

## SUPPLEMENTAL MATERIAL

### Antagonistic effects of the cytotoxic molecules granzyme B and TRAIL in the immunopathogenesis of sclerosing cholangitis

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## SUPPORTING MATERIALS AND METHODS

**Determination of liver injury.** Heart blood was drawn from individual mice and liver injury was quantified by automated measurement of plasma activities of alanine transaminase (ALT) using a COBAS Mira System (Roche Diagnostic, Mannheim, Germany).

**Hydroxyproline assay.** The concentration of the collagen component hydroxyproline was determined in liver tissue by a spectrophotometric assay as previously described.<sup>[1]</sup>

**Immunohistochemistry.** Sirius red staining was done with 3  $\mu$ m paraffin-embedded liver sections as previously described<sup>[2]</sup> to determine collagen fibers. To identify cholangiocytes, 3  $\mu$ m paraffin-embedded liver sections were stained with an anti-CK19 antibody (TROMA-III; DSHB, Iowa City, IA, USA). A biotin-conjugated secondary antibody was used (Abcam, Cambridge, UK). For antigen detection, the DAB+ Substrate Chromogen System (Agilent Technologies, Santa Clara, CA, USA) was used. To stain myofibroblasts, 3  $\mu$ m paraffin-embedded liver sections were incubated with an anti- $\alpha$ SMA antibody (1A4; Abcam, Cambridge, UK). After incubation with a biotin-conjugated secondary antibody (ImmunoReagents, Raleigh, NC, USA), liver sections were stained with the DAB+ Substrate Chromogen System. CK19 and  $\alpha$ SMA staining was evaluated per  $\mu$ m<sup>2</sup> using the Keyence BZ-II Analyzer software.

**Phospho-protein staining.** Following extracellular staining, cells were fixed with 4% PFA. Cells were permeabilized with ice-cold methanol and stained with anti-phospho-Zap-70 (65E4, PE) and anti-phospho-PLC $\gamma$ 1 antibodies (D6M9S, Alexa Fluor 488; both Cell Signaling, Danvers, MA, USA).

***Annexin V/7-AAD staining.*** To analyze apoptotic immune cells, cells were stained with Annexin V and 7-AAD using the PE Annexin V Apoptosis Detection Kit with 7-AAD (BioLegend) according to manufacturer's protocol.

***Determination of GzmB in culture supernatants.*** To measure GzmB concentrations in supernatants of re-stimulated NPCs, an enzyme-linked immunosorbent assay (ELISA) was carried out according to manufacturer's protocol (R&D Systems, Minneapolis, MN). GzmB levels were measured using a Tecan Infinite plate reader (Tecan, Männedorf, Switzerland) and Magellan Data Analysis Software (V6.5; Tecan).

***Quantitative real-time PCR analysis.*** Total RNA was isolated from shock-frozen liver tissue using the NucleoSpin RNA Midi Kit (Macherey-Nagel, Düren, Germany). Genomic DNA was digested using the DNA-free Kit (ThermoFisher Scientific). 1 µg RNA was transcribed into cDNA using the Verso cDNA Synthesis Kit (ThermoFisher Scientific) and the Biometra TAdvanced thermal cycler (Analytik Jena, Jena, Germany). Quantitative RT-PCR was performed using the PowerUp SYBR Green Master Mix (ThermoFisher Scientific) and the QuantStudio 7 Flex (Applied Biosystems, Waltham, USA). Relative mRNA levels were calculated using the  $\Delta\Delta CT$  method after normalization to the reference gene  $\beta$ -actin. Quantification was shown in x-fold changes to the corresponding control cDNA. Primers were designed for detection of exon overlapping amplicons and were obtained from Metabion (Martinsried, Germany). Sequences of the primers are listed in Table S2.

