T Cells Expressing a Chimeric Antigen Receptor That Binds 1 Hepatitis B Virus Envelope Proteins Control Virus Replication in 2 Mice 3

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Supplemental Figures 5

Figure S1 6

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S-CAR IL-12

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Figure S1. Comparison of IL-2 and IL-12 preconditioning of T cells before adoptive transfer. 9 Stimulation and transduction was carried out either in the presence of IL-2 for total 10 splenocytes or IL-12 for positively selected CD8⁺ T cells. (A) Cell surface expression of the 11 S-CAR on primary murine CD4⁺ (upper row) and CD8⁺ (lower row) T cells was quantified by 12 flow cytometry. n.d.= not determined. (B) Recovery of transferred cells (CD45.1⁺) from blood 13 and liver 6 days after transfer of 3x10⁶ CAR⁺ T cells into HBVtg mice. Cells from 4 mice per 14 15 group were pooled and numbers of transferred cells are given relative to total isolated mononuclear cells. 16



Figure S2. Adoptive transfer of IL-12 stimulated and CAR-grafted CD8⁺ T cells. (A) Schematic representation of the S Δ -CAR, which combines the extracellular part of the S-CAR with the intracellular domain of the nerve growth factor receptor (NGFR). (B) Cell surface expression of CARs on murine CD8⁺ T cells quantified by flow cytometry. Numbers at the bottom indicate the percentage of CD8⁺ T cells; numbers at the top indicate the percentage of CAR-expressing cells. (C-H) Characterization of leukocytes in the liver analyzed by histological staining on day 12 after adoptive T cell transfer. Arrows indicate single positive cells. Tissue was cryo-preserved (D,E) or paraffin-embedded (C,F,G,H). F4/80 stains macrophages, B220 stains B-cells, Ki-67 stains proliferating cells. Scale bars: 100 µm.

35 Figure S3



Figure S3. PD-1 expression and IFN-γ response to *ex vivo* antigen stimulation of adoptively
transferred CD8⁺ T cells. Cells were stimulated with plate-bound, recombinant HBsAg for 20h
and subsequently stained for CD8, CD45.1, S-CAR, PD-1 and IFN-γ. Left: CD8⁺ T cells
before transfer. Right: Liver associated lymphocytes isolated at d12 post transfer from HBVtg
or wild-type C57BL/6 recipient mice. Cells were first gated on CD45.1⁺ CD8⁺ transferred T
cells and then on expression of S-CAR as indicated on the X-axis. Data represent mean
values, n=4. LAL from C57BL/6 mice were pooled for analysis.



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Figure S4. Preparation of S-CAR-grafted and natural HBV-specific CD8⁺ T cells. (*A*) Cell surface expression of S-CAR on murine CD8⁺ T cells quantified by flow cytometry. Numbers indicate the percentage of CAR-expressing cells. (*B*) CD45.1⁺ mice were immunized with HBsAg and MVA-S and intracellular cytokine staining assessed numbers of S-specific T cells. Four representative FACS dot plots show IFN- γ and TNF- α production of CD8⁺ T cells after stimulation with HBV S190 and S208 peptides, an irrelevant peptide or no peptide. (*C*) HBV-specific cytokine production of CD8⁺ T cells after immunization.

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62 Supplemental Methods

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Cultivation of hepatoma cells. Hepatoma cells were kept in full DMEM medium (DMEM, 10 64 65 % FCS, 1 % Pen/Strep, 1 % sodium pyruvate, 1 % NEAA; Life Technologies, Darmstadt, Germany). For co-culture experiments 5x10⁴ target cells per well were seeded in collagen-66 67 coated (Serva, Heidelberg, Germany) 96-well flat bottom plates. When cells had reached confluence, differentiation medium was applied (Williams medium, 5 % FCS, 1 % Pen/Strep, 68 69 1 % sodium pyruvate, 1 % NEAA (Life Technologies) and 0.5 % DMSO (Sigma-Aldrich, 70 Taufkirchen, Germany). Target cells were used for experiments after 10 to 14 days of differentiation. 71

Cultivation of murine cells. Primary murine cells were maintained in murine T cell medium
(RPMI dutch modified, 10 % FCS, 1 % glutamine, 1 % Pen/Strep, 1 % sodium pyruvate and
50 μM β-mercaptoethanol; Life Technologies).

