## **Additional Tables**

Gene	Sequence
GAPDH Fp	ACAACTTTGGTATCGTGGAAGG
GAPDH Rp	GCCATCACGCCACAGTTTC
SAMHD1 Fp	CCCAAAGTATTGCTAGACGTGA
SAMHD1 Rp	TGCATTCCATAATCCATGTTG
MRE11 Fp	ATCGGCCTGTCCAGTTTGAAA
MRE11 Rp	TGCCATCTTGATAGTTCACCCAT
CtIP Fp	AGCGTTTGTGGAGCCGTAT
CtIP Rp	GTTGGGTGGAATGTAGCG
RAD50 Fp	TAAGTGTGCAGAAATTGACCGAG
RAD50 Rp	GACGTACCTGCCGAAGTGTT
NBS1 Fp	AGGTGGGGAAGCTAGGTTGAT
NBS1 Rp	CACCGCCAATCCAATTTCTGC
IFI16 Fp	AGACTGAAGACTGAACCTGAAGA
IFI16 Rp	GAACCCATTGCGGCAAACATA
STING Fp	AGCATTACAACAACCTGCTACG
STING Rp	GTTGGGGTCAGCCATACTCAG
IFNβ Fp	AACTTGCTTGGATTCCTACAAAG
IFNβ Rp	TATTCAAGCCTCCCATTCAATTG
CCL5 Fp	CGCTGTCATCCTCATTGCTA
CCL5 Rp	CCAGACTTGCTGTCCCTCTC

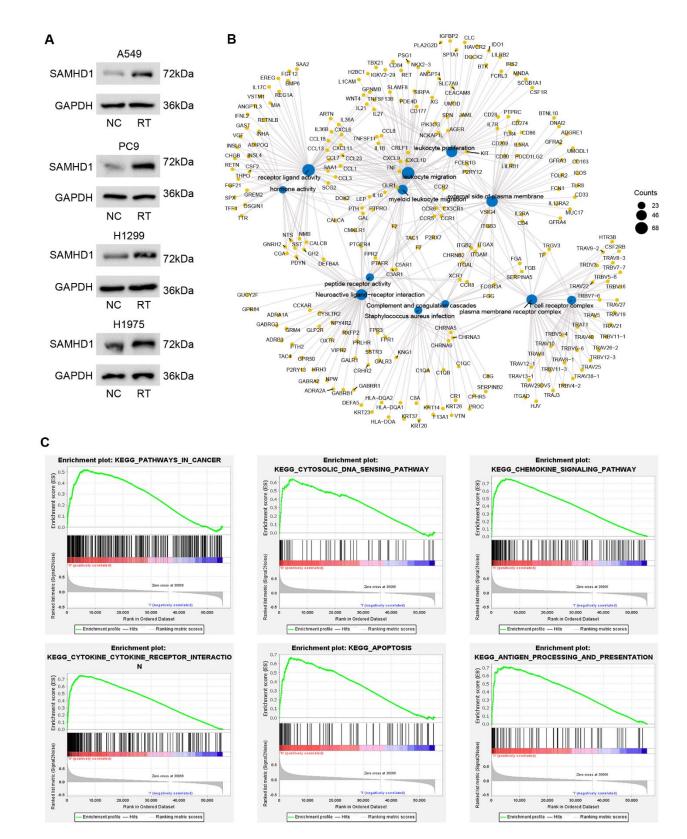
Table S1. Primer sequences used for amplification and the targeting siRNA sequences.

