Supplemental Figure 1 (cont). Confirmation of HMGB1 deletion by Mx1Cre and Vav1Cre. A-C. Mx1Cre-mediated Hmgb1 deletion results in almost complete loss of HMGB1 protein in the liver and bone marrow, partial loss in other tissues and almost complete loss of HMGB1 secretion in LPS-stimulated macrophages (n=4 per group). D. Vav1Cre-mediated Hmgb1 deletion leads to strong reduction of protein levels in bone marrow and spleen but not liver (n=4 per group). E. Vav1Cre efficiently deletes HMGB1 in F4/80-positive macrophages in the liver but only to a small extent in endomucin-positive endothelial cells (arrows pointing to endothelial cells or macrophages expressing HMGB1). F. Similarly, most endothelial cells in the kidney and heart retain HMGB1 expression (arrows point towards endothelial cells with HMGB1 expression). Statistical significance assessed via two-tailed unpaired t-test (C). ** p<0.01. Scale bars 100 µm (B) and 50 µm (E, F).
Supplementary Figure 2.
Supplementary Figure 2 (cont.). *Hmgb1Δhep* mice exhibit effective deletion of HMGB1 in hepatocytes and biliary epithelial cells but not in other hepatic cell populations. A-E. Confocal microscopy of liver sections co-stained for HMGB1 and markers for liver cell subpopulations including hepatocytes (HNF-4α, A), biliary epithelial cells (cytokeratin 19, B), Kupffer cells (F4/80, C), endothelial cells (endomucin, D) and hepatic stellate cells (desmin, E) in *Hmgb1*+/+ and *Hmgb1Δhep* mice, showing deletion in hepatocytes and biliary epithelial cells, and preserved expression in Kupffer cells, endothelial cells and hepatic stellate cells of *Hmgb1Δhep* mice (arrows indicate HMGB1 expression in cells positive for each of these markers). F. Western blot analysis reveals strongly reduced HMGB1 expression in the liver, and preserved HMGB1 expression in kidneys, lungs and spleens of *Hmgb1Δhep* mice (n=4 per group). Scale bars 50 µm.
Supplemental Figure 3. HMGB1 status does not affect acetaminophen metabolization, but early neutrophil recruitment to the liver. 

