

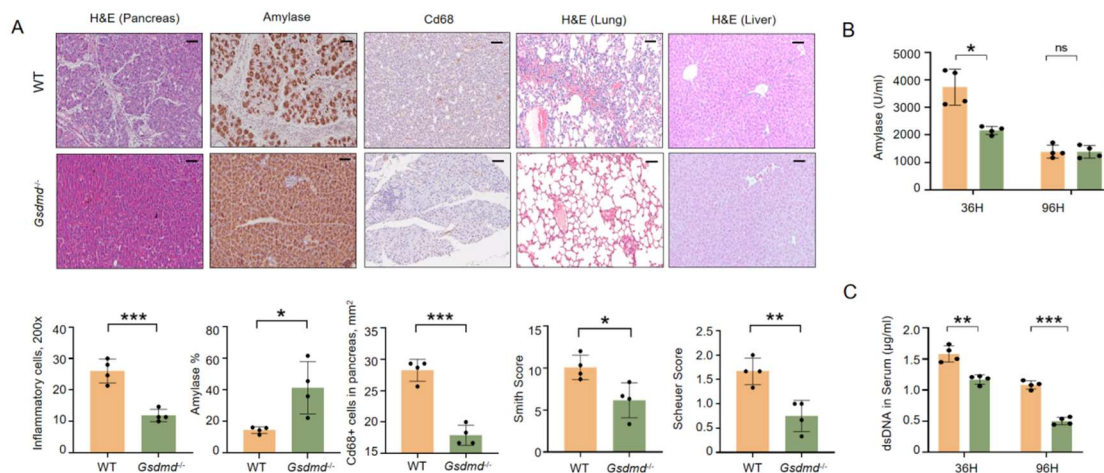
## Supplementary Materials

### I: Antibodies and dilution

Primary/Secondary antibody	Company	Cat. No	Dilution
NLRP3	CST	15101	1:1000
GSDMD	Abcam	ab219800	1:1000
N-GSDMD	Affinity	DF13758	1:1000 for WB, 1:500 for IHC
AMYLASE	CST	3796	1:1000 for WB, 1:500 for IHC
ACTIN	Proteintech	66009-1-Ig	1:1000
MPO	Abcam	ab208670	1:1000 for WB, 1:500 for IF
H3CIT	Abcam	ab281584	1:1000
NE	Abcam	ab131260	1:1000
Caspase4	Proteintech	11856-1-AP	1:1000
Cleaved-Caspase-1	Affinity	AF4005	1:1000
IL-33 (FOR WB)	Abclonal	A8096	1:1000
IL-33 (FOR IHC)	Abways	CY5105	1:100
ST2	Abcam	ab194113	1:500
P-STAT6	Affinity	AF3301	1:500
MMP12	Dreambio	YM-Y23544	1:1000
LY6G	Abclonal	A22270	1:100
GATA3	Affinity	AF6233	1:100
TBET	Affinity	DF7759	1:100
CD68	Abcam	ab303565	1:500
CD68	Abcam	ab53444	1:500
CD86	Abways	DF6332	1:500
CD206	Proteintech	18704-1-AP	1:500
TBK1	Abways	CY5145	1:1000
P-TBK1	Abways	CY8364	1:1000
STING	Proteintech	19851-1-AP	1:1000
P-STING	Affinity	AF7416	1:1000
IRF3	Abways	CY5779	1:1000
P-IRF3	Proteintech	29528-1-AP	1:1000
FITC Anti-Mouse CD4 Antibody[RM4-5]	Elabscience	E-AB-F1353C	1:20
APC Anti-Mouse Foxp3 Antibody[3G3]	Elabscience	E-AB-F1238E	1:20
APC Anti-Mouse IL-4 Antibody[11B11]	Elabscience	E-AB-F1204E	1:20
PE Anti-Mouse F4/80	Elabscience	E-AB-F0995D	1:20