CXCL10 FpCTGTACGCTGTACCTGCATCACXCL10 RpTTCTTGATGGCCTTCGATTCActin FpGGCTGTATTCCCCTCCATCGActin RPCCAGTTGGTAACAATGCCATGTSamhd1 FpGAGAATCGTGGTTTCCGAGAGSamhd1 RpCTCCAAGGAACTTACTCCCAGAIfhβ FpCAGCTCCAAGAAAGGACGAACIfhβ RpGGCAGTGTAACTCTTCTGCATCcl5 FpGCTGCTTTGCCTACCTCCCCxel10 FpCCAAGTGCTGCAGCGTCATTTCCxel10 RpGGCTCGCAGGGATGATTTCAAMhc-2 FpTCAGTCGCAGACGGTGTTTATMhc-2 RpGGGGGCTGGAATCTCAGGTCd86 FpTTGTTTCCGTGGAGACGCAAGsiSAMHD1-1CCUCGUCCGAAUCAUUGAUTTsiSAMHD1-2GCAGAUGACUACAUAGAGATTsiSAMHD1-3GCAGAUGACUACAUAGAGATTsiSAMHD1-3GCAGAUGACUACAUAGAGATTsiSTINGGGUCAUAUUACAUGGAUATT		
Actin FpGGCTGTATTCCCCTCCATCGActin RPCCAGTTGGTAACAATGCCATGTSamhd1 FpGAGAATCGTGGTTTCCGAGAGSamhd1 RpCTCCAAGGAACTTACTCCCAGAIfnβ FpCAGCTCCAAGAAAGGACGAACIfnβ RpGGCAGTGTAACTCTTCTGCATCcl5 FpGCTGCTTTGCCTACCTCCCCcl10 FpCCAAGTGCTGCGTCATTTCCxcl10 RpGGCTGCCAGGGATGATTTCAAMhc-2 FpTCAGTCGCAGACGGTGTTATMhc-2 RpGGGGGCTGGAATCTCAGGTCd86 FpTTGAGCCTTTGTAAATGGGCAsiSAMHD1-1CCUCGUCCGAAUCAUUGAUTTsiSAMHD1-2GCGUAUUUGUGCUAGAGATTsiSAMHD1-3GCAGAUGACUACAUAGAGATTshSamhd1CGATGAGGATCGTCTGGAA	CXCL10 Fp	CTGTACGCTGTACCTGCATCA
Actin RPCCAGTTGGTAACAATGCCATGTSamhdl FpGAGAATCGTGGTTTCCGAGAGSamhdl RpCTCCAAGGAACTTACTCCCAGAIfhβ FpCAGCTCCAAGAAAGGACGAACIfhβ RpGGCAGTGTAACTCTTCGCATCcl5 FpGCTGCTTTGCCTACCTCCCCcl5 RpTCGAGTGACAAACACGACTGCCxcl10 FpCCAAGTGCTGCCGTCATTTCCxcl10 RpGGCTCGCAGGGATGATTCAAMhc-2 FpTCAGTCGCAGGAACGGTGTTATCd86 FpTGTTTCCGTGGAAACGCAAGCd86 RpTTGAGCCTTTGTAAATGGGCAsiSAMHD1-1CCUCGUCCGAAUCAUUGAUTTsiSAMHD1-2GCGUAUUUGUGCUAGAGATTsiSAMHD1-3GCAGAUGACUACAUAGAGATTshSamhd1GGATGAGGATCGTCTGGAA	CXCL10 Rp	TTCTTGATGGCCTTCGATTC
