

T Cell Receptor Grafting allows Virological Control of Hepatitis B Virus Infection

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31 **Abstract**

32 T cell therapy is a promising means to treat chronic HBV infection and HBV-associated
33 hepatocellular carcinoma. T cells engineered to express an HBV-specific T cell receptor (TCR)
34 may achieve cure of HBV infection upon adoptive transfer. We investigated the therapeutic
35 potential and safety of T cells stably expressing high affinity HBV envelope- or core-specific
36 TCRs recognizing European and Asian HLA-A2 subtypes. Both CD8⁺ and CD4⁺ T cells from
37 healthy donors and from chronic hepatitis B patients became polyfunctional effector cells when
38 grafted with HBV-specific TCRs and eliminated HBV from infected HepG2-NTCP cell
39 cultures. A single transfer of TCR-grafted T cells into HBV-infected, humanized mice
40 controlled HBV infection and virological markers declined 4-5 log or below detection limit.
41 When – as in a typical clinical setting - only a minority of hepatocytes were infected,
42 engineered T cells specifically cleared infected hepatocytes without damaging non-infected
43 cells. Cell death was compensated by hepatocyte proliferation and alanine amino transferase
44 levels peaking at day 5-7 normalized again thereafter. Co-treatment with the entry inhibitor
45 Myrcludex B ensured long-term control of HBV infection. Thus, T cells stably transduced with
46 highly functional TCRs have the potential to mediate clearance of HBV-infected cells causing
47 limited liver injury.

48

49 **Keywords:** chronic hepatitis B; hepatocellular carcinoma; adoptive T-cell therapy; functional
50 cure; humanized mice, Myrcludex B

51 **Introduction**

52 Worldwide, more than 250 million humans suffer from chronic hepatitis B (CHB), which
53 accounts for an estimated 880.000 deaths per year caused by secondary complications like liver
54 cirrhosis or hepatocellular carcinoma (HCC) (1). Current therapeutic regimens based on the
55 use of nucleos(t)ide analogues (NUCs) can efficiently suppress viral replication, but are unable
56 to eradicate the virus. The so-called covalently closed circular DNA (cccDNA) of HBV persists
57 as a transcription template in the nucleus of infected cells and re-initiates HBV replication
58 when antiviral treatment is discontinued (2). Hence, therapeutic options are needed, in which
59 cccDNA is eliminated or at least strongly reduced and controlled by the immune system to
60 allow functional HBV cure and to prevent relapse (3).

61 During acute, resolving hepatitis B a strong immune response is mounted with T cells being
62 key to clear the virus (4). In CHB by contrast, the scarce and oligoclonal T-cell response against
63 HBV fails to control the virus and to prevent disease progression (5). T cells have been shown
64 to kill infected hepatocytes and to secrete cytokines that control virus replication in a non-
65 cytolytic fashion by silencing (6-8) but also by destabilizing cccDNA (9, 10). In addition,
66 killing of HBV-infected cells will promote compensatory cell proliferation that in turn favors
67 further cccDNA loss by cell division (11).

68 Restoring a potent T-cell response by adoptive T-cell therapy is an interesting therapeutic
69 option (12, 13). The concept of transferring adaptive immunity to control HBV has already
70 been applied successfully in CHB patients who underwent stem cell transplantation and
71 received bone marrow from HBV-immune donors (14-16). Since allogeneic stem cell
72 transplantation is limited by its severe side effects like graft-versus-host disease and a high
73 mortality, alternative approaches are required. Genetic engineering of autologous T cells to
74 express HBV-specific receptors is an attractive alternative to treat CHB, to prevent HBV-
75 related complications or to treat HBV-related HCC (17).

76 We have already demonstrated that T cells expressing a chimeric antigen receptor binding the
77 antigenic loop within the “S”-domain of all HBV envelope proteins selectively eliminated
78 HBV-infected and thus cccDNA-positive target cells (18, 19). It has also been shown that T
79 cells from HBV-infected patients can be engrafted with T cell receptors (TCR) against the
80 envelope or the core protein and become activated upon recognition of HBV peptides (20, 21).
81 Moreover, significant reduction of HBV infection in humanized mice has recently been
82 demonstrated after repeated adoptive transfers of human T cells engineered to express HBV-
83 specific TCRs via mRNA electroporation (22). However, due to the transient expression of the
84 TCR and the high numbers of infected cells present in that experimental setting, the antiviral
85 effects remained limited and HBV rebound was observed within ten days after the last T cell
86 injection. Most likely, a more sustained T cell activity, ideally combined with strategies aiming
87 at blocking new infection events (23), is required to achieve a more profound and durable
88 control of HBV infection. We recently reported the cloning, characterization and permanent
89 expression of a set of eleven HBV-specific TCRs and determined their functional avidity (21).
90 This allowed us to identify those most promising TCRs for adoptive T cell therapy.

91 In the present study, we aimed at determining the potential of T cells grafted by retroviral
92 vector transduction with HBV-specific TCRs to clear HBV-infected cells without damaging
93 non-infected neighboring cells. Furthermore, we combined T cell therapy with the virus entry
94 inhibitor Myrcludex B (MyrB) (24) to restrict new virus spread after T-cell control of HBV
95 infection, and showed that T cells grafted with selected, high avidity TCRs were able to control
96 HBV-infection both in cultured hepatocytes and in vivo in livers of HBV-infected humanized
97 mice.

98

99 **Results**

100 ***Core- and S-specific TCRs confer HBV-specificity upon retroviral transduction***

101 HLA-A2 restricted HBV envelope- or core-specific TCRs were cloned as gene-optimized
102 constructs into a retroviral vector. TCR-transgenic T cells were generated by retroviral
103 transduction resulting in high expression of both TCRs on CD4⁺ as well as CD8⁺ T cells (Figure
104 1A). Functional avidities of eleven TCRs were compared in extensive analyses resulting in a
105 functionality rating as described in detail in Wisskirchen et al. (21) (Supplemental table 1). The
106 core₁₈₋₂₇ (C₁₈)-specific TCR 6K_{C18} and the S₂₀₋₂₈ (S₂₀)-specific TCR 4G_{S20} were selected for
107 comprehensive testing of their antiviral activity. Transduced T cells killed stable HBV-
108 replicating hepatoma cells in co-cultures at effector to target (E:T) ratios as low as 1:12. At an
109 E:T ratio of 1:3, core-specific, 6K_{C18}-grafted T cells eliminated 50% of HBV-replicating cells
110 after 6-7 hours. S-specific, 4G_{S20}-grafted T cells showed slower kinetics and needed about 20
111 hours (Figure 1B). Thus, endogenously processed peptides were readily recognized by both
112 receptors and activated T cell effector functions.

113

114 **TCR-grafted T cells efficiently target HBV-infected cells in vitro**

115 Our next step was to assess the antiviral capacity of TCR-grafted T cells on HepG2 cells stably
116 expressing the HBV entry receptor NTCP (HepG2-NTCP) and infected with HBV. Based on
117 titration experiments (Supplemental Figure 1A-F), we incubated the HBV-infected cells with
118 TCR-grafted T cells at an effector to target (E:T) cell ratio of 1:2 and tested whether this would
119 be sufficient to eliminate HBV-infected cells. After 6 and 10 days of co-culture, viral HBsAg
120 and HBeAg were not detected anymore in cell culture media, respectively (Figure 2A,B), while
121 secreted and intracellular viral rcDNA were largely reduced (Figure 2C,D). Most importantly,
122 the persistence form of the viral DNA – cccDNA – became undetectable by qPCR after ten
123 days (Figure 2E). A more prominent effect on cccDNA than on rcDNA was expected, since

124 rcDNA is protected from DNase activity within the HBV capsid (18). The amount of
125 extracellular rcDNA even increased temporarily when infected cells were lysed by HBV-
126 specific T cells (Figure 2C), likely because of the release of non-enveloped DNA-containing
127 capsids (25).

128 To assess whether pre-treatment with antivirals would influence antiviral T cell activity, we
129 treated HBV-infected cells with the NUC Entecavir (ETV) for three weeks before adding TCR-
130 grafted T cells. While killing of ETV-treated target cells within 72 hours was reduced (Figure
131 2F), the overall antiviral effect of HBV-specific T cells remained equally pronounced
132 compared to that without NUC treatment (Figure 2G-J). Thus, both core- and S-specific T cells
133 generated by genetic engineering were capable of eliminating HBV-infected cells even after
134 treatment with NUCs.

135

136 ***HBV-specific TCRs mediate redirection of T cells from patients with chronic hepatitis B***

137 Adoptive T cell therapy imposes the challenge of creating an autologous T-cell product from a
138 patient that has high viral antigens circulating and chronic inflammatory liver disease.
139 Therefore, we used PBMC from two CHB patients, grafted T cells with the two selected TCRs
140 and evaluated their antiviral potency. T cells could be transduced as efficiently as T cells from
141 healthy donors (Figures 3A, 1A). T cell expansion was more than 200-fold, starting from less
142 than a million cells, irrespective of the donor or the TCR being expressed (Figure 3B). 4G_{S20}-
143 and 6K_{C18}-grafted T cells killed infected cells (Figure 3C) and secreted up to 10 ng/ml of IFN-
144 γ within 2 days (Figure 3D). After 10 days of co-culture, secreted HBeAg became negative -
145 with the exception of 4G_{S20} T cells in donor 2 where very low levels were still detected - and
146 cccDNA was not detectable anymore (Figure 3E-F). Importantly, T cells from CHB patients
147 did not inherently contain a relevant number of functional HBV-specific T cells before TCR-
148 grafting, as no antiviral activity of mock-transduced T cells was observed.