## Tables

Table S1: Antibodies used for cell surface staining.

| Target        | Fluorochrome    | Clone       | Distributor                            |
|---------------|-----------------|-------------|--|
| CD4           | BV 711          | RM4-5       | BioLegend                              |
| CD4           | APC-Cy7         | GK1.5       | BioLegend                              |
| CD8           | BV 785, BV 650  | 53-6.7      | BioLegend                              |
| CD25          | PE              | PC-61       | BioLegend                              |
| CD69          | APC             | H1.2F3      | BioLegend                              |
| TCR $\beta$   | PE-Cy7          | H57-597     | BioLegend                              |
| TCR $\delta$  | PerCP-Cy5.5     | GL3         | BioLegend                              |
| CD1d-Tetramer | Alexa Fluor 647 |             | NIH Tetramer Core Facility, Atlanta GA |
| NKp46         | APC, BV 711     | 29A1.4      | BioLegend                              |
| NK1.1         | BV 605          | PK136       | BioLegend                              |
| CD107a        | FITC            | 1D4B        | BioLegend                              |
| TRAIL         | PerCP-Cy5.5, PE | N2B2        | BioLegend                              |
| Sca-1         | BV 421          | D7          | BioLegend                              |
| KLRG1         | BV 785          | 2F1/KLRG1   | BioLegend                              |
| ICOS          | FITC            | 7E.17G9     | eBioscience                            |
| CD11c         | FITC            | N418        | BioLegend                              |
| MHC-II        | APC-Cy7         | M5/114.15.2 | BioLegend                              |

Table S2: Sequences of the primer used for mRNA analysis.

