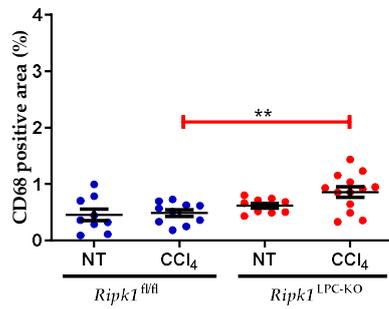
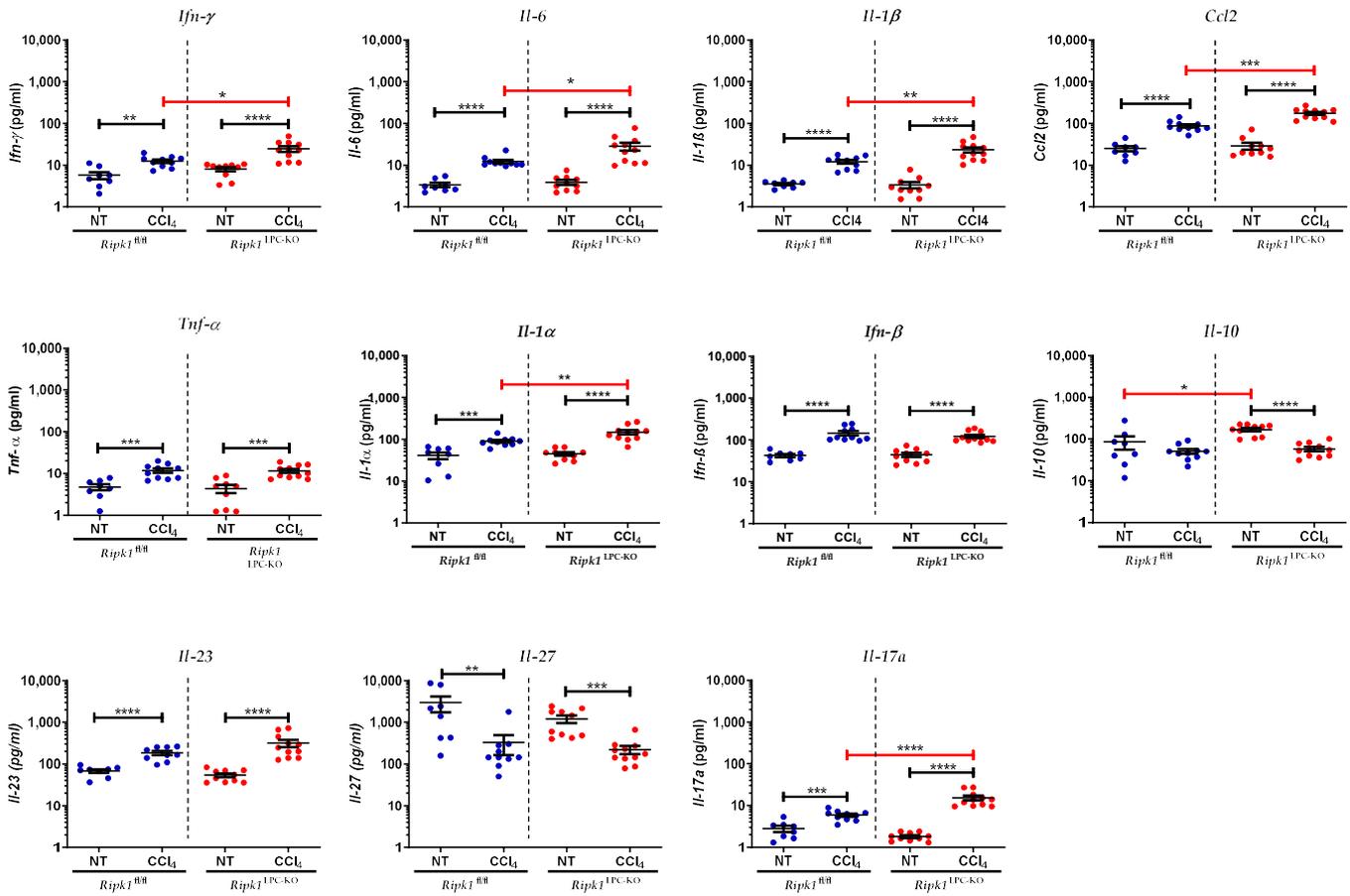


