

# An 85-aa segment of the GB virus type C NS5A phosphoprotein inhibits HIV-1 replication in CD4<sup>+</sup> Jurkat T cells

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GB virus type C (GBV-C) is an apparently nonpathogenic virus that replicates in T and B lymphocytes and is a common cause of persistent human infection. Among HIV-1-infected individuals, persistent coinfection with GBV-C is associated with prolonged survival, and infection of blood mononuclear cells or CD4<sup>+</sup> T cells with GBV-C and HIV *in vitro* results in significantly reduced HIV-1 replication. To date, the viral protein(s) that lead to HIV inhibition have not been identified. The GBV-C nonstructural phosphoprotein (NS5A) is predicted to have pleiotropic effects on cells, including interactions with the IFN-induced dsRNA-activated protein kinase (PKR). We studied GBV-C NS5A to determine whether it is involved in inhibition of HIV replication. GBV-C NS5A protein from an isolate that was cleared by IFN therapy did not inhibit PKR, whereas NS5A from an isolate that was not cleared by IFN-inhibited PKR function in a yeast genetic system. Both of these GBV-C NS5A proteins were expressed in a CD4<sup>+</sup> T cell line (Jurkat), and both induced a potent, dose-dependent inhibition of HIV-1 replication, thus the effect was independent of PKR inhibition. NS5A induced the release of the chemokine SDF-1 and decreased surface expression of the HIV coreceptor CXCR4, potentially explaining the HIV inhibition. Deletion mapping of the NS5A protein found that an 85-aa region between amino acids 152 and 237 inhibits HIV-1 replication. Thus, GBV-C NS5A protein alters the cellular milieu necessary for HIV-1 replication and may provide a previously undescribed therapeutic approach for anti-HIV therapy.

chemokine receptor | chemokines | nonstructural proteins

**G**B virus C (GBV-C) is a human flavivirus that is present in 1–3% of healthy U.S. blood donors (reviewed in ref. 1). Because of shared modes of transmission, up to 86% of HIV-positive individuals have evidence of active (39.6%) or prior (46%) GBV-C infection (1, 2). Although viremia may persist for decades in some individuals, most immune-competent hosts clear GBV-C infection concurrently with the development of antibodies to the envelope glycoprotein E2 (3, 4), which appear to confer some, although not complete, protection against reinfection (4). Although no disease entity has yet been associated with GBV-C infection (reviewed in refs. 1 and 5), numerous studies found a significant association between persistent GBV-C viremia and prolonged survival among HIV-infected individuals (6; reviewed in ref. 1).

The GBV-C genome organization is similar to that of hepatitis C virus (HCV) (7–9). The single-strand, positive-sense RNA genome encodes a long ORF that is translated into a polyprotein of ≈3,000 aa (10). Based on comparisons with HCV, the structural proteins (E1 and E2) are thought to be cleaved from the polyprotein by cellular signal peptidases, whereas the nonstructural (NS) proteins are predicted to be cleaved by two viral proteases (NS2 and NS3; Fig. 1A) (11). A major difference between GBV-C and HCV is the apparent tissue tropism, because GBV-C appears to replicate in bone marrow cells and both T and B lymphocytes rather than in hepatocytes (10, 12–15).

HIV-1 attachment to cells involves interaction between HIV gp120 and the CD4 receptor, whereas entry into target cells requires interaction with a coreceptor, almost always the chemokine receptor CCR5 or CXCR4 (16, 17). HIV-1 isolates transmitted *in vivo* generally use CCR5 as their coreceptor and replicate in monocytes, macrophages, and primary CD4<sup>+</sup> T cells (R5 viruses) (17, 18). In contrast, HIV-1 isolates that use CXCR4 as their entry coreceptor (X4 viruses) frequently emerge in HIV-infected people later in the course of infection and replicate in T cell lines (17, 18). The natural ligands for CCR5 are the chemokines RANTES, MIP-1 $\alpha$ , and MIP-1 $\beta$ , whereas SDF-1 is the ligand for CXCR4 (19, 20). These chemokines inhibit HIV-1 by competing for binding to the chemokine receptors and, thus, inhibiting HIV-1 entry into the cell (19), and by inducing postentry inhibition of HIV reverse transcription (21–23).

