

Supplementary figures

Direct antiviral properties of TLR ligands against HBV replication in immune-competent hepatocytes

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\$ equal contribution

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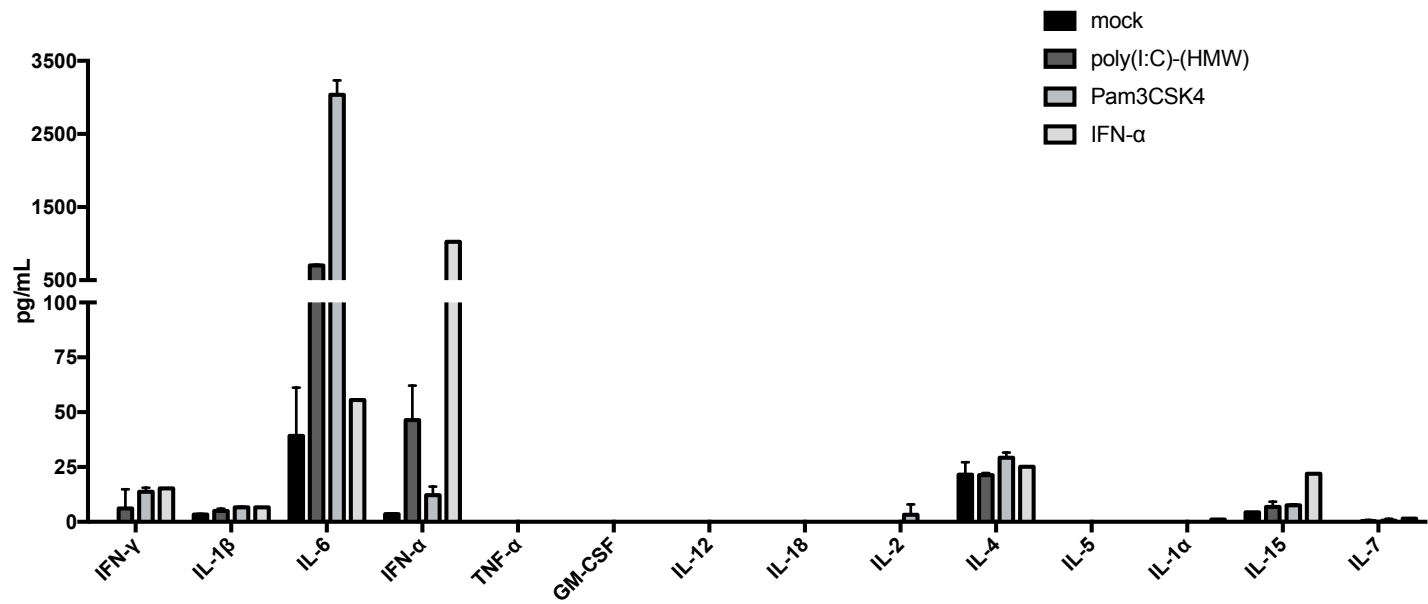
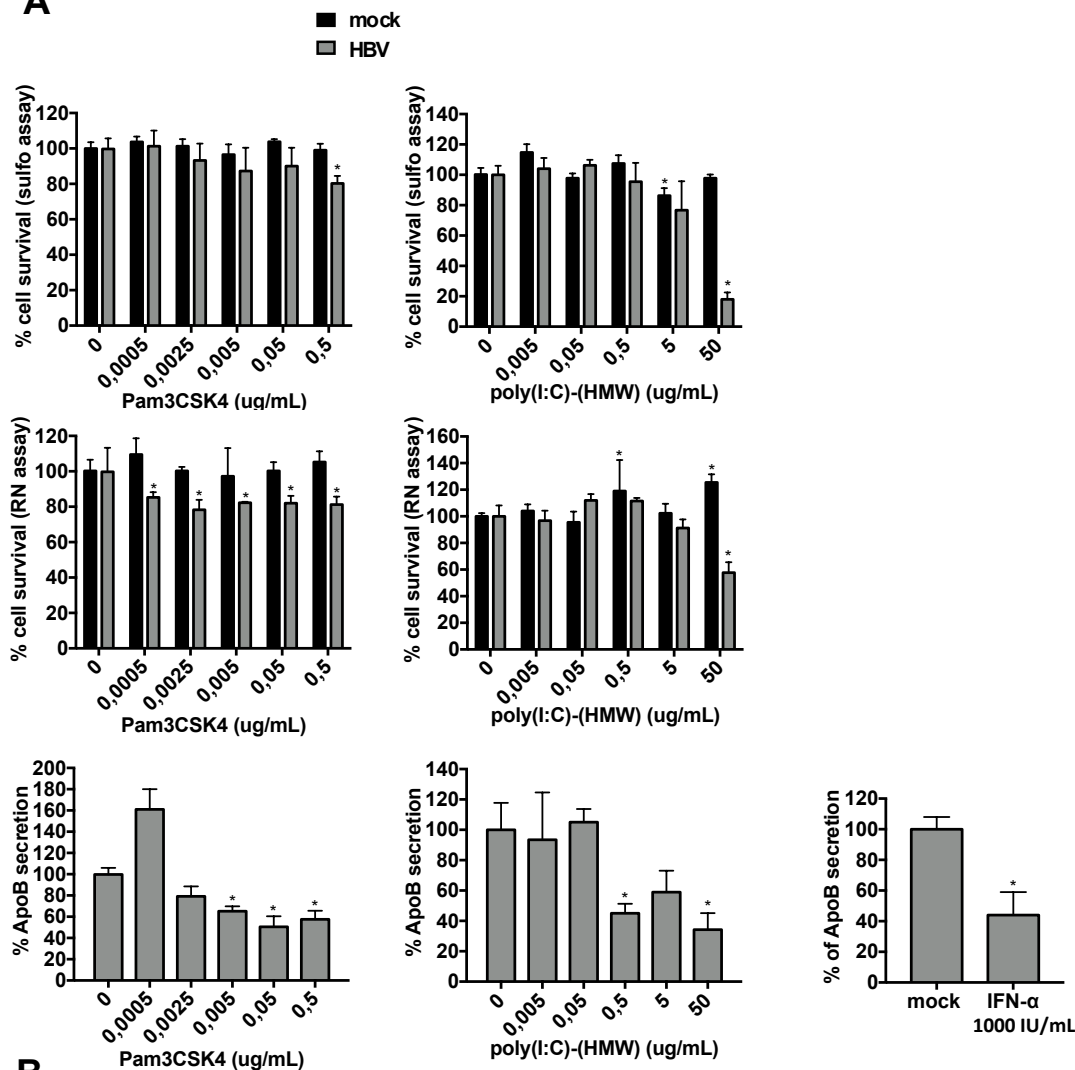