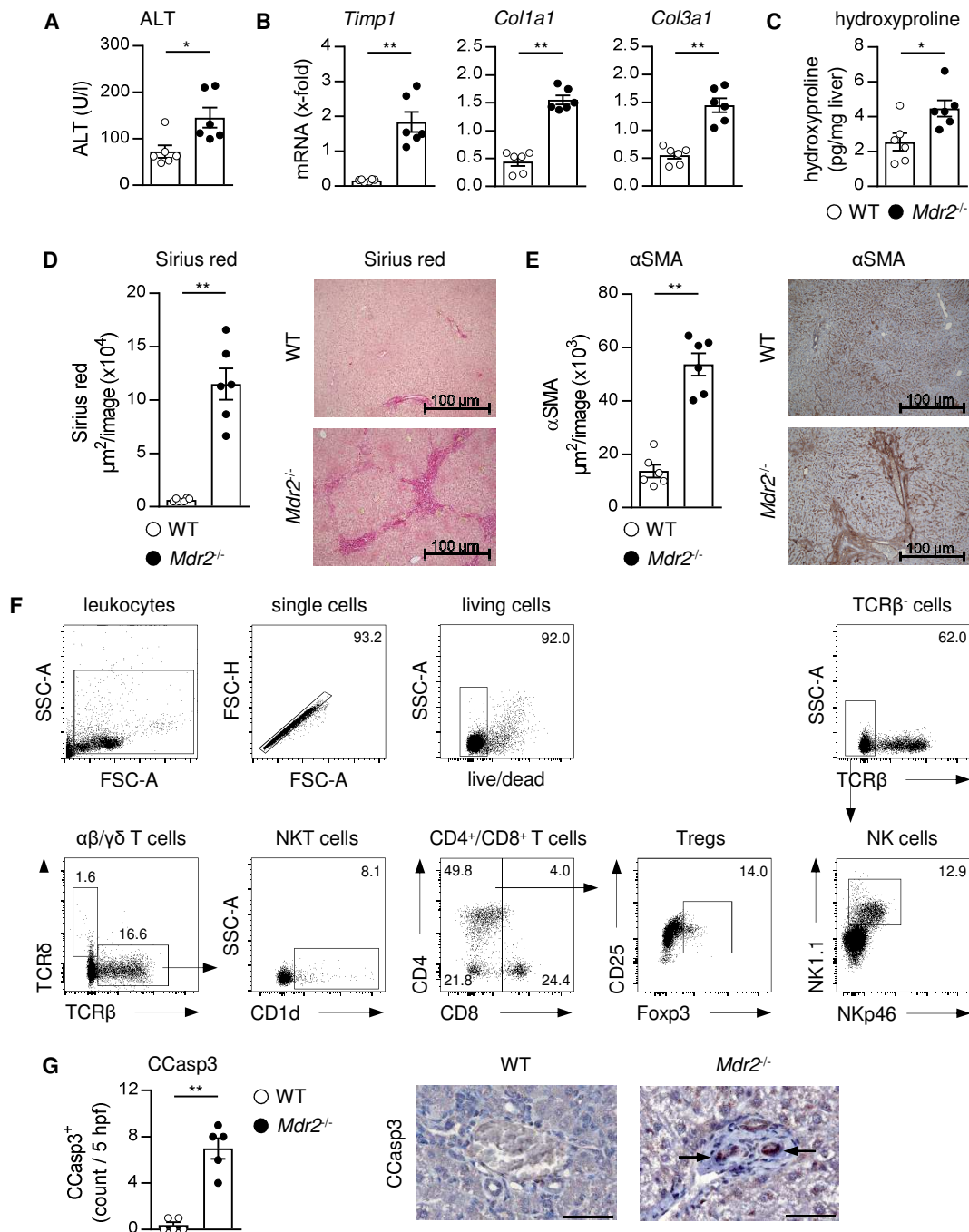
| Target | Forward primer<br>Reverse primer                           | Amplicon length | Annealing temperature |
|--------|--|-----------------|-----------------------|
| Timp1  | 5'-CGGACCTGGATGCTAAAAGGA-3'<br>3'-ACTCTCCAGTTTGCAAGGGA-5'  | 300 bp          | 60 °C                 |
| Col1a1 | 5'-GAGCGGAGAGTACTGGATCG-3'<br>3'-TACTCGAACGGGAATCCATC-5'   | 204 bp          | 60 °C                 |
| Col3a1 | 5'-GTCCACGAGGTGACAAAGGT-3'<br>3'-GATGCCCACTTGTTCCATCT-5'   | 204 bp          | 60 °C                 |
| Ccl2   | 5'-AGCTGTAGTTTTTGTACCAAGC-3'<br>3'-GTGCTGAAGACCTTAGGGCA-5' | 155 bp          | 60 °C                 |
| Cxcl2  | 5'-AGGCTACAGGGGCTGTTGT-3'<br>3'-TTCAGGGTCAAGGCAAACCTT-5'   | 176 bp          | 60 °C                 |
| Cxcl1  | 5'-CCCAAACCGAAGTCATAGCCA-3'<br>3'-CTCCGTTACTTGGGGACACC-5'  | 118 bp          | 60 °C                 |
| Ccl4   | 5'-CTGTGCAAACCTAACCCCGA-3'<br>3'-AGGGTCAGAGCCCATTGGT-5'    | 118 bp          | 60 °C                 |
| IL-12  | 5'-AGACCCTGCCCATTGAAGT-3'<br>3'-GGCGGGTCTGGTTTGATGAT-5'    | 99 bp           | 60 °C                 |

|              |  |        |       |
|--------------|--|--------|-------|
| IFN $\gamma$ | 5'-ACAGCAAGGCGAAAAAGGATG-3'<br>3'-TCTTCCCCACCCCGAATCA-5'     | 171 bp | 60 °C |
| Cxcl10       | 5'-GCCGTCATTTTCTGCCTCAT-3'<br>3'-TGCAGCGGACCGTCCTT-5'        | 76 bp  | 60 °C |
| Ccl28        | 5'-GCCTCACCTGAGTCATTGCC-3'<br>3'-CCATGGGAAGTATGGCTTCTGA-5'   | 121 bp | 60 °C |
| Cxcl16       | 5'-TTGGACCCTTGTCTCTTGCG-3'<br>3'-CCAGTTCCACACTCTTTGCG-5'     | 292 bp | 60 °C |
| Actb         | 5'-TATTGGCAACGAGCGGTTCC-3'<br>3'-GGCATAGAGGTCTTTACGGATGTC-5' | 180 bp | 60 °C |