IL-12 production. Murine recombinant IL-12 was produced by using a single polycistronic
 vector containing IL12p40 and IL-12p35 obtained from Dr. W.J. Storkus (1) and subsequently
 affinity purified.

Co-culture with hepatoma cell lines and HBsAg stimulation. The number of effector T cells/well was adjusted according to the transduction efficiency of each receptor to identical numbers of CAR-expressing cells. 5-10 μ g /ml HBsAg (kindly provided by RheinBiotech-Dynavax, Düsseldorf, Germany) was coated on 96-well flat bottom plates for 7 h at RT or overnight at 4°C. Plates were washed with PBS before cells were added. Supernatants of co-cultures and stimulation with recombinant antigen were subjected to a murine IFN- γ ELISA (BD Biosciences).

Proliferation assay. S-CAR-grafted T cells were incubated with 2.5µM Carboxyfluorescein
succinimidyl ester (CFSE, Life Technologies) at 37°C for 10'. Cells were washed two times
and then 5x10⁵ T cells were co-cultured in a 24-well plate with HBV-replicating cell lines

(HepG2.2.15) or the parental HBV-negative cell line HepG2 for 3 days. Dilution of CFSE in
dividing cells was assessed by flow cytometry. Cells were additionally stained with anti-CD8
and anti-human IgG to discriminate between S-CAR positive and S-CAR negative cells.

ALT measurement. Serum alanin amino transferase activity was determined in 32 µl of
serum using the Reflotron system (Roche Diagnostics, Mannheim, Germany).

93 Histopathology. Liver tissue was either embedded in Tissue-Tek O.C.T. (Sakura, Staufen, Germany) and frozen or fixed with 4 % paraformaldehyde and embedded in paraffin. 94 95 Cryo sections (5 µm) were fixed in aceton and stained with various antibodies: rabbit-anti-96 CD3 (Neomarkers, Fremont, USA), rat-anti-B220, rat-anti-CD8a, rat-anti-CD4 (all BD 97 Biosciences, Heidelberg, Germany), mouse-anti-CD45.1 FITC, mouse-anti-CD45.2 FITC 98 (both eBioscience, Frankfurt, Germany), goat-anti-humanlgG FITC (Sigma-Aldrich), rat-anti-99 CD68 (Serotec, Oxford, UK), goat-anti-rabbit Alexa488, goat-anti-rat Alexa594, goat-anti-100 FITC Alexa488 (all Life Technologies), rabbit-anti-rat AP (Bond Polymer Refine Red 101 Detection, Leica, Newcastel Upon Tyne, UK). Paraffin sections (2 µm) were either stained 102 with haematoxylin / eosin or automated immunohistochemistry staining was performed using 103 rabbit-anti-HBcAg (Diagnostic Biosystems, Pleasanton, USA), rat-anti-F4/80 (BMA 104 Biomedicals, Augst, Switzerland), rabbit-anti-CD3, rat-anti-B220 (both BD Biosciences), 105 rabbit-anti-Ki67 (Neomarkers), rabbit-anti-cleaved caspase 3 (Cell Signaling, Danvers, USA), 106 rabbit-anti-rat HRP and anti-rabbit-poly HRP (Leica Bond Polymer Refine Detection 107 DS9800). Image acquisition was performed on the DotSlide BX51 (Olympus), SCN400 108 (Leica), Axio Z1 (Zeiss) or BX53 (Olympus) microscope.