Figure S1: **Cytokines secretion by dHepaRG stimulated with poly(I:C)-(HMW) or Pam3CSK4.** dHepaRG cells were infected by HBV at a multiplicity of infection of 100 viral genome equivalent (vge)/cell for seven days, treated or not with Pam3CSK4 (0,5 ug/mL), poly(I:C)-(HMW) (5 ug/mL) or PEG-IFN-α (1000 IU/mL) for 24h. Supernatants were collected and amount of the indicated cytokines were measured by Luminex assay. Results are the mean +/- SD one experiment performed with two biological replicate.

dHepaRG

A



B

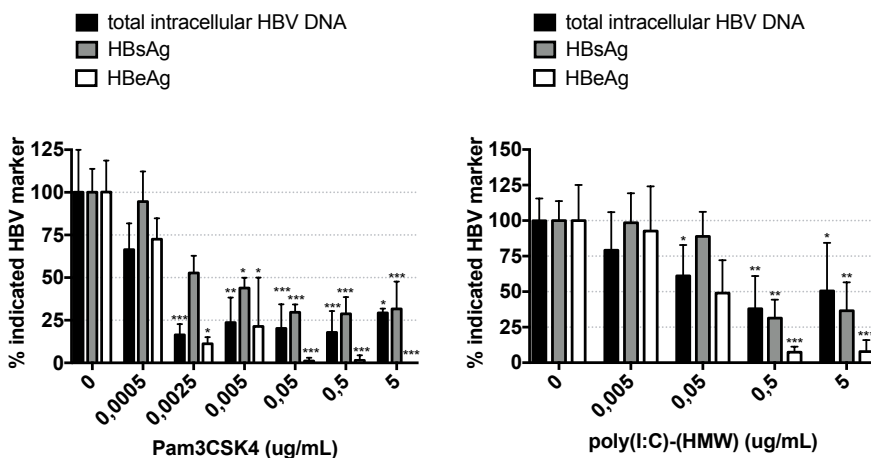
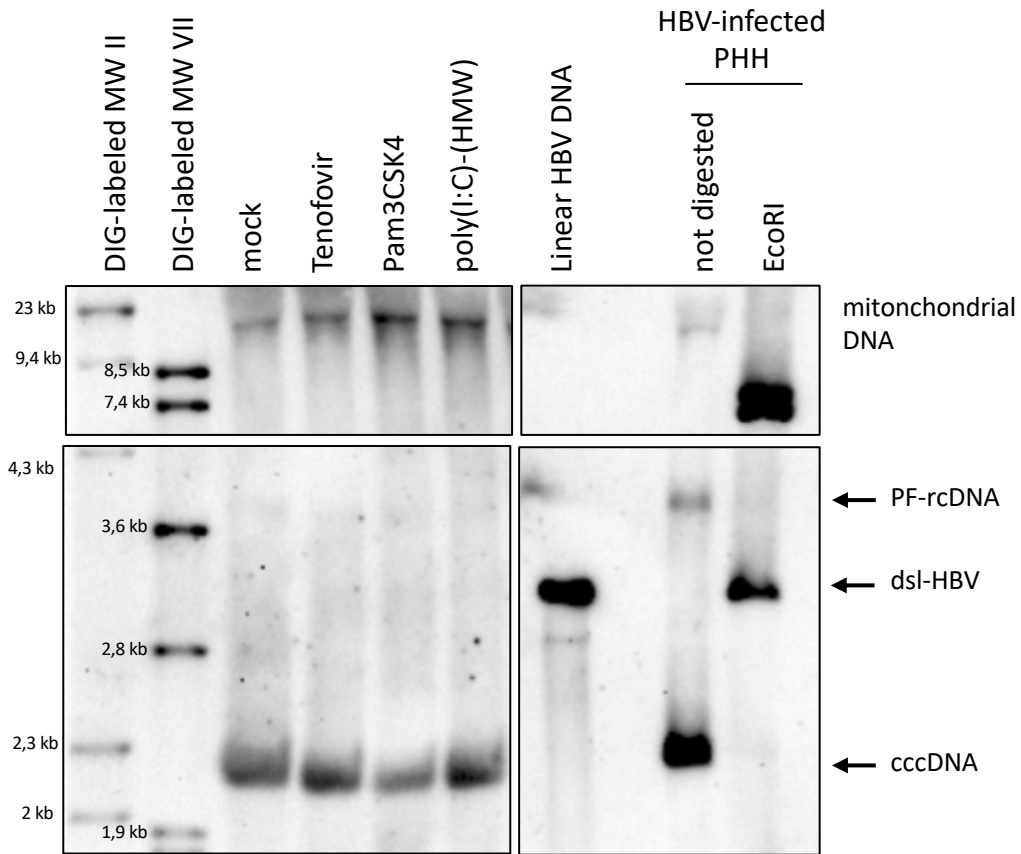


