

Serum amyloid A contributes to radiation-induced lung injury by activating macrophages through FPR2/Rac1/NF- κ B pathway

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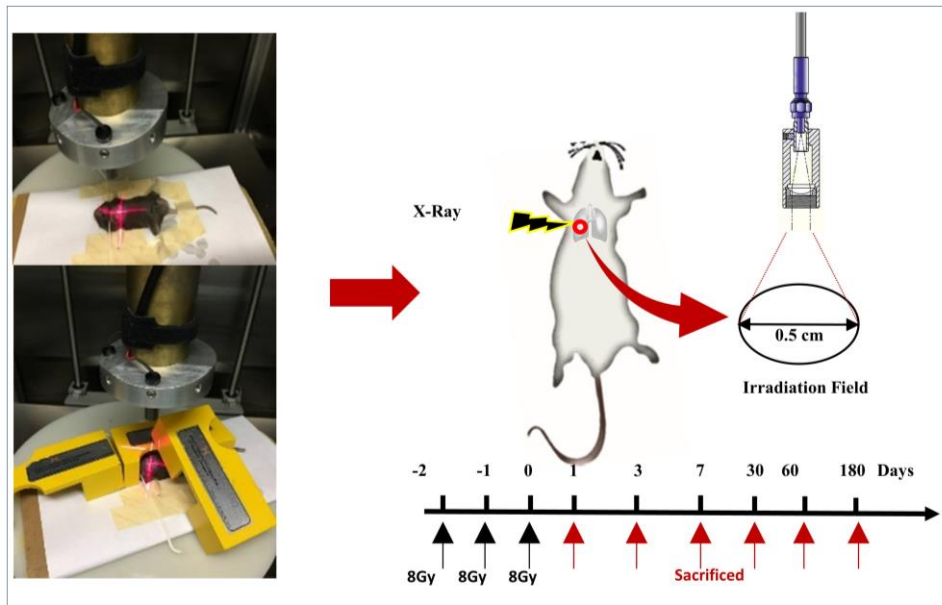
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Supplementary figures

A



B

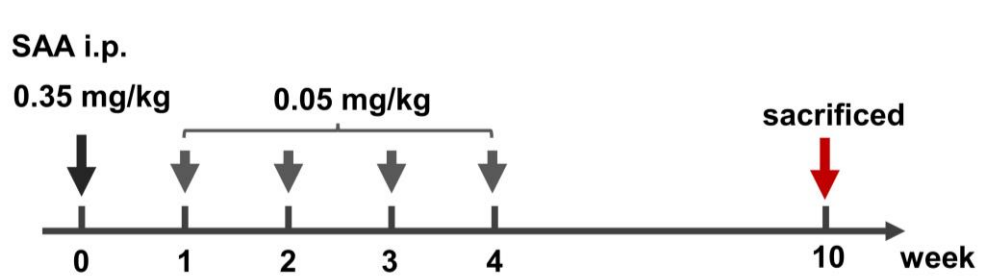


Figure S1. **A** Thoracic irradiation model in mice. The right lung of the mice was irradiated by X-rays with 8 Gy each day for three days through a circular collimator. **B** Schematic flow of SAA-induced lung injury mouse model. The mice were intraperitoneally injected with 0.35 mg/kg SAA on the first day and then with 0.05 mg/kg SAA repeated once a week for 4 weeks, lung tissues were collected at 10 weeks after the first SAA injection.