149 Adoptive T cell therapy using T cell receptors entails that the TCR is customized to fit to the
150 patient's HLA-type. Therefore, we asked whether our high-affinity TCRs would recognize
151 peptide presented on different HLA-A*02 subtypes (Supplemental Figure 2). In total, TCR
152 4G_{S20} or 6K_{C18} recognized their cognate peptide on 9/12 or 7/12 tested HLA-A*02-subtypes,
153 respectively, including subtypes A*02:03, A*02:06 and A*02:07 which are most frequently
154 found among the Asian population. Taken together, these TCRs conferred HBV-specificity to
155 T cells from donors with CHB with a performance comparable to T cells from healthy donors
156 and can be applied to patients with different HLA-A*02 subtypes.

157

158 ***TCR-grafted CD4⁺ and CD8⁺ T cells show antiviral activity***

159 To quantify the contribution of CD4⁺ T cells in controlling and eliminating HBV, infected cells
160 were first co-cultured either with a mixture of CD4⁺ and CD8⁺ TCR-grafted T cells or with
161 TCR-grafted CD4⁺ T cells only. The combination of CD4⁺ and CD8⁺ T cells grafted with either
162 TCR killed infected cells even at a starting E:T ratio as low as 1:64 (Figure 4A). CD4⁺ T cells
163 were also able to kill infected cells, although less efficiently requiring a 4-fold higher number
164 of transduced T cells i.e. a higher E:T ratio (Figure 4B). Cytotoxic effector function of core-
165 specific T cells started immediately after onset of the co-culture while that of S-specific T cells
166 only started 1 or 2 days later. Secreted HBeAg declined 2-3 days after the onset of cytotoxicity
167 (Supplemental Figure 3A,B). A direct comparison of both T cell types revealed that CD8⁺ T
168 cells mainly produced IFN- γ and CD4⁺ T cells TNF- α in particular when using TCR 4G_{S20}
169 (Figure 4C,D and Supplemental Figure 3C). Overall, CD8⁺ T cells showed a stronger antiviral
170 effect, and CD4⁺ T cells that carried TCR 6K_{C18} were more efficient than CD4⁺ T cells grafted
171 with the TCR 4G_{S20} (Figure 4E,F and Supplemental Figure 3D,E). Taken together, not only
172 CD8⁺, but also CD4⁺ T cells were able to kill HBV-infected cells when expressing a TCR with
173 a high functional avidity.

174

175 ***TCR-grafted T cells clear HBV infected cells mainly by direct cytotoxicity***

176 Both cytotoxicity and secretion of cytokines by T cells play a role in viral clearance. To
177 determine the contribution of non-cytotoxic, cytokine-mediated antiviral activity of TCR-
178 grafted T cells, cytokine-depleting antibodies were employed in our in vitro infection model.
179 The antibodies reduced the amount of IFN- γ by about 60 and 90% for 4G_{S20}- and 6K_{C18}-
180 expressing T cells, respectively, and almost completely depleted TNF- α from the cell culture
181 medium (Figure 4C,D and Supplemental Figure 4A,B). Reduction of secreted HBeAg
182 remained similar when cytokines were deprived (Figure 4E, Supplemental Figure 4C). While
183 cccDNA levels remained strongly reduced when co-cultures with CD8⁺ T cells were treated
184 with cytokine-blocking antibodies, cytokine removal reduced the capacity of CD4⁺ T cells to
185 eliminate cccDNA (Figure 4F). A detailed comparison of 4G_{S20}- and 6K_{C18}-expressing T cells
186 showed that cccDNA clearance was reduced by a factor of 1.5-2 when IFN- γ and TNF- α were
187 neutralized (Figure 4G). This indicated that TCR-grafted CD8⁺ T-cells mainly clear HBV by
188 direct cytotoxicity, while TCR-grafted CD4⁺ T cells elicit both a cytotoxic and a prominent
189 non-cytotoxic, cytokine-mediated effect.

190 The high cytolytic T cell activity in vitro raised the concern of potential bystander killing of
191 non-infected cells. To address this concern, HBV-infected HepG2-NTCP cells MOI 100, about
192 half of the cells productively infected with HBV (26)) were mixed with non-infected cells at
193 different ratios. Interestingly, 4G_{S20}-specific T cells were not activated when only 20% of cells
194 were derived from the infection cell batch while 6K_{C18} T cells still were (Supplementary Figure
195 5A-C). For both receptors, 4G_{S20} and 6K_{C18}, killing increased directly proportionally to the
196 percentage of cells from the infected batch (Supplementary Figure 5A,B). When T cells were
197 activated for 24 hours by co-culture with HepG2-NTCP cells, which had been infected at MOI

198 of 500 and transferred to non-infected cell cultures, no bystander killing was observed while T
199 cells remained active on infected cells (Supplementary Figure 5D). From these data we did not
200 get any indication of non-specific activity of our TCR-grafted T cells.

201

202 ***Transfer of TCR grafted T cells results in strong reduction of HBV markers in vivo***

203 For the clinical success of adoptive T cell therapy of CHB it is important that transferred T
204 cells migrate to the liver, exert their effector function and eliminate HBV while liver function
205 is maintained despite a loss of infected hepatocytes. In a first set of experiments we employed
206 human liver chimeric USG mice harboring HLA-A*02-positive human hepatocytes to assess
207 the antiviral activity of T cells stably expressing the selected HBV-specific TCRs. Mice were
208 infected with HBV for 12 weeks and displayed median viral titers of 1.4×10^8 (2.4×10^6 to
209 2.2×10^9) HBV-DNA copies/ml before they received one single injection of either 2×10^6 mock
210 transduced or 1×10^6 6K- together with 1×10^6 4G-grafted T cells. They were followed-up either
211 for a short time of 15-20 days or a longer time of 55 days (Figure 5A). Two additional mice (1
212 treated and 1 mock control) received a second injection of T cells at day 5 and were sacrificed
213 at day 15. Alanine aminotransferase levels (ALT) were increased between day 3 and 7 post T-
214 cell transfer in all mice that had received HBV-specific T cells indicating transient liver damage
215 (Figure 5B). Liver damage was accompanied by a <10% transient body weight reduction
216 (Supplementary Figure 6A). Mock-treated animals had ALT levels comparable to untreated
217 liver chimeric mice indicating that no alloreaction was caused by the transferred T cells. In
218 these high viremic, HBV-infected mice treated with TCR-grafted T cells, human serum
219 albumin (HSA) decreased substantially (average 5.7-fold) (Figure 5C). Nevertheless, HSA
220 levels started to slowly rebound in some animals (Figure 5C). Within the first 3 weeks after T-
221 cell transfer, viremia decreased by $\approx 4 \log_{10}$ (Figure 5D), and HBeAg and HBsAg dropped below
222 the limit of detection in most animals (Figure 5E, F). At sacrifice, HBeAg proved non-reactive

223 in 6/7 and HBsAg in 3/7 mice treated with TCR-grafted T cells, and reduction of HBV DNA
224 viremia was confirmed (Supplemental Figure 6B-D).

225 In line with the serological results, intrahepatic analyses of mice treated with effector T cells
226 showed significantly lower levels (median: $-3\log_{10}$) of intracellular HBV RNA transcripts
227 (Figure 5G) and rcDNA (Figure 5H) in comparison to mock-treated mice. In all treated mice
228 intrahepatic cccDNA dropped to very low levels (median: $-2\log_{10}$) 3 weeks after T cell transfer
229 and became undetectable after 8 weeks of treatment (Figure 5I). The second T cell injection
230 did not have any further effect, as one single injection of TCR-grafted T cell was already
231 sufficient to achieve strong and sustained antiviral effects.

232 HBV pgRNA was not detectable by RNA in situ hybridization in liver tissues of mice that were
233 sacrificed at day 18 (Figure 6A). The discrepancy between in situ and qPCR detection may be
234 explained by the different sensitivity levels of the respective assays and by the fact that in situ
235 analysis only reflects the viral state within a limited area of a liver section, since DNA and
236 RNA for PCR analysis were extracted from a larger piece of liver tissue. Immunofluorescence
237 co-staining for HBV core protein and human cytokeratin 18 (Figure 6B) revealed that nearly
238 all human hepatocytes were HBV-positive in control mice with high viral titers, whereas
239 animals with 10-fold lower viral titers (e.g. 2×10^8 HBV-DNA copies/ml) had fewer positive
240 cells (Figure 6B, mock-treated group). T-cell injection provoked a massive elimination of
241 infected hepatocytes over time in mice with high infection rates (Figure 6B, upper panel), while
242 in mice with intermediate infection rates at baseline, a large proportion of human hepatocytes
243 survived (Figure 6B, lower panel). Human CD45⁺ lymphocytes were still detected in liver
244 tissue at day 19, but no longer after 8 weeks (Figure 6C). Nevertheless, human T cells,
245 especially CD8⁺ T cells, were still present in the spleen of mice 8 weeks post T-cell transfer,
246 although the proportion of TCR⁺ T cells had decreased compared to what had been injected in
247 the mice (Supplemental Figure 7A, B). This could be attributed to downregulation of the TCR

248 after T cell activation or contraction of the population of HBV-specific T cells after most of
249 the infection had been cleared. Of note, Ki67-staining showed the potential of human
250 hepatocytes to proliferate and hence their ability to compensate for the immune-mediated cell
251 loss (Figure 6D). Taken together, these experiments demonstrated that a single injection of T
252 cells grafted with TCRs of high avidity can efficiently reduce HBV infection by promoting the
253 clearance of HBV-infected hepatocytes in vivo.