Figure S2: **Pam3CSK4 and poly(I:C)-(HMW) inhibit HBV in HBV-infected dHepaRG in a dose dependent manner without toxicity.** dHepaRG were infected by HBV at a multiplicity of infection of 100 viral genome equivalent (vge)/cell for seven days and treated with the indicated molecules at the indicated concentrations for seven more days (2 treatments). At the end of the experiments, (A) cells survival rates was assessed by sulforhodamin or neutral red assays, (B) total DNAs were extracted and total intracellular HBV DNA amounts were evaluated by qPCR analyses. Supernatant were collected and (A) ApoB, (B) HBeAg or HBsAg levels were assessed by ELISA. Results are the mean +/- standard deviation (SD) of three independent experiments each performed with three biological replicate.

dHepaRG

A



B cccDNA quantification by Southern Blot on Hirt DNA extraction

C cccDNA quantification by qPCR on total DNA extractions

D Total intracellular HBV DNA quantification by qPCR on total DNA extractions

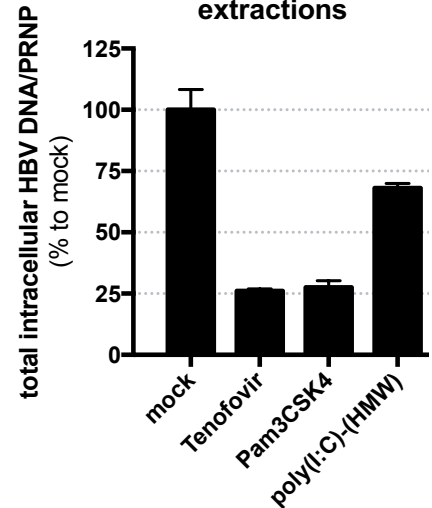
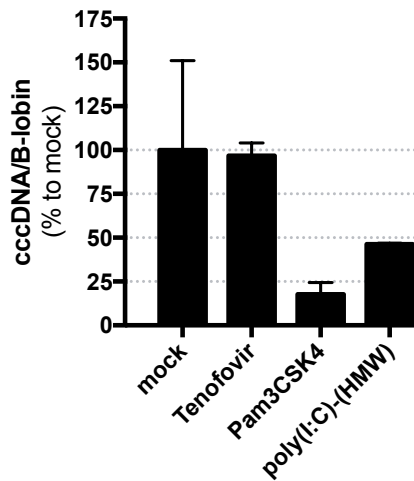
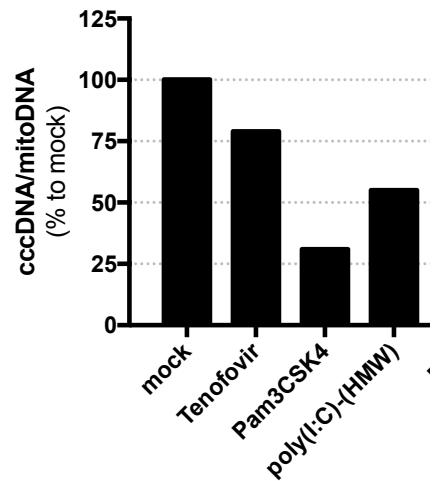
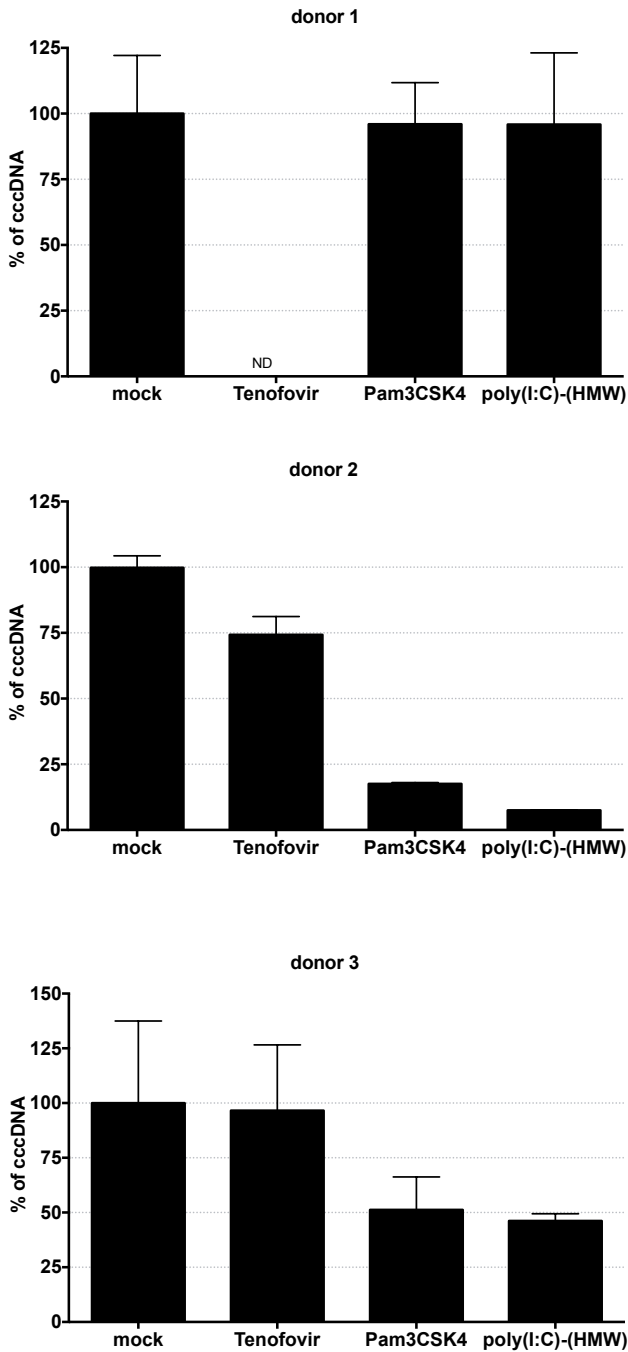