Samhdl FpGAGAATCGTGGTTTCCGAGAGSamhdl RpCTCCAAGGAACTTACTCCCAGAIfnβ FpCAGCTCCAAGAAAGGACGAACIfnβ RpGGCAGTGTAACTCTTCTGCATCcl5 FpGCTGCTTTGCCTACCTCCCCcl5 RpTCGAGTGACAAACACGACTGCCxcl10 FpCCAAGTGCTGCCGTCATTTCAAMhc-2 FpTCAGTCGCAGGATGATTCAAGMhc-2 RpGGGGGCTGGAATCTCAGGTCd86 FpTTGAGCCTTTGTAAATGGGCAsiSAMHD1-1CCUCGUCCGAAUCAUUGAUTTsiSAMHD1-2GCGAUGACUACAUAGAGATTsiSAMHD1-3GCAGAUGACUACAUAGAGATTshSamhdlGGATGAGGATCGTCTGGAA	Actin Fp	GGCTGTATTCCCCTCCATCG
YCTCCAAGGAACTTACTCCCAGASamhdl RpCAGCTCCAAGAAAGGACGAACIfnβ FpCAGCTCCAAGAAAGGACGAACIfnβ RpGGCAGTGTAACTCTTCGCATCcl5 FpGCTGCTTTGCCTACCTCTCCCcl5 RpTCGAGTGACAAACACGACTGCCxcl10 FpCCAAGTGCTGCCGTCATTTTCCxcl10 RpGGCTCGCAGGGATGATTTCAAMhc-2 FpTCAGTCGCAGAGGTGTTTATCd86 FpTGTTTCCGTGGAAACGCAAGGCd86 RpTTGAGCCTTTGTAAATGGGCAsiSAMHD1-1CCUCGUCCGAAUCAUUGAUTTsiSAMHD1-2GCGUAUUUGUGCUAGAGATTsiSAMHD1-3GCAGAUGACUACAUAGAGATTshSamhdlGGATGAGATCGTCTGGAA	Actin RP	CCAGTTGGTAACAATGCCATGT
Ifnβ FpCAGCTCCAAGAAAGGACGAACIfnβ RpGGCAGTGTAACTCTTCTGCATCcl5 FpGCTGCTTTGCCTACCTCCCCcl5 RpTCGAGTGACAAACACGACTGCCxcl10 FpCCAAGTGCTGCCGTCATTTCCxcl10 RpGGCTCGCAGGGATGATTCAAMhc-2 FpTCAGTCGCAGAGACGGTGTTATCd86 FpTGTTTCCGTGGAGACGCAAGcd86 RpTTGAGCCTTTGTAAATGGGCAsiSAMHD1-1CCUCGUCCGAAUCAUUGAUTTsiSAMHD1-2GCGUAUUUGUGCUAGAGAUTTsiSAMHD1-3GCAGAUGACUACAUAGAGATshSamhd1GGATGAGGATCGTCTGGAA	Samhd1 Fp	GAGAATCGTGGTTTCCGAGAG
Imp r pEncorrent function funct	Samhd1 Rp	CTCCAAGGAACTTACTCCCAGA
Cel5 FpGCTGCTTTGCCTACCTCTCCCel5 RpTCGAGTGACAAACACGACTGCCxel10 FpCCAAGTGCTGCCGTCATTTTCCxel10 RpGGCTCGCAGGGATGATTTCAAMhe-2 FpTCAGTCGCAGAGCGTGTTTATMhe-2 RpGGGGGCTGGAATCTCAGGTCd86 FpTGTTTCCGTGGAGACGCAAGCd86 RpTTGAGCCTTTGTAAATGGGCAsiSAMHD1-1CCUCGUCCGAAUCAUUGAUTTsiSAMHD1-2GCGUAUUUGUGCUAGAGAUTTsiSAMHD1-3GCAGAUGACUACAUAGAGATTshSamhd1GGATGAGGATCGTCTGGAA	Ifnβ Fp	CAGCTCCAAGAAAGGACGAAC