254

255 ***TCR grafted T cells have the potential for clinical application***

256 To assess the specificity with which effector T cells stably expressing HBV-specific TCRs
257 target the infected hepatocytes in vivo without provoking damage of non-infected neighboring
258 hepatocytes, we used mice in which only a minority of the human hepatocytes was infected.
259 This mimics the clinical situation more closely, where typically only a low percentage of cells
260 is infected and expresses HBV core and envelope proteins (27). To obtain partially HBV-
261 infected humanized livers, we stopped HBV spreading at 5 weeks post virus inoculation by
262 applying the HBV entry inhibitor MyrB. One week later, five mice received TCR-grafted T
263 cells, whereas three animals served as controls and were sacrificed either 2 weeks (n=2) or 13
264 weeks after injection of mock T cells (Figure 7A). To assess whether HBV may relapse after
265 T cell therapy, we stopped MyrB application in 2 mice 3 weeks after T cell transfer and
266 monitored viremia levels for additional 10 weeks (i.e. until week 19 post infection).

267 In line with our previous experiments, a single injection of HBV-specific T cells caused a
268 transient ALT elevation (Figure 7B) and concomitant reduction of HSA levels (Figure 7C) in
269 all treated mice while their body weight remained stable (data not shown). Compared to high-
270 titer HBV-infected mice (Figure 5C), HSA drop was less pronounced in the partially infected
271 animals and HSA levels rebounded to baseline within 2-3 weeks (Figure 7C), indicating that

272 hepatocyte loss was limited and promptly compensated by human hepatocyte proliferation.
273 HBV viremia, as well as circulating HBeAg and HBsAg dropped to borderline detection levels
274 in all animals within two weeks after T cell transfer (Figure 7 D-F). When mice received MyrB
275 throughout the experiment, intrahepatic levels of HBV transcripts and DNA were ≈ 4 -log lower
276 3 and 13 weeks after transfer of TCR-grafted compared to mock T cells (Figure 7G-I). When
277 MyrB treatment was stopped 3 weeks after T cell injection, all serological HBV markers started
278 to rebound 9 weeks after injection of TCR-grafted T cells (Figure 7D-F) and reached baseline
279 levels again at 13 weeks (Figure 7G-I). Co-staining of HBcAg and human CK18 revealed that
280 only a minority of human hepatocytes was HBV-positive in mice that received MyrB at 5
281 weeks post infection and sacrificed 2 weeks post mock T-cell treatment (Figure 7J, left panel).
282 HBV-positive cells were detected neither 3 nor 13 weeks after transfer of HBV-specific T cells
283 under continuous MyrB treatment, whereas HBV-positive hepatocytes were detected in
284 animals that discontinued MyrB treatment (Figure 7I, right panel). These results show that
285 TCR-grafted T cells were able to target HBV-infected cells with high efficiency also when only
286 a minority of human hepatocytes was infected. Lack of off-target effects also indicated high
287 specificity of TCR-redirected T cells. Most importantly, HBV infection was fully controlled
288 after adoptive transfer of HBV-specific T cells when MyrB was continuously administered.
289

290 **Discussion**

291 T cell therapy of CHB aims at supplementing a patient's lacking or functionally exhausted
292 HBV-specific T cell repertoire to achieve HBV control. In this study, we demonstrate that T
293 cells genetically engineered to stably express HBV-specific TCRs with high functional avidity
294 (21) are able to eliminate HBV-infected cells with high efficiency and specificity. Although
295 our TCRs were cloned from European donors, testing them on different HLA-A2 subtypes
296 showed a broad applicability irrespective of the patient's ethnicity. HBV-specific T cells
297 obtained using either T cells of healthy volunteers or CHB patients expanded to clinically
298 relevant numbers, were equally efficient and able to eliminate HBV-infected cells. Importantly,
299 a single adoptive T cell transfer combined with administration of the HBV entry inhibitor MyrB
300 was able to achieve long-term control of HBV infection in HBV-infected humanized mice.

301 Currently approved polymerase inhibitors are well tolerated and effectively suppress HBV
302 replication. However, they do not directly affect the HBV persistence form, the cccDNA
303 demanding additional means to cure hepatitis B (28). T cells grafted with HBV-specific
304 receptors become activated by HBV-expressing hepatoma cell lines (20, 29) or HBV-infected
305 hepatocytes (30) and target HBV persistence (18).

306 Our TCRs proved to be functional not only on CD8⁺ but also on CD4⁺ T cells, and grafting
307 onto healthy volunteers' or CHB patients' T cells lead to elimination of viral antigens as well
308 as cccDNA when co-cultured with HBV-infected cells. Both cell types showed cytotoxic and
309 non-cytotoxic effector function while CD8⁺ T cells mainly produced IFN γ and CD4⁺ T cells
310 TNF α . Thus, the manufacturing of a T cell product for clinical application can be simplified as
311 purification of CD8⁺ T cells will not be essential to generate sufficient numbers of antiviral T
312 cells. As it has been shown that the inclusion of CD4⁺ T cells grafted with a chimeric antigen
313 receptor has a synergistic antitumor effect (31), CD4⁺ T cells engrafted with our MHC-I-
314 restricted high affinity TCRs that are fully functional without CD8 co-receptor binding (21)

315 will also very likely contribute to the success of adoptive T cell therapy either by direct antiviral
316 activity or by helper function.

317 Several cytokines, secreted by adoptively transferred T cells or after bystander activation of
318 other immune cells have been reported to influence HBV gene expression and replication in a
319 non-cytolytic fashion (32) by a number of different means (summarized in (33)). In our
320 experiments, however, killing of infected cells was the dominant antiviral mechanism
321 eliminating HBV in cell culture and in vivo. Since IFN γ , however, could only partially be
322 blocked by antibodies, the non-cytolytic effect of TCR-grafted T cells that has recently been
323 reported (29, 34) may have been underestimated in our experiments.

324 In our first in vivo experimental setting, most human hepatocytes (>90%) were infected with
325 HBV and mice were highly viremic after the virus had had 12 weeks to spread in the
326 immunodeficient animals. In these animals, a large proportion of the human cells was
327 eliminated upon T cell transfer, cccDNA was reduced by >95% and neither HBV core protein
328 nor viral RNAs could be detected anymore in situ in surviving human hepatocytes.
329 Nevertheless, non-cytolytic activity of T cells (29, 35) may have contributed to clear HBV
330 infection, since some human hepatocytes may well have lost cccDNA by cell division (11), as
331 demonstrated by positive Ki67 staining, and T-cell derived cytokines may have contributed to
332 purging (10, 29) or silencing (6) of cccDNA molecules.

333 A concern about HBV-specific T-cell therapy is the loss of a significant part of functional liver
334 cells, particularly in patients with liver cirrhosis and end-stage liver disease. After adoptive T
335 cell therapy, adverse events have been observed when the TCR was specific for a tumor-
336 associated (self-) antigen and bound “off-target” to related peptides (36), or when it recognized
337 the cognate antigen “off-organ” on healthy tissue (37). Both scenarios seem unlikely to happen
338 during T-cell therapy of CHB, as the viral antigens we are targeting are very distinct from “self-
339 antigens” and are only expressed in hepatocytes. This is supported by the notion that no severe

340 side effects have been noted after the transfer of HBV-specific immune cells to CHB patients
341 with hematological malignancies and normal liver function, in whom bone marrow
342 transplantation led to viral clearance while only causing moderate liver toxicity (14-16). In this
343 regard, we found that despite the capacity of our engineered T cells to clear infected cells,
344 cytotoxicity of the stably transduced T cells was limited, since ALT indicating hepatocyte death
345 increased only transiently in the first week after T cell transfer. Serum HSA reflecting
346 hepatocyte function dropped in correlation with the amount of intrahepatic infection detected.
347 Humanized mice in which - as in a typical clinical setting - the liver was only partially infected
348 neither lost weight nor showed any other signs of distress. Of note, TCR-grafted T-cells did
349 not cause any measurable damage of neighboring non-infected cells, since large hepatocyte
350 areas were maintained and HSA rebound to baseline levels emphasizing the capacity of human
351 hepatocytes to proliferate and compensate for the immune-mediated cell loss.

352 The situation was different when virtually all human hepatocytes were infected. Here, we found
353 a significant drop in HSA indicating a significant reduction in liver function. A fully infected
354 liver, however, is not expected to reflect the situation of the majority of CHB patients (27).
355 Another limitation of the humanized mouse model is the mismatch of human cytokines and
356 murine receptors that may underestimate the effect of a cytokine storm. Huang et al. (38)
357 showed in a study including more than one hundred HBeAg-negative patients, that in patients
358 with a viral load below 1×10^6 copies/ml the amount of HBcAg⁺ hepatocytes is below 20%.
359 Hence, patients, which would be considered as candidates for the clinical application of T cell
360 therapy, should be carefully selected for their viremia and antigenemia. Nevertheless a liver
361 biopsy should be obtained to assess the number of HBV-infected cells as well as the extend of
362 pre-existing liver inflammation and fibrosis.

363 Treatment with NUCs only blocks the viral polymerase and production of new viral progeny,
364 but not transcription of HBV RNA and antigen production per se. Thus, the presentation of

365 HBV antigen is not expected to change substantially and HBV-specific T cells should remain
366 able to clear infected cells. Our in vitro study proved that recognition and clearance capacity
367 of TCR-grafted T cells was maintained upon Entecavir treatment. Pretreatment with NUCs
368 would even be preferred, since long term NUC treatment is associated with a lower extend of
369 intrahepatic HBV infection (27). As in our experiments with partially infected mice, we would
370 expect a less pronounced hepatocyte loss in patients that have received NUCs for some years
371 and this should not impair liver function in non-cirrhotic or child A patients. Moreover,
372 treatment with polymerase inhibitors will also reduce inflammation and thus increase safety
373 of adoptive T-cell therapy. Safety could further be increased if a safeguard molecule would
374 be co-transduced and co-expressed with the TCR to allow for rapid depletion of transferred
375 T cells. To limit the circulation of HBV-specific T cells, the application of T cells that only
376 transiently express the HBV-specific TCR after RNA electroporation is explored (22).
377 However, in the same humanized mouse model TCR-electroporated T cells led to a
378 comparable increases of ALT, but despite several T-cell re-injections did not achieve the same
379 strong antiviral effect as their retrovirally transduced counterparts (22), and was even lower
380 when resting, non-cytolytic T cells were used (34).