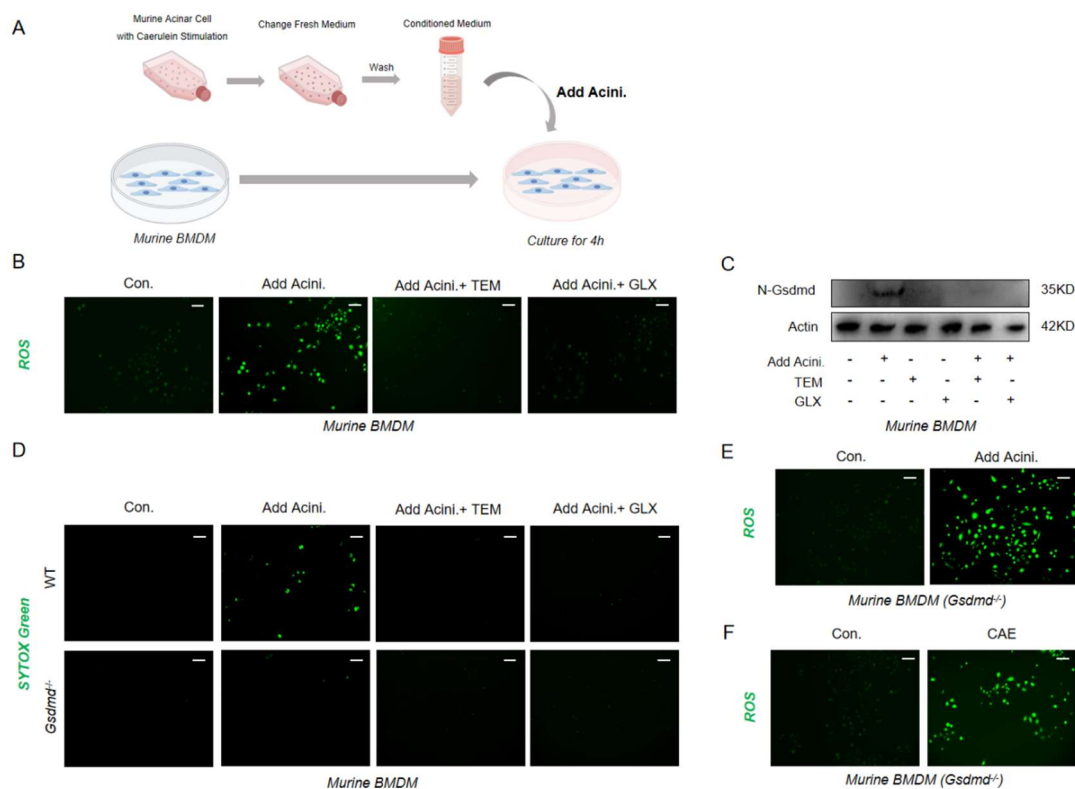
Antibody[CI:A3-1]			
APC Anti-Mouse CD206/MMR Antibody[C068C2]	Elabscience	E-AB-F1135E	1:20
PE Anti-Mouse IFN- $\gamma$ Antibody[XMG1.2]	Elabscience	E-AB-F1101D	1:20
Alexa Fluor 488-labeled Goat Anti-Rabbit IgG(H+L)	Beyotime	A0423	1:200
Alexa Fluor 488-labeled Goat Anti-Mouse IgG(H+L)	Beyotime	A0473	1:200
Cy3-labeled Goat Anti-Rat IgG(H+L)	Beyotime	A0507	1:200
Cy3-labeled Goat Anti-Rabbit IgG (H+L)	Beyotime	A0562	1:200
FITC-labeled Goat Anti-Mouse IgG (H+L)	Beyotime	A0568	1:200
Anti-rabbit IgG, HRP-linked Antibody	CST	7074	1:5000
Anti-mouse IgG, HRP-linked Antibody	CST	7076	1:5000
HRP-labeled Goat Anti-Rabbit IgG(H+L)	Beyotime	A0208	1:200
HRP-labeled Rabbit Anti-Mouse IgG(H+L)	Beyotime	A0354	1:200