Cel5 RpTCGAGTGACAAACACGACTGCCxel10 FpCCAAGTGCTGCCGTCATTTCCxel10 RpGGCTCGCAGGGATGATTTCAAMhe-2 FpTCAGTCGCAGACGGTGTTTATMhe-2 RpGGGGGCTGGAATCTCAGGTCd86 FpTGTTTCCGTGGAGACGCAAGcd86 RpTTGAGCCTTTGTAAATGGGCAsiSAMHD1-1CCUCGUCCGAAUCAUUGAUTTsiSAMHD1-2GCGUAUUUGUGCUAGAGATTsiSAMHD1-3GCAGAUGACUACAUAGAGATTshSamhd1GGATGAGGATCGTCTGGAA	Ifnβ Rp	GGCAGTGTAACTCTTCTGCAT
Cxcl10 FpCCAAGTGCTGCCGTCATTTCCCxcl10 RpGGCTCGCAGGGATGATTTCAAMhc-2 FpTCAGTCGCAGACGGTGTTTATMhc-2 RpGGGGGCTGGAATCTCAGGTCd86 FpTGTTTCCGTGGAGACGCAAGCd86 RpTTGAGCCTTTGTAAATGGGCAsiSAMHD1-1CCUCGUCCGAAUCAUUGAUTTsiSAMHD1-2GCGUAUUUGUGCUAGAGAUTTsiSAMHD1-3GCAGAUGACUACAUAGAGATTshSamhd1GGATGAGGATCGTCTGGAA	Ccl5 Fp	GCTGCTTTGCCTACCTCTCC
Cxcl10 RpGGCTCGCAGGGATGATTTCAAMhc-2 FpTCAGTCGCAGACGGTGTTTATMhc-2 RpGGGGGCTGGAATCTCAGGTCd86 FpTGTTTCCGTGGAGACGCAAGCd86 RpTTGAGCCTTTGTAAATGGGCAsiSAMHD1-1CCUCGUCCGAAUCAUUGAUTTsiSAMHD1-2GCGUAUUUGUGCUAGAGAUTTsiSAMHD1-3GCAGAUGACUACAUAGAGATTshSamhd1GGATGAGGATCGTCTGGAA	Ccl5 Rp	TCGAGTGACAAACACGACTGC
Mhc-2 FpTCAGTCGCAGACGGTGTTTATMhc-2 RpGGGGGCTGGAATCTCAGGTCd86 FpTGTTTCCGTGGAGACGCAAGCd86 RpTTGAGCCTTTGTAAATGGGCAsiSAMHD1-1CCUCGUCCGAAUCAUUGAUTTsiSAMHD1-2GCGUAUUUGUGCUAGAGAUTTsiSAMHD1-3GCAGAUGACUACAUAGAGATTshSamhd1GGATGAGGATCGTCTGGAA	Cxcl10 Fp	CCAAGTGCTGCCGTCATTTTC
Mhc-2 RpGGGGGCTGGAATCTCAGGTCd86 FpTGTTTCCGTGGAGACGCAAGCd86 RpTTGAGCCTTTGTAAATGGGCAsiSAMHD1-1CCUCGUCCGAAUCAUUGAUTTsiSAMHD1-2GCGUAUUUGUGCUAGAGAUTTsiSAMHD1-3GCAGAUGACUACAUAGAGATTshSamhd1GGATGAGGATCGTCTGGAA	Cxcl10 Rp	GGCTCGCAGGGATGATTTCAA
ICd86 FpTGTTTCCGTGGAGACGCAAGCd86 RpTTGAGCCTTTGTAAATGGGCAsiSAMHD1-1CCUCGUCCGAAUCAUUGAUTTsiSAMHD1-2GCGUAUUUGUGCUAGAGAUTTsiSAMHD1-3GCAGAUGACUACAUAGAGATTshSamhd1GGATGAGGATCGTCTGGAA	Mhc-2 Fp	TCAGTCGCAGACGGTGTTTAT