Figure S3: Pam3CSK4 and poly(I:C)-(HMW) treatments decreased cccDNA levels in HBV-infected dHepaRG. dHepaRG were infected by HBV at a multiplicity of infection of 100 viral genome equivalent (vge)/cell for seven days and treated with Tenofovir (10 μ M), Pam3CSK4 (0,5 μ g/mL) or poly(I:C)-(HMW) (0,5 μ g/mL) for seven more days (2 treatments). At the end of the experiments, cells were harvested, (A, B) DNAs were extracted following a Hirt procedure or (C, D) a total DNA extraction procedure and submitted to (A) Southern blot analyses using HBV-DIG labeled probes or (C, D) qPCR analyses to detect (C) cccDNA or (D) total intracellular HBV DNAs. (B) cccDNA and mitochondrial DNA were quantified on the southern blot analyses using the « ImageLab » software from Bio-Rad.

PHH

A



B

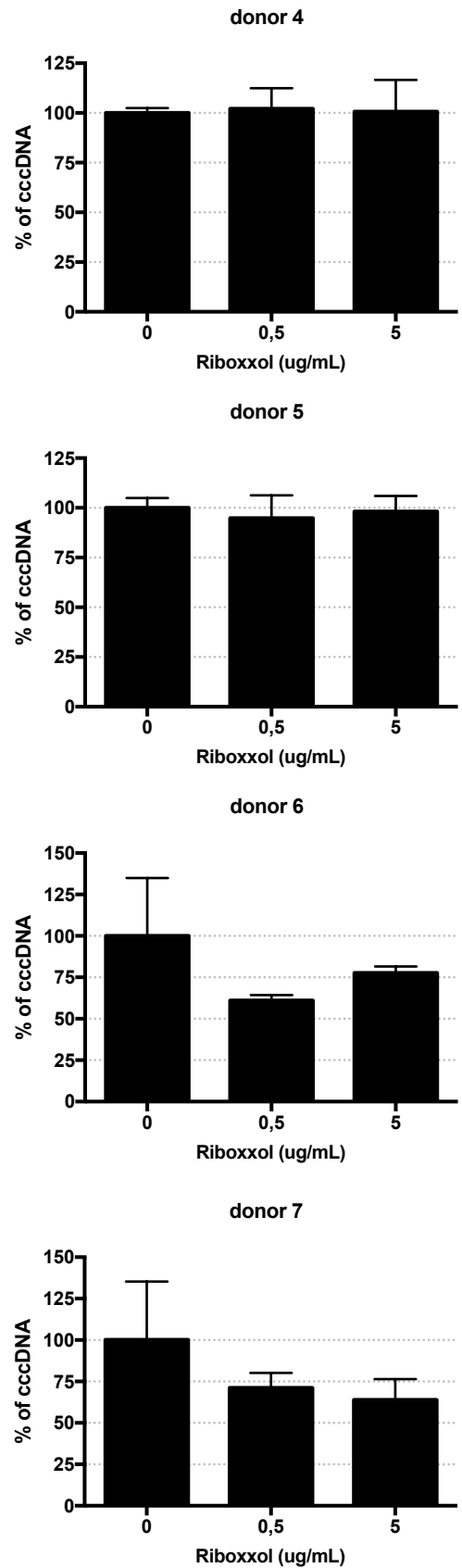


Figure S4: **Effect of Pam3CSK4, poly(I:C)-(HMW) and riboxsol treatments on cccDNA levels in HBV-infected PHH.** PHH were infected by HBV at a multiplicity of infection of 100 viral genome equivalent (vge)/cell for four days and treated with Tenofovir (10 uM), Pam3CSK4 (0,5 ug/mL), poly(I:C)-(HMW) (0,5 ug/mL) or Riboxsol (indicated concentrations) for seven more days (2 treatments). At the end of the experiments, cells were harvest, total DNAs were extracted and cccDNA amounts were evaluated by qPCR analyses. For each donor, results are the mean +/- SD one experiment performed with three biological replicate.