## II: Supplementary Figures and Figure Legend

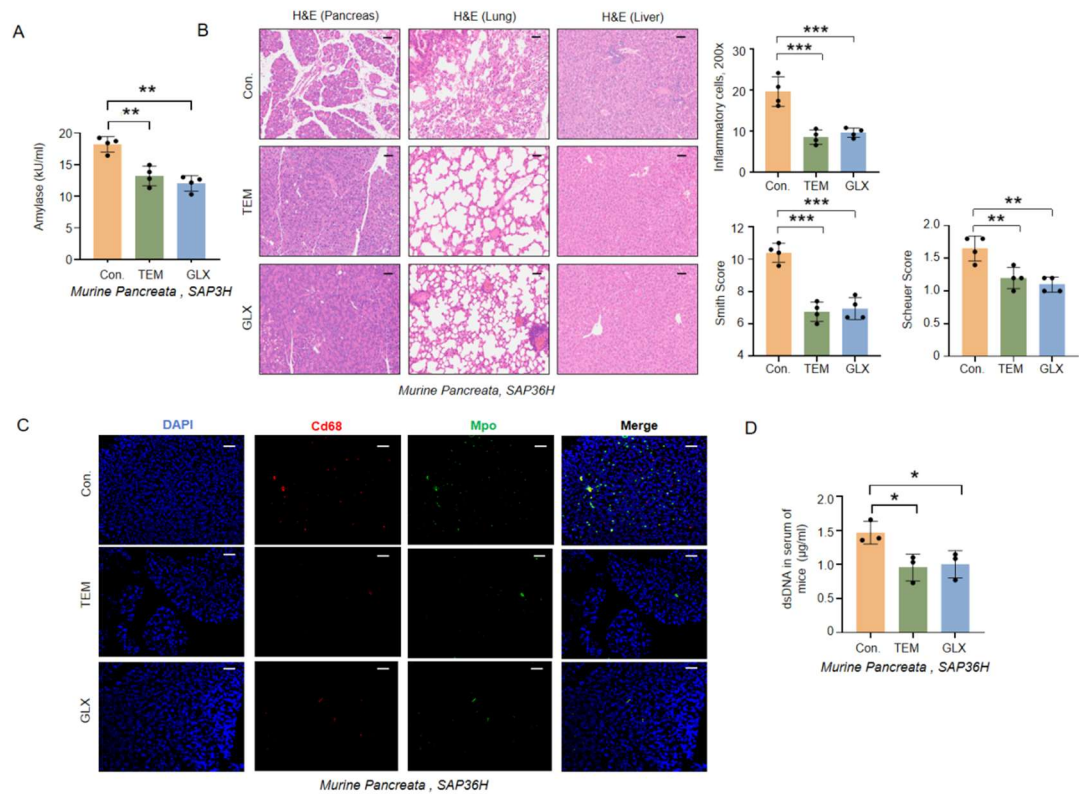


**Figure S1. GSDMD Knockout Alleviates SAP in Mice** A) Representative images and quantitative assessments show pancreatic inflammation cell infiltration, amylase staining of pancreata, pancreatic immunostaining for Cd68<sup>+</sup> cells, as well as histological scoring of lung and liver tissues in wild-type and *Gsdmd*<sup>-/-</sup> mice after SAP induction (n=4 mice per group).

Scale bar=50μm. **B)** Serum amylase measured in wild-type and *Gsdmd*<sup>-/-</sup> mice after SAP induction (n=4 mice per group). **C)** dsDNA levels measured in wild-type and *Gsdmd*<sup>-/-</sup> mice after SAP induction (n=4 mice per group). Statistical significance for **A**, **B** and **C** was determined using unpaired Student's t-tests. Data are presented as mean ± SD. Statistical significance: \*P < 0.05, \*\*P < 0.01, \*\*\*P < 0.001, ns. not significant.