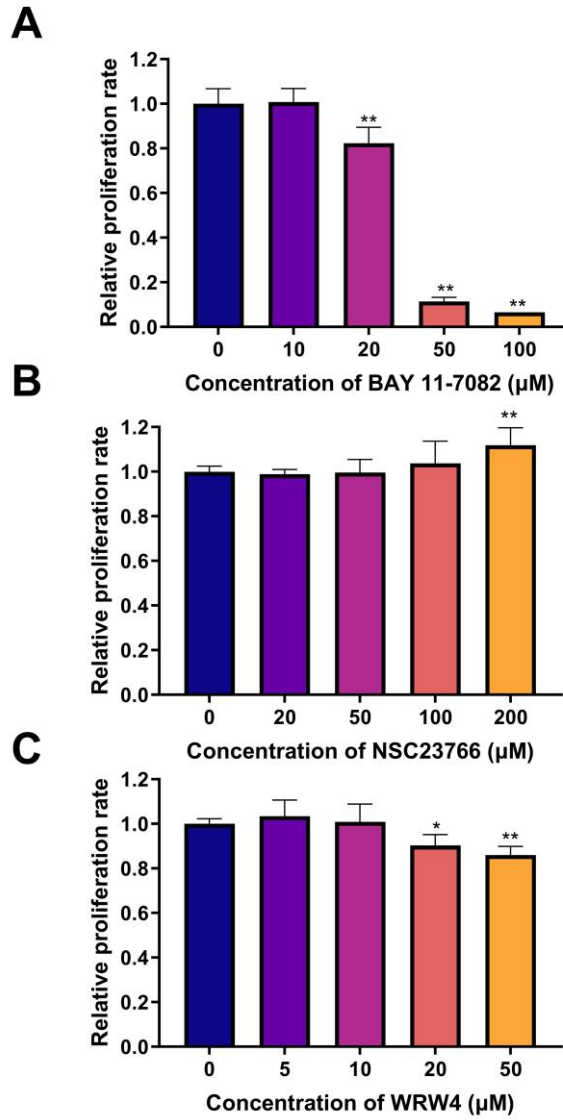


Figure S2. The effect of inhibitors on cell proliferation. A-C Relative proliferation level of RAW264.7 macrophages treated with BAY11-7082 (NF- κ B inhibitor), NSC33766 (Rac1 inhibitor), and WRW4 (FPR2 inhibitor). * $p < 0.05$, ** $p < 0.01$ between indicated groups.