Cd86 RpTTGAGCCTTTGTAAATGGGCAsiSAMHD1-1CCUCGUCCGAAUCAUUGAUTTsiSAMHD1-2GCGUAUUUGUGCUAGAGAUTTsiSAMHD1-3GCAGAUGACUACAUAGAGATTshSamhd1GGATGAGGATCGTCTGGAA	Mhc-2 Rp	GGGGGCTGGAATCTCAGGT
siSAMHD1-1CCUCGUCCGAAUCAUUGAUTTsiSAMHD1-2GCGUAUUUGUGCUAGAGAUTTsiSAMHD1-3GCAGAUGACUACAUAGAGATTshSamhd1GGATGAGGATCGTCTGGAA	Cd86 Fp	TGTTTCCGTGGAGACGCAAG
siSAMHD1-2 GCGUAUUUGUGCUAGAGAUTT siSAMHD1-3 GCAGAUGACUACAUAGAGATT shSamhd1 GGATGAGGATCGTCTGGAA	Cd86 Rp	TTGAGCCTTTGTAAATGGGCA
siSAMHD1-3 GCAGAUGACUACAUAGAGATT shSamhd1 GGATGAGGATCGTCTGGAA	siSAMHD1-1	CCUCGUCCGAAUCAUUGAUTT
shSamhd1 GGATGAGGATCGTCTGGAA	siSAMHD1-2	GCGUAUUUGUGCUAGAGAUTT
	siSAMHD1-3	GCAGAUGACUACAUAGAGATT
siSTING GGUCAUAUUACAUGGAUATT	shSamhd1	GGATGAGGATCGTCTGGAA
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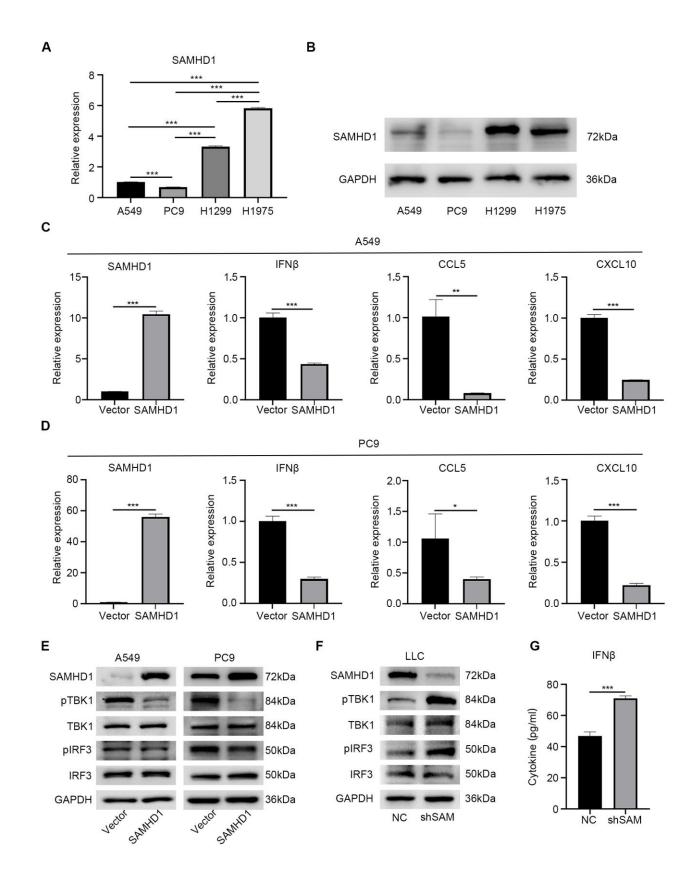
Antibody	Company	Catalog number
GAPDH	Proteintech	10494-1-AP
ssDNA	Sigma-Aldrich	MAB3868
IFI16	Abcam	Ab169788
SAMHD1	Origene	TA502024
STING	Proteintech	19851-1-AP
IRF3	Proteintech	11312-1-AP
p-IRF3	Cell Signaling Technology	4947S
TBK1	Cell Signaling Technology	30138
p-TBK1	Cell Signaling Technology	54838
Dylight 549 Goat Anti-Mouse	Abbkin	A23310
FITC Goat Anti-Rabbit IgG	Proteintech	SA00003-2
HRP Goat Anti-Rabbit IgG	Proteintech	SA00001-2
HRP Goat Anti-Mouse IgG	Proteintech	SA00001-1
Fixable Viability Stain 700	BD Pharmingen	564997
Ms CD45 PerCP-Cy5.5	BD Pharmingen	550994
Ms CD3 APC-Cy7	BD Pharmingen	560590
PE-Cy7 Anti-mouse CD4	BD Pharmingen	552775
Flour647 Anti-mouse CD8a	BD Pharmingen	557682
Ms CD86 PE-Cy7 GL1	BD Pharmingen	560582
CD11b APC-Cy7	BD Pharmingen	557657
Ms F4/80 BV421	BD Pharmingen	565411
Ms I-A/I-E BV480	BD Pharmingen	746669

