

Extended Data

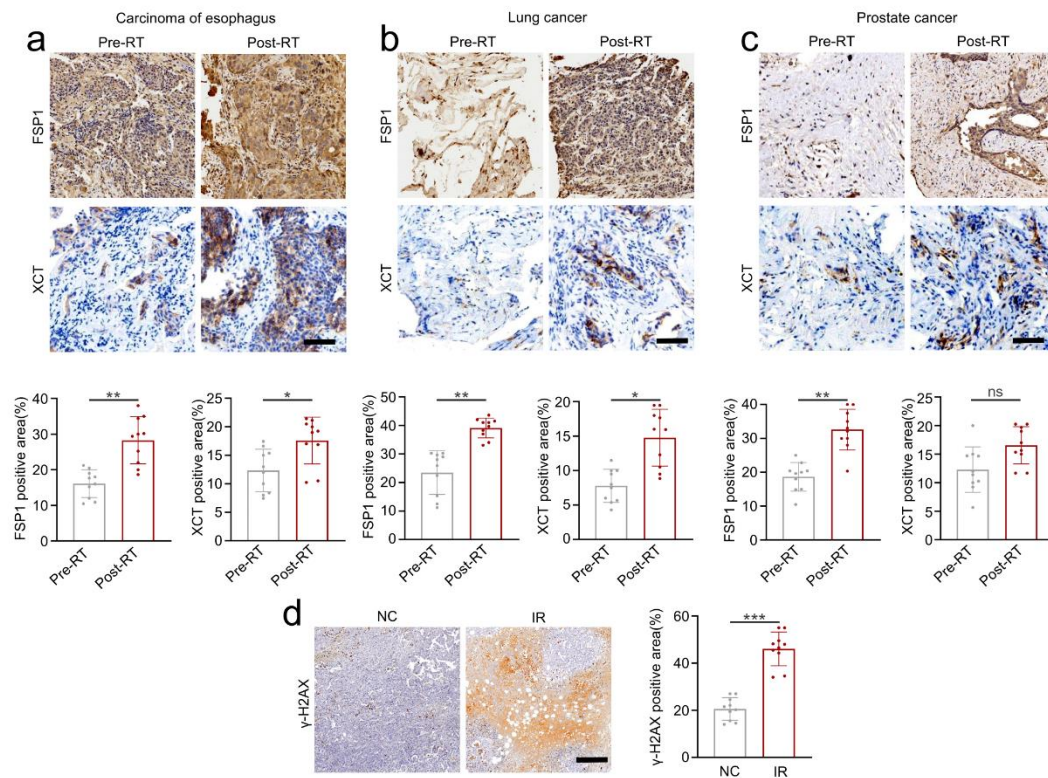


Figure 1 | Expression of ferroptosis-related genes across malignancies post-radiotherapy

a-c Representative IHC staining images (scale bar: 100 μ m) of FSP1 and xCT in pan-cancer patients before and after radiotherapy. **d** IHC staining showing γ -H2AX expression in tumor tissues of a mouse PCa subcutaneous xenograft model pre- and post-irradiation (representative images; scale bar: 100 μ m).

Data: Mean $\pm$ SD; ns = not significant, * p < 0.05, ** p < 0.01, *** p < 0.001.

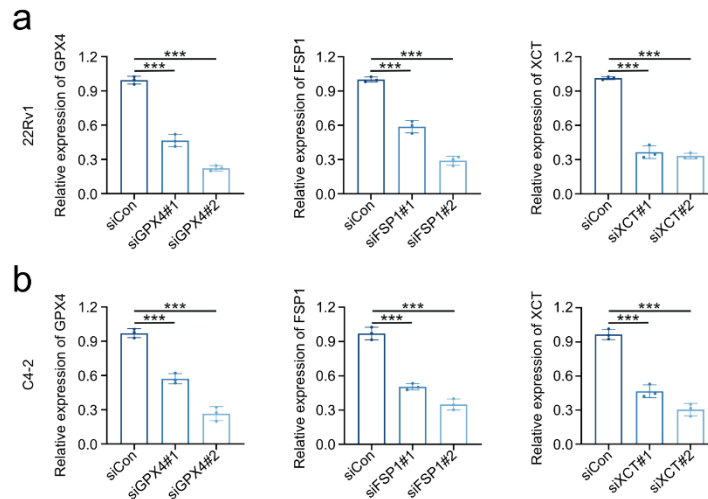


Figure 2 | Screening optimal FINs for radiosensitization

a-b Validation of the knockdown efficiency of two siRNA pairs targeting multiple ferroptosis-related genes using qPCR.