Coinfection of peripheral blood mononuclear cells (PBMCs) with GBV-C and HIV-1 results in inhibition of HIV-1 replication (2, 24–27). PBMCs infected with GBV-C release significantly more RANTES, MIP-1 $\alpha$ , MIP-1 $\beta$ , and SDF-1 into culture supernatants and have significantly less CCR5 on their surface than do control cells (24, 25), suggesting that modulation of chemokines and chemokine receptors may be a mechanism by which GBV-C influences HIV-1 disease progression. No specific GBV-C viral protein has been implicated in the inhibitory effect of GBV-C on HIV-1 isolates that use the CXCR4 coreceptor.

Based on comparisons with HCV, the GBV-C NS5A protein is thought to be anchored at the N terminus in the endoplasmic reticulum and necessary for RNA replication (10, 28–30). The HCV NS5A appears to have three structured domains, one of which contains a zinc-binding motif that is conserved in GBV-C (31). Like HCV NS5A (10), there are two phosphorylated forms of GBV-C NS5A, which represent basal and hyperphosphorylated forms of the protein (32). HCV NS5A appears to modulate host antiviral responses at least, in part, by inhibiting dsRNA-activated protein kinase (PKR)-mediated phosphorylation of the eukaryotic initiation factor eIF-2 $\alpha$  (33, 34). HCV NS5A also has been shown to inhibit apoptosis and induce oxidative stress (35–38). GBV-C NS5A is also phosphorylated and inhibits

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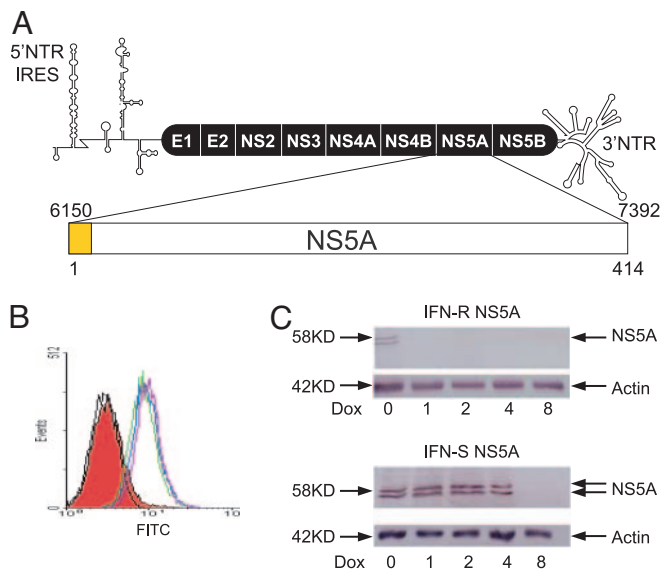
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Abbreviations: GBV-C GB Virus type C; HCV hepatitis C virus; IFN-R, IFN-resistant; IFN-S IFN-sensitive; PKR, dsRNA-activated protein kinase; VC, vector control; VC-GFP, vector control expressing GFP; X4, HIV isolates that utilize CXCR4 as their entry coreceptor.

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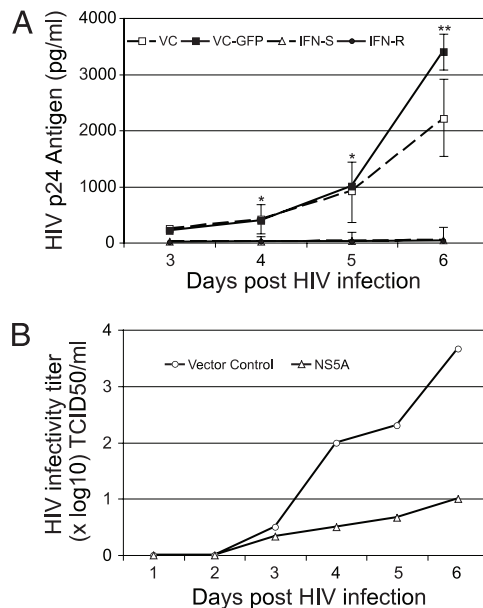
**Fig. 1.** GB virus C NS5A expression. (A) The predicted genome organization of GBV-C is shown, including the 5' nontranslated region [containing an internal ribosomal entry site (IRES)], envelope glycoproteins (E1 and E2) nonstructural proteins (NS), and 3' nontranslated region. The predicted 414-aa-long NS5A protein is encoded by nucleotides 6150–7392 (GenBank accession no. AF121950) with the putative N-terminal membrane anchoring domain shown in yellow. (B) GFP expression was detected by flow cytometry in Jurkat cells alone (black) or stably transfected with VC (red), VC-GFP (blue), or expressing the NS5A from an IFN-S isolate (GenBank accession no. DQ177420 in green) and an IFN-R isolate (DQ177421) in purple. (C) Immunoreactive GBV-C NS5A proteins from both the IFN-S and IFN-R isolates were observed in GFP-positive clones, and NS5A expression was reduced when cells were maintained in doxycycline for 48 h before preparing cell lysates.