Table S2. Antibodies used in this study.

## **Additional Figures**



## **Figure S1. Functional enrichment analysis to confirm the association of SAMHD1 with immunity.** (A) The SAMHD1 protein levels after radiation were detected by immunoblotting in LUAD cell lines. **(B)** GO and KEGG enrichment analysis. **(C)** GSEA analysis.



**Figure S2. SAMHD1 inhibited TBK1-IRF3-IFN-I pathway in LUAD cells. (A, B)** The mRNA and protein levels of SAMHD1 in LUAD cell lines. **(C, D)** The mRNA levels of IFNβ, CCL5 and CXCL10 in A549 and PC9 cells upon SAMHD1 overexpression were detected by qPCR. (E) The phosphorylation levels of TBK1 and IRF3 after SAMHD1 overexpression in A549 and PC9 cells were detected by immunoblotting. **(F)** TBK1-

IRF3 pathway protein levels in LLC cells were detected by immunoblotting. (G) Supernatant levels of IFN $\beta$  in LLC cells were detected by ELISA. N = 3; \*, *P* < 0.05; \*\*, *P* < 0.01; \*\*\*, *P* < 0.001.

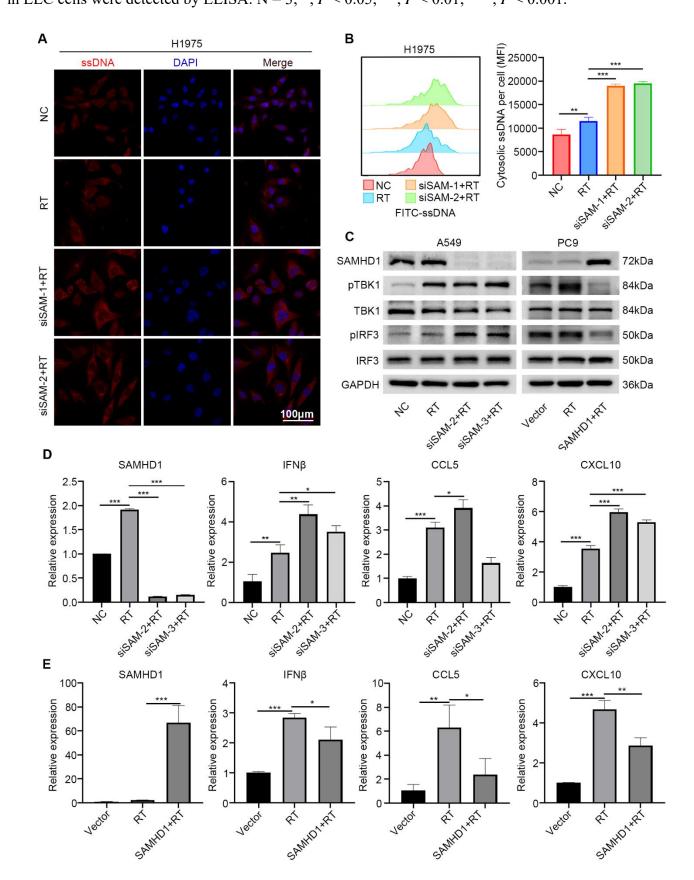
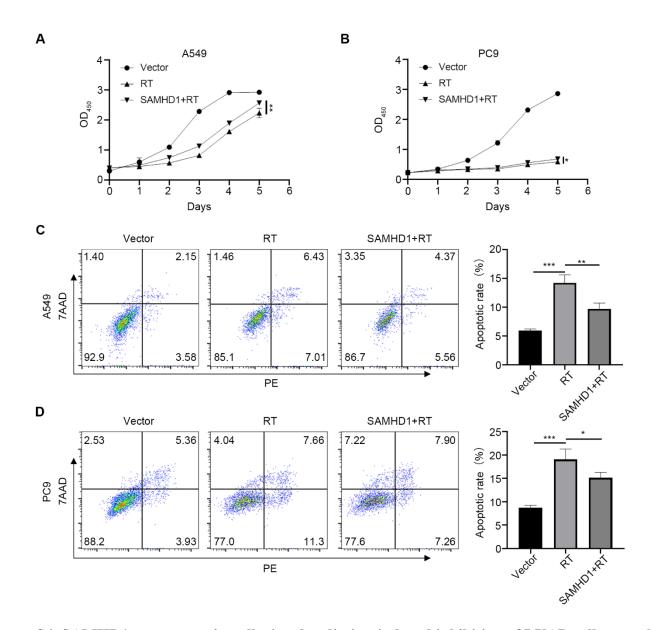


