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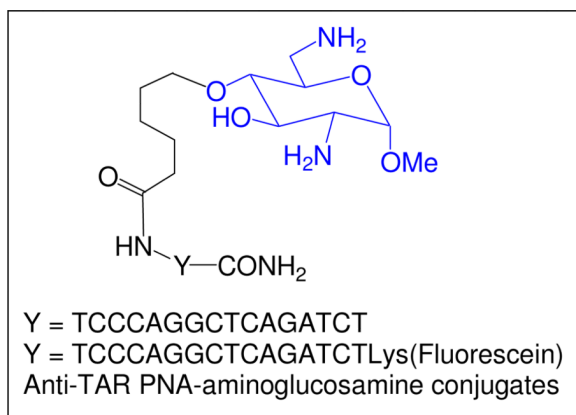
## A Peptide Nucleic Acid-Aminosugar Conjugate Targeting Transactivation Response Element of HIV-1 RNA Genome Shows a High Bioavailability in Human Cells and Strongly Inhibits Tat-mediated Transactivation of HIV-1 Transcription

Indrajit Das<sup>†</sup>, Jérôme Désiré<sup>†,§</sup>, Dinesh Manvar<sup>‡</sup>, Isabelle Baussanne<sup>†</sup>, Virendra N. Pandey<sup>‡,\*</sup>, and Jean-Luc Décout<sup>†,\*</sup>

<sup>†</sup>Université de Grenoble I/CNRS, UMR 5063, Département de Pharmaco-chimie Moléculaire, ICMG FR 2607, 470 rue de la Chimie BP 53 F-38041 Grenoble, France

<sup>‡</sup>Center for the Study of Emerging and Re-emerging Pathogens, UMDNJ-New Jersey Medical School, Department of Biochemistry and Molecular Biology, 185 South Orange Avenue, Newark, New Jersey 07103, USA

### Abstract



The 6-aminoglucosamine ring of the aminoglycoside antibiotic neomycin B (ring II) was conjugated to a 16 mer peptide nucleic acid (PNA) targeting HIV-1 TAR RNA. For this purpose we prepared the aminoglucosamine monomer **15** and attached it to the protected PNA prior to its cleavage from the solid support. We found that the resulting PNA-aminoglucosamine conjugate is stable under acidic condition, efficiently taken up by the human cells and fairly distributed in both cytosol and nucleus without endosomal entrapment since co-treatment with endosome-disrupting agent had no effect on its cellular distribution. The conjugate displayed very high target specificity *in vitro* and strongly inhibited Tat mediated transactivation of HIV-1 LTR transcription in cell culture system. The unique properties of this new class of PNA conjugate suggest it to be a potential candidate for therapeutic application.

\*Jean-Luc Décout: Phone +33 4 76 63 53 17, Fax +33 4 76 63 52 98, Jean-Luc.Decout@ujf-grenoble.fr. <sup>†</sup>Virendra N. Pandey: Phone 973-972-0660, Fax 973-972-8657 / 5594, pandey@umdnj.edu.

<sup>§</sup>Present address: Université de Poitiers/CNRS, UMR 7285, Institut de Chimie des Milieux et Matériaux de Poitiers, 4 rue Michel Brunet, 86022 Poitiers, France

### ASSOCIATED CONTENTS

Supporting Information Available: General information for the synthesis, <sup>1</sup>H and <sup>13</sup>C NMR spectra of the glucosamine derivatives and HPLC profiles of the conjugates **5a** and **5b**. This material is available free of charge via the Internet at <http://pubs.acs.org>.

## Keywords

Peptide nucleic acids; polyamide nucleic acids; aminosugar; aminoglycoside; 6-aminoglucosamine; neosamine; neamine; antiviral activity; HIV; cellular penetration; TAR; Tat

## INTRODUCTION

Aminoglycosides such as neomycin B **1** (Figure 1) are highly potent and broad-spectrum pseudo-oligosaccharidic antibiotics which act by binding to 16S rRNA, causing mRNA decoding errors, mRNA and tRNA translocation blockage, ribosome recycling inhibition and in fine protein synthesis alteration.<sup>1–5</sup> The strong affinities of aminoglycosides for other RNA targets such as several ribozymes<sup>6–16</sup> and important RNA motifs of HIV, the Transactivation Response Element (TAR), the Rev Responsive Element (RRE) and the Dimerisation Initiation Site (DIS),<sup>17–20</sup> led to intensive works for their modifications in the search for RNA-targeting drugs.<sup>21–25</sup> Amphiphilic aminoglycosides have been also reported as antimicrobial agents<sup>26–34</sup> targeting bacterial ribosomal RNA and/or membranes<sup>33,34</sup> and as efficient gene<sup>35,36</sup> and siRNA<sup>37</sup> delivery vectors.