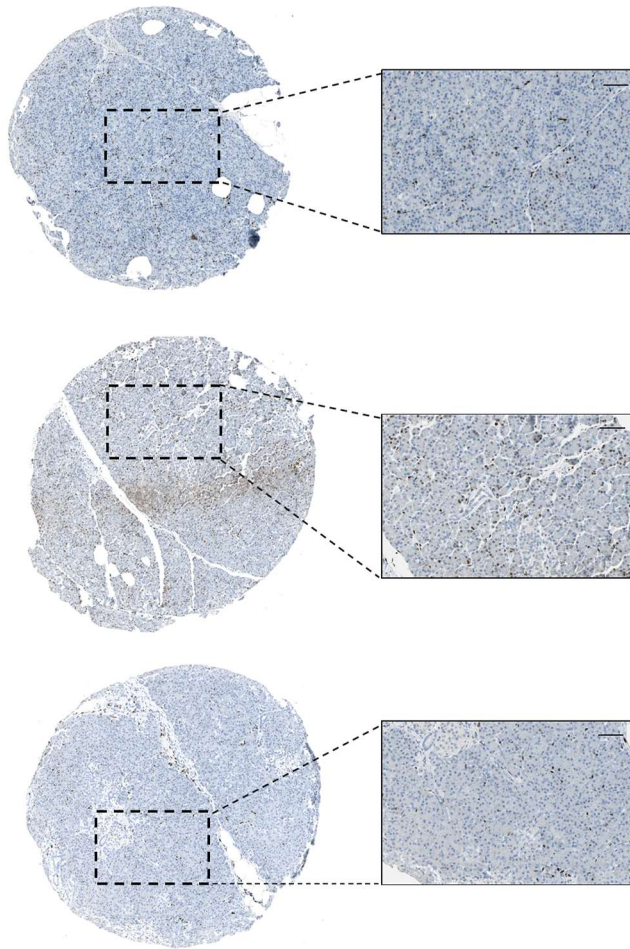
**Figure S2. Injured pancreatic acinar cells mediate METs release via the ROS-GSDMD axis.** **A)** Schematic of the procedure for obtaining conditioned medium from injured acinar cells. Created with Figdraw.com. **B)** Representative images of reactive oxygen species (ROS) in wild-type bone marrow-derived macrophages (BMDMs) treated with conditioned medium from injured acinar cells for 4 hours, with or without co-treatment with Tempol (20 nM) and GLX481304 (20 nM). Scale bar=50μm. **C)** Western blot analysis of N-terminal Gasdermin D (N-Gsdmd) in BMDMs treated with conditioned medium from injured acinar cells, Tempol (20 nM), GLX481304 (20 nM), or combinations of these factors. **D)** Representative images of SYTOX Green in wild-type or *Gsdmd* knockout BMDMs treated with conditioned medium from injured acinar cells for 4 hours, with or without co-treatment with Tempol (20 nM) and GLX481304 (20 nM). Scale bar=50μm. **E, F)** Representative images of ROS in *Gsdmd* knockout BMDMs treated with conditioned medium from injured acinar cells or caerulein (500 pM) for 4 hours. Scale bar=50μm.



**Figure S3. Pharmacological Inhibition of ROS Suppresses METs Formation and Alleviates Pancreatic Injury.** **A)** Serum amylase levels in SAP mice treated with Tempol(TEM) or GLX481304(GLX) (n=4 mice per group). **B)** Representative images and quantitative analysis of H&E staining of mouse pancreas, lung and liver. (n=4 mice per group). **C)** Immunofluorescence images of DAPI (blue), Cd68 (red), and Mpo (green) in pancreata of SAP mice treated with Tempol or GLX481304 (n=4 mice per group). **D)** Serum dsDNA levels in SAP mice treated with Tempol or GLX481304 (n=3 mice per group). Statistical significance for **A**, **B** and **D** was determined using a one-way ANOVA with multiple comparisons test. Data are presented as mean  $\pm$  SD. Statistical significance: \*P < 0.05, \*\*P < 0.01, \*\*\*P < 0.001. Scale bar=50µm.