Data: Mean $\pm$ SD; ***p < 0.001.

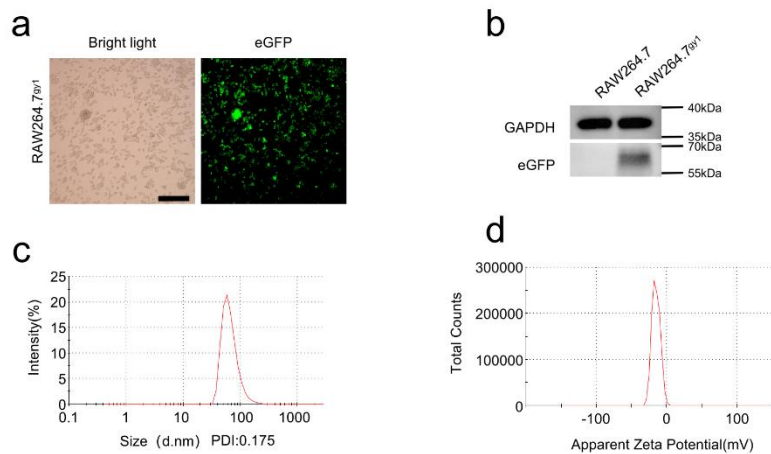


Fig. 3 | Fabrication and functional characterization of Au-MOF/RSL3@M-gy1

a. Fluorescence microscopy of RAW264.7 macrophages expressing PSMA-targeting gy1 antibody (green: eGFP signal; scale bar: 100 μ m). **b.** Western blot validation of eGFP-tagged gy1 expression in engineered RAW264.7 cells. **c.** Hydrodynamic size distribution of Au-MOF nanoparticles by DLS. **d.** Zeta potential analysis of Au-MOF.

