Shanghai, China). Female BALB/c mice (8 weeks old, ~20 g) were from Hunan SJA Laboratory Animal Co. And maintained under controlled conditions of temperature (22-24 °C) and humidity (40-60 %) with free access to water and standard feed. 7-AAD Viability

Staining Solution, FITC anti-mouse CD80, PE anti-mouse CD86, APC anti-mouse CD11c, PE anti-mouse CD8a, FITC anti-mouse CD4, and APC-Cy7 anti-mouse CD3 antibodies were purchased from BioLegend. Mouse TNF- $\alpha$ , IFN- $\gamma$ , and IL-6 were part of ELISA kits obtained from mlbio (China). Anti-HMGB1, anti-CRT, anti- $\gamma$ H2AX, anti-Areg, anti-Phospho-Egfr (Anti-P-Egfr, Tyr1068), anti-Egfr, anti-Bcl-2, anti-Caspase3, and anti- $\beta$ -actin were acquired from Proteintech (Wuhan, China). Milli-Q water was prepared using a water purification system (Millipore, Burlington, MA, USA) and used in all the tests. All mouse experimental protocols were approved by the Laboratory Animal Ethics Committee.

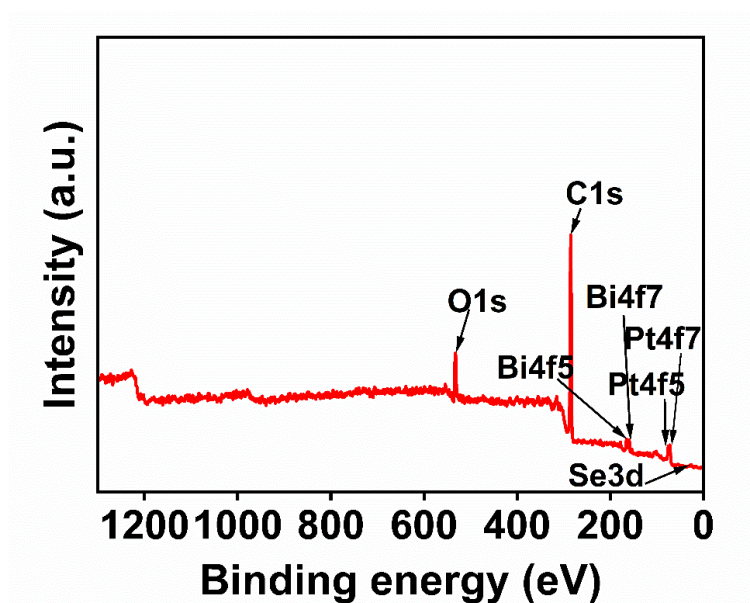
### **Characterization**

TEM images of the PBR nanoassembly and high-angle annular dark-field scanning transmission electron microscopy images were obtained using a FEI TALOS G2 high-resolution transmission electron microscope operating at 200 kV. The  $\zeta$ -potential and size of the PBR were measured using a Malvern Zetasizer NANO ZS90. FTIR spectroscopy characteristics were recorded using a Thermo Fisher Scientific Nicolet iS20 FT-IR spectrometer. Elemental analysis was performed using an iCAP TQ system (Thermo Fisher Scientific). Moreover, X-ray photoelectron spectroscopy was performed using a Thermo Fisher Scientific ESCALAB 250Xi spectrometer. The crystalline morphology of the nanostructures was characterized through XRD using an X-ray diffractometer (XRD-7000; Shimadzu, Japan) with Cu K- $\alpha$  radiation ( $\lambda = 1.5406$  Å). Ultraviolet-visible absorbance measurements were performed using a UV-3600 UV-visible spectrophotometer (Shimadzu). The X-ray parameters used were: 160 kVp and 25 mA, with a 0.3 mm copper filter. BS and B were analyzed by thermogravimetric analysis.

## Supplementary data



**Figure S1.** High-resolution transmission electron microscopy (TEM) image of  $\text{Bi}_2\text{Se}_3$ .



**Figure S2.** XPS pattern of full spectrum of PBR.

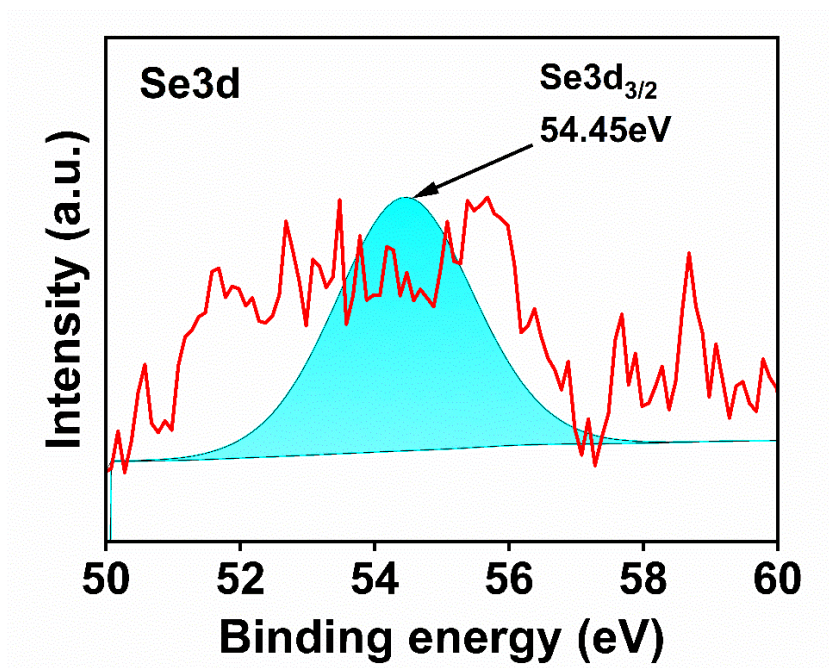


Figure S3. XPS spectrum spectra of Se3d in PBR.