381 The efficacy of our retrovirally transduced T cells carrying high-avidity TCRs was striking in
382 vivo, especially when compared to previous studies (22, 34). These results point out how a
383 different technology used to engineer effector T cells (i.e. careful selection of TCRs (21) and
384 stable versus transient TCR expression) helps to achieve a higher efficacy of T cell therapy.
385 We needed only a single T cell injection to achieve a sustained drop of HBsAg and HBV
386 DNA (> 4log) in the serum of mice, whereas multiple injections of transiently transduced T
387 cells were needed to achieve a reduction of HBV viremia by a median 1 log, and HBsAg
388 levels barely changed (22).

389 However, in the absence of adaptive immune responses, even the smallest HBV reservoir
390 persisting eventually leads to viral rebound in our mice. To avoid HBV rebound, we combined
391 T cell therapy with administration of the entry inhibitor MyrB and by this mean succeeded in
392 maintaining control of HBV infection for the entire period of three months after T cell transfer.
393 Although in CHB patients - in contrast to our immune incompetent mouse model – long term
394 survival of transferred T cells is more likely and may even be supported by reconstitution of
395 the patient's own anti-HBV immune response, inhibition of new infection events will have a
396 supportive role also in the clinical setting.

397 Adoptive T cell therapy of HBV-related diseases is already proceeding towards the clinics.
398 Bertoletti and colleagues recently reported T cell therapy of an HLA-A2-positive patient who
399 had received an HLA-A2-negative liver transplant because of HCC and developed metastases
400 from his original HBV S⁺, HLA-A2⁺ HCC. This patient was treated with retrovirally TCR-
401 grafted, S₂₀-specific T cells. The therapy was safe and circulating HBsAg decreased (17).
402 Although these study results are very encouraging numbers of TCR-grafted, transferred HBV-
403 specific T cells were low (<2.5% within 3.9x10⁸ infused cells). By improved retroviral
404 transduction we can now generate high numbers of HBV-specific T cells stably expressing
405 TCRs (>75% within 2.5x10⁸ cells) from less than one million cells from healthy donors but -
406 as shown here - also from CHB patients even allowing to spare leukapheresis.

407 Taken together, we show that high-affinity HBV-specific T cells can be generated by TCR-
408 grafting irrespective of the donor being HBV-positive or -negative prior to T cell therapy. TCR-
409 grafted T cells have a strong antiviral capacity in cell culture and in vivo, most strongly
410 reducing cccDNA, which is regarded as a hallmark of virological cure (3). Thus, adoptive T-
411 cell therapy of CHB, mimicking T-cell responses in self-limiting HBV infection, may result in
412 functional cure of HBV infection.

413

414 **Materials and Methods**

415 ***Retroviral transduction of T cells.*** Stable 293GP-R30 (RD114-pseudotype) producer cells
416 were generated by transduction with cell culture supernatant from 293GP-GLV9 cells, both
417 provided by BioVec Pharma (39) that had been transfected with TCR plasmids as described
418 earlier (21). T cells were enriched using human T activator CD3/CD28 Dynabeads (Thermo
419 Fisher Scientific, Waltham, MA, USA) and pre-stimulated for 2 days in T-cell medium (TCM):
420 RPMI, 10 % human serum, 1 % pen/strep, 1 % glutamine, 1 % sodium pyruvate, 1 % non-
421 essential amino acids, 0,01M HEPES (all Thermo Fisher Scientific), supplemented with 300
422 U/ml IL-2. 0.45 µm-filtered retrovirus cell culture supernatant from stable producer cell lines
423 was centrifuged at 2000xg, 32°C for 2 hours on non-treated culture plates coated with 20 µg/ml
424 RetroNectin (Takara, St. Germain en Laye, France). Retrovirus cell culture supernatant was
425 removed and T cells were spinoculated onto the retrovirus-coated plate at 1000xg for 10
426 minutes. A second transduction was performed after 24 hours. TCR expression was determined
427 by flow cytometry. Staining was done for 30 minutes on ice in the dark, using the primary
428 antibodies anti-human CD4_{APC} (clone OKT4, #17-0048-42, eBioscience, Frankfurt,
429 Germany), anti-human CD8_{PB} (clone DK25, #PB984, Dako, Waldbronn, Germany) and anti-
430 mouse TCRβ_{PE} (clone H57-597, #553172, BD Biosciences, Heidelberg, Deutschland)
431 diluted in FACS Buffer (0.1% BSA/PBS). Cells were analyzed using a FACSCanto II flow
432 cytometer (BD Biosciences) and data were analyzed with FlowJo 9.2 software.

433 ***Co-culture with HBV-infected cells.*** HepG2 cells expressing the sodium-taurocholate
434 cotransporting polypeptide (HepG2-NTCP K7, generated by our group (26)) were seeded in
435 DMEM (10% FCS, 1% penicillin/streptomycin, 1% glutamine, 1% NEAA) on collagen-coated
436 plates. At 90% confluency, 2.5% DMSO was added to the medium. Cells were infected 2-6
437 days later with HBV genotype ayw purified via heparin column affinity chromatography
438 followed by sucrose gradient ultracentrifugation from HepAD38 cell culture supernatant in the

439 presence of 4.8% PEG overnight at the indicated number of virions per cell (MOI), and
440 maintained with 1% FCS as a monolayer. For T-cell co-culture, medium was changed to
441 DMEM 10% FCS / 2% DMSO. T cells were added in equal amounts of TCM (final
442 concentration of 1% DMSO in co-culture) at the indicated ratio of 6K_{C18} (specific for C18-27:
443 FLPSDFFPSV), or 4G_{S20} (specific for S20-28: FLLTRILTI) T cells to target cells (E:T). For
444 Figure 2 F-J, HepG2-NTCP cells were infected with an MOI of 500. One week after infection
445 cells were treated with 0.1 μ M of Entecavir (ETV) twice a week for a duration of three weeks.
446 For Figure 4 C-F, CD8⁺ and CD4⁺ T cells were separated using MACS beads for positive
447 selection of the respective cell type (Miltenyi Biotech, Bergisch-Gladbach, Germany) and
448 added at an E:T ratio of 1:2. Cytokine-blocking antibodies against IFN- γ (10ng/ml, clone B27,
449 #506513, Biolegend, San Diego, CA, USA) or TNF- α (5ng/ml, clone D1B4, #7321, Cell
450 Signaling, Danvers, MA, USA) were given every other day when medium was exchanged.

451 ***Analyses of co-cultures.*** Cytokines in the cell culture supernatant were detected using ELISA
452 kits for IFN- γ and IL-2 (BioLegend, San Diego, USA) or for TNF- α (BD Biosciences). HBeAg
453 in the cell culture supernatant was measured using the Enzygnost HBe monoclonal assay
454 (Siemens Healthcare Diagnostics, Eschborn, Germany). Total DNA was extracted from cells
455 using the NucleoSpin tissue kit (Macherey-Nagel, Düren, Germany) or from cell culture
456 supernatant using a Tecan extraction robot (40). Viral DNA forms were amplified and detected
457 by PCR as described previously (10) and quantified with a standard obtained from accredited
458 assays in our diagnostics department. 7.5 μ l DNA were digested with 5 units of T5 exonuclease
459 (NEB, Ipswich, USA) for 30 minutes at 37°C to eliminate rcDNA before cccDNA
460 quantification.

461 ***Real-time cytotoxicity measurement.*** 5x10⁴ HepG2 (ATCC®) or HepG2.2.15 cells (kind gift
462 of Heinz Schaller, University of Heidelberg, Germany) prepared as described previously (21),
463 or 4x10⁴ infected or non-infected HepG2-NTCP cells were seeded onto 96-well electronic

464 microtiter plates. T cells were added 1-2 days later when target cells had reached confluence.
465 The impedance, which reflects adherence of the target cells to the bottom of the plate, was
466 measured every 15-30 minutes using an xCELLigence® SP real-time cell analyzer (ACEA
467 Biosciences, San Diego).

