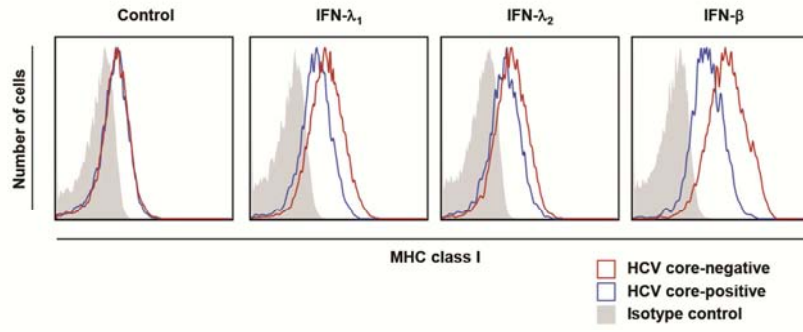
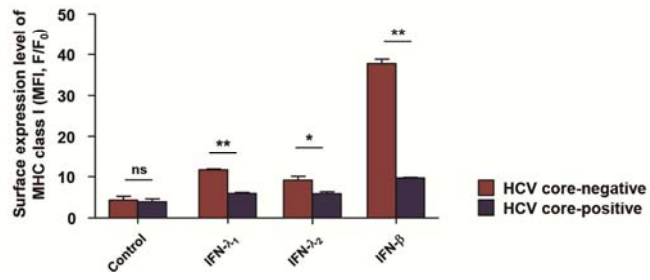


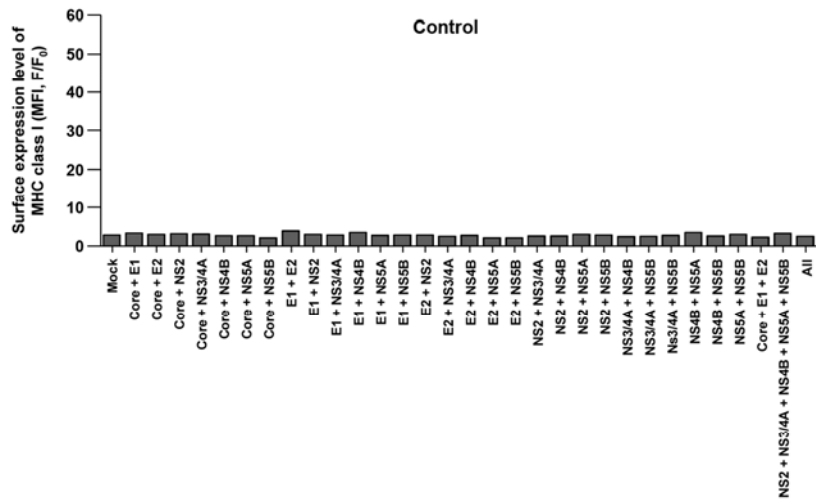
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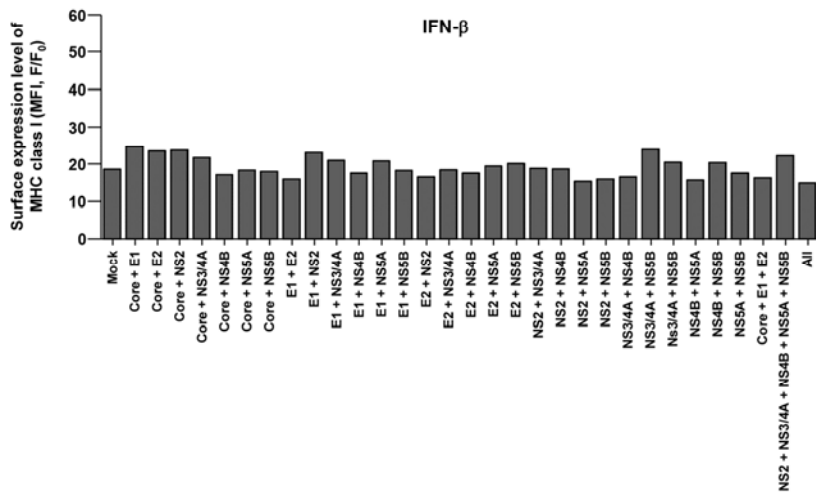
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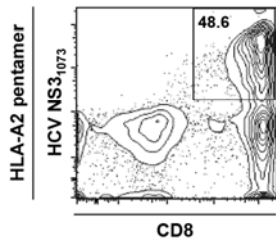
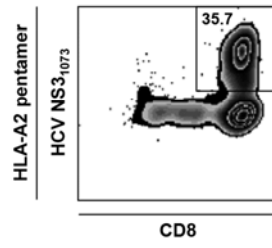