## References

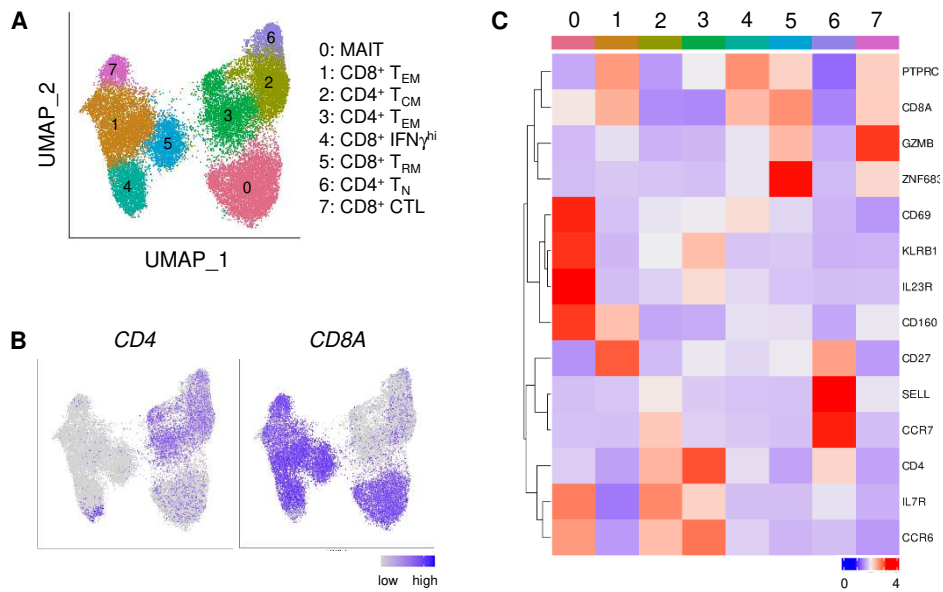
- [1] Uchinami H, Seki E, Brenner DA, D'Armiento J. Loss of MMP 13 attenuates murine hepatic injury and fibrosis during cholestasis. *Hepatology*. 2006;44:420-9.
- [2] Segnani C, Ippolito C, Antonioli L, Pellegrini C, Blandizzi C, Dolfi A, et al. Histochemical detection of collagen fibers by sirius red/fast green is more sensitive than van Gieson or sirius red alone in normal and inflamed rat colon. *PLoS One*. 2015;10:e0144630.