In the antisense approach for RNA targeting,<sup>38,39</sup> the conjugation of aminoglycoside to oligonucleotides or peptide (polyamide) nucleic acids (PNA) is attractive since the strong binding of the covalently attached aminoglycoside may enhance hybridization and direct the drug design to shorter oligonucleotide sequences. Aminoglycosides are able to stabilize DNA- and RNA-triplexes, DNA-RNA hybrid triplexes and hybrid duplexes.<sup>40–42</sup> Improvements in the cellular uptake also may be provided since neomycin has been shown to assist the lipid-mediated delivery of oligonucleotides.<sup>43</sup> Due to the presence of protonated and unprotonated amino groups in the neamine core, an acido-basic catalysis of hydrolysis of the target RNA could be expected.<sup>44</sup> A number of aminoglycoside antibiotics, and in particular neomycin B, were demonstrated to promote strand cleavage of RNA oligonucleotides (minimized HIV-1 TAR element and prokaryotic ribosomal A-site) by binding and causing sufficient distortion to the RNA backbone to render it more susceptible to intramolecular transphosphorylation.<sup>45–46</sup> Regarding the conjugation of aminoglycosides to oligo-2'-deoxyribonucleotides (ODN), neomycin has been covalently attached to the 5'-end of an ODN complementary to a seven-bases long  $\alpha$ -sarcin loop RNA sequence, and its ability to enhance duplex formation has been studied.<sup>47</sup> A set of neamine- and ribostamycin-2'-O-methyl oligoribonucleotide conjugates has been prepared in order to study their nuclease activity.<sup>48</sup> Neamine and paromamine-ODN conjugates were also synthesised using click chemistry and DNA duplex stabilization was observed with a paromamine conjugate.<sup>49,50</sup> More recently, aminoglycoside (neomycin, ribostamycin, and methyl neobiosamine) conjugated to the 3'-end of 2'-O-methyl oligoribonucleotides were synthesised to target a <sup>19</sup>F labeled HIV-1 TAR RNA model which allows monitoring of the invasion by <sup>19</sup>F NMR spectrometry.<sup>51</sup> A remarkably enhanced invasion, compared to that resulting from the corresponding unmodified 2'-O-methyl oligoribonucleotide was observed with the neomycin conjugate. The conjugation of neamine to dinucleotides has been reported to remarkably decrease affinity to HIV-1 TAR RNA due to electrostatic repulsion between the phosphates, whereas conjugates of the corresponding PNA show about 2-fold binding affinities compared to that of neamine.<sup>52</sup> In the antisense approach with ODN conjugates, the strong binding of aminoglycosides to nucleic acids also probably leads to intramolecular and intermolecular charge-charge interaction between the protonated aminoglycoside core and the phosphodiester backbone of the ODN that can disturb the binding to the RNA target.

Peptide nucleic acids (PNAs) are a class of antisense DNA analogues first synthesized by Nielsen and colleagues in 1991.<sup>53</sup> The PNA molecules, devoid of sugar phosphate backbone and charges under physiological conditions, have been shown to display high affinity for complementary sequences on RNA and DNA both in single and double stranded forms.<sup>54–56</sup> PNA are highly stable and remained uncleaved when incubated with blood or cell lysate from human and bacterial cells.<sup>57</sup> Initial expectations held that PNAs would quickly enter the field of antisense as genespecific, nontoxic, and nonimmunogenic agents. However, problems associated with solubility and poor cellular uptake of this class of compounds hampered developments in this direction.<sup>58</sup> The synthesis of modified PNAs or PNA conjugates presents new means of improving their solubility and cellular uptake.<sup>58</sup>

A number of strategies have been used for improving the biodelivery of naked or modified forms of PNA in cell culture systems or in vivo.<sup>58–61</sup> The major advance in biodelivery of PNA was made by conjugating cