Supplementary Materials for

Vaccination to prevent T cell subversion can protect against persistent hepacivirus infection

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Supplemental Figure 1. Gating strategy for quantification of intracellular cytokines. Shown are intrahepatic mononuclear cells after 5-hr PMA/Ionomycin stimulation in the presence of brefeldin A.

Supplemental Table 1. Core-specific MHC class II epitopes.

Peptide	Position	Sequence	
P9	Core ₅₇	AKRRRRHRRDQGGWRRSP	
P12	Core ₇₈	VDPYVRQGLQILLPSAAY	
P13	Core ₈₅	GLQILLPSAAYPVRDPRR	
P17	Core ₁₁₃	GTLGWTADLLHHVPLVGP	
P18	Core ₁₂₀	DLLHHVPLVGPLVGHPAR	
P20	Core ₁₃₄	HPARLICRAVRACEDGIN	
P22	Core ₁₄₈	DGINSFTGIAGVHLFLIC	

Supplemental Table 2. List of PCR primers used in study.

Name	Sequence (5'→3')		
GNN-F1	GATTTGTGGAAATAACACGGTGTGCATATTTG		
GNN-R1	AACCAGCTGGGTGACGCG		
5'Kpnl_NS35B_F	GATGGGGTACCATGGCCCCCTTCAACTTAAG		
3'EcoRV_NS35B_R	GGTGGTAAGATATCGAGGTTGAGGGATTTACC		
RHV_NS3amino-F	CCGCTGAGGACACTGGTTTCA		
RHV_NS3amino-R	CGGCATGGCACTCGTCGC		
RHV_NS3carboxy-F	CCATGATAACGGCTCAAGGCTC		
RHV_NS3carboxy-R	GTCGCCCGTCAAGTCCCAGG		
RHV_NS3amino-F2	CCGCTGAGGACACTGGTTTCA		
RHV_NS3amino-R2	CGTGGCCGTCGCAAGAAGC		
RHV_NS4B-F	GCTTCCAGCATCCAGGGGTG		
RHV_NS4B-R	CGGGTCGGCAGGCGTAGAGT		
RHV_NS5A-F	CAAGCCACCACACTCAGACAAAC		
RHV_NS5A-R	TGGGATTACGCATCAAGCCG		
RHV_NS5Bamino-F	GCCCACAGACCCAGACACTTTCA		
RHV_NS5Bamino-R	GCTTCATGGCCACTGCGAAT		
RHV_NS5Bcarboxy-F	TTACATGACAACCTGTATGCTGG		
RHV_NS5Bcarboxy-R	ACATCAGGAGAAGATGAAGGCTAT		
NS3aseq-F1	CCCTAAAGGGCTTACCAGTG		
NS3aseq-F2	TCTTTCTAAGGTGCAGGAGC		
NS3cseq-F1	CATGTTCCTAGCGAAGAACC		
NS3cseq-F2	TGATGCCAACATTCAAGCTG		
NS4Bseq-F1	GAAGAATACTTTGCTGAGACGG		
NS4Bseq-F2	CTACAAAGCTAACAACACAGACG		
NS5Aseq-F1	AAGTGGCTGGAAACAGCCTC		
NS5Aseq-F2	CTGTAGGCTGGACTAGTATGGC		
NS5Aseq-F3	TGCACACATGAAAGACATCTCACG		
NS5Aseq-F4	GTGCATCCTGTTCCAGAATC		
NS5Baseq-F1	AGTCGTCTGGATCATGGACC		
NS5Baseq-F2	TGGTGGAGAAGATGGTGCTC		
NS5Bcseq-F1	CATGGTGATGCAAGGTGCTG		
NS5Bcseq-F2	GTGTGTGCAATTCTTACAGCAG		



Supplemental Figure 2. Gating strategies for analysis of tetramer positive cells. Analysis of tetramer positive cells after staining with markers for CD44H and granzyme B (a) or Annexin V (b).



Supplemental Figure 3. Absence of in vivo killing during acute RHV infection. (a) Representative FACS plots of CFSE⁺ cells in infected liver at 10 days post infection. (b) Summary of percent specific lysis of RHV-rn1 peptide-loaded cells at days (d) 10 and 14 post infection in livers (grey circles) and spleens (red squares).