## Figure S1



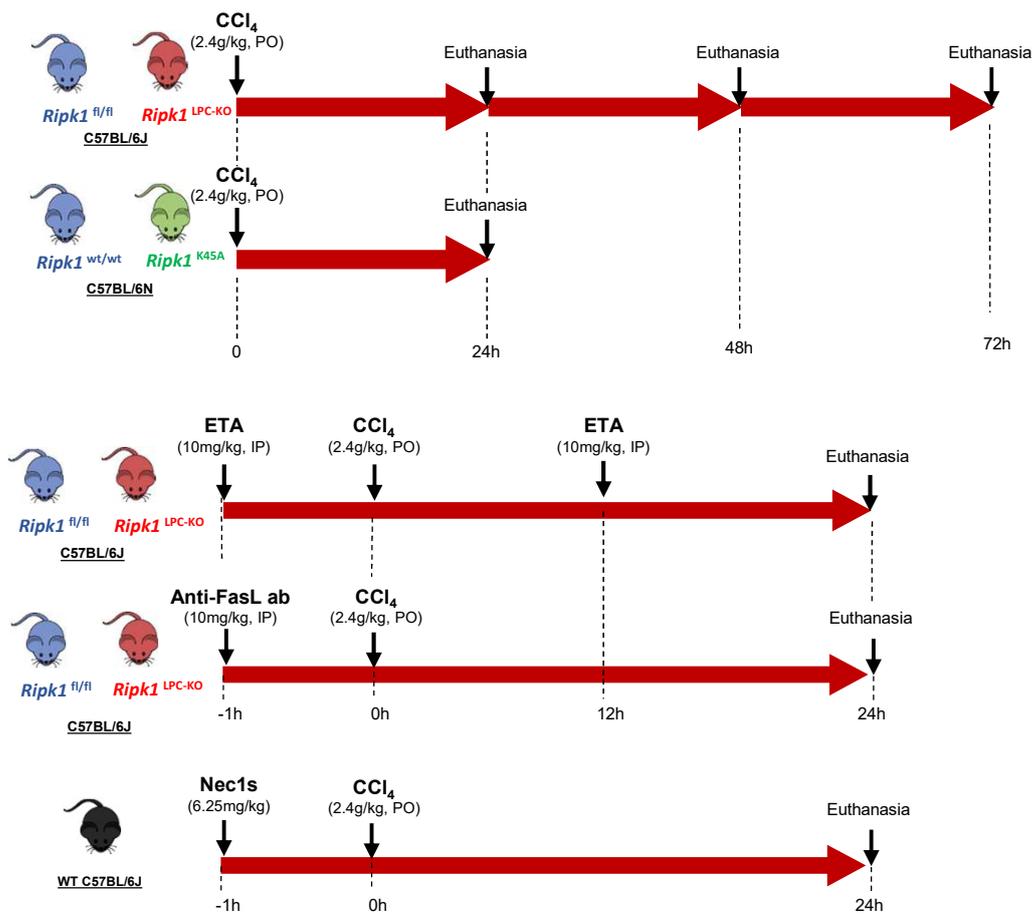
**Figure S1: Macrophage infiltration analysis in non-treated and CCl<sub>4</sub>-treated *Ripk1<sup>fl/fl</sup>* and *Ripk1<sup>LPC-KO</sup>* mice.** *Ripk1<sup>fl/fl</sup>* and *Ripk1<sup>LPC-KO</sup>* mice were force-fed either with olive oil alone (control; NT for non-treated) or containing CCl<sub>4</sub> (dose of 2.4 g/kg body weight), 24 h before sample collection. Signal quantification of anti-CD68 positive area. For the graph, each dot represents an individual and errors bars are expressed as means ± SEM (\*\* p < 0.01).

## Figure S2



**Figure S2: Serum levels of cytokines in non-treated and CCl<sub>4</sub>-treated *Ripk1<sup>fl/fl</sup>* and *Ripk1<sup>LPC-KO</sup>* mice.** *Ripk1<sup>fl/fl</sup>* and *Ripk1<sup>LPC-KO</sup>* mice were force-fed either with olive oil alone (control; NT for non-treated) or containing CCl<sub>4</sub> (dose of 2.4 g/kg body weight), 24 h before serum collection. Murine cytokines were captured by a bead-based immunoassay (BioLegend's LEGENDplex™ Mouse Inflammation Panel, multi-analyte flow assay kit) and quantified on a LSR X-20 Fortessa™ flow cytometer (BD Biosciences). For all graphs, each dot represents an individual and errors bars are expressed as means ± SEM (\* p < 0.05; \*\* p < 0.01; \*\*\* p < 0.001 and \*\*\*\* p < 0.0001).