PKR-mediated phosphorylation of eIF-2 $\alpha$  in a yeast genetic system (32). To characterize the effect of GBV-C NS5A protein on lymphocytes, stable CD4<sup>+</sup> Jurkat T cell lines were selected that expressed NS5A in a tetracycline repressible system, and the effect of GBV-C NS5A protein on HIV-1 replication was examined in these cells.

## Results

**Effect of GBV-C NS5A Protein on HIV Replication.** The predicted full-length GBV-C NS5A coding sequences from two isolates were expressed in Jurkat cells under the control of tetracycline. One of the isolates was obtained from a patient whose GBV-C viremia cleared during IFN therapy administered for coexisting HCV infection (IFN-sensitive, IFN-S; GenBank accession no. DQ177420), and the other NS5A coding sequence was obtained from a patient who did not clear GBV-C viremia (IFN-resistant, IFN-R; GenBank accession no. DQ177421) (32). NS5A was expressed in a bicistronic vector that also expressed GFP contained on the same transcript, with translation of NS5A by using capped mRNA, whereas translation of the GFP was directed by the EMC internal ribosomal entry site (IRES). GFP expression was monitored by flow cytometry (Fig. 1B). Cell lines were cloned, and those containing the NS5A sequences demonstrated two immunoreactive proteins with estimated molecular masses of 56 kDa and 58 kDa (Fig. 1C) as described in ref. 31. Expression of IFN-S NS5A was greater than that of IFN-R NS5A; however, expression of both proteins was decreased to undetectable levels when cells were incubated in doxycycline (Fig. 1C). NS5A expression was confirmed by immunofluorescence microscopy (Fig. 7, which is published as supporting information on the PNAS web site).

To determine whether GBV-C NS5A influenced HIV-1 rep-



**Fig. 2.** Expression of GBV-C NS5A in Jurkat cells inhibits HIV-1 replication. (A) Cells containing IFN-R or IFN-S GBV-C NS5A or VC or VC-GFP were infected with HIV-1 (isolate no. 1073) after doxycycline was removed for 4 days. HIV-1 p24 antigen released into culture supernatants was statistically greater in control cells on days 4–6 after HIV-1 infection. \*,  $P < 0.03$ ; \*\*,  $P < 0.001$ . Infections were performed in triplicate, and SE for IFN-R NS5A and vector control cells are shown. (B) The HIV-1 infectious titer (TCID<sub>50</sub> in PBMC culture) of culture supernatants was determined for cells expressing the VC-GFP or the IFN-S NS5A.

lication, cells were maintained in doxycycline to suppress NS5A expression. Doxycycline was removed 4 days before infection with an X4-tropic HIV-1 isolate (NIH repository no. 1073). HIV-1 replication (p24 antigen release into culture supernatants) was reduced by >99% in cells expressing either the IFN-R or the IFN-S NS5A protein compared with the vector control expressing GFP (VC-GFP) or the vector control without GFP (VC) (Fig. 2A). Previous studies demonstrated that the IFN-R NS5A protein inhibited PKR-mediated phosphorylation of eIF-2 $\alpha$ , whereas IFN-S NS5A did not (32). Both IFN-R and IFN-S NS5A expression inhibited the replication of all HIV-1 isolates tested, including clade B and clade D isolates (NIH repository nos. 317, 2969, and 2521; Fig. 8, which is published as supporting information on the PNAS web site). To ensure that NS5A was not affecting the HIV p24 antigen assay, HIV-1 infectivity in culture supernatants was measured, confirming the HIV-1 replication inhibition (Fig. 2B).

Control and NS5A expressing cells were maintained in various concentrations of doxycycline for 96 h to decrease NS5A and GFP expression before infection with HIV. Fig. 3A demonstrates that the doxycycline did not alter HIV replication in VC-GFP Jurkat cells; however, dose-dependent inhibition of HIV-1 replication was observed in the NS5A expressing Jurkat cells maintained in doxycycline (Fig. 3B). Similar results were observed in the IFN-R NS5A-expressing cells (data not shown). When IFN-S NS5A expressing cells were maintained in doxycycline (5  $\mu$ g/ml) for several passages, NS5A was not detectable by immunoblot; however, HIV replication still was inhibited, although to a lesser extent (73% inhibition on day 4; data not shown).