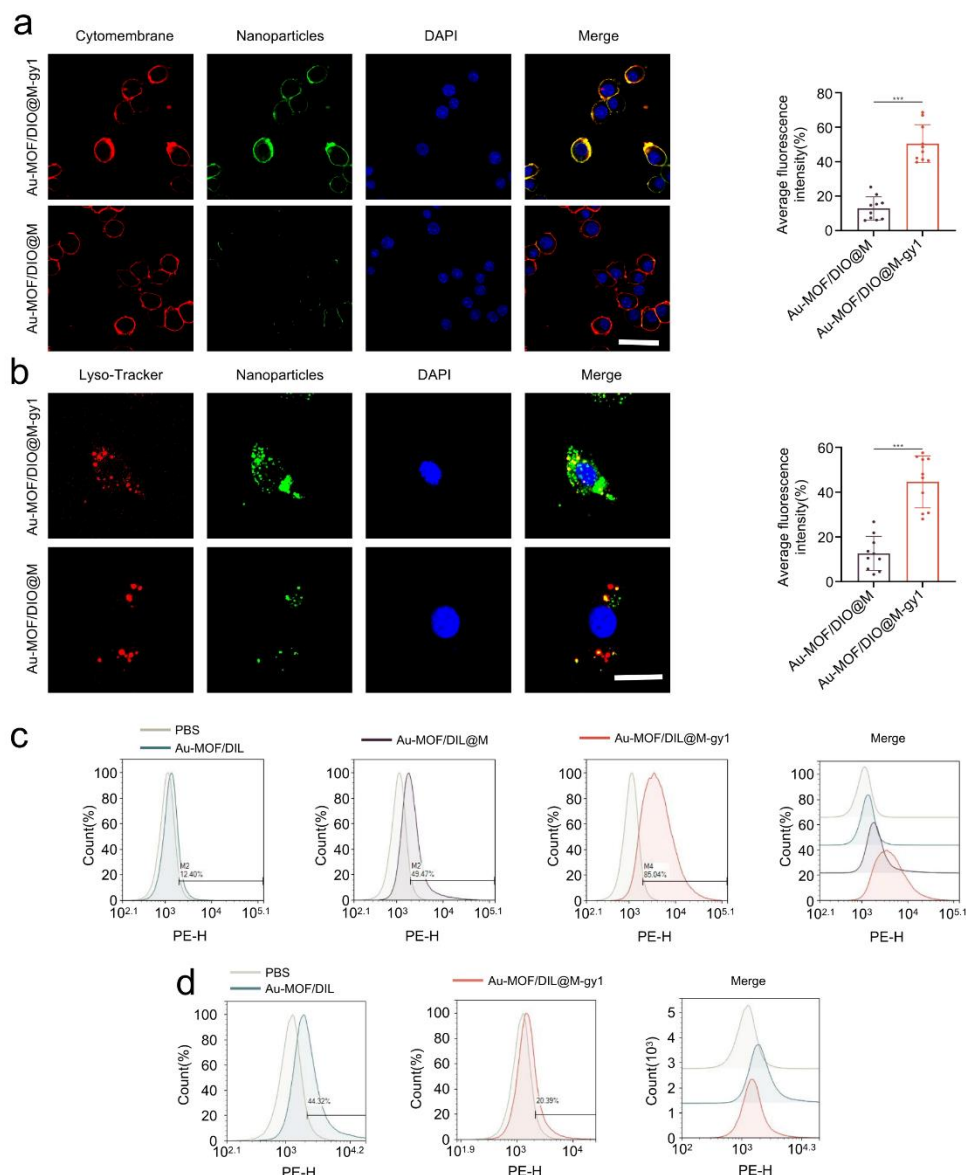


Fig. 4 | The targeting capability of Au-MOF/RSL3@M-gy1

a-c) Evaluation of targeting specificity in C4-2 cells: **a.** Immunofluorescence co-localization analysis of Au-MOF@M-gy1 binding to target cells (Nanoparticles: green; scale bars: 40 μ m). **b.** Intracellular internalization efficiency of Au-MOF@M-gy1 visualized by confocal microscopy (Nanoparticles: green; scale bars: 20 μ m). **c.** Quantitative flow cytometry analysis of cellular uptake. **d.** Phagocytic efficiency of target cells (macrophages) towards Au-MOF@M-gy1 analyzed by flow cytometry.

Data: Mean $\pm$ SD; ***p < 0.001.