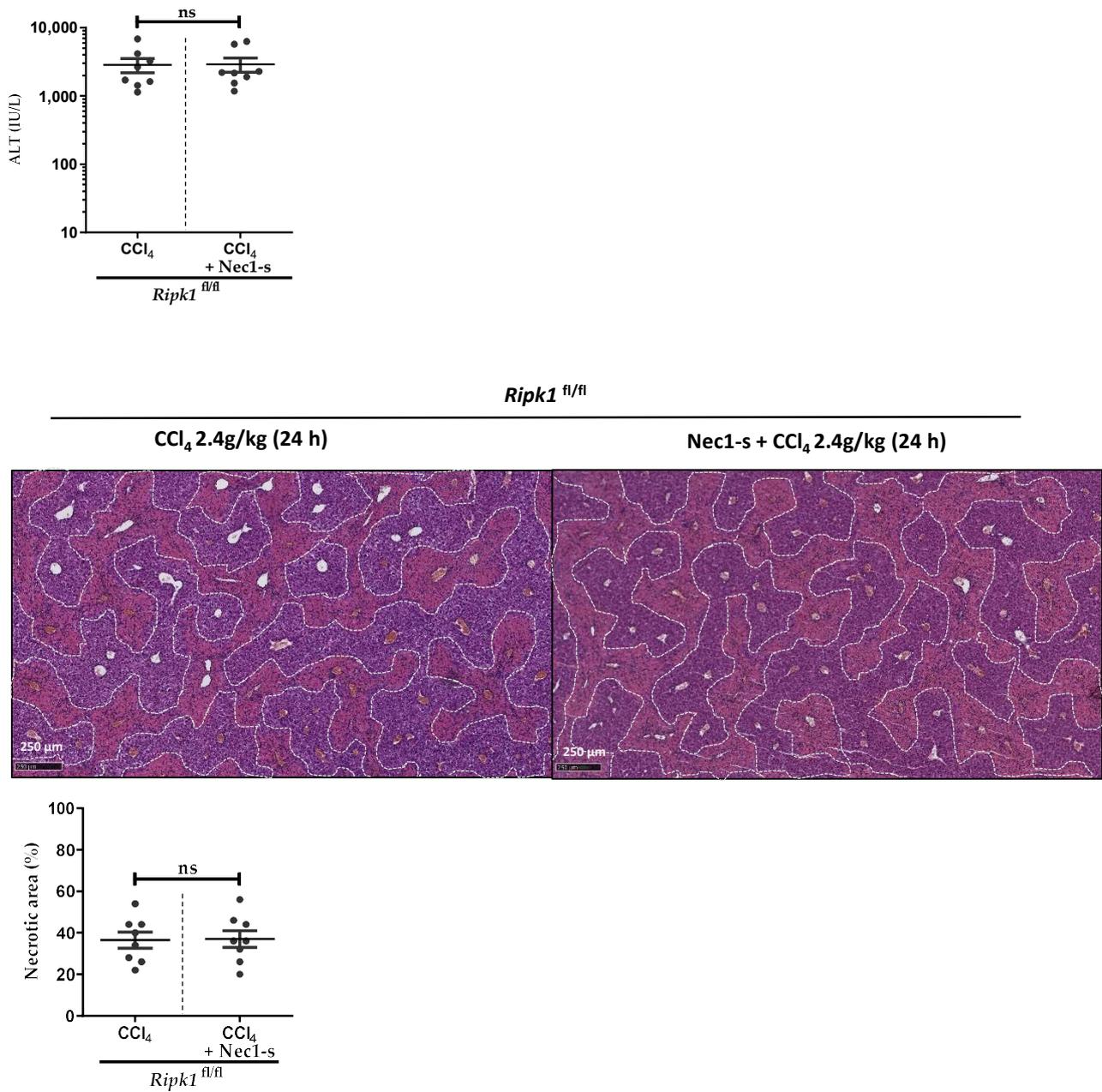
**Figure S3**



LPC : Liver parenchymal cells  
 K45A : Kinase Dead (KD)

**Figure S3: Experimental design.** *Ripk1*<sup>LPC-KO</sup> (red; LPC-KO, Liver Parenchymal Cell – Knock Out) and *Ripk1*<sup>K45A</sup> (green; Kinase Dead, KD) C57BL/6 mice along with their respective WT littermates (blue; *Ripk1*<sup>fl/fl</sup> [fl, floxed] and *Ripk1*<sup>wt/wt</sup> [wt, wild-type]) were force-fed with olive oil alone (non-treated, NT) or containing CCl<sub>4</sub> with a final administered dose of 2.4g/kg (PO, Per Os). When used, etanercept (ETA) or anti-FasL antibody (ab) or vehicle (PBS) were intraperitoneally (IP) injected at 10 mg/kg, 1 hour before CCl<sub>4</sub> treatment, and 12 hour later for a second dose of ETA. In another experiment, Nec-1s or vehicle (PBS-DMSO 6%) was intravenously injected at 6.25 mg/kg 1 hour before CCl<sub>4</sub> treatment. Analysis were conducted 12, 24, 48 or 72 hours post CCl<sub>4</sub> administration as shown in the diagram.