To determine whether the amount of NS5A protein expression in Jurkat cells is comparable with that found in GBV-C-infected PBMCs,  $2 \times 10^6$  PBMCs were infected with either a clinical GBV-C isolate (39) or a mock-control preparation, and





increased release of SDF-1 into culture supernatants and by decreased expression of CXCR4 on the cell surface (Fig. 4). Although SDF-1 contributes to HIV-1 replication inhibition, neutralizing antibodies to SDF-1 only partially blocked this inhibitory effect, suggesting that NS5A inhibits HIV-1 replication through additional mechanisms.

HIV replication inhibition required NS5A amino acids 152–237, a region that includes differing sequence polymorphisms in IFN-S and IFN-R GBV-C isolates (32). IFN-R and IFN-S GBV-C NS5A proteins were shown to have different inhibitory effects on PKR function, and only the IFN-R NS5A inhibited PKR-mediated phosphorylation of eIF-2 $\alpha$  (32). The fact that both IFN-R and IFN-S NS5A proteins inhibited HIV-1 to a similar extent indicates that the effect of NS5A on PKR function is distinct from the HIV-1 inhibitory effect. It has been proposed that GBV-C genotypes present in diverse geographic regions may have differential effects on HIV-1 (43). The IFN-S NS5A protein comes from a genotype 3 isolate (32), whereas the IFN-R NS5A is genotype 2, suggesting that genotypic differences in NS5A do not appear to influence HIV-1 inhibition. The full-length GBV-C NS5A protein and the 152–237 NS5A deletion protein both appear to be good phosphorylation substrates in Jurkat cells (Figs. 1 and 10). Because of the large number of potential phosphorylation sites in the 85 NS5A amino acids that inhibit HIV (15 for IFN-S and 18 for IFN-R), it is tempting to speculate that phosphorylation of this peptide interferes with chemokine or chemokine receptor pathways or other intracellular pathways involved in HIV replication. Thus, identification of cellular proteins that interact with NS5A, and determination of the effects of the protein on cellular gene expression, appears warranted.

In previous studies, we and others were not able to demonstrate reduced CXCR4 levels in GBV-C-infected PBMCs (24, 25). During GBV-C infection of PBMCs, NS5A is expressed only in cells supporting active GBV-C replication, and the protein is not released from cells, nor is it part of the virus particle. Hence, the effect of NS5A may be limited to actively infected cells. Because <5% of PBMCs are actively infected with GBV-C and *in vitro* GBV-C replication in PBMCs is not robust (39), the effect of GBV-C on CXCR4 may be insufficient to be detected by flow cytometry. However, PBMCs infected with GBV-C *ex vivo* demonstrated similar levels of NS5A expression, as did the Jurkat cells in which HIV replication was inhibited (Fig. 3C). By controlled expression of NS5A in a stably transformed population of Jurkat cells, we were able to demonstrate that NS5A leads to decreased surface density of CXCR4 (Fig. 4). This decreased surface expression was not related to blockade of CXCR4 detection by SDF-1, because maintaining the cells in neutralizing anti-SDF-1 antibodies did not increase the detection of CXCR4 on the cell surface.

HIV-positive people with acute dengue or measles viruses infection have marked reductions in plasma HIV RNA concentration, suggesting viral interference (44, 45). Although measles viruses has been shown to inhibit HIV replication independent of chemokines *in vitro* (46), several members of the *Flavivirus* family (HCV, Japanese Encephalitis Virus and Dengue Virus) alter the cellular release of cytokines and chemokines (47–49). Thus, chemokine and cytokine modulation may be a common feature of *Flavivirus* replication, and the effects observed with GBV-C NS5A protein may be found in related viruses as well. Nevertheless, a more prominent HIV inhibitory effect would be expected for GBV-C infection compared with other members of the *Flaviviridae* because of its propensity to result in persistent viremia (present in up to 39% of HIV-positive people; ref. 2) and because it replicates in CD4<sup>+</sup> and CD8<sup>+</sup> T lymphocytes and B lymphocytes (13). Thus, modulation of cellular chemokines occurs in the local milieu of HIV replication. In addition, GBV-C appears to be nonpathogenic (5), and, therefore, there does not appear to be a risk of comorbidity. These data provide support for the hypothesis that GBV-C replication is related causally to the observed epidemiological associa-

tion between persistent GBV-C viremia and prolonged survival in HIV-infected people (6).