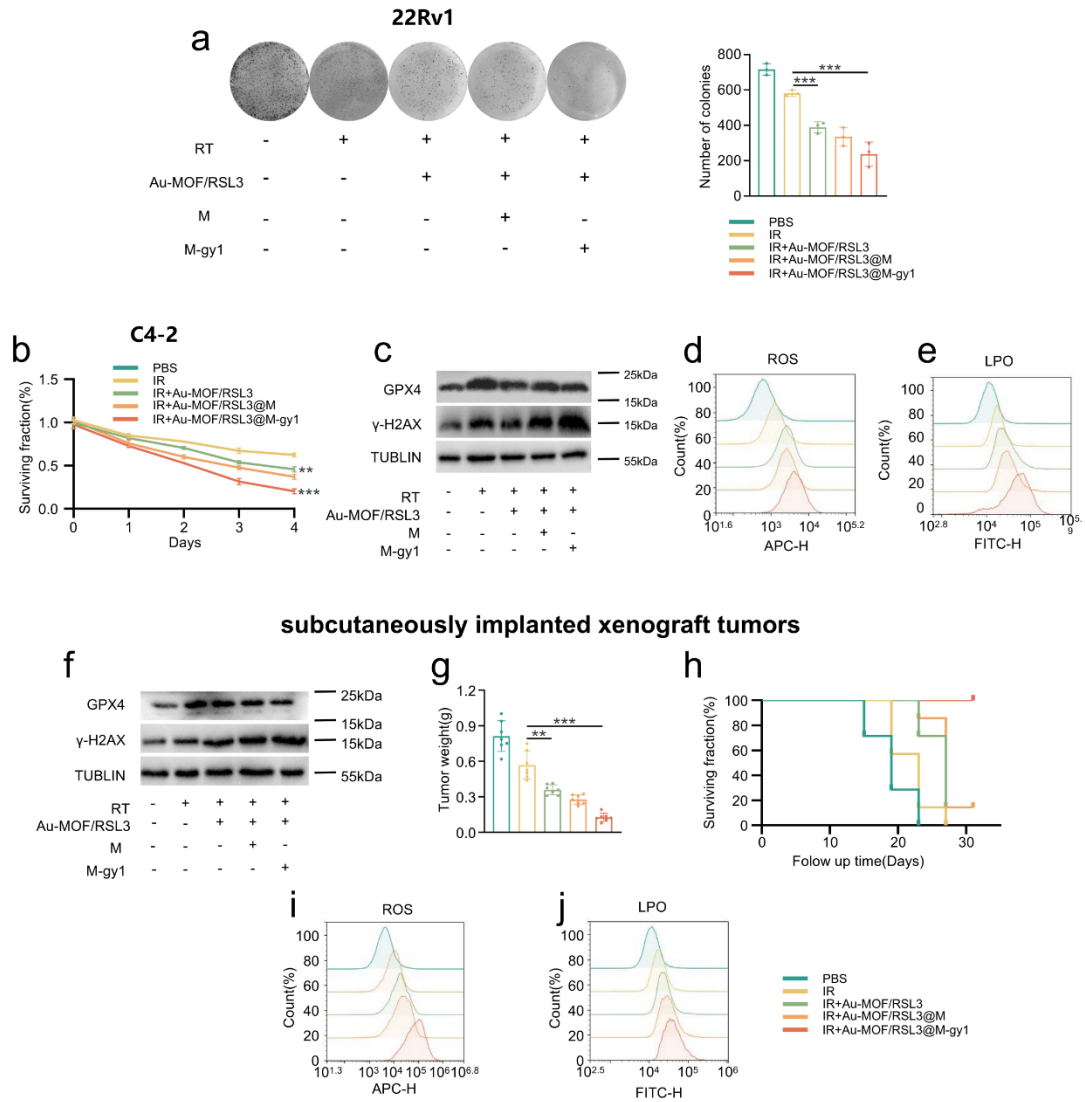


Fig. 5 | Evaluation of the radiosensitization effect of Au-MOF/RSL3@M-gy1.

a. Clonogenic assay demonstrating radiosensitization efficacy in 22Rv1 prostate cancer cells.

b-e). Validation in C4-2 cells: **b.** CCK-8 viability assay under X-ray irradiation (2 Gy). **c.** Western blot analysis of GPX4 and γ-H2AX. **d-e.** Flow cytometry quantification of ROS and LPO.

f-j). In vivo assessment in xenografted mice: **f.** Western blot of tumor lysates for GPX4、γ-H2AX expression. **g.** Tumor weight comparison across treatment groups. **h.** Survival curves of mice in different treatment groups. **i-j.** Flow cytometry analysis of tumor tissue ROS、LPO levels.

Data: Mean ± SD; **p < 0.01, ***p < 0.001.

c-d) Validation in C4-2 cells: **c.** Live/dead cell staining assay under radiation treatment (representative images; scale bar: 100 μ m). **d.** Immunofluorescence analysis of γ -H2AX expression (representative images; scale bar: 40 μ m).
 Data: Mean $\pm$ SD; **p < 0.01, ***p < 0.001.