468 ***Adoptive T cell transfer in humanized mice.*** Human liver chimeric mice were generated using
469 male or female, 3-week-old uPA/SCID/beige/IL-2R γ ^{-/-} (USG) mice (41). They originated from
470 IL-2R γ ko mice (JAX Mice stock number 003169; C.129S4-Il2rg^{-/-}/J) were backcrossed
471 10 times on uPA/SCID mice, which originated from crossing uPA mice (Jax Stock JR2214 (not
472 available anymore); B6SJL-TgN(Alb1Plau)144Bri) for 10 generations on SCID beige mice
473 (Taconic model: CBSCBG; C.B-Igh-1b/GbmsTac-Prkdcscid-Lystbg N7. After transplanting
474 1×10^6 thawed human HLA-A2⁺ hepatocytes the repopulation levels were determined by
475 measuring human serum albumin (HSA) in mouse serum using the human Albumin ELISA
476 quantitation kit (Bethyl Laboratories, Biomol GmbH, Hamburg, Germany). To establish HBV
477 infection, animals received a single intra-peritoneal injection of HBV-infectious serum (1×10^7
478 HBV-DNA copies/mouse; genotype D) 10 weeks after transplantation. For adoptive T cell
479 transfer, T cells were thawed and cultured overnight in AIM-V medium (Gibco/Thermo Fisher
480 Scientific), 2% human AB⁺-serum and 180 IU/ml rIL-2. 2×10^6 TCR-grafted T cells (1×10^6 with
481 6K_{C18} + 1×10^6 with 4G_{S20}) or equal numbers of mock-treated T cells were injected
482 intraperitoneally into HBV-infected mice. In the first experiment, mice were monitored until day
483 15-19 (short-term) or day 55 (long-term) after transfer. In the short-term follow-up groups, one
484 mouse of each group (both HBeAg 1.5 (S/CO) on day 3 post transfer) received a second injection
485 of 1×10^6 TCR-grafted or mock T cells on day six (mice are specified in Figure 5) and one mouse
486 received MyrB, 5 days before sacrifice. In the second experiment, all mice received MyrB
487 (100 μ l; 2mg/kg) subcutaneously (11, 24) 5 weeks post infection to block the viral spreading and
488 mimic a partial infection in these mice. Within the first 5 days, mice received MyrB daily. After

489 5 days, mice were treated every other day to avoid re-or new-infection of non-infected
490 hepatocytes. Blood was taken for analyses of viral antigens, ALT and viremia as indicated in
491 results.

492 ***Virological measurements.*** DNA and RNA were extracted from liver specimens using the
493 Master Pure DNA purification kit (Epicentre, Madison, USA) and RNeasy RNA purification kit
494 (Qiagen), respectively. Intrahepatic total viral loads were quantified with the help of primers and
495 probes specific for total HBV-DNA, pgRNA and cccDNA while the human housekeeping gene
496 *GAPDH* was used for normalization (42). HBsAg and HBeAg quantification were performed on
497 the Abbott Architect platform (Abbott, Ireland, Diagnostic Division) at indicated serum dilution.
498 HBeAg results are displayed as signal to noise ratio (S/CO).

499 ***Analysis of biochemical parameters.*** ALT was measured by using the Roche Cobas c111
500 System (Roche, Basel, Switzerland). For the measurements 5µl of serum of the mice was used.

501 ***Preparation and staining of splenocytes.*** Spleens were dissected in cold mTCM
502 (RPMI/10%FCS supplemented with Glutamin, 1% Pen/Strep, 0,1% beta-Mercaptoethanol and
503 1% sodium pyruvate, passed through a 100 µm strainer and homogenized through a 20 G needle.
504 Splenocytes were centrifuged at 300xg for 5' at 4°C and incubated in TAC-buffer for 2' at 37°C
505 to lyse erythrocytes. Reaction was stopped with mTCM. Cells were counted after repeat
506 centrifugation and subsequently stained for anti-human CD8-PE-Cy7 (clone RPA-T8, #557746,
507 BD Biosciences,), anti-human CD4-Pacific Blue (clone RPA-T4, #558116, BD Biosciences)
508 and anti-mouse TRBC-PE (clone H57-597, #553172, BD Biosciences). Flow cytometry data
509 were acquired on the BD FACS LSRII System.

510 ***Immunofluorescence and RNA in situ hybridization.*** Human hepatocytes were identified in
511 frozen mouse liver sections using a monoclonal anti-human cytokeratin-18 mouse antibody
512 (clone DC-10, #11-107-C100, Exbio, Praha, Czech Republic) or monoclonal anti-human
513 calnexin rabbit antibody (clone C5C9, #2679, Cell Signaling Technology, Massachusetts, USA).

514 HBcAg staining was detected with anti-rabbit anti-HBcAg antibodies (polyclonal, #B0586,
515 Dako Diagnostika, Glostrup, Denmark) and specific signals were visualized with Alexa 488- or
516 Alexa 555-labelled secondary antibodies (polyclonal, #A21429, #A11029, # A11034,
517 Invitrogen, Darmstadt, Germany) or the TSA Fluorescein System (Perkin Elmer, Jügesheim,
518 Germany). Nuclear staining was achieved by Hoechst 33258 (Invitrogen, Eugene, USA). Human
519 CD45 staining was performed using a monoclonal anti-mouse CD45 antibody (clone
520 2B11+PD7/26, # IS751, Dako). Proliferation of human hepatocytes and immune cells was
521 determined using antibodies to human Ki-67 (polyclonal, # ABIN152984, antibodies-online,
522 Aachen, Germany). RNA in situ hybridization was performed on paraformaldehyde-fixed, cryo-
523 preserved liver sections using the RNAScope Fluorescent Multiplex Kit (Advanced Cell
524 Diagnostics, ACD, Hayward, CA, USA) with target probes recognizing the pregenomic HBV
525 (assay 442741-C2) and human beta2-microglobulin-RNA (assay 478171-C3). Stained sections
526 were analysed by fluorescence microscopy (Bioevo BZ-9000, Keyence, Osaka, Japan) using
527 the same settings for all groups.

528 ***Study approval.*** The use of healthy volunteer PBMC was approved by the local ethics board of
529 the University Hospital rechts der Isar, Munich (G 548/15S), and written informed consent was
530 obtained from all blood donors. PHH were isolated from rejected explant livers using protocols
531 approved by the Ethical Committee of the city and state of Hamburg (OB-042/06) in accordance
532 with the principles of the Declaration of Helsinki. Animals were housed under specific pathogen-
533 free conditions according to institutional guidelines under authorized protocols. All animal
534 experiments were approved by the City of Hamburg, Germany (G118/16), conducted in
535 accordance with the European Communities Council Directive (86/EEC) and the ARRIVE
536 standard.

537

538 **Author contributions:** K.W. and U.P. designed in vitro experiments. K.W., A.M. and T.A.
539 performed in vitro experiments and K.W. analyzed data. K.W., J.K., M.D., and U.P. wrote the
540 manuscript. T.B. provided and analyzed patient material. A.M. established transduction
541 protocols. J.W. produced and purified HBV stocks. J.K., M.D., K.W and U.P. designed in vivo
542 experiments. J.K. conducted in vivo experiments and acquired data. J.K. and K.W. analyzed
543 the data. L.A. and T.V. generated human chimeric liver mice. M.L. and L.A. established RNA
544 in situ hybridization assays. L.A. and J.K. performed RNA in situ hybridization assay. S.U.
545 conceptually contributed to entry inhibition experiments and provided MyrB for the in vivo
546 experiments. All authors discussed data and corrected the manuscript.

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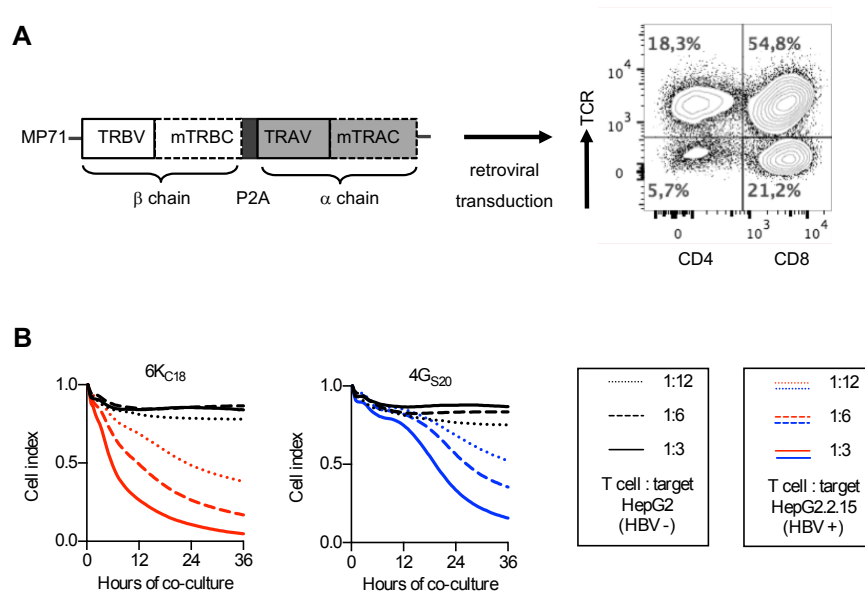
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679

680 **Figure 1. Genetic engineering and analysis of HBV-specific T cells.** (A) Codon-optimized
 681 TCR α (TRAV) and β (TRBV) chains of high-affinity TCRs were cloned into the retroviral
 682 vector MP71. Murine constant domains (mTRBC, mTRAC) and insertion of additional
 683 cysteines were used to increase pairing. After retroviral transduction T cells were stained for
 684 mTRBC and TCR expression was quantified by flow cytometry. (B) Functional comparison
 685 scored eleven TCRs directed against HBV peptides Core₁₈₋₂₇, S₂₀₋₂₈ and S₁₇₂₋₁₈₀ (21). TCRs
 686 6K_{C18} and 4G_{S20} were identified to have the highest functional avidity and were therefore
 687 chosen for further analyses: Parental HepG2 or HBV+ HepG2.2.15 target cells were co-
 688 cultured with increasing numbers of TCR-grafted T cells. Killing of target cells was determined
 689 by detachment from the plate using a real-time cell analyzer (XCelligence™) and is given as
 690 normalized cell index relative to the starting point of the co-culture. HBV⁻ target cells co-
 691 cultured with TCR-grafted T cells are shown as black lines. HBV⁺ target cells co-cultured with
 692 6K_{C18}-transduced T cells are shown in red and 4G_{S20}-transduced T cells in blue. Data are
 693 presented as mean values of quadruplicate co-cultures (n=4).