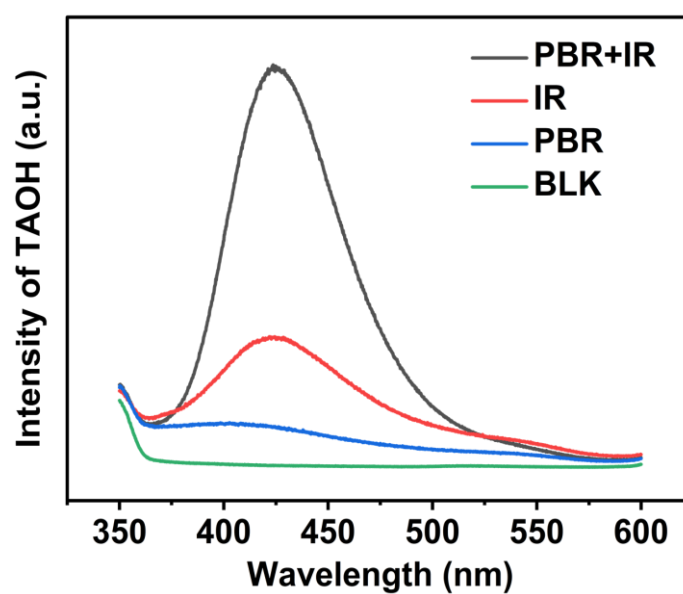
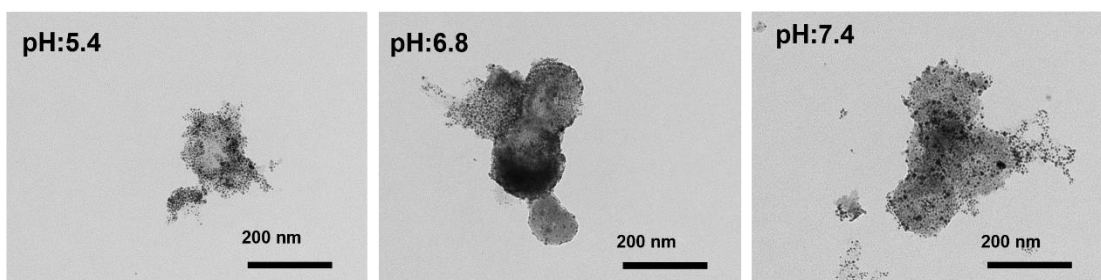
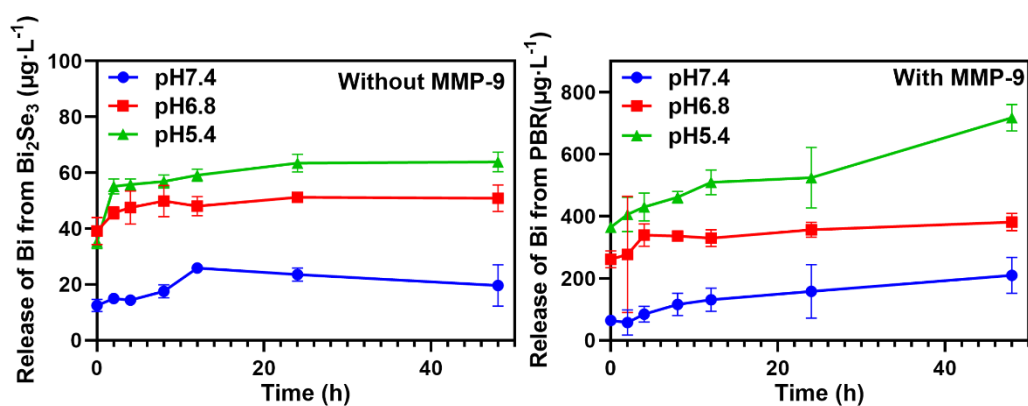


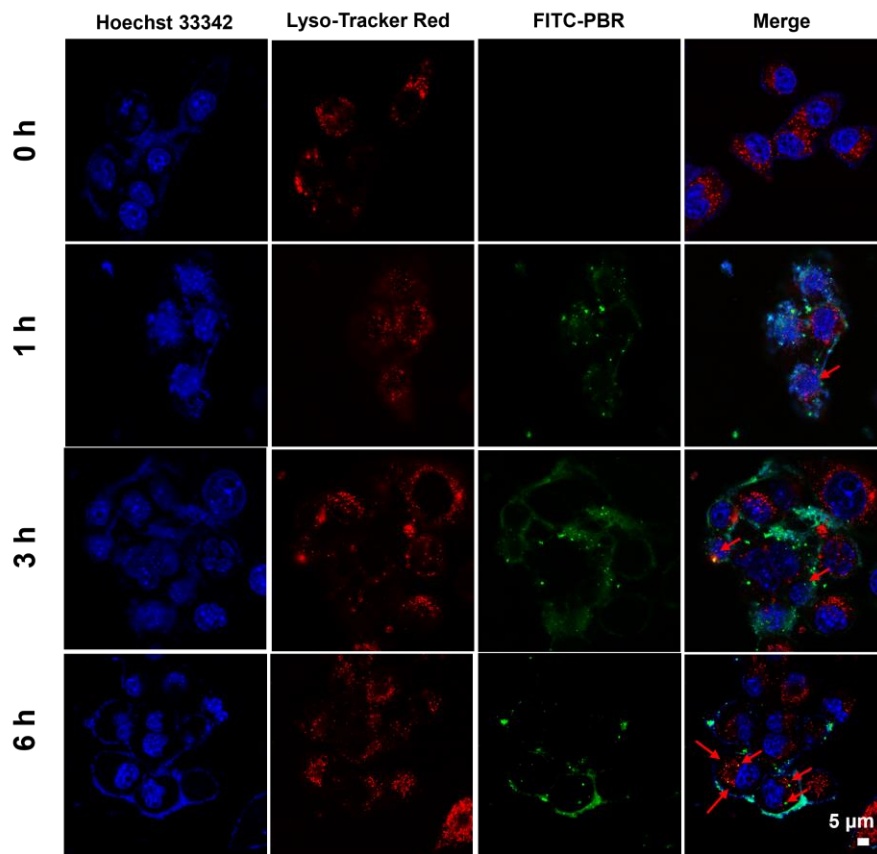
Figure S4. TPA as a tracing agent for  $\bullet\text{OH}$  generated from PBR in the presence of irradiation (4 Gy).



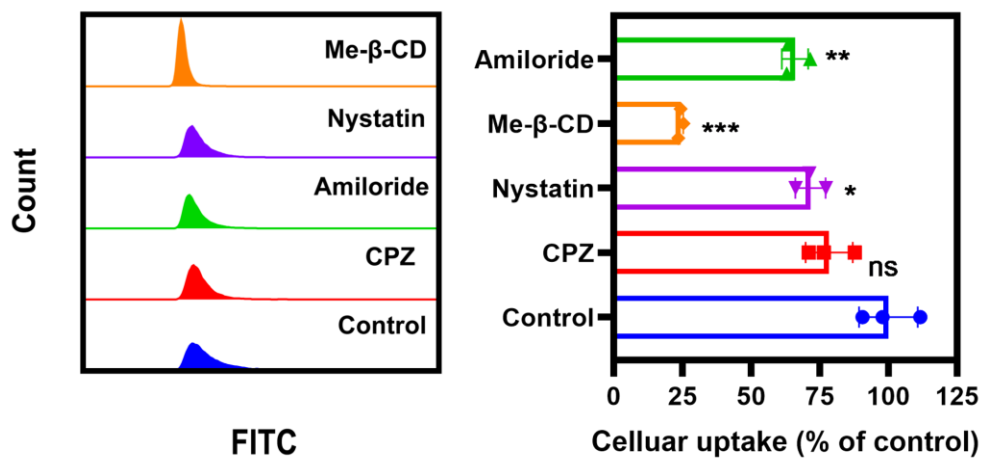
**Figure S5.** *In vitro* simulated biodegradation of PBR in Phosphate buffer solution of different pH (pH=5.4, 6.8, and 7.4) with Cathepsin B.