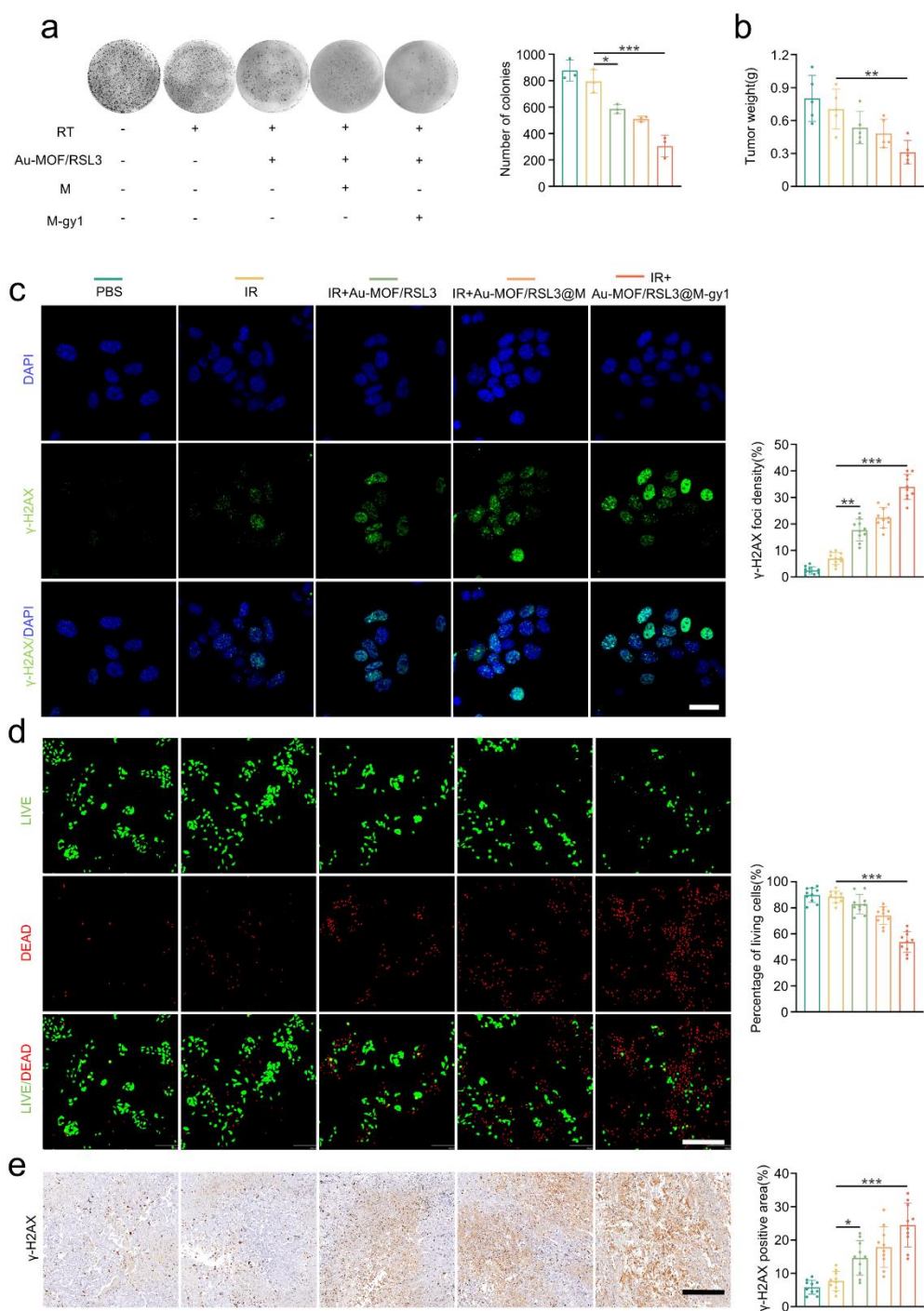


Fig. 7 | Radiosensitization efficacy validation of Au-MOF/RSL3@M-gy1 in radioresistant prostate cancer models
a. Clonogenic assay quantifying radiosensitization effects in radiation-resistant 22Rv1-RR

cells. **b.** Subcutaneous tumor mass quantification in radioresistant murine xenografts (n = 5). **c.** Immunofluorescence co-localization analysis of γ -H2AX foci across treatment groups (representative images; scale bar: 40 μ m). **d.** Live/dead cell staining demonstrating the radiosensitization of Au-MOF/RSL3@M-gy1 in 22Rv1-RR cells (representative images; scale bar: 100 μ m). **e.** Immunohistochemical staining of γ -H2AX expression in radioresistant tumor tissues (representative images; scale bar: 100 μ m).
Data: Mean $\pm$ SD; *P < 0.05, **P < 0.01, ***P < 0.001.