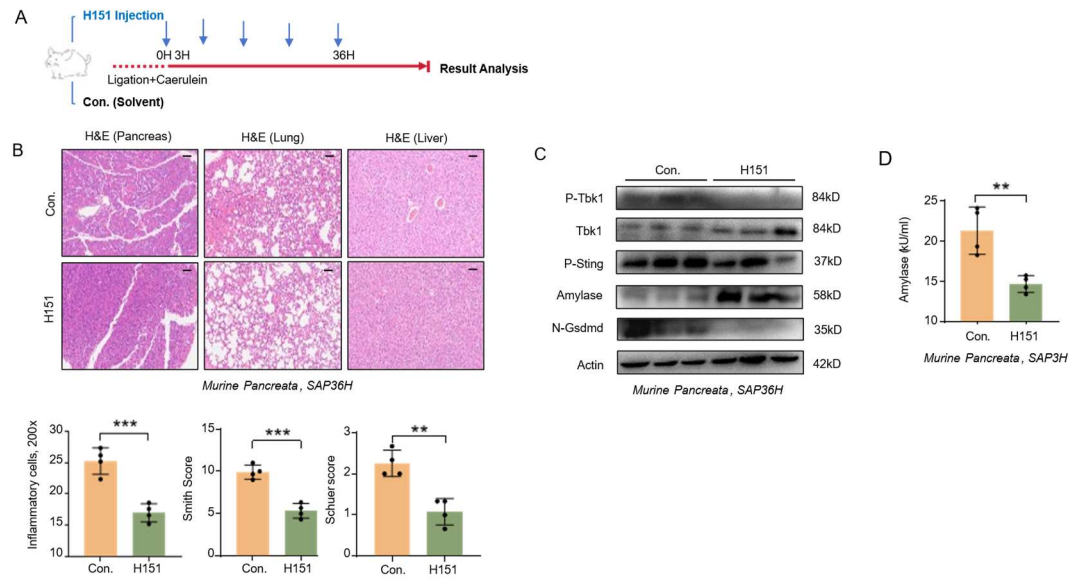
A

*Normal Pancreas*



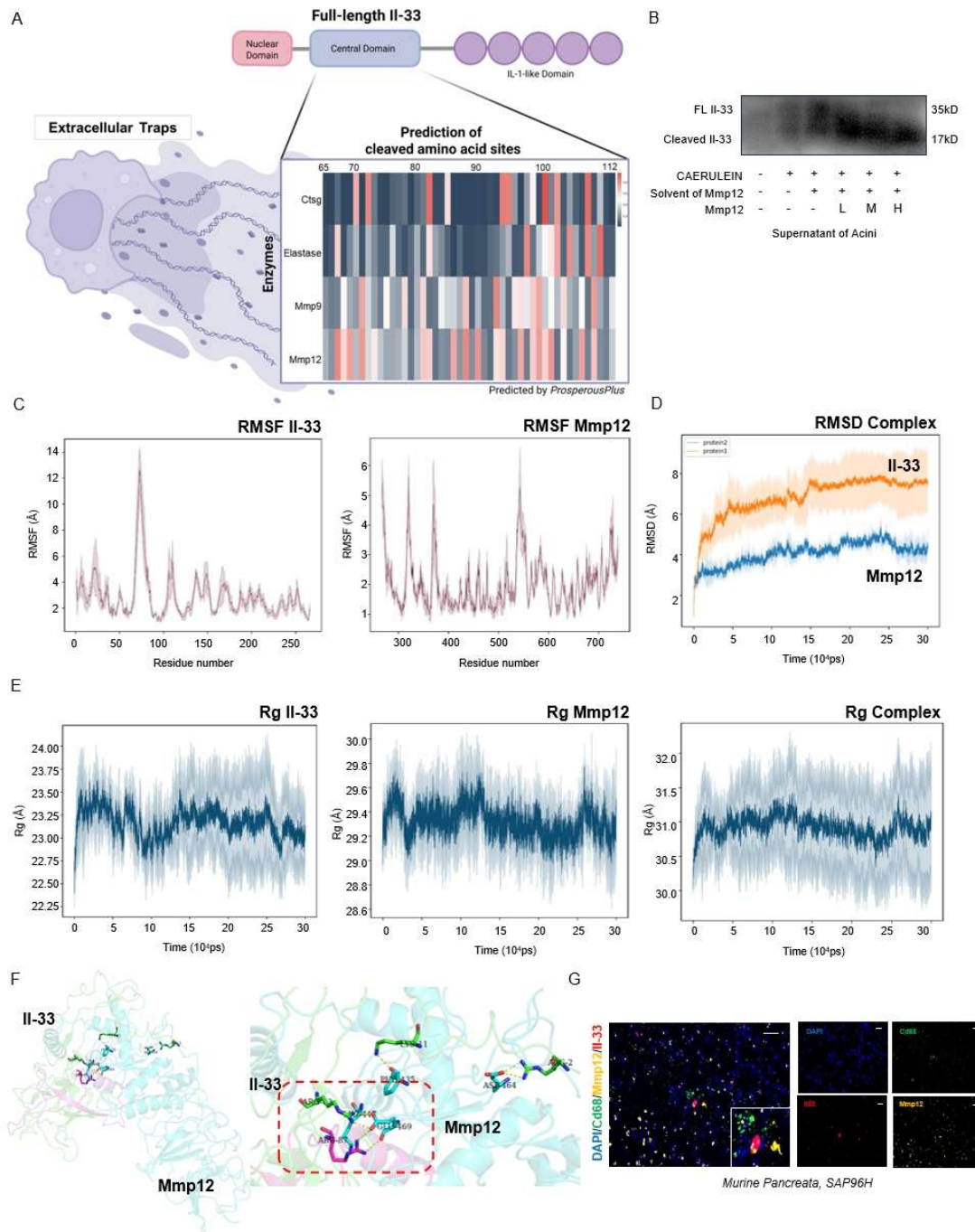
*HPA DATABASE*

**Figure S4. Expression of IL-33 in Human Normal Pancreas.** A) Immunohistochemistry data from the Human Protein Atlas (HPA) database show that IL-33 is expressed in the nucleus of acinar cells in the normal human pancreas from three patients. Scale bar=50 $\mu$ m.



**Figure S5. Pharmacological Inhibition of STING Signaling Pathway Alleviates Pancreatic Injury during SAP.** **A)** Schematic diagram illustrating the establishment of the SAP mouse model and H151 administration procedures. **B)** Representative images and quantitative analysis of H&E staining of mouse tissues (n=4 mice per group). Scale bar=50µm. **C)** Western blot analysis of Amylase, N-Gsdmd, and phosphorylated forms of Tbk1 and Sting in the pancreata of SAP mice with and without H151 treatment (n=3 mice per group). **D)** Serum amylase levels in SAP mice with and without H151 treatment (n=4 mice per group). Statistical significance for **B** and **D** was determined using unpaired Student's t-tests. Data are presented as mean  $\pm$  SD. Statistical significance: \*\*P < 0.01, \*\*\*P < 0.001.





**Figure S6. Prediction of the Cleavage of IL-33 by MMP12.** **A)** The potential cleavage sites in the central domain of IL-33 were predicted using the *ProsperousPlus*. Created with Biorender.com. **B)** Western blot analysis demonstrating that addition of Mmp12 results in cleavage of IL-33 in injured acinar supernatants [L=low dose(10pM), M=medium dose(100pM), H=high dose(1nM)]. **C)** Root mean square fluctuation (RMSF) plots for individual residues of IL-33 (left) and Mmp12 (right) throughout the simulation. **D)** Root mean square deviation (RMSD) over time for the IL-33-Mmp12 complex, indicating stability during the simulation period. **E)** Radius of gyration (Rg) plots for IL-33, Mmp12, and their complex, illustrating changes in compactness over time. All molecular dynamics simulations

were performed in triplicate to ensure reliability. **F)** Molecular dynamics simulation snapshots showing the interaction between Il-33 (green) and Mmp12 (blue). The binding interface is highlighted by a red dashed box. **G)** Immunofluorescence staining of pancreatic tissues following SAP induction showing colocalization of Cd68 (green), Mmp12 (gold), and Il-33 (red). Nuclei are stained with DAPI (blue); scale bar=25 $\mu$ m.