A. Serum HMGB1 levels were determined after treating mice with 300 mg/kg i.p. acetaminophen ("APAP", n=4 per group). B. Hepatic neutrophil infiltration was assessed in Hmgb1\textsuperscript{f/f} (n=7) and Hmgb1\textsuperscript{Δhep} (n=7) mice 3h after treatment with acetaminophen (300 mg/kg). C. Hepatic Cyp2e1 mRNA levels were determined in Hmgb1\textsuperscript{f/f} and Hmgb1\textsuperscript{Δhep} (n=5 per group) mice by qPCR. D. Early hepatic GSH depletion was determined in Hmgb1\textsuperscript{f/f} and Hmgb1\textsuperscript{Δhep} (n=3 per group) mice 60 minutes after treatment with acetaminophen (300mg/kg). E. Hepatic acetaminophen adducts were determined in untreated control mice (n=2) and Hmgb1\textsuperscript{f/f} (n=4) and Hmgb1\textsuperscript{Δhep} (n=4) mice treated with acetaminophen (300 mg/kg) for 3h. F. Acetaminophen adducts were determined in untreated primary hepatocytes (n=2) and primary hepatocytes isolated from Hmgb1\textsuperscript{f/f} (n=2) and Hmgb1\textsuperscript{Δhep} (n=2; representative results from three separate isolations), after treatment with acetaminophen (10 mM) for 3h. Statistical significance assessed via one-way ANOVA followed by Tukey’s Multiple Comparison Test (A, B, D), or two-tailed unpaired t-test (C), respectively. * p<0.05, ** p<0.01. n.s.= non-significant. Scale bars 200 µm (B).
Supplemental Figure 4. Chemokines are not regulated by HMGB1 status at early time points of hepatic injury, and do not regulate neutrophil migration towards necrotic liver tissue. A. Hepatic expression of Ccl2 and Cxcl2 mRNA was assessed in Hmgb1fl/fl (n=7) and Hmgb1Δhep (n=7) mice 3h after treatment with acetaminophen (300 mg/kg). B. Neutrophil migration towards necrotic liver extracts was determined in the presence or AMD3100, an inhibitor of CXCL2 receptor CXCR4, or vehicle (PBS), or a neutralizing antibody against CXCL2 or isotype-matched control antibody (n=3 per group, shown is a representative of three independent experiments, each performed in triplicate). Statistical significance assessed via one-way ANOVA followed by Tukey’s Multiple Comparison Test. *** p<0.001. n.s. = non-significant.
Supplemental Figure 5. HMGB1 in bone marrow-derived cells does not alter acetaminophen-induced inflammation and injury. A-D. Vav1Cre-negative \( Hmgb1^{ff} \) mice (\( n=5 \)) and Vav1Cre-positive \( Hmgb1^{ff} \) mice (\( Hmgb1^{\Delta BM} \), \( n=7 \)) were injected with acetaminophen (300 mg/kg), followed by determination of hepatic neutrophil infiltration (A), hepatic inflammatory gene expression (B), hepatic H&E staining and quantification of necrosis (C) and serum ALT levels (D), 24h later. Statistical significance assessed via two-tailed unpaired t-test (A-D). n.s. = non-significant. Scale bars 200 \( \mu \)m (A, C).
Supplemental Figure 6. RAGE status does not affect acetaminophen metabolism or acetaminophen-induced injury in primary hepatocytes. A. Acetaminophen adducts were determined in untreated primary hepatocytes (n=2) and primary hepatocytes isolated from Ager^{+/+} (n=2) or Ager^{-/-} (n=2, representative result from three separate isolations) mice, after treatment with acetaminophen (10 mM) for 3h. B-C. Primary hepatocytes, isolated from wt or Ager^{-/-} mice (n=2 per group), were treated with various concentrations of acetaminophen for 24h and stained with propidium iodide (red) for cell death determination. Shown are representative images (B) and quantification of acetaminophen-induced cell death (C), representative results from three separate isolations. Statistical significance assessed via two-tailed unpaired t-test (C). n.s.= non-significant. Scale bars 100 µm (B).
Supplemental Figure 7. Reconstitution of hepatic macrophages and spleen by transplantation of Ager<sup>-/-</sup> bone marrow. Reconstitution of mice with Ager<sup>-/-</sup> BM was demonstrated by GFP fluorescence (expressed in Ager<sup>+/+</sup> mice) which overlapped with F4/80 staining in the liver, and GFP fluorescence in the spleen. Scale bars 50 µm.
Supplemental Figure 8. Acetaminophen-induced neutrophil recruitment into the liver does not require macrophages. Mice were treated with vehicle liposomes (n=7) or liposomal clodronate (n=8), followed by injection of acetaminophen (300 mg/kg) three days later. A-B. Livers were stained with F4/80 (A) and Ly6B (B), and the numbers of F4/80-positive macrophages and Ly6B-positive neutrophils, respectively, was quantified. C. Hepatic expression of inflammatory genes was determined by qPCR and normalized to untreated control livers. Statistical significance assessed via one-way ANOVA followed by Tukey’s Multiple Comparison Test (B), or via two-tailed unpaired t-test (C), respectively. ** p<0.01, n.s.=non-significant. Scale bars 200 µm (A, B).
Supplemental Figure 9. Pharmacologic HMGB1 inhibition ameliorates acetaminophen-induced liver injury. A-C. Mice were pretreated with glycyrrhizin (50 mg/kg, n=9) or vehicle (n=8) followed by injection of acetaminophen (300 mg/kg) 2h later. Mice were sacrificed 24h later, followed by determination of hepatic neutrophil infiltration (A), hepatic H&E staining and quantification of necrosis (B) and serum ALT levels (C). D-E. Mice were injected with acetaminophen (300 mg/kg), followed by treatment with either glycyrrhizin (50 mg/kg, n=8) or vehicle (NaCl, n=7) 2h later. Mice were sacrificed 24h after acetaminophen injection, followed by hepatic H&E staining and quantification of necrosis (D) and determination of serum ALT levels (E). Statistical significance assessed via two-tailed unpaired t-test (A-E). *p<0.05, **p<0.01, n.s.=non-significant. Scale bars 200 µm (A, B, D).
Supplemental Figure 10. Human neutrophils require HMGB1 for migration towards tissue lysates. Migration of human neutrophils, isolated from peripheral blood, towards towards $Hmgb1$-floxed and $Hmgb1$-deleted ($Hmgb1^{del}$) liver extracts (induced by Mx1Cre), was determined in Boyden chambers. Statistical significance assessed via one-way ANOVA followed by Tukey’s Multiple Comparison Test. ** p<0.01.
Supplemental Figure 11. HMGB1 promotes neutrophil migration towards cardiac tissue extracts. Neutrophil migration towards Hmgb1-floxed and Hmgb1-deleted (Hmgb1del) cardiac tissue extracts (achieved by MHC-Cre-mediated deletion; insert showing immunoblot confirming HMGB1 deletion) was determined in a Boyden chamber. Statistical significance assessed via one-way ANOVA followed by Tukey's Multiple Comparison Test. ** p<0.01.
All Western blot exposed on Kodak Image station (digital)

FULL UNEDITED GEL FOR Fig.2A

FULL UNEDITED GEL FOR Fig.3A

FULL UNEDITED GEL FOR Fig.3B

FULL UNEDITED GELS FOR Suppl. Fig.1A

FULL UNEDITED GELS FOR Suppl. Fig.1D