Figure S3. SAMHD1 silencing synergized with radiotherapy to induce ssDNA accumulation and activate TBK1-IRF3-IFN-I signaling. (A) Confocal images were performed to detect the accumulation of

ssDNA in cytoplasm in H1975 cells after radiation and transfecting siSAMHD1. (**B**) The accumulation of ssDNA in cytoplasm after radiation and siSAMHD1 transfection was detected by flow cytometry. The cytosolic ssDNA was evaluated by MFI. (**C**) TBK1-IRF3 pathway protein levels in A549 and PC9 cells after transfection and radiation were detected by immunoblotting.(D) The mRNA levels of SAMHD1, IFN $\beta$ , CCL5 and CXCL10 after SAMHD1 silencing and radiation in A549 cells were detected by qPCR. (E) The mRNA levels of SAMHD1, IFN $\beta$ , CCL5 and CXCL10 after SAMHD1 overexpression and radiation in PC9 cells were detected by qPCR. N = 3; \*, *P* < 0.05; \*\*, *P* < 0.01; \*\*\*, *P* < 0.001.



**Figure S4. SAMHD1 overexpression alleviated radiation-induced inhibition of LUAD cells growth. (A, B)** CCK8 assays were performed to evaluate the cell growth inhibition of SAMHD1 overexpression and radiation in A549 and PC9 cells. **(C, D)** The effects of SAMHD1 overexpression and radiation on apoptosis

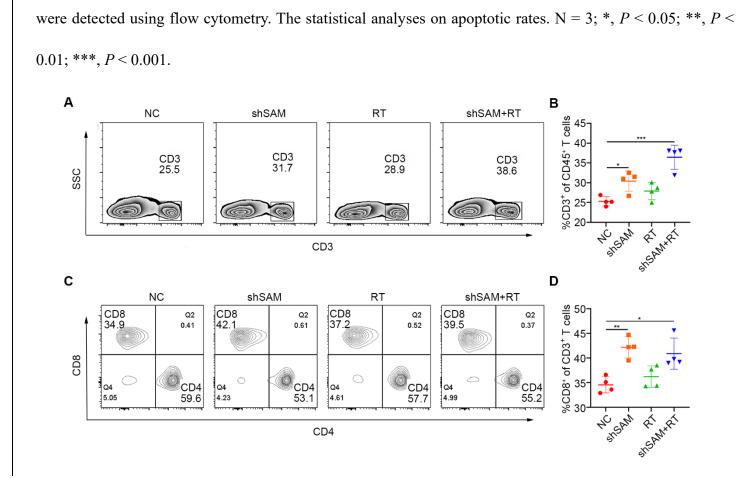


Figure S5. SAMHD1 silencing synergized with radiotherapy to regulate CD3<sup>+</sup> and CD8<sup>+</sup> T cell infiltration in spleen. (A) Representative flow cytometry of CD3<sup>+</sup> T cells in spleens. (B) Quantitative analysis of CD3<sup>+</sup> T cells in spleens. (C) Representative flow cytometry of CD4<sup>+</sup> T cells and CD8<sup>+</sup> T cells in spleens. (D) Quantitative analysis of CD8<sup>+</sup> T cells in spleens. N = 4; \*, P < 0.05; \*\*, P < 0.01; \*\*\*, P < 0.001.