Supplemental Figure 4. Immunogenicity of Ad-NSmut and characterization of vaccineelicited T cell repertoire. Lewis rats were vaccinated intramuscularly with 5 x 10⁸ ifu Ad-NSmut. Two weeks after vaccination, splenocytes were harvested for analysis of cellular immunity. **(a)** IFNγ ELISpot responses against vaccine insert (NS3, NS4; 2 µg/mL) and backbone (Hexon; 0.6 nmol/mL) peptides. Data from n=3 rats per group is shown (mean ± SEM). **(b-e)** Splenocytes from four rats were pooled together at equal ratios, magnetically enriched for CD8^{pos} and CD8^{neg} cells, and used to map MHC class I and II epitopes in the NS3-5B insert by IFNγ ELISpot assay. MHC class II epitopes (b and c) were identified by testing individual NS3-4B peptides and NS5B matrix pools. MHC class I epitopes (d and e) for the RT1-A⁷ protein of the Lewis rat were screened and confirmed using the SYFPEITHI prediction algorithm. Panels b and d show representative ELISpot responses of identified epitopes in duplicate.



Supplemental Fig 5. Serum ALT values in vaccinated rats. Levels of serum ALT (sALT) in (a)

resolvers and (b) non-resolvers after RHV infection.



Supplemental Fig. 6. Mapping of NS3₉₇₄ class I epitope. Flow cytometric analysis of intracellular IFN γ production by CD8⁺ T cells in immune splenocytes following 5-hr stimulation with untruncated or truncated NS3₉₇₄ peptide.



Extended Figure 7. RHV-specific T cell response in Ad-null vaccinated rats after virus challenge. Lewis rats were vaccinated intramuscularly with 5 x 10⁸ ifu Ad-null. Three weeks after vaccination, rats were challenged intravenously with 10⁶ genomes RHV. At days 14 and 100 post infection, virus-specific T cell responses were quantified in spleens and livers by flow cytometric analysis following 5-hr stimulation with a pool of MHC class I and II epitopes (5 μ g/mL). (a,b) Representative flow plots of intracellular IFN γ and TNF α cytokine production in CD8⁺ (a) and CD4⁺ (B) T cells from livers of vaccinated rats. (c,d) Calculated frequencies of total and dual positive cytokine producers in CD8⁺ (c) and CD4⁺ (d) T cell compartments following RHV-specific peptide stimulation. Liver, grey circles; spleen, red squares. Data from n=3 rats per group are shown (mean ±SEM).



Supplemental Figure 8. RHV-specific CD4⁺ T cell responses in depleted rats after virus challenge. Lewis rats were vaccinated intramuscularly with 5 x 10⁸ ifu Ad-NSmut. Three weeks after vaccination, rats were challenged intravenously with 10⁶ genomes RHV. At -2 and 5 days post infection, rats were depleted of CD8α⁺ cells by antibody or isotype control. At day 9 post infection, CD4⁺ T cell responses were quantified in liver by flow cytometric analysis following 5-hr stimulation with a pool of MHC class I and II epitopes (5 µg/mL). **(a,b)** Representative flow plots of intracellular IFNγ and TNFα cytokine production by intrahepatic CD4⁺ T cells in (a) mock and (b) CD8α⁺ cell-depleted rats. **(c)** Calculated frequencies of total and dual positive cytokine producing cells in mock and CD8α⁺ cell-depleted rats. IFNγ, red circles; TNFα, blue squares; IFNγ,TNFα, green triangles. Data from n=3 rats per group are shown (mean ±SEM). **(d)** Relative frequency of intrahepatic CD4⁺ T cells producing dual or single IFNγ or TNFα cytokines. Data from n=3 rats per group are shown (mean ±SEM).

day 21	NS3 ₉₇₄	NS3 ₁₄₉₇	NS4A ₁₅₇₈
	SICVIGTPL	YTYLYAAQY	CVFMAIDLF
506		506 5	06
507		507 5	07
508		508 5	08
509		509 5	09

Supplemental Figure 9. Class I epitope evolution in CD4-depleted rats. Serum viral RNA in CD4-depleted rats at 21 days post infection was analyzed for presence of mutations in class I-restricted epitopes by consensus PCR sequencing. A single non-synonymous V \rightarrow I mutation was identified at position 4 in the NS3₉₇₄ epitope of serum virus recovered from rat 508.