GBV-C inhibits multiple HIV-1 clades (Fig. 8; ref. 25) and a zidovudine-resistant HIV-1 isolate (isolate no. 1073). Because the effect of the GBV-C NS5A protein appears to occur at the cellular level and not by targeting an HIV-1 replication enzyme or receptor binding domain, the barrier for HIV-1 to develop resistance to this effect would be expected to be greater than that seen with current treatment strategies. Thus, further characterization of this inhibitory effect may lead to previously undescribed approaches for antiretroviral therapy.

## Materials and Methods

**Expression of NS5A Proteins in Jurkat Cells.** GBV-C nucleotide sequences predicted to encode the full-length NS5A protein were amplified from plasma of individuals with GBV-C viremia as described in refs. 10 and 32. NS5A sequences were subcloned into the pTRE2-Hgy plasmid (Clontech, Mountain View, CA) modified to include a stop codon after NS5A, followed by the EMC internal ribosomal entry site element directing the translation of GFP. NS5A deletion mutants were generated by using convenient restriction sites or by PCR mutagenesis and subcloned into the pTRE2-Hgy plasmid. All sequences were confirmed by automated fluorescent dye terminator cycle sequencing (University of Iowa DNA Core Facility; automated DNA sequencer 373A, Applied Biosystems, Foster City, CA). Tet-Off Jurkat cells (Clontech) were transfected with NS5A-containing plasmids or either the VC or with VC-GFP by using the Amaxa (Gaithersburg, MD) nucleofection method. After selection in hygromycin and neomycin (200  $\mu$ g/ml), cell lines were selected for hygromycin and G418 resistance and cloned at least twice by terminal dilution.

Cells were maintained in RPMI medium 1640 containing 10% FCS with or without doxycycline to regulate NS5A protein expression. To monitor NS5A expression, Jurkat cells were lysed in RIPA buffer containing protease and phosphatase inhibitors, clarified (13,000  $\times$  g, 2 min and 4°C) and subjected to SDS/PAGE before transfer to nitrocellulose membranes (Bio-Rad, Hercules, CA) as described in ref. 32. Immunoreactive proteins were identified by using the GE3 anti-GBV-C NS5A rabbit serum kindly provided by Jungsook Kim (Genelabs Technologies, Redwood City, CA), which was generated against GBV-C nucleotide sequences 6615–6977 expressed in *Escherichia coli* (50) or by an anti-NS5A peptide sera as described in ref. 32. Alternatively, NS5A was identified in cells by indirect immunofluorescence microscopy by using the GE3 antibody (51).

**HIV Infections.** Three HIV-1 isolates (X4) representing clade B (NIH AIDS Research and Reference Reagent Program, catalog nos. 1073, 317, and 2969) and one X4 isolate representing clade D (catalog no. 2521) were studied. HIV-1 was applied to Jurkat cells (200 pg of HIV p24 antigen per 10<sup>6</sup> cells) containing NS5A sequences (with or without doxycycline to repress expression) or vector control cells with or without doxycycline. After 3 h of attachment, cells were washed and maintained in fresh media. Culture supernatants were obtained at various time points to measure HIV-1 replication and GBV-C NS5A expression. All infections were performed in duplicate or triplicate and repeated at least three times with similar results. HIV-1 replication was determined either by measuring HIV p24 antigen in culture supernatants (Retro-Tek HIV-1 p24 antigen ELISA kits; Zeptometrix, Buffalo, NY) or by measuring infectivity of culture supernatants in PBMC cultures determining the TCID<sub>50</sub> as described in refs. 2 and 24.

**Characterization of Cell Surface Receptors and Chemokine Release.** CD4, CCR5, and CXCR4 expression on the surface of Jurkat cells (with and without NS5A and doxycycline) were determined by flow cytometry as described in refs. 2 and 24. Polyclonal rabbit anti-CD4 (PE-conjugated), anti-CCR5 (FITC-conjugated), and

anti-CXCR4 (PE-conjugated) antibodies (BD Pharmingen, San Jose, CA) were used in these studies. Flow cytometry was performed by using a FACScan (Becton Dickinson, San Jose, CA). SDF-1, RANTES, MIP-1 $\alpha$ , and MIP-1 $\beta$  were detected in culture supernatant fluids by ELISA (R & D Systems, Minneapolis, MN) as described in ref. 24. Neutralizing anti-SDF-1 antibody was obtained from R & D Systems.

**Statistics.** All statistics were performed by using SigmaStat software V2.03S (Jandel Scientific, Chicago, IL). Comparisons of two samples used *t* tests, and comparisons of more than three samples used ANOVA.

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