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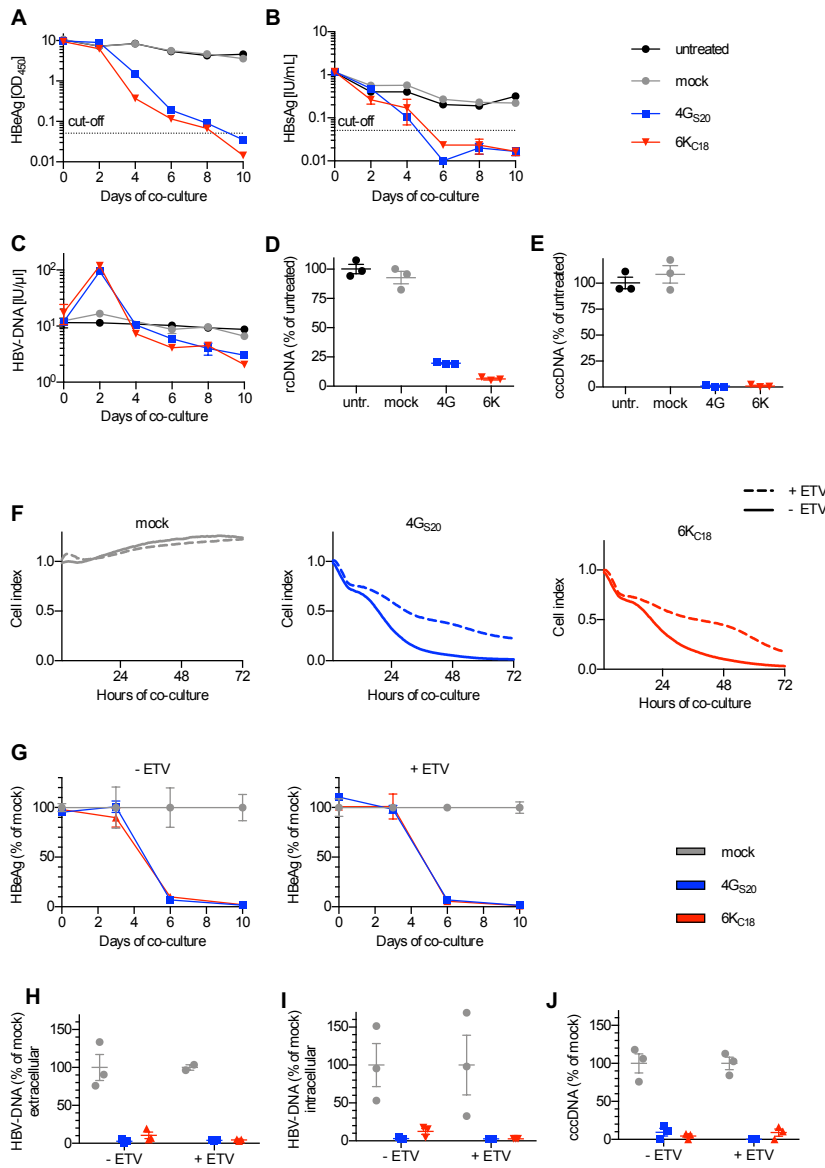
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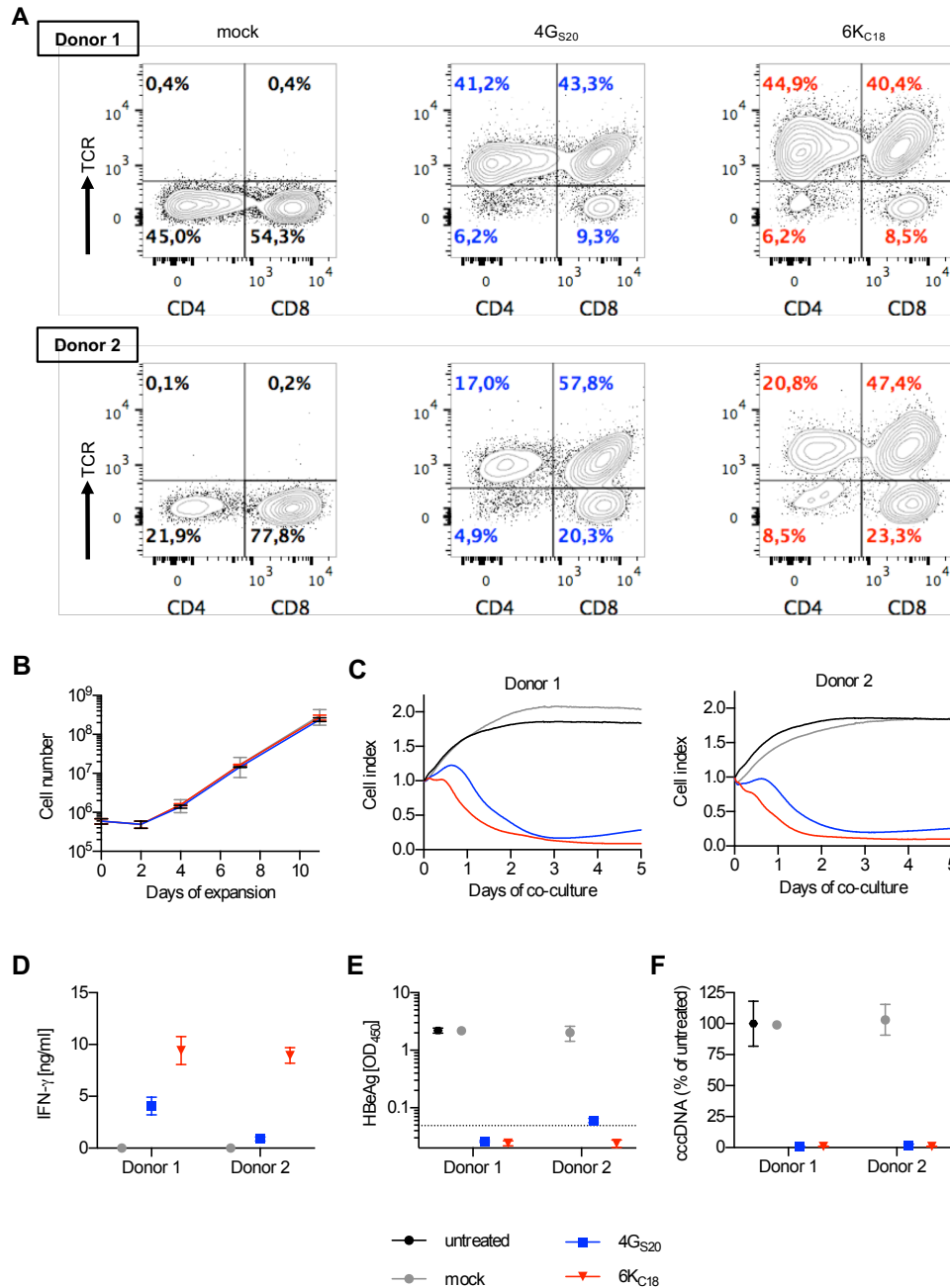
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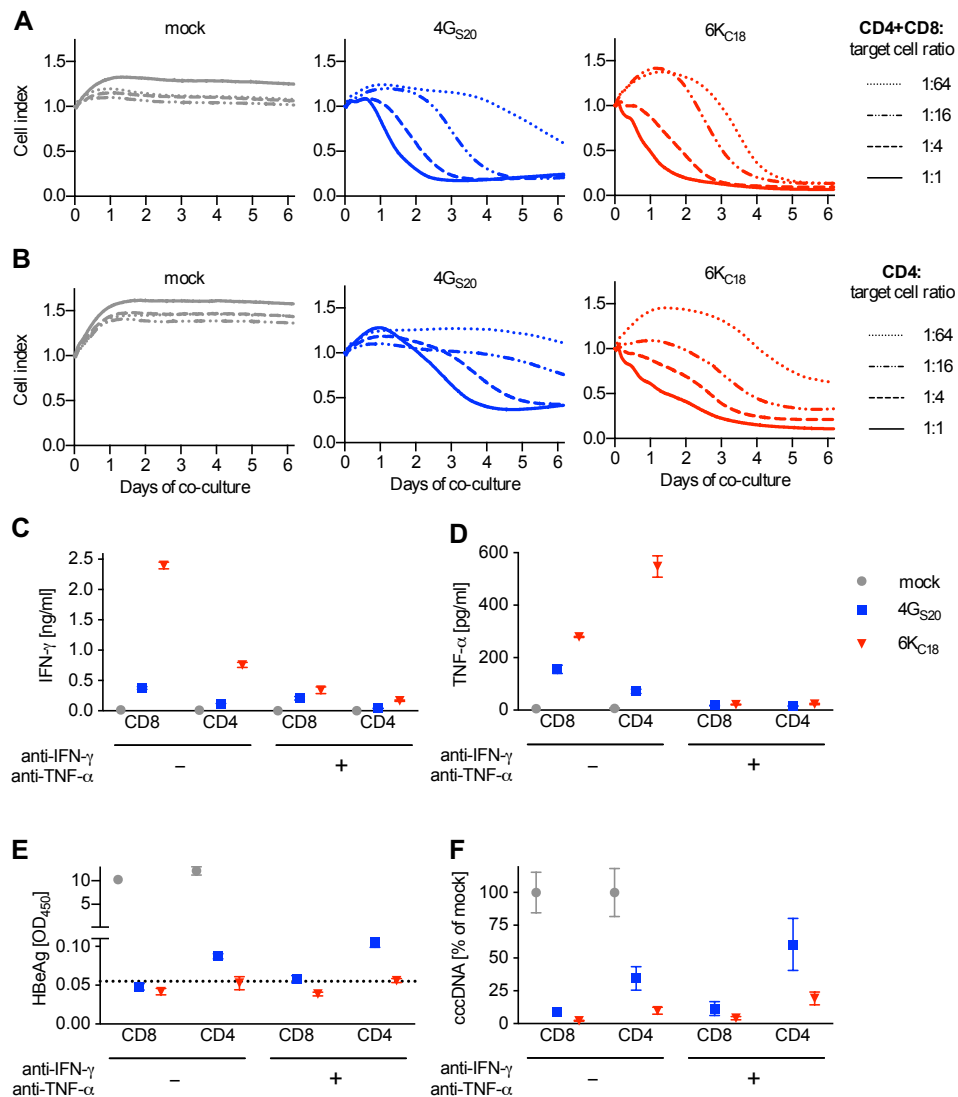
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701 **Figure 2. Antiviral effect of TCR-grafted T cells on HBV-infected cells.** HepG2-NTCP cells
 702 were infected with HBV at an MOI of 100. After two weeks, T cells grafted with HBV S-
 703 specific TCR 4G_{S20} (blue squares) or HBV core-specific TCR 6K_{C18} (red triangles) or non-
 704 transduced T cells (mock, grey circles) were added for ten days at an E:T ratio of 1:2. Medium
 705 was changed every other day and used to determine (A) HBeAg and (B) HBsAg, by diagnostic
 706 ELISA. (C) HBV relaxed circular (rc)DNA contained in virions that had been secreted was
 707 extracted from cell culture supernatant every other day and DNA extracted from cell lysates on
 708 day ten was used to determine (D) intracellular HBV rcDNA and (E) nuclear cccDNA using
 709 qPCR. (F-J) Cells were infected with an MOI of 500. One week after infection cells were
 710 treated with 0.1 μM of Entecavir (ETV) twice a week for a duration of three weeks. (F) Killing
 711 of target cells was measured using a real-time cell analyzer and is given as normalized cell
 712 index relative to the starting point of the co-culture. E:T 1:1. (G-J) Medium was changed every
 713 3-4 days and values are normalized for co-cultures treated with mock T cells. (G) HBeAg in
 714 supernatant of co-cultures without (left) or with (right) ETV pre-treatment (H, I) HBV relaxed
 715 circular (rc)DNA contained in virions secreted into the cell culture medium or extracted from
 716 cell lysates on day ten and (J) nuclear cccDNA determined using qPCR. Data are presented as
 717 mean values from triplicate co-cultures (n=3).