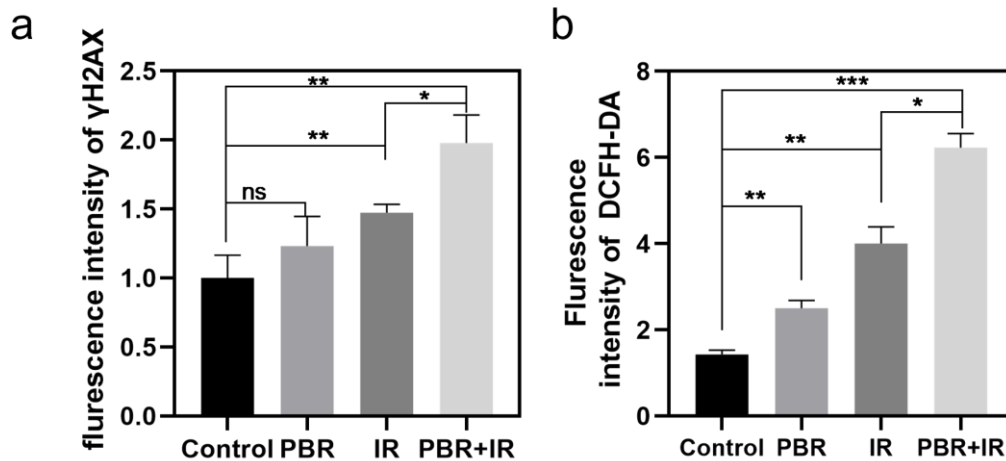
**Figure S6.** The release of Bi element from Bi<sub>2</sub>Se<sub>3</sub> and PBR at different pH values detected by ICP-MS.



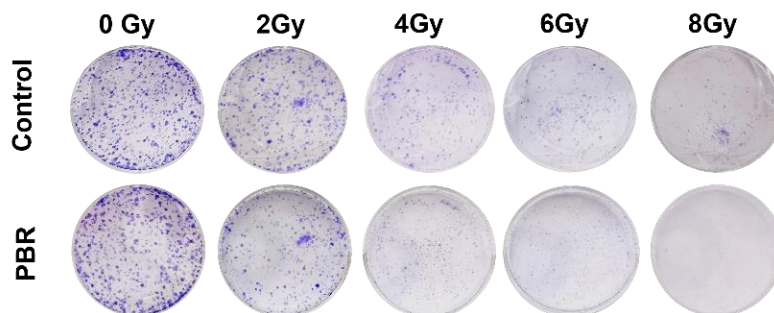
**Figure S7.** Confocal images showing the subcellular localization of PBR after various incubation time.



**Figure S8.** Flow cytometry analysis of cellular uptake of PBR in 4T1 cells after treated with different endocytosis inhibitors for 6h.

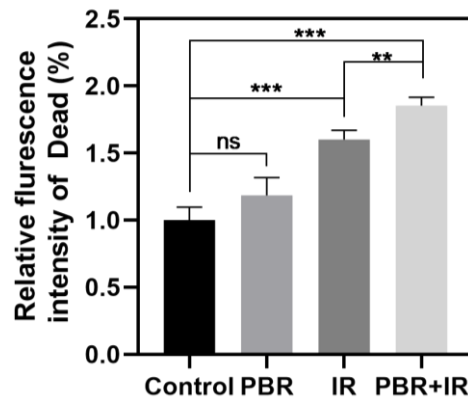
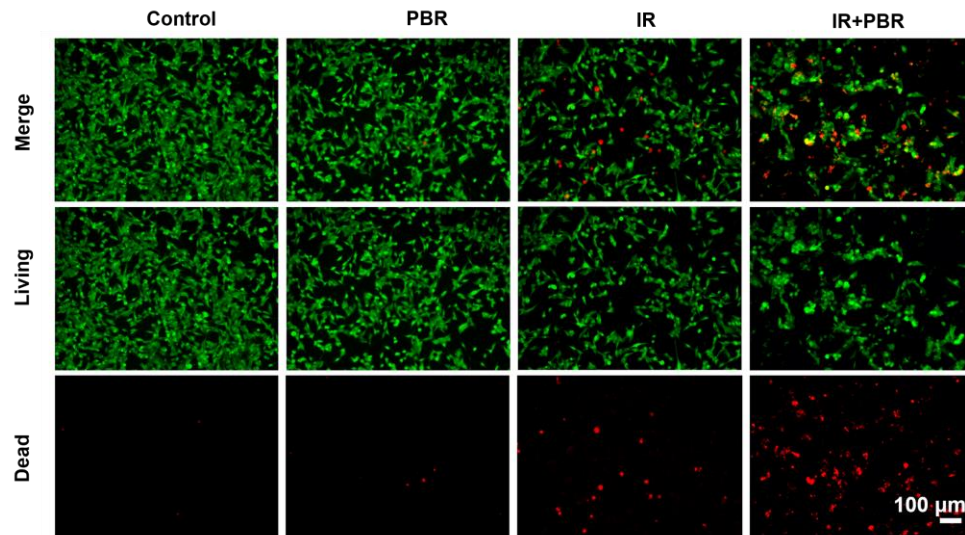


**Figure S9** Fluorescence intensity of (a)  $\gamma$ H2AX, (b) DCFH-DA. Statistical analysis was performed using a *t*-test. Data are presented as the mean  $\pm$  SD. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$  and \*\*\*\* $P < 0.0001$ .