dHepaRG

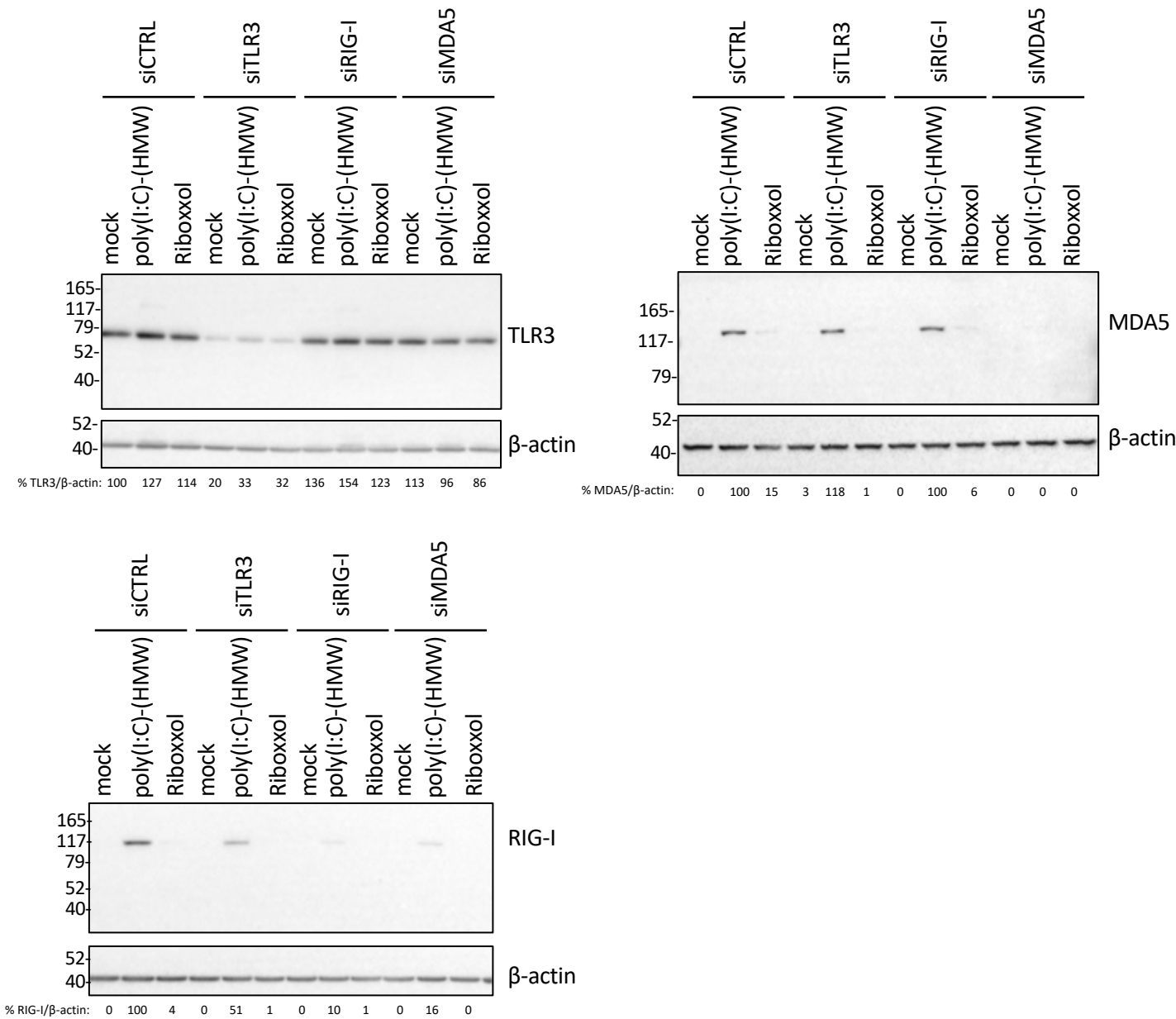


Figure S5: TLR3, MDA5 and RIG-I protein levels after siRNA transfection. Uninfected dHepaRG cells were transfected with the indicated siRNA. 14 days later, cells were stimulated with 10 μ g/ml of the indicated ligands. Cells were lysed 24h later and TLR3, MDA5 or RIG-I protein levels were analyzed by western blot. Blots presented here are representative of three independent experiments.