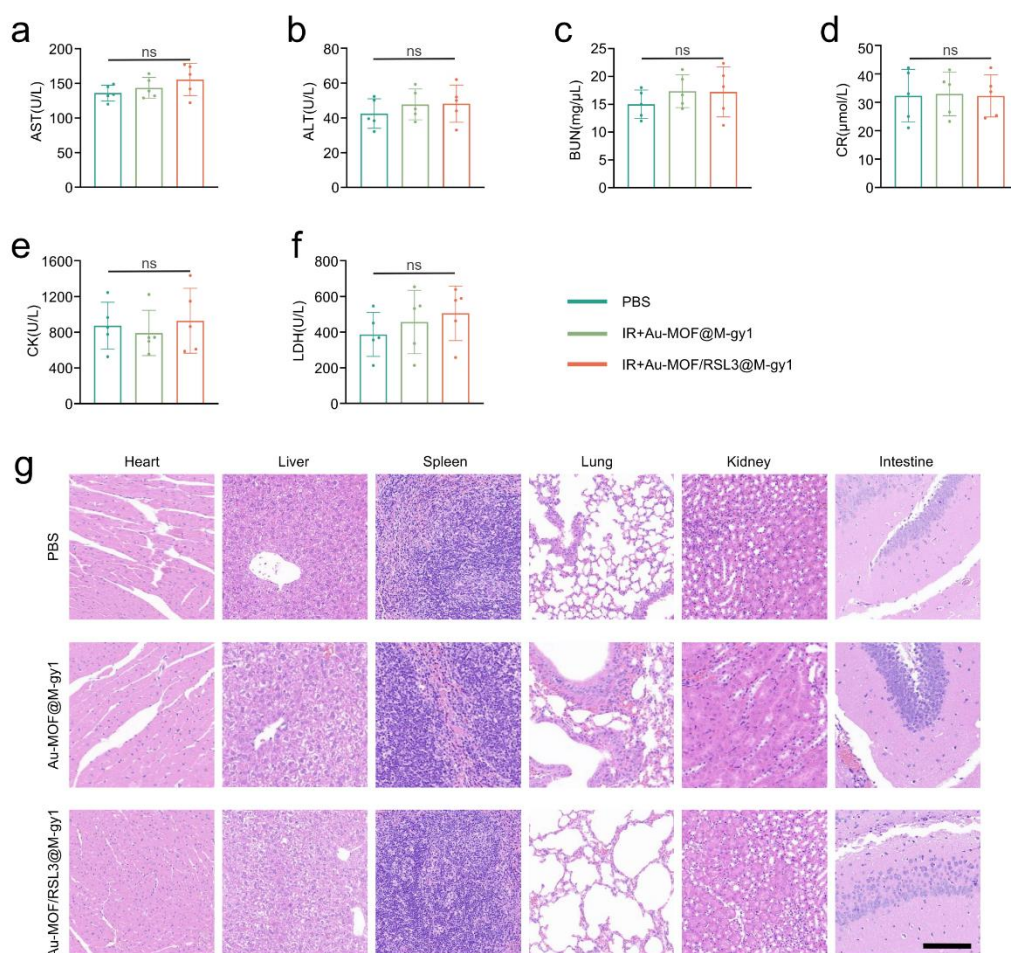


Fig. 8 | Biosafety evaluation of Au-MOF/RSL3@M-gy1

a-f). Serum biochemical profiling: **a.** Aspartate aminotransferase (AST) levels. **b.** Alanine aminotransferase (ALT) levels. **c.** Blood urea nitrogen (BUN) levels. **d.** Creatinine (CR) levels. **e.** Creatine kinase (CK) levels. **f.** Lactate dehydrogenase (LDH) levels. **g.** H&E staining of major organ paraffin sections (representative images; scale bar: 100 μ m).
Data: Mean $\pm$ SD; ns = not significant.