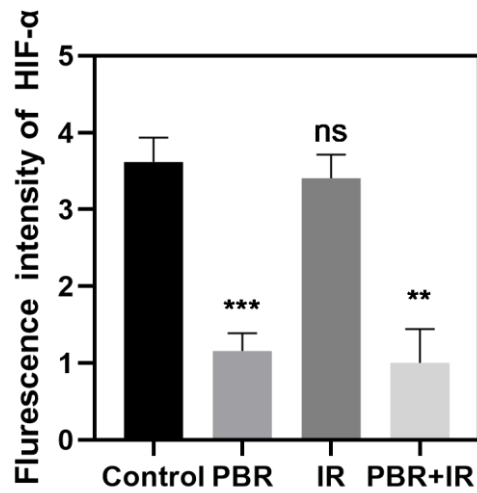


**Figure S10.** Colony formation of 4T1 cells incubated with or without PBR ( $25 \mu\text{g}\cdot\text{mL}^{-1}$ ) and different irradiation doses ( $n = 5$ ).

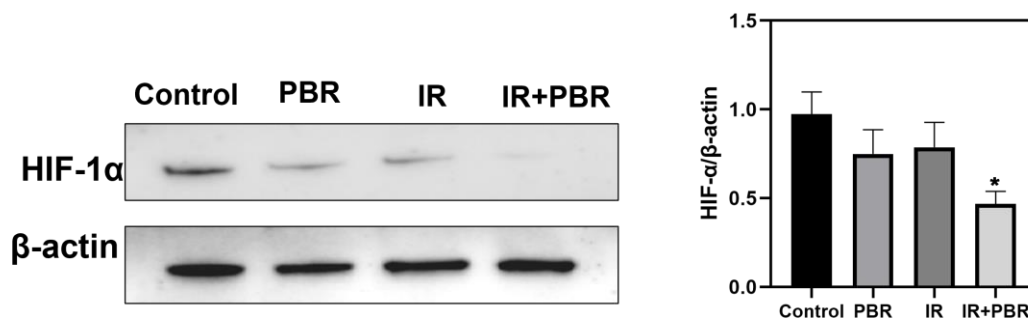




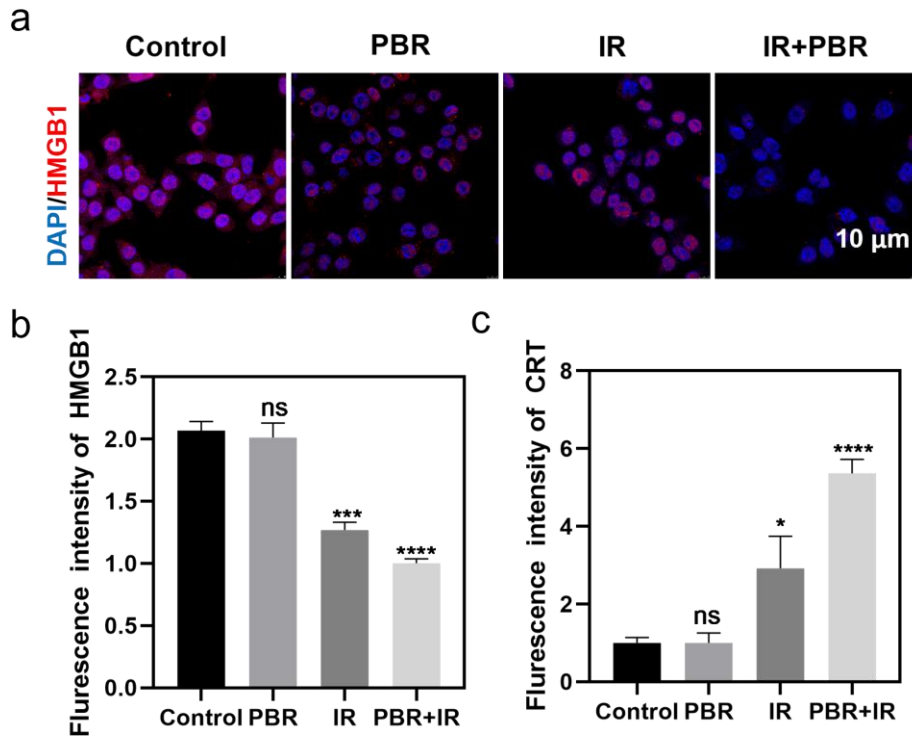
**Figure S11.** Cell death of 4T1 cells after PBR treatment with or without ionizing radiation (Ionizing radiation: 4 Gy, PBR incubation time: 48 h). Statistical analysis was performed using a *t*-test. Data are presented as the mean  $\pm$  SD. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ .



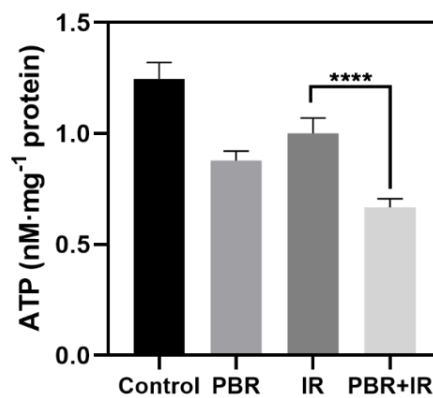
**Figure S12.** Fluorescence intensity of HIF-1 $\alpha$ . Statistical analysis was performed using a *t*-test. Data are presented as the mean  $\pm$  SD. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ .



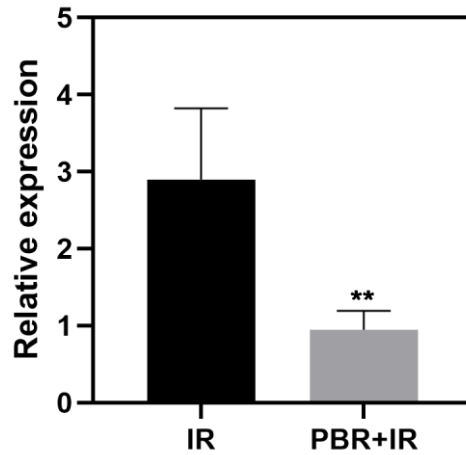
**Figure S13.** Western blot analysis and semi-quantitative analysis evaluating changes in the expression of HIF-1 $\alpha$  in 4T1 cells following different treatments. Statistical analysis was performed using a *t*-test. Data are presented as the mean  $\pm$  SD. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ .



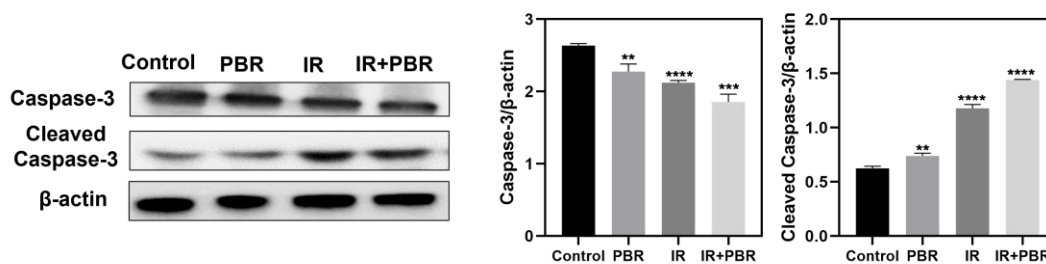
**Figure S14.** Fluorescence intensity of HGB1 (a-b) and CRT (c). Statistical analysis was performed using a *t*-test. Data are presented as the mean  $\pm$  SD. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$  and \*\*\*\* $P < 0.0001$ .



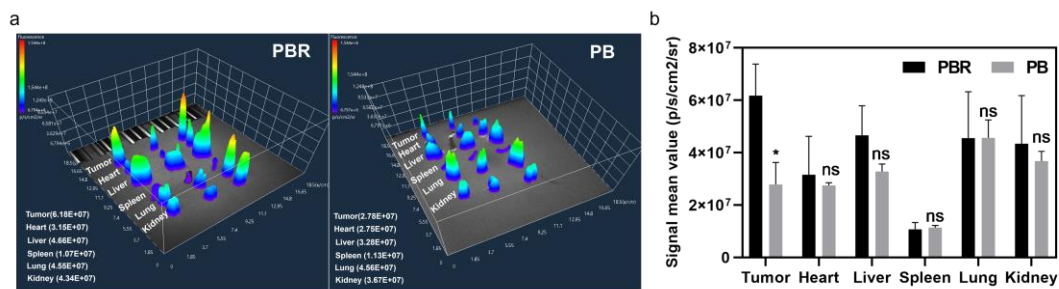
**Figure S15.** Intracellular ATP level changes after different treatments. Statistical analysis was performed using a *t*-test. Data are presented as the mean  $\pm$  SD. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$  and \*\*\*\* $P < 0.0001$ .



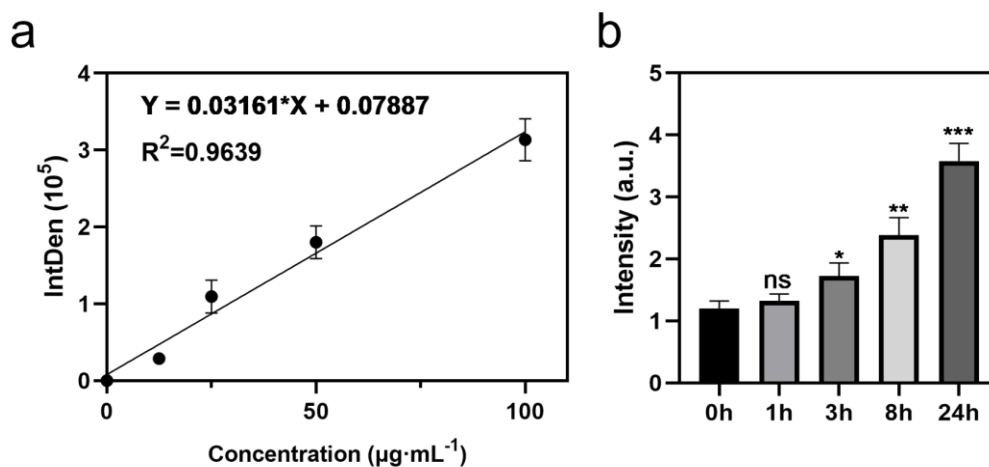
**Figure S16.** Transcription level of Areg was assessed by using real-time quantitative PCR (qPCR). Statistical analysis was performed using a *t*-test. Data are presented as the mean  $\pm$  SD. \**P* < 0.05, \*\**P* < 0.01, \*\*\**P* < 0.001.



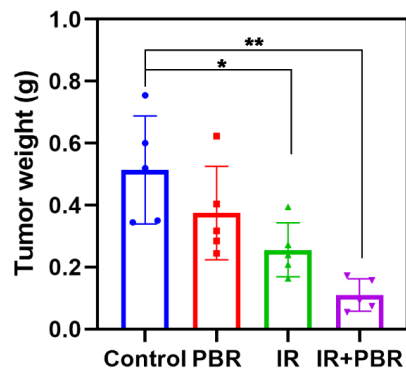
**Figure S17.** Western blot analysis and semi-quantitative analysis evaluating changes in the expression of Caspase-3 and Cleaved Caspase-3 in 4T1 cells following different treatments. Statistical analysis was performed using a *t*-test. Data are presented as the mean  $\pm$  SD. \**P* < 0.05, \*\**P* < 0.01, \*\*\**P* < 0.001 and \*\*\*\**P* < 0.0001.



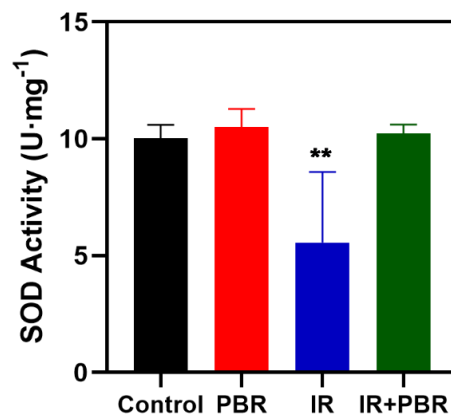
**Figure S18.** Fluorescence imaging of 4T1 tumor-bearing mice injected with IR783 labeled PBR or PB ( $20 \text{ mg} \cdot \text{k}^{-1}$ ), and fluorescence distribution in excised organs and tumors 24 h after injection. Statistical analysis was performed using a *t*-test. Data are presented as the mean  $\pm$  SD. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ .



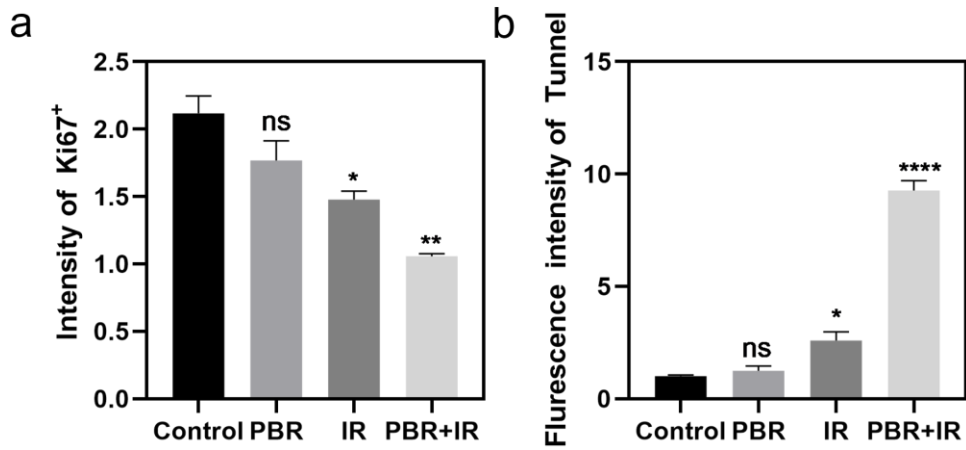
**Figure S19.** (a) The relationship between the photoacoustic imaging signal intensity and concentration of PBR. (b) Semi quantitative statistics of photoacoustic imaging signal intensity in tumor areas. Statistical analysis was performed using a *t*-test. Data are presented as the mean  $\pm$  SD. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ .



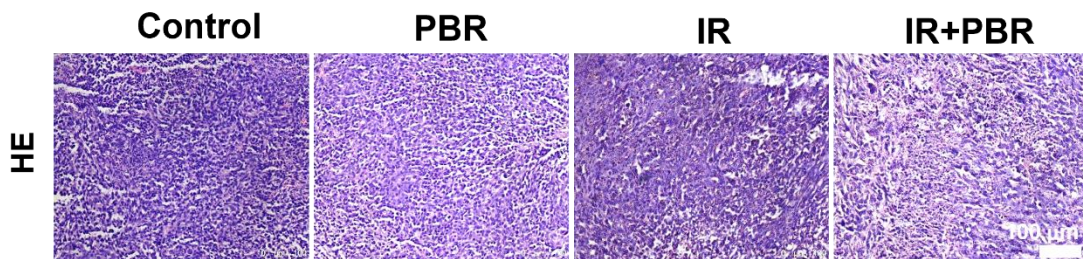
**Figure S20.** Weight of dissected solid tumors after two weeks post-treatment. Statistical analysis was performed using a *t*-test. Data are presented as the mean  $\pm$  SD. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ .



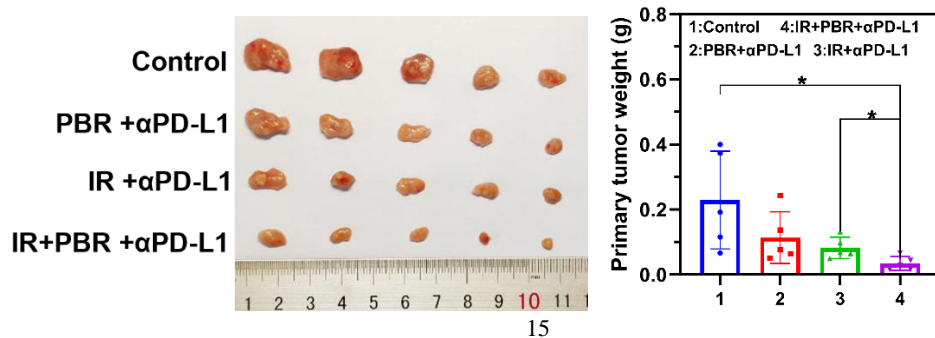
**Figure S21.** Serum levels of superoxide dismutase (SOD) activity in mouse after two weeks post-treatment. Statistical analysis was performed using a *t*-test. Data are presented as the mean  $\pm$  SD. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ .



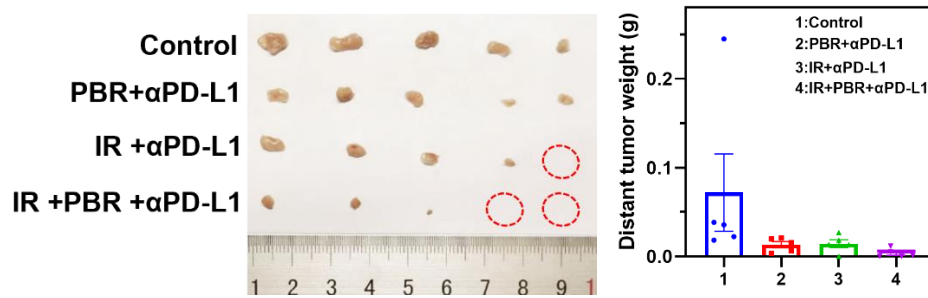
**Figure S22.** The Intensity of tumor sections by Ki67 immunohistochemistry (a) and TUNEL immunofluorescence staining (b) Statistical analysis was performed using a *t*-test. Data are presented as the mean  $\pm$  SD. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$  and \*\*\*\* $P < 0.0001$ .



**Figure S23.** Histological analysis of tumor sections via Hematoxylin and Eosin (H&E) staining on day 14.

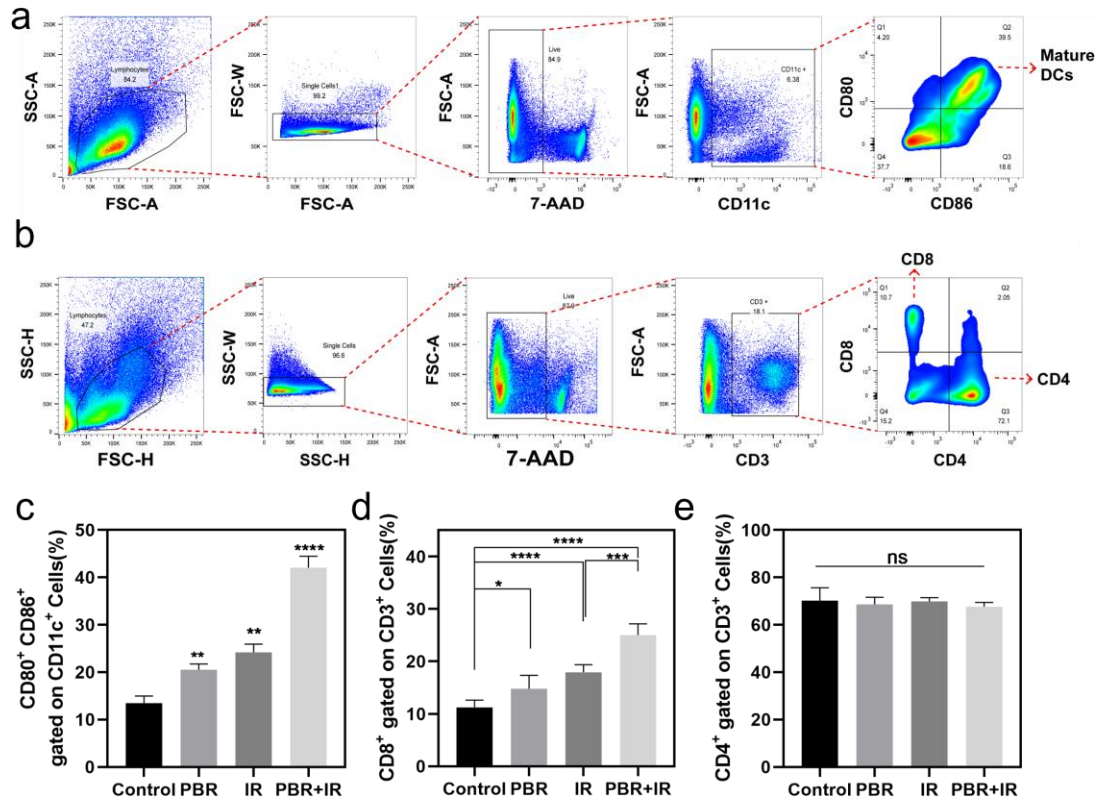


**Figure S24.** Graphs and weight of dissected primary solid tumor on day 14 after two weeks post-treatments. Statistical analysis was performed using a *t*-test. Data are presented as the mean  $\pm$  SD. \**P* < 0.05, \*\**P* < 0.01, \*\*\**P* < 0.001.

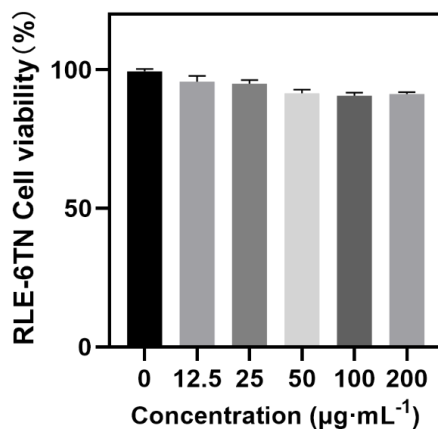


**Figure S25.** Graphs and weight of dissected distant solid tumor on day 14 after two weeks post-treatments. Statistical analysis was performed using a *t*-test. Data are presented as the mean  $\pm$  SD. \**P* < 0.05, \*\**P* < 0.01, \*\*\**P* < 0.001.

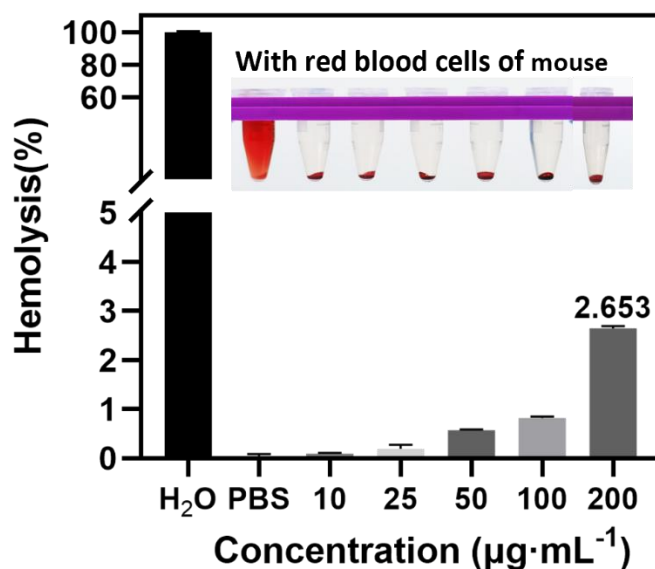




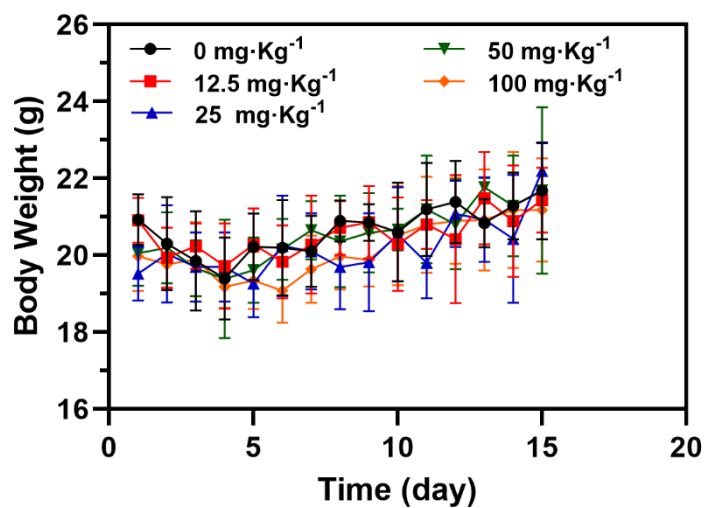
**Figure S26.** Representative FCM images of DC cells and T cells. (a) Gating strategies of CD80<sup>+</sup> and CD86<sup>+</sup> T cells in lymph tissues. (b) Gating strategies of CD3<sup>+</sup> and CD4<sup>+</sup> T cells in spleen tissues. CD80<sup>+</sup> and CD86<sup>+</sup> T cells of (c) lymph, and CD8<sup>+</sup> and CD4<sup>+</sup> T cells of (d-e) spleen tissues in the Control, PBR, IR, PBR +IR groups detected by FCM (n = 5 mice). Statistical analysis was performed using a *t*-test. Data are presented as the mean ± SD. \**P* < 0.05, \*\**P* < 0.01, \*\*\**P* < 0.001 and \*\*\*\**P* < 0.0001.