## Figure S4



**Figure S4: No contribution of RIPK1 kinase activity during CCl<sub>4</sub>-induced hepatotoxicity.** C57BL/6J *Ripk1*<sup>fl/fl</sup> mice were forced with olive oil containing CCl<sub>4</sub> (dose of 2.4 g/kg body weight), 24 hours before sample collection. Levels of serum alanine aminotransferase (ALT). Representative pictures of liver tissue sections stained by H&E with necrotic area quantifications. For all graphs, each dot represents an individual and errors bars are expressed as means ±SEM (ns, non-significant).

**Table S1**

Mouse Gene	Forward	Reverse
18S	5'-CGCCGCTAGAGGTGAAATTC-3'	5'-TTGGCAAATGCTTTCGCTC-3'
<i>Hmox-1</i>	5'-AGGTACACATCCAAGCCGAGA-3'	5'-CATCACCAGCTTAAAGCCTTCT-3'
<i>Nfe2l2</i> (Nrf-2)	5'-TAGATGACCATGAGTCGCTTGC-3'	5'-GCCAACTTGCTCCATGTCC-3'
<i>Cybb</i> (Nox-2)	5'-AGTGCGTGTGCTCGACAA-3'	5'-GCGGTGTGCAGTGCTATCAT-3'
<i>Nqo1</i>	5'-AGGATGGGAGGTACTIONCGAATC-3'	5'-AGGCGTCCTTCTTATATGCTA-3'
<i>Il-6</i>	5'-CGATGATGCACTTGCAGA-3'	5'-CTCTGAAGGACTCTGGCTTTG-3'
<i>Ifn-γ</i>	5'-AAGACAATCAGGCCATCAGC-3'	5'-CAGCAGCGACTCCTTTTCC-3'
<i>Il-1b</i>	5'-GCCACCTTTTGACAGTGATGAG-3'	5'-GACAGCCCAGGTCAAAGTT-3'
<i>Ccl2</i>	5'-TCCCAATGAGTAGGCTGGAG-3'	5'-TCTGGACCCATTCTTCTTG-3'
<i>Rela</i> (NFκBp65)	5'-GACCCCTGCTCTCACATCCG-3'	5'-CAGCTCCAGAGTCCGGTT-3'
<i>Nfkbia</i> (IκBα)	5'-TGAAGGACGAGGAGTACGAGC-3'	5'-TTCGTGGATGATTGCCAAGTG-3'
<i>Tnfaip3</i> (A20)	5'-TGCCCAGTCTGTAGTCTTCG-3'	5'-AGTTGTTAGCCATGGTCCT-3'
<i>Cflar</i> (cFLIP)	5'-GAGGCCAGCTCTCTTTTGCT-3'	5'-TCCACGGTTGCTTTGTCTGT-3'
<i>Ikkb-γ</i> (Nemo)	5'-GGTGGAGAGACTGAGCTTGG-3'	5'-CCTCTAAAGCTTGCCGATCC-3'
<i>Saa1</i>	5'-TGTTACGAGGCTTTCCAAG-3'	5'-GTCCTTGCCGAAGAATTCC-3'
<i>Cxcl1</i>	5'-CGCCTATCGCCAATGAGC-3'	5'-GAACCAAGGGAGCTTCAGG-3'
<i>Ccl20</i>	5'-TCTGCTTCTTCTGCTTTGG-3'	5'-TCACCCAGTTCTGCTTTG-3'
<i>Tnf-α</i>	5'-TAGTCCCAGAAAAGCAAGC-3'	5'-TTTTCTGGAGGGAGATGTGG-3'
<i>Tnfrsf1a</i> (TNFR1)	5'-ATCGAGAGGCTGGAGATGC-3'	5'-GGCGGGATTTCTCAGAGC-3'
<i>Tnfrsf1b</i> (TNFR2)	5'-CGCTGGTCTTCAACTGC-3'	5'-CAGGAGGACACTTAGCACAGC-3'
<i>Fas</i>	5'-CTCCGAGTTAAAGCTGAGG-3'	5'-TGTACTIONTCCCTTCTGTGC-3'
<i>Fasl</i>	5'-GCAGCAGCCCATGAATTACC-3'	5'-GGTAATTCATGGGCTGCTGC-3'
<i>Tnfsf10</i> (TRAIL)	5'-TGAGAACCTTTCAGGACACC-3'	5'-GGCCTAAGGTCTTCCATCC-3'
<i>Tnfrsf10b</i> (TRAIL-R2)	5'-CCGGGCAGATCACTACACC-3'	5'-AGACAGGGTCTCTTGATGG-3'

**Table S1: Primer sequences**