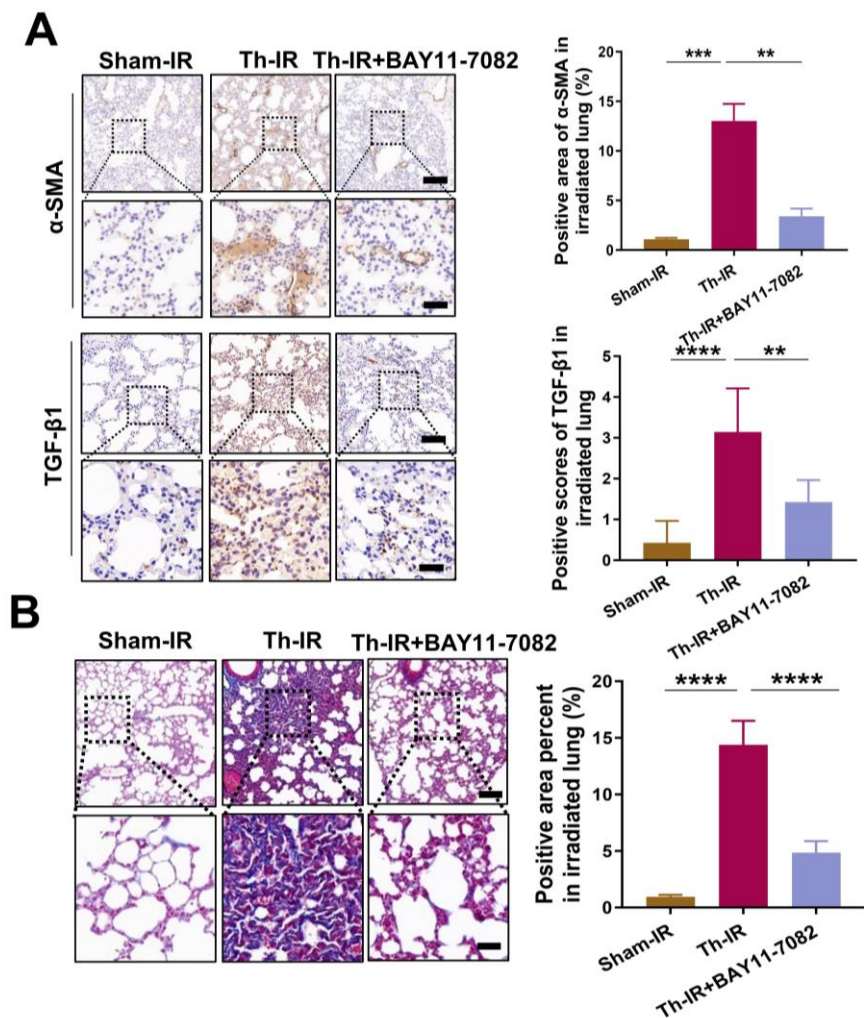


Figure S3. NF- κ B mediated lung injury induced by Th-IR. **A** IHC staining for α -SMA and TGF- β 1 expression in irradiated lung tissues at week 8 after Th-IR. **B** Masson trichrome staining of irradiated lung tissues at week 8. Scale bar=100 μ m, magnifying scale bar=50 μ m. Data were presented as mean \pm SEM. ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$, and ns, no significant difference between indicated groups.