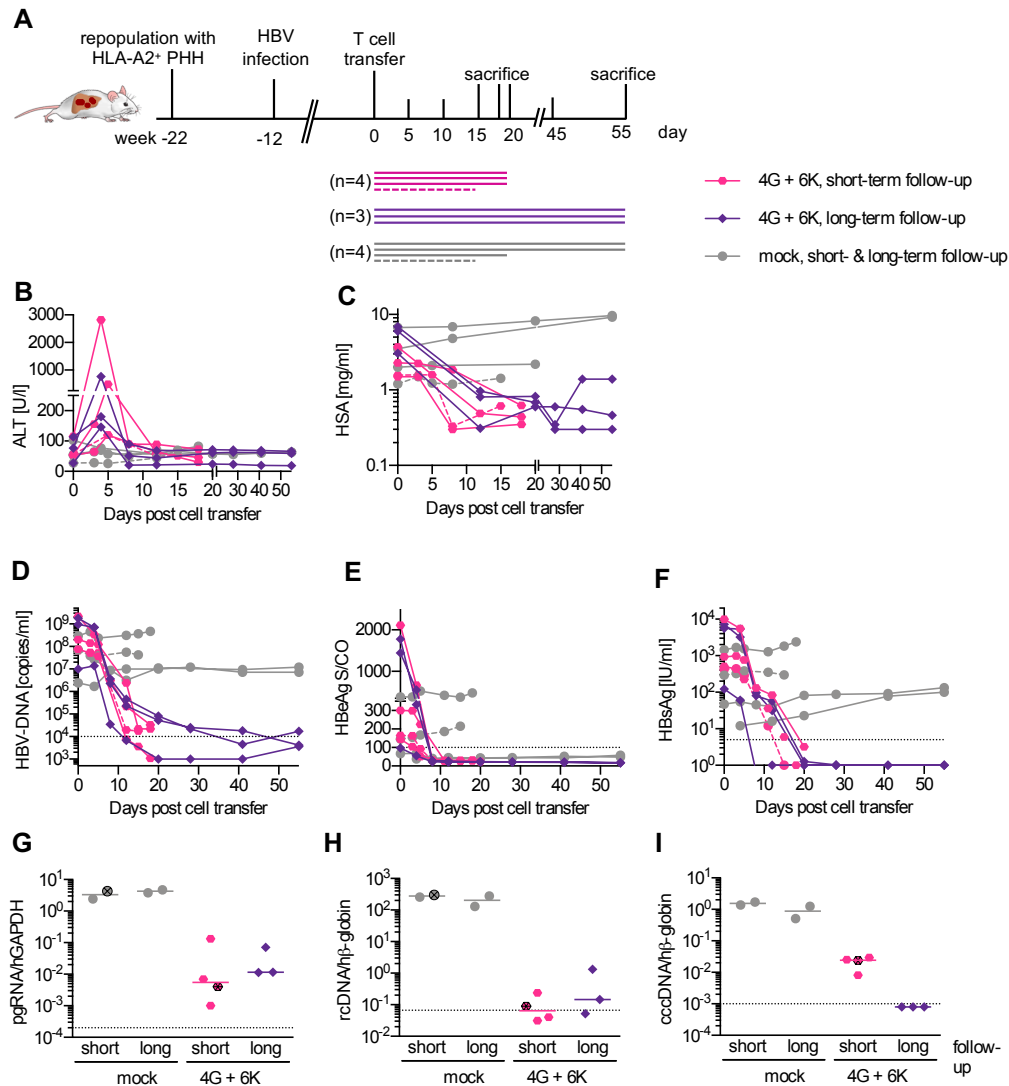
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719 **Figure 3. Antiviral activity of T cells from patients with CHB.** (A) CD3⁺ T cells were
 720 isolated from two donors with CHB and transduced to express HBV-specific TCRs 4G_{S20}
 721 (blue) and 6K_{C18} (red). TCR expression was quantified by flow cytometry. (B) Expansion of T
 722 cells during retroviral transduction with CD3/CD28 T activator Dynabeads and IL-2. (C-F)
 723 HepG2-NTCP cells were infected with HBV at an MOI of 500 three weeks prior to co-culture
 724 with TCR-grafted T cells from CHB donors at an E:T ratio of 1:2. Since the ratio of CD4⁺ and
 725 CD8⁺ T cells varied strongly between the donors, effector cell number was calculated on the
 726 basis of TCR⁺ CD8⁺ T cells only. (C) Killing of target cells was measured using a real-time
 727 cell analyzer and is given as normalized cell index relative to the starting point of the co-culture.
 728 E:T 1:2.7. (D) IFN-γ was determined in cell culture medium on day 2. Secreted HBeAg (E)
 729 and intracellular HBV cccDNA (F) were measured after 10 days of co-culture. Data are
 730 presented as mean values of triplicate co-cultures (n=3).



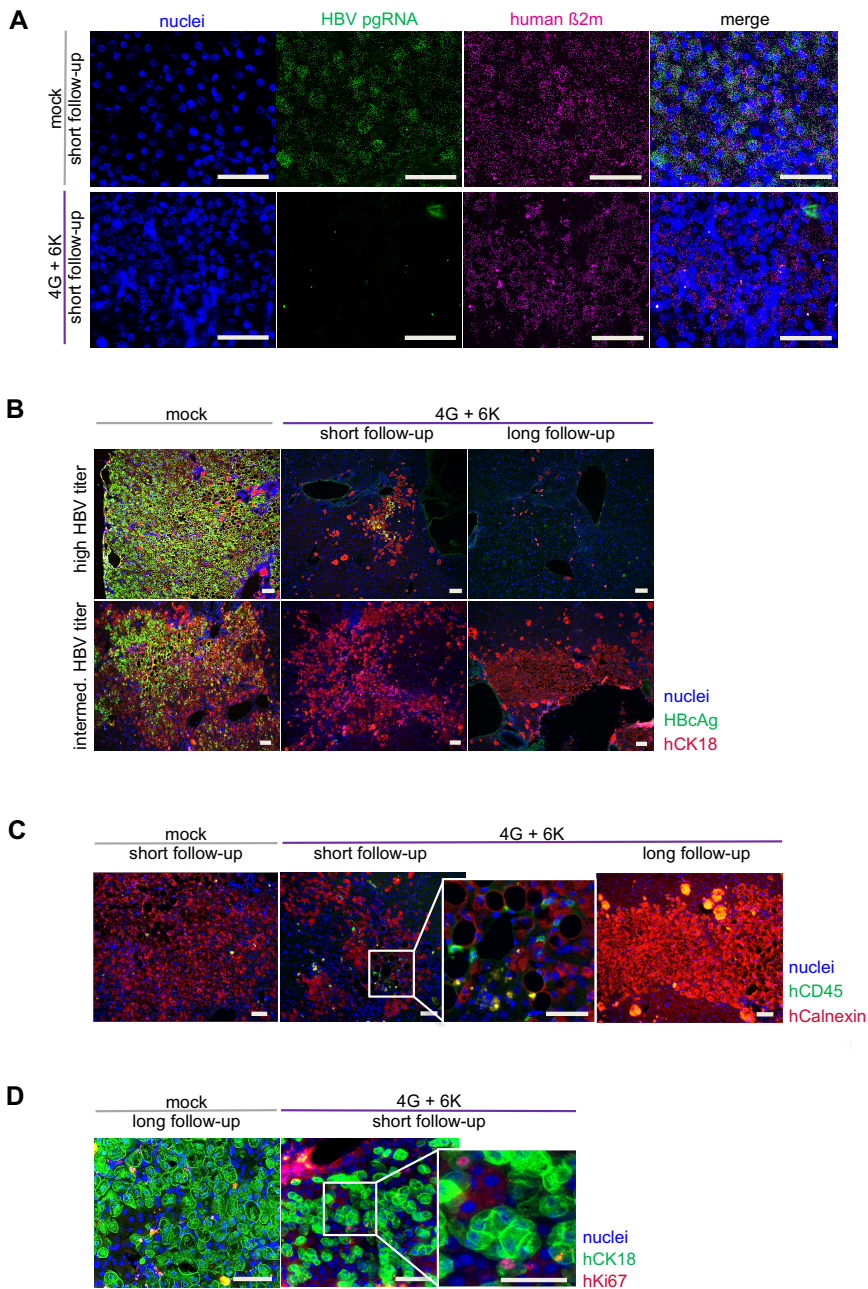
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 732 **Figure 4. Antiviral activity of different TCR-grafted T cell subsets.** HepG2-NTCP cells
 733 were infected with HBV at an MOI of 500. After two-three weeks, T cells grafted with TCR
 734 4G_{S20} (blue squares) or 6K_{C18} (red triangles) or non-transduced T cells (mock, grey circles)
 735 were added at decreasing E:T ratios. (A,B) Killing of target cells determined by detachment
 736 from the bottom of the 96-well plate was measured in real-time (XCelligence™) and is given
 737 as normalized cell index relative to the starting point of the co-culture with CD4⁺ and CD8⁺ T
 738 cells (A) or CD4⁺ T cells only (B). (C-F) CD8⁺ and CD4⁺ T cells were separated by positive
 739 magnetic cell sorting and added at an E:T ratio of 1:2. Cytokine-blocking antibodies against
 740 IFN- γ (10ng/ml) or TNF- α (5ng/ml) were given every other day when medium was exchanged.
 741 (C, D) IFN- γ and TNF- α were measured in the cell culture medium after 2 days. (E) Secreted
 742 HBeAg and (F) intracellular HBV cccDNA were measured after 10 days of co-culture. Data
 743 are presented as mean values (A,B) or mean values \pm SEM (C-F) of triplicate co-cultures (n=3).

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Figure 5. Antiviral activity of TCR-grafted T cells in HBV-infected humanized mice. (A) USG mice were repopulated with HLA-A*02-matched PHH, infected with 1×10^7 HBV virions, followed until a stable viremia had established (week 12-14) and injected with 2×10^6 TCR-grafted T cells (1×10^6 with 6K_{C18} plus 1×10^6 with 4G_{S20}; colored symbols, n=7) or equal numbers of mock-treated human T cells (grey circles, n=4). 4 mice were sacrificed within 3 weeks (short-term follow-up, pink hexagons) and 3 mice 8 weeks (long-term follow-up, purple diamonds) after T cell transfer, respectively. 2/11 mice received a second dosage of either effector cells or mock cells and were sacrificed on day 15 (presented by broken lines in A-F and crossed dots in G-I). (B) ALT activity and progression of (C) HSA or (D) HBV-DNA in sera. (E, F) HBeAg and HBsAg determined using immunoassays. (G-I) Intrahepatic HBV RNA and DNA transcripts were quantified by qPCR. (G) Levels of pregenomic HBV RNA were normalized to human GAPDH RNA. (H, I) rcDNA and cccDNA were quantified relative to an HBV plasmid standard curve and normalized to human beta globin. Each data point or longitudinal line represents one mouse. Dotted line represents the technical cut-off of the respective test. For DNA and RNA analyses dotted lines indicate the lower limit of detection = LLoD (defined as 10 HBV rcDNA or cccDNA copies per ≥ 1000 human beta globin copies).



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Figure 6. In situ analysis of antiviral effects of TCR-grafted T cells in humanized livers.

Liver tissue of HBV-infected mice treated with mock-transduced T cells or 2×10^6 HBV-

specific T cells (1×10^6 with 6K_{C18} plus 1×10^6 with 4G_{S20}) were used from day 18-19 (short

follow-up) or days 55 post T cell transfer (long follow-up). (A) In situ RNA hybridization for

HBV pregenomic RNA (green) and human β 2-microglobulin (magenta) against nuclei staining

(blue) was performed to determine the occurrence of HBV specific RNA transcripts. (B) As

indicated mice were distinguish in the level of infection into high (10^9 - 10^{10} copies/ml) or low

(10^7 - 10^8 copies/ml) HBV titer. Representative immunohistochemistry staining for cell nuclei

(blue), HBV core protein (green), and human cytokeratin 18 (red). (C) Human CD45 positive

cells (green) were stained against human Calnexin (red) as marker for human hepatocytes and

cell nuclei (blue). (D) To determine potential proliferation in mice treated with T cells, staining

for cell nuclei (blue), human cytokeratin 18 (green) and Ki67 (red) was employed. Scale bar of

liver tissue sections: 50 μ m.

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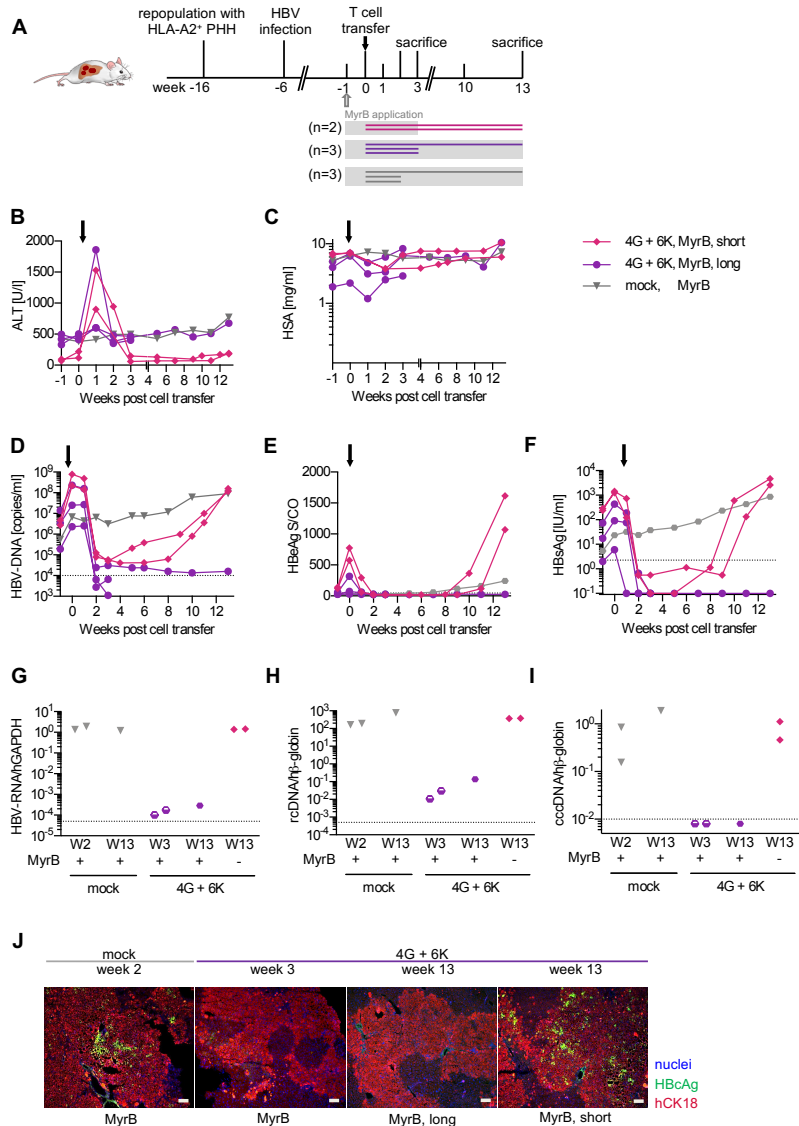
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786 **Figure 7. Long-term follow-up of mice partially infected with HBV and treated with**
787 **HBV-specific T cells and an entry inhibitor. (A)** USG mice were repopulated with HLA-
788 A*02-matched PHH, infected with 1×10^7 HBV virions. Viral spreading was stopped at week 5
789 after infection by administration of MyrB (see methods) (grey blocks). After 1 week of MyrB
790 application, 2×10^6 TCR-grafted T cells (1×10^6 with 6K_{C18} plus 1×10^6 with 4G_{S20}, n=5) or mock
791 T cells (grey triangles, n=3) were transferred. To address the question whether mice could be
792 re-infected after treatment with effector T cells, MyrB application was stopped in 2/5 mice
793 after 3 weeks (MyrB short, pink diamonds, n=2) and followed up till week 13. MyrB was
794 administered continuously in 3/5 mice (MyrB long, purple hexagons, n=3). Mice were
795 sacrificed at week 2 or 3 during short-term follow-up or at week 13 during long-term follow-
796 up. Time course of ALT activity (B), HSA (C), HBV viremia (D) HBeAg (E) and HBsAg (F)
797 in sera followed until week 3 or 13, respectively. (G) Levels of pregenomic RNA were
798 normalized to human GAPDH RNA. (H, I) rcDNA and cccDNA were quantified relative to an
799 HBV plasmid standard curve and normalized to human beta globin. Each data point or
800 longitudinal line represents one mouse. Dotted lines represent the technical cut-off of the
801 respective test. For DNA and RNA analyses dotted lines indicate the lower limit of detection
802 = LLoD (defined as 10 HBV DNA or cccDNA copies per ≥ 1000 human beta globin copies).
803 Dots below LLoD symbolize undetectable measurements. (J) Representative staining of liver
804 tissue slides from mice either treated with mock or 4G+6K effector T cells.