## SUPPLEMENTAL FIGURE 1



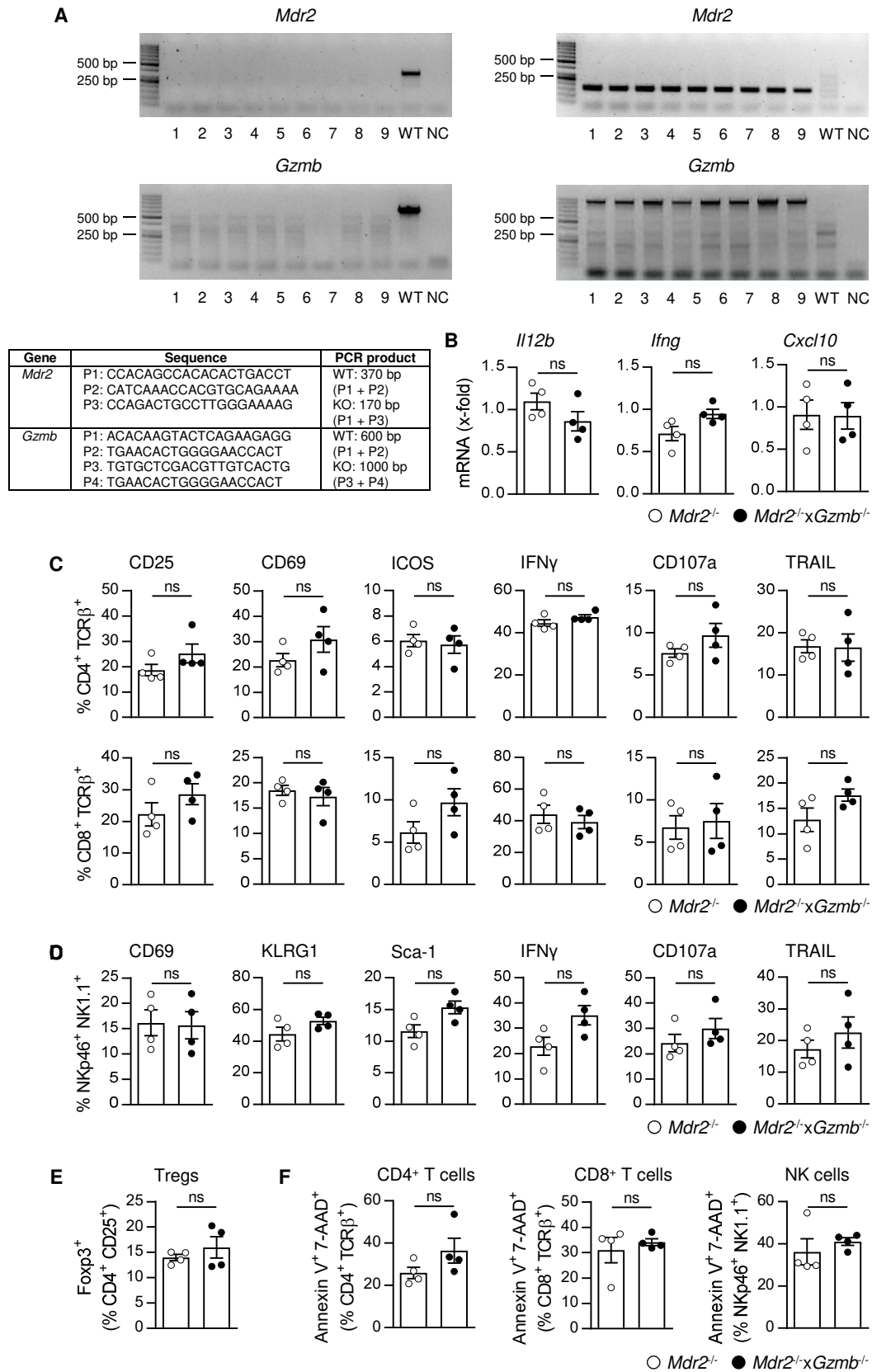
**Figure S1. Disease pathology and leukocyte populations in *Mdr2*<sup>-/-</sup> mice.** (A) Plasma ALT activities were determined in 12 weeks old *Mdr2*<sup>-/-</sup> and C57BL/6 WT mice. (B) Hepatic mRNA expression was analysed in *Mdr2*<sup>-/-</sup> mice and normalized to WT mice. (C) Hydroxyproline concentrations in liver tissue. (D) Sirius red and (E)  $\alpha\text{SMA}$  stainings were quantified in liver sections. (F) Leukocytes were isolated from livers of 12 weeks old *Mdr2*<sup>-/-</sup> mice and stained for TCR $\beta$ , TCR $\delta$ , CD4, CD8, CD1d, NKp46 and NK1.1 to identify CD4<sup>+</sup> TCR $\beta$ <sup>+</sup> T cells, CD8<sup>+</sup> TCR $\beta$ <sup>+</sup> T cells, CD25<sup>+</sup> Foxp3<sup>+</sup> CD4<sup>+</sup> Tregs, CD1d<sup>+</sup> TCR $\beta$ <sup>+</sup> NKT cells, NKp46<sup>+</sup> NK1.1<sup>+</sup> TCR $\beta$ <sup>-</sup> NK cells and TCR $\delta$ <sup>+</sup>  $\gamma\delta$  T cells. Representative frequencies and dot plots are shown. (G) CCasp3 was stained in liver sections and CCasp3-expressing cholangiocytes were counted. Arrows mark CCasp3<sup>+</sup> cholangiocytes. Bars represent 50  $\mu\text{m}$ . Mean  $\pm$  SEM of one out of two experiments are shown. \* $p \leq 0.05$ , \*\* $p \leq 0.01$ .