109 RNA extraction and quantification of mRNA levels. Approximately 20 mg of liver tissue 110 was homogenized in 1 ml of Trizol (Life Technologies) using the Ultrathorax T10 (IKA, 111 Staufen, Germany). The RNA was extracted according to the manufacturer's instruction. 112 Residual DNA was DNase digested (Roche Diagnostics, Mannheim, Germany) and 1 µg of 113 RNA was reverse transcribed into cDNA using the Superscript III Kit (Life Technologies). 114 Cytokine expression levels were determined relative to GAPDH expression by a quantitative

115 PCR on a LightCycler480 using SybrGreen (both Roche Diagnostics) and the primers 116 described before (2) or TNF α -fw 5'-ACGTCGTAGCAAACCAC-3' and TNF α -rev 5'-117 AGATAGCAAATCGGCTG-3'. Expression levels were determined relative to GAPDH and 118 normalized to the CEA-CAR control group.

DNA extraction and quantification of viral DNA. Approximately 20 mg of liver tissue was homogenized in 1 ml of Proteinase-K buffer (Tris pH 8.5 100 mM, EDTA 5 mM, SDS 0.2 %, NaCl 200 mM) followed by a phenol-chloroform extraction of the DNA. Intracellular HBV replicative intermediates were detected by Southern blot analysis of 75 µg of total liver DNA digested with HindIII (Fermentas, St. Leon-Rot, Germany) and separated through a 0.85 % agarose gel.

Quantitative PCR was performed using p-fwHBV 5'-TACTAGGAGGCTGTAGGCATA-3' and p-revHBV 5'-GGAGACTCTAAGGCTTCCC-3'. Data were normalized to the cell number determined by the single copy gene *Nid2*; mNid2Fw 5'-ATCCTACCGGGAAACAG-3' and mNid2Rev 5'-CCCAAGGTCTTCGTTGA-3'. Viral DNA from serum was extracted using the High pure viral nucleic acid Kit (Roche Diagnostics) and analyzed using fwHBV and revHBV primers.

131 **Cytokines.** Serum cytokines were measured using the Cytometric Bead Array Mouse 132 Inflammation Kit (BD Biosciences). Briefly, 15 μ I of undiluted or 1:4 diluted serum were 133 incubated with 15 μ I of mixed capture beads and 15 μ I of PE detector for 2 h at RT. 5000 134 beads were acquired on a BD FACS Canto II and analyzed with FCAP Array Software 3.0 135 (BD Biosciences).

Immunoglobulin screening. 100 µl of 1:1200 diluted serum were coated on Nunc Maxisorp
ELISA plates (Sigma) at 4 °C overnight. HRP-labeled detection antibodies (Southern
Biotech, Birmingham, USA) were diluted 1:1000 (anti-IgM and anti-IgG1), 1:500 (anti-IgA)
and 1:250 (anti-IgG2a, anti-IgG2b, anti-IgG3). ELISA was performed according to the manual
of the SBA Clonotyping System/HRP (Southern Biotech).

Immunization. CD45.1⁺ mice were immunized s.c. on day 0 with a volume of 50 µl vaccine 141 142 preparation containing 20 µg rHBsAg (Rheinbiotech, Düsseldorf, Germany), 0.1 mM CpG 143 1668 (Molbiol Syntheselabor GmbH, Berlin, Germany) and 50 µg Polyphosphazene (PCEP); 144 kindly provided by G. Mutwiri (Vaccine and Infectious Disease Organization/International 145 Vaccine Center, University of Saskatchewan, Saskatoon, Canada). On day 28 mice were boosted i.p. with 1x10⁸ MVA pH5-S expressing HBs in 0.9 % NaCl. On day 35, splenocytes 146 147 were isolated and numbers of HBV-specific T cells were determined by intracellular cytokine 148 staining. For stimulation of cells from immunized mice, we used peptides &Gal 96-103, S190-149 197 and S208-215 at a concentration of 1 µg/ml as well as three peptide pools of 18 15mers 150 overlapping by 11 amino acids covering the complete S protein (pool S1: S1-S83, S2: S73-151 S155, S3: S144-S226). After overnight resting, CD8⁺ T cells were isolated using magnetic 152 cell sorting (Miltenyi) and processed as described for CAR-transduced T cells. The number of HBV-specific CD8⁺ T cells was estimated to be approx. 4%. 153

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155 **References**

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161