dHepaRG

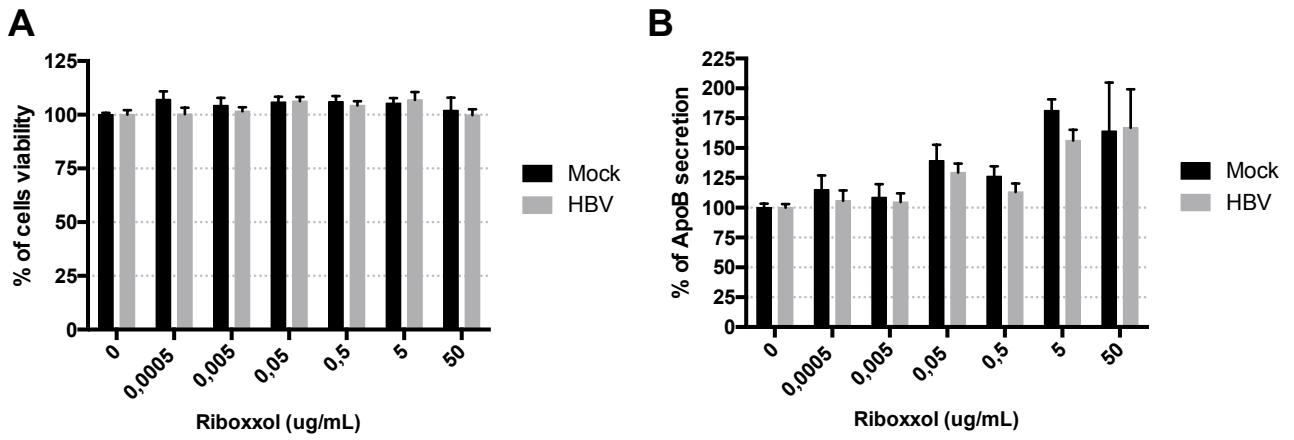


Figure S6: **Treatments with Riboxzol did not impact on dHepaRG survival and differentiation status.** dHepaRG were infected by HBV at a multiplicity of infection of 100 viral genome equivalent (vge)/cell for seven days and treated with the riboxzol at the indicated concentrations for seven more days (2 treatments). At the end of the experiments, (A) cells survival rates was assessed by sulforhodamin assay, (B) supernatant were collected and ApoB levels were assessed by ELISA. Results are the mean +/- standard deviation (SD) of three independent experiments each performed with three biological replicate.

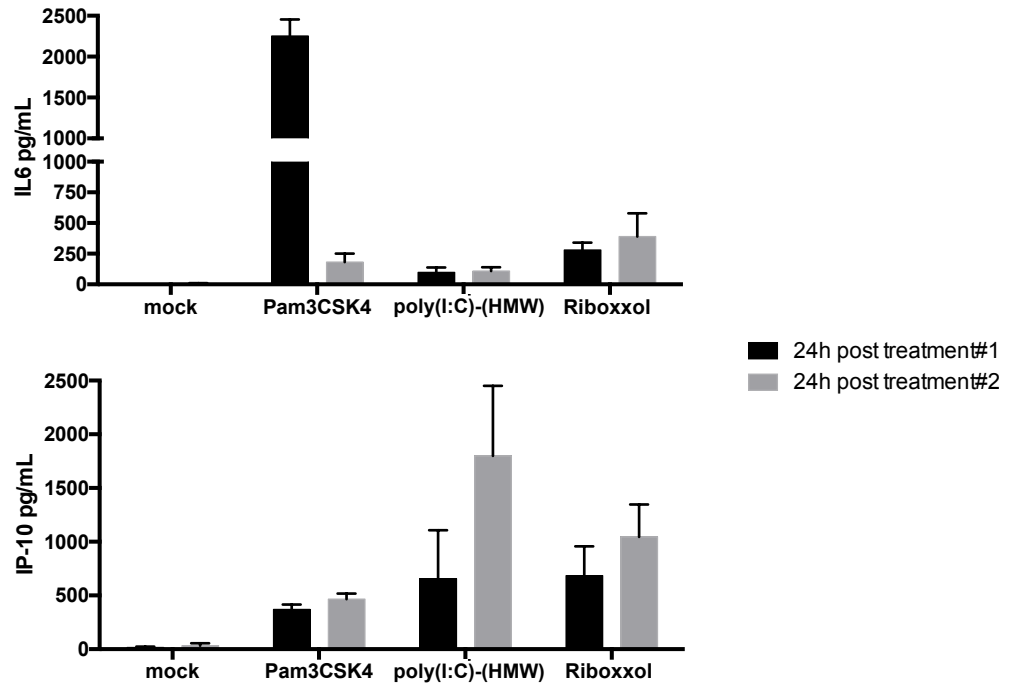


Figure S7: **Riboxxol induced secretion of both IL-6 and IP-10.** dHepaRG cells were infected by HBV at a multiplicity of infection of 100 viral genome equivalent (vge)/cell for seven days, treated or not with Pam3CSK4 (0,5 ug/mL), poly(I:C)-(HMW) (5 ug/mL) or Riboxxol (50 ug/mL) for seven days (2 treatments). Supernatants were collected and amount of IL6 and IP-10 were quantified by ELISA. Results are the mean +/- SD of three independent experiments each performed with three biological replicate.

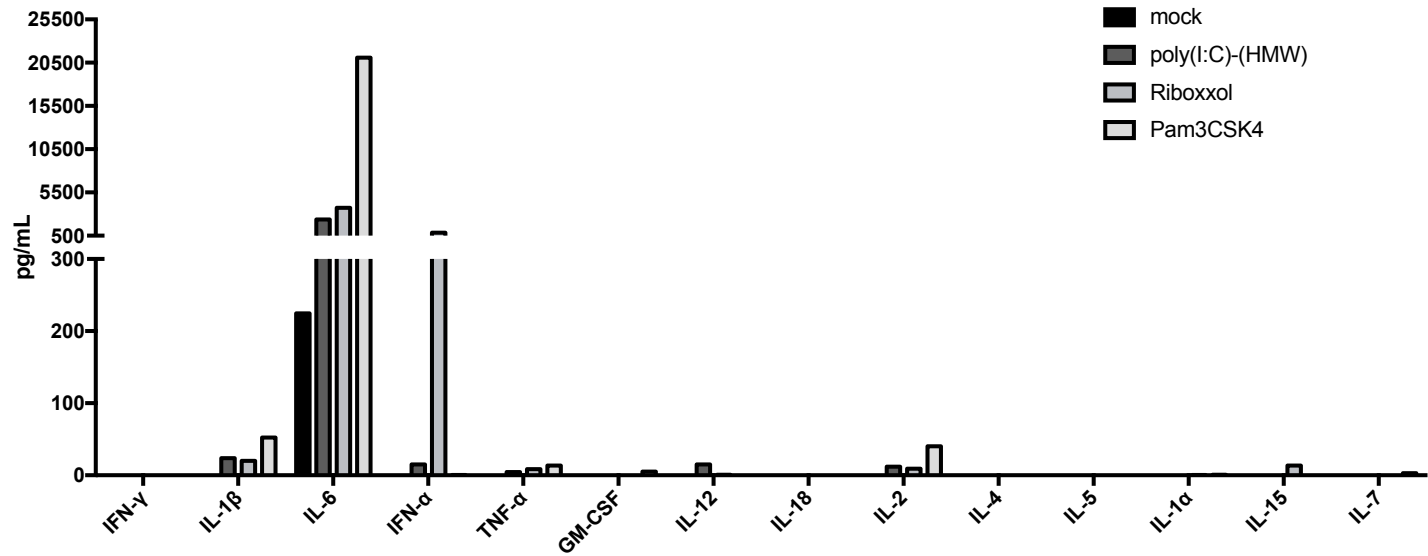
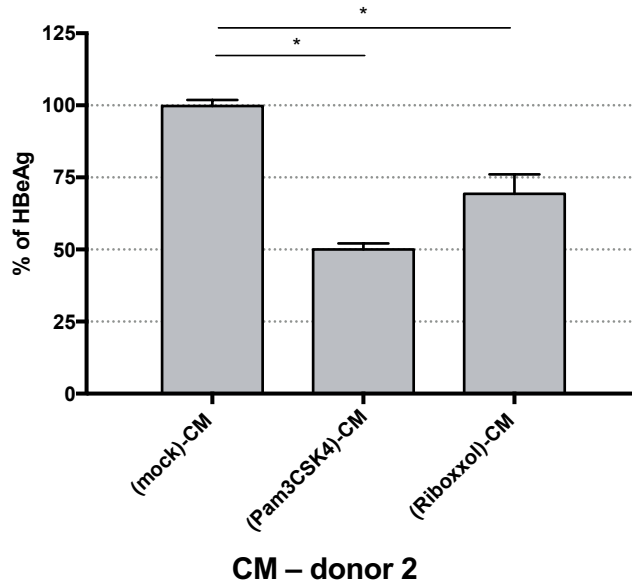
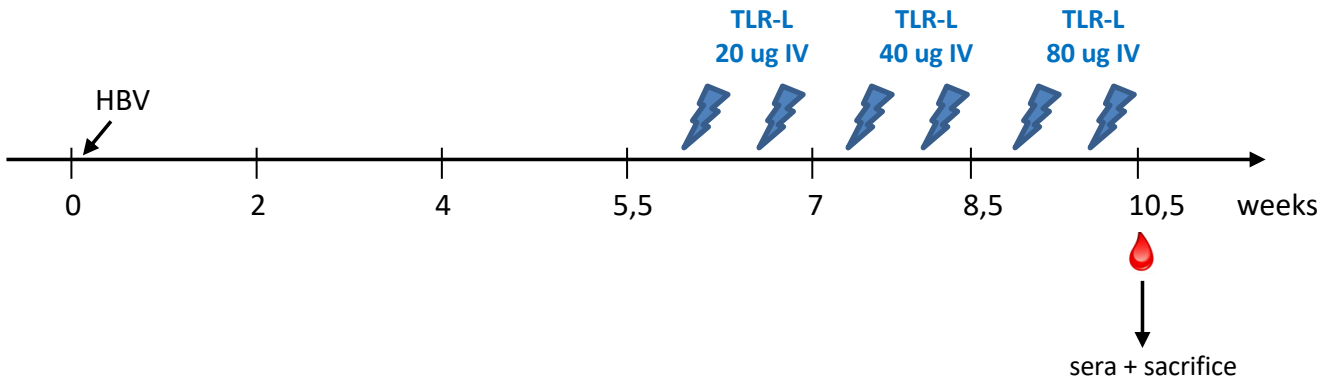
A**PBMC – donor 2****B****dHepaRG**

Figure S8: **TLR2 and 3 ligands lead to production of inflammatory cytokines by immune blood cells.** (A) Fresh PBMC from a healthy donor were cultivated and stimulated or not with Pam3CSK4 (5 $\mu\text{g}/\text{mL}$), poly(I:C)-(HMW) (5 $\mu\text{g}/\text{mL}$) or Riboxzol (5 $\mu\text{g}/\text{mL}$) for 24h. Supernatant were collected and cytokines content was analyzed with Luminex Assay. (B) dHepaRG cells were infected by HBV at a multiplicity of infection of 100 viral genome equivalent (vge)/cell for seven days and treated or not during 10 days with the indicated conditioned media (CM) diluted 1/100. Supernatant were collected and HBeAg levels were assessed by ELISA.

HBV-infected HuHep mice

A



B

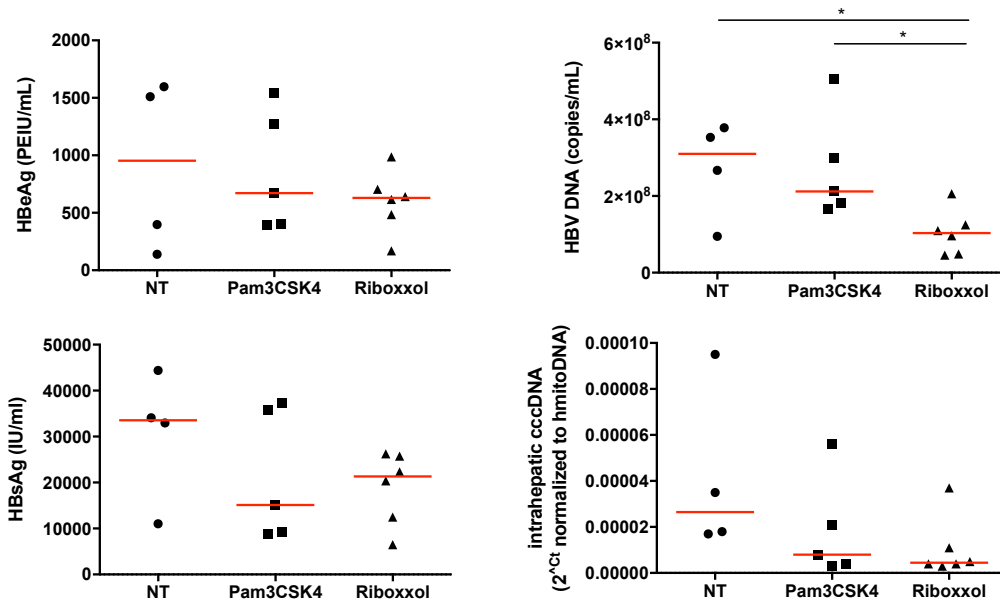


Figure S9: Effect of Pam3CS4K and Riboxsol in HBV-infected HuHep mice. HuHep mice were infected IP with HBV and treated IV with TLR-L (Pam3CSK4 or Riboxsol) or not (NT; no treatment) (A) according to the scheme. Mice were sacrificed after 3 weeks of treatment and (B) levels of HBV markers (HBeAg, HBSAg, viremia and intrahepatic HBV cccDNA) were assessed by ELISA, qPCR or RT-qPCR. Each dot represents one animal and red lines the median from the 4-5 animals.

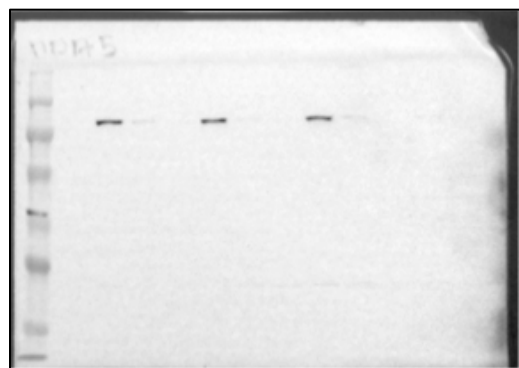
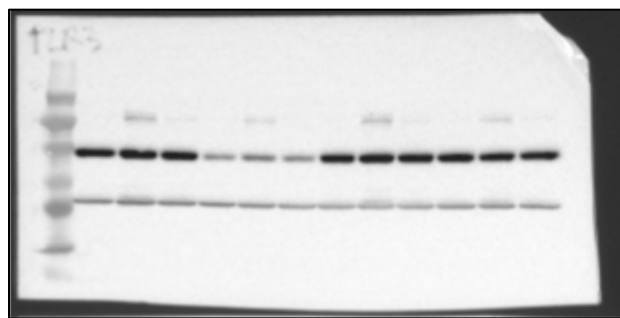
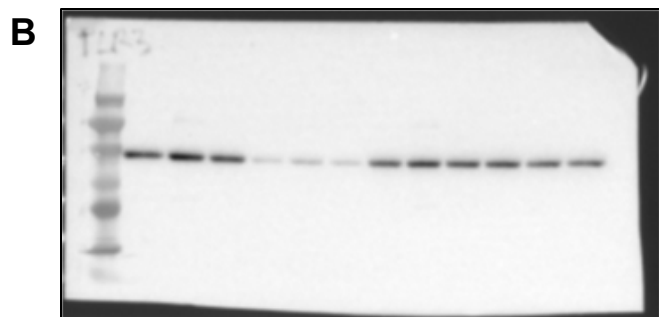
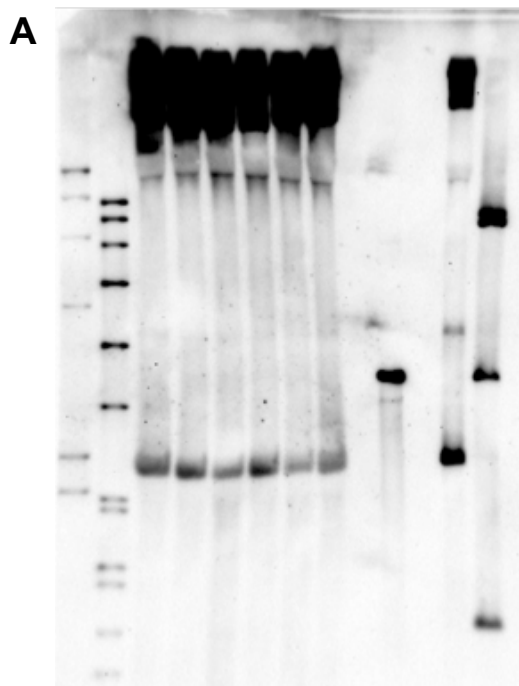


Figure S11: **Full blots** from (A) fig S3A, (B) fig S5