## SUPPLEMENTAL FIGURE 2



**Figure S2. Hepatic T cell subsets in patients with PSC.** (A) UMAP plot shows clustering of CD3<sup>+</sup> T cell subsets in livers of PSC patients undergoing liver transplantation. (B) UMAP plots depict CD4<sup>+</sup> and CD8<sup>+</sup> T cells clusters. (C) Heat map shows expression of subset-defining genes within the T cell clusters.

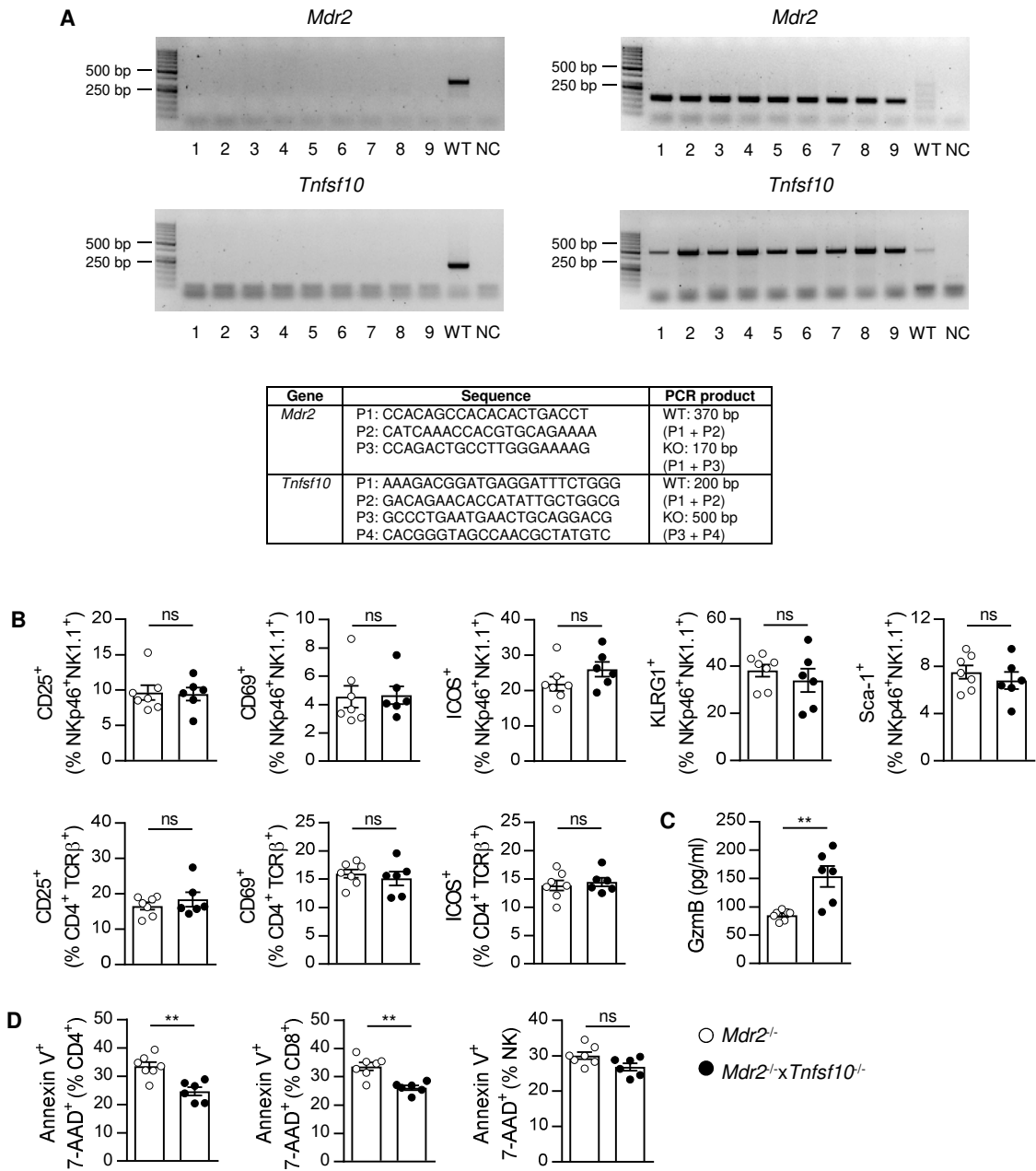
## SUPPLEMENTAL FIGURE 3



**Figure S3. Phenotype of hepatic T cells and NK cells in *Mdr2*<sup>-/-</sup> × *Gzmb*<sup>-/-</sup> mice.** (A) Genotyping of *Mdr2*<sup>-/-</sup> × *Gzmb*<sup>-/-</sup> mice is shown. (B) Hepatic mRNA expression was analysed in *Mdr2*<sup>-/-</sup> × *Gzmb*<sup>-/-</sup> mice and normalised to *Mdr2*<sup>-/-</sup> mice. (C) Phenotype of hepatic T cells and (D) NK cells was analysed. (E) Frequencies of hepatic Tregs and (F) apoptotic T cells and NK cells. Mean ± SEM of one out of three experiments is shown. ns: not significant.

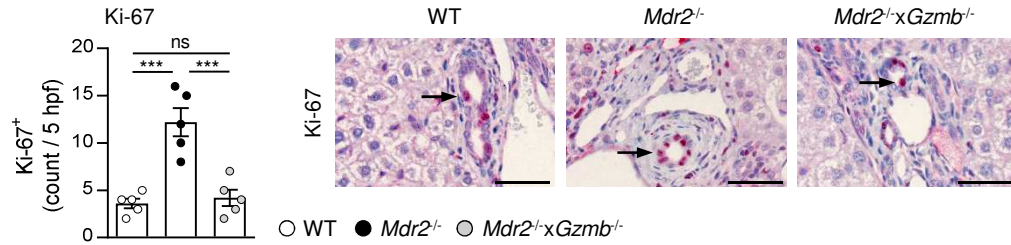