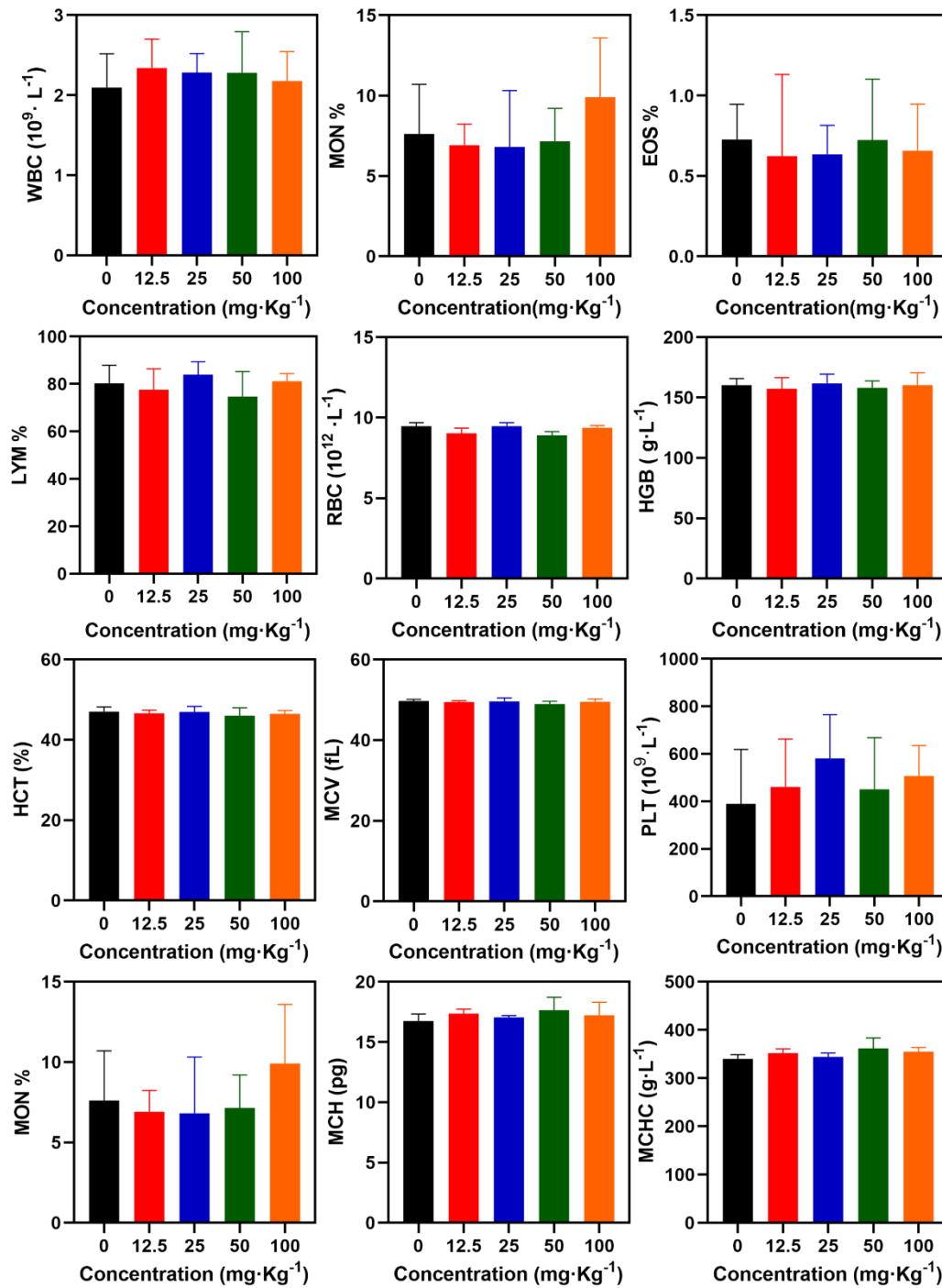
**Figure S27.** Viability of RLE-6TN cells after incubation with various concentrations of PBR for 48h.



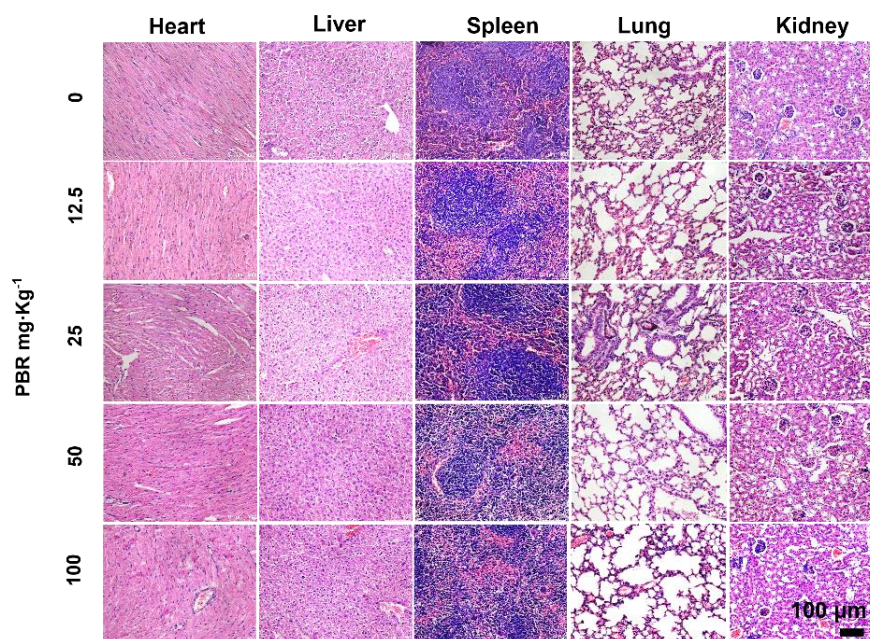
**Figure S28.** Hemolytic rate of erythrocytes after incubation with PBR at various concentrations at 37 °C for 3 h. DI water served as the positive control (100% of hemolysis rate). Inset: photograph of the centrifuge tubes containing various samples.



**Figure S29.** Variation of BALB/c mouse body weight after intravenous injection with PBR with different dosages during the treatment period of 15 days.



**Figure S30.** Primary indicators of blood routine test on day 15 after BALB/c mice being intravenously injected with PBR at different dosages.



**Figure S31.** H&E staining of major organs (heart, liver, spleen, lung and kidney) harvested from BALB/c mice on day 15 after intravenous injection of PBR at different dosages.

**Table S1.** Differential gene statistics table

compare	all	up	down	shold
IR+PBR vs IR	1135	618	517	$ \log_2\text{FoldChange}  > 1 \& \text{pvalue} < 0.05$

**Table S2.** Reactome Pathway Enrichment Analysis Results Table (top minimum 5

padj PathWays)

Description	GeneRatio	BgRatio	pvalue	p.adjust	qvalue	geneNAME	Count
GRB2 events in EGFR signaling	4/76	12/8846	2.36E-06	0.000793	0.000719	Egf/Areg/Btc/Hbegf	4
SHC1 events in EGFR signaling	4/76	13/8846	3.39E-06	0.000793	0.000719	Egf/Areg/Btc/Hbegf	4
EGFR downregulation	5/76	30/8846	4.94E-06	0.000793	0.000719	Egf/Areg/Btc/Rps27a/Hbegf	5
GAB1 signalosome	4/76	17/8846	1.10E-05	0.001326	0.001201	Egf/Areg/Btc/Hbegf	4
Signaling by EGFR	5/76	45/8846	3.83E-05	0.003691	0.003345	Egf/Areg/Btc/Rps27a/Hbegf	5