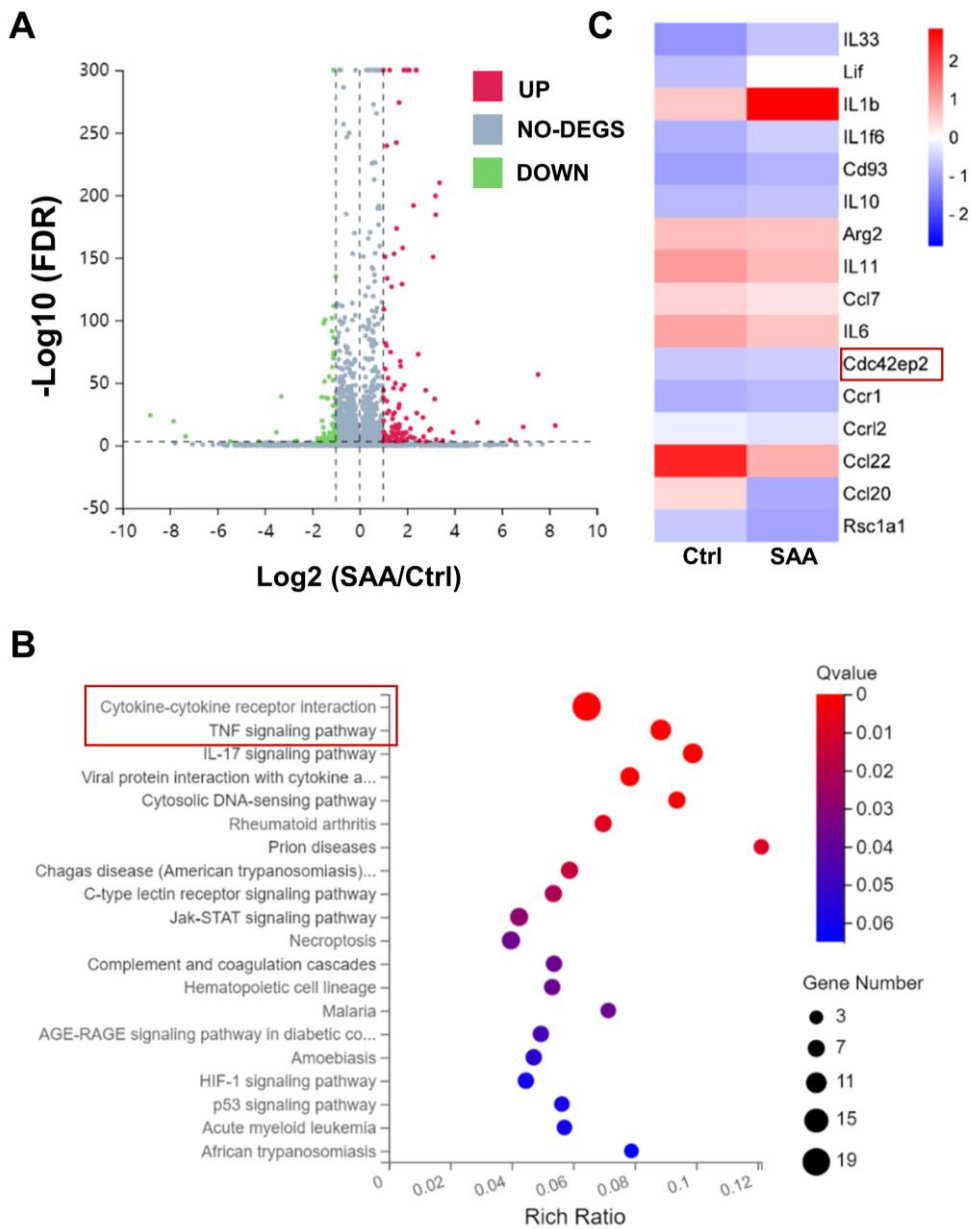


Figure S4. RNA sequencing transcriptome analysis on macrophages. **A** Volcano plots presenting the differentially expressed genes (DEGs) between 5 µg/mL SAA-treated group and control group (fold change > 2, $p < 0.01$). **B** Functional pathway analysis of DEGs. **C** Heatmap for differentially expressed cytokine and chemokine genes induced by SAA.

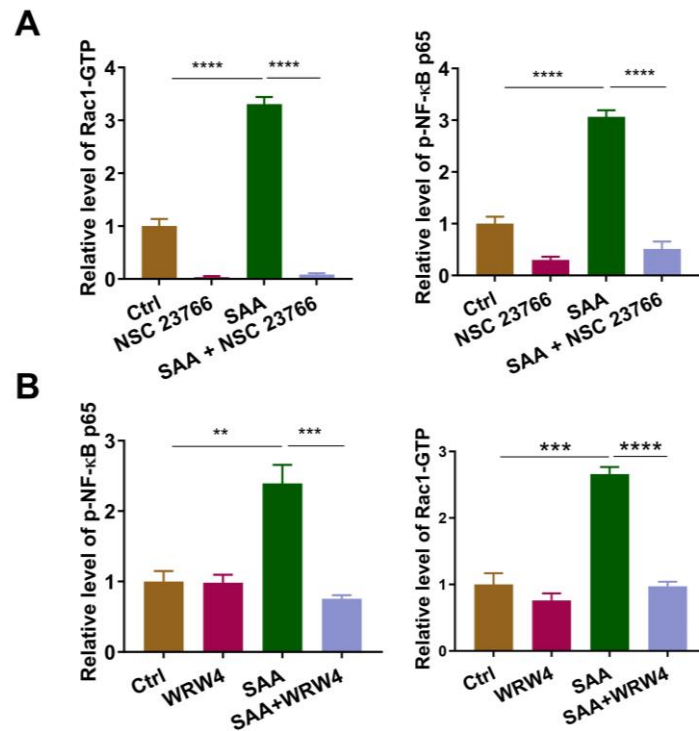


Figure S5. The statistical graphs for western blotting in Figure 4D-E. A-B The relative expression levels of p-NF-κB p65 and Rac1-GTP in macrophages after the cells being treated by NSC23766 (A) and WRW4 (B), respectively, in Figure 4D-E. $**p < 0.01$, $***p < 0.001$, and $****p < 0.0001$ compared with the indicated group.

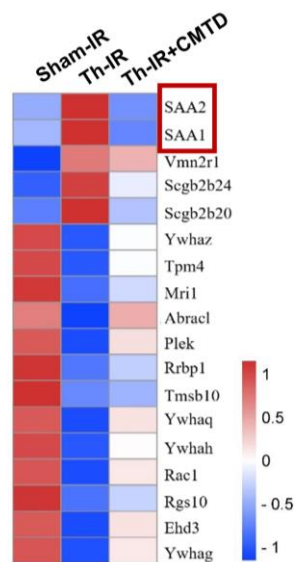


Figure S6 Heatmap of differentially expressed proteins between Sham-IR, Th-IR and Th-IR+CMTD group at day 7 after Th-IR.