## SUPPLEMENTAL FIGURE 4



**Figure S4. *Mdr2*<sup>-/-</sup> × *Tnfsf10*<sup>-/-</sup> mice.** (A) Genotyping of *Mdr2*<sup>-/-</sup> × *Tnfsf10*<sup>-/-</sup> mice is shown. (B) Phenotype of hepatic NK cells and CD4<sup>+</sup> T cells was analysed in 12 weeks old *Mdr2*<sup>-/-</sup> × *Tnfsf10*<sup>-/-</sup> and *Mdr2*<sup>-/-</sup> mice. (C) GzmB levels were determined in culture supernatants of hepatic non-parenchymal cells from *Mdr2*<sup>-/-</sup> × *Tnfsf10*<sup>-/-</sup> and *Mdr2*<sup>-/-</sup> mice. (D) Frequency of apoptotic hepatic CD4<sup>+</sup> and CD8<sup>+</sup> T cells and NK cells is shown. Mean ± SEM of one out of three experiments is shown. \*\*p ≤ 0.01 ; ns: not significant.

## SUPPLEMENTAL FIGURE 5



**Figure S5. Proliferation of cholangiocytes in sclerosing cholangitis.** Ki-67 was stained in liver sections of 12 weeks old WT, *Mdr2*<sup>-/-</sup> and *Mdr2*<sup>-/-</sup>*xGzmb*<sup>-/-</sup> mice. Ki-67-expressing cholangiocytes were counted. Bars represent 50  $\mu$ m. Arrows mark Ki-67<sup>+</sup> cholangiocytes. Mean  $\pm$  SEM of one out of two experiments is shown.