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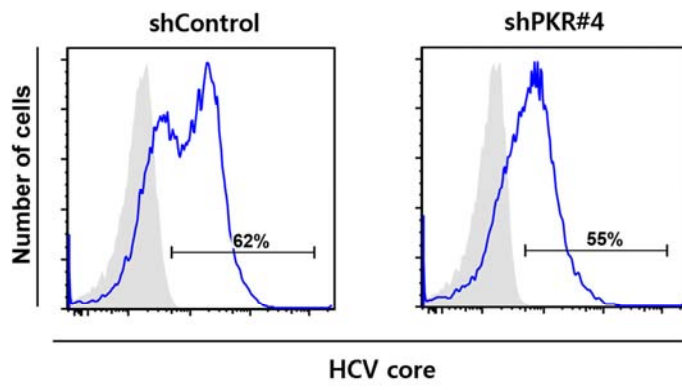


B



A**B**

ACCEPTED MANUSCRIPT



1 **Supplementary Materials and Methods**

2

3 ***Cell culture***

4 Huh-7.5 cells were cultured at 37°C with 5% CO₂ in Dulbecco's Modified Eagle Medium (DMEM;
5 Welgene) supplemented with 10% fetal bovine serum (FBS; Welgene), 4.5 g/L glucose, L-glutamine, and
6 1% penicillin/streptomycin (Invitrogen). Stable HLA-A2-transfectants were maintained in complete
7 medium supplemented with 1 mg/mL G418 (A.G. Scientific, San Diego, CA). PKR-silenced cell lines
8 were maintained in complete medium supplemented with 1 µg/mL puromycin (Sigma-Aldrich). HCV-
9 specific CD8⁺ T cells were expanded in RPMI 1640 (Welgene) supplemented with 5% FBS, 1%
10 penicillin/streptomycin in the presence of relevant cytokines as described below. For HCV RNA
11 transfection studies, Huh-7.5 cells were transfected with *in vitro*-transcribed, replication-competent JFH-1
12 using DMRIE-C reagent (Invitrogen,).

13

14 ***Confocal microscopy***

15 The antibodies used for confocal imaging included mouse monoclonal anti-HCV core IgG₁ (Clone C7-50),
16 rabbit monoclonal anti-MHC class I (Clone EP1395Y), AlexaFluor 594-conjugated donkey anti-mouse
17 IgG (Invitrogen), and AlexaFluor 488-conjugated donkey anti-rabbit IgG (Invitrogen). HCV-infected or
18 uninfected Huh-7.5 cells were plated on 4-well Lab-Tek Chamber slides (Nunc, Naperville, IL) and
19 treated with IFN-β for 24 h. Cells were fixed with 4% paraformaldehyde in PBS, permeabilized with 0.2%
20 (v/v) Triton-X100, and incubated with primary antibodies (1:300 dilution). After washing with PBS, the
21 slides were incubated with secondary antibody conjugates (1:500 dilution). After nuclear staining with
22 DAPI and mounting, images were acquired on a LSM-710 confocal laser scanning microscope (Carl
23 Zeiss, Jena, Germany).

24

1 ***PKR shRNA clones***

2 Five validated clones of MISSION[®] shRNA bacterial glycerol stocks with lentiviral constructs expressing
3 PKR shRNA were purchased from Sigma-Aldrich: NM_002759.x-2288s1c1 (TRCN0000001379),
4 NM_002759.x-1409s1c1 (TRCN0000001381), NM_002759.x-2010s1c1 (TRCN0000001382),
5 NM_002759.1-1870s1c1 (TRCN0000196400), and NM_002759.1-1580s1c1 (TRCN0000197012).

6
7 ***TaqMan real-time PCR***

8 Total RNA isolation, cDNA synthesis, and TaqMan real-time PCR were performed as previously
9 described²². In brief, total RNA was isolated with the RNeasy Mini kit (Qiagen, Valencia, CA), and
10 cDNA was synthesized using the First-Strand cDNA Synthesis kit (Origene/Marligen Biosciences,
11 Ijamsville, MD). TaqMan Gene Expression Assays (Applied Biosystems, Foster City, CA) were used to
12 determine mRNA levels of HLA-A. Target mRNA levels were normalized to an endogenous reference
13 gene (β -actin).

14
15 ***Statistical analysis***

16 Data are presented as mean + standard error of mean. Two-tailed Mann-Whitney U test or unpaired t-test
17 was performed where appropriate. Statistical analyses were performed with GraphPad Prism version 5.01
18 (GraphPad Software, San Diego, CA). A *p* value of less than 0.05 was considered statistically significant.

19

20

1 **Supplementary Figure legends**

2

3 **Supplementary Figure 1. IFN- λ -induced MHC class I expression is attenuated by HCV infection.**

4 Huh-7.5 cells were inoculated with HCVcc at 0.01 MOI and cultured for 3 days, with an infection rate of
5 30-45%. Cells were then treated with 100 ng/mL IFN- λ_1 , 100 ng/mL IFN- λ_2 or 3 ng/mL IFN- β or were
6 left untreated (control). (A) The expression level of MHC class I in HCV-infected (blue) or uninfected
7 (red) cells was analyzed by co-staining for HCV core and MHC class I and flow cytometry. (B) Data from
8 three independent experiments are presented as a bar graph (mean + SEM). *, $P < 0.05$; **, $P < 0.01$; ns,
9 not significant.

10

11 **Supplementary Figure 2. Effect of HCV gene transfection on IFN-induced MHC class I expression.**

12 Huh-7.5 cells were transfected with pairs of HCV gene plasmids in 28 combinations. In addition,
13 transfection was performed with all structural protein genes (core, E1 and E2), all non-structural protein
14 genes (NS2, NS3/4A, NS4B, NS5A and NS5B) or all HCV genes (core, E1, E2, NS2, NS3/4A, NS4B,
15 NS5A and NS5B). The transfected cells were treated with 3 ng/mL IFN- β for 24 h. The expression level
16 of MHC class I was analyzed by flow cytometry (A), and the MFI of MHC class I expression is presented
17 (B). F, fluorescence intensity of stained sample; F_0 , fluorescence intensity of isotype control.

18

19 **Supplementary Figure 3. HCV-specific CD8⁺ T cell lines used for co-culture experiments.**

20 NS31073 (CINGVCWTV)-specific CD8⁺ T cells and core35 (YLLPRRGPRRL)-specific CD8⁺ T cells
21 were expanded from PBMC of HLA-A2⁺ patients with acute HCV infection (genotype 1a) as described
22 in Materials and Methods. The frequency of NS31073-specific CD8⁺ T cells (A) and core35-specific
23 CD8⁺ T cells (B) was determined with MHC class I pentamers.

1
2 **Supplementary Figure 4. HCV infection rates in PKR-silenced and unsilenced cells.** shControl-
3 transduced Huh-7.5/P_(HLA-A2)-(HLA-A2) (left) or shPKR#4-transduced Huh-7.5/P_(HLA-A2)-(HLA-A2) (right)
4 cells were infected with HCVcc at 0.1 MOI, cultured for 3 days, and treated with 3 ng/mL IFN- β for 24 h.
5 Then, the infection rate was determined with anti-HCV core immunostaining and flow cytometry.
6 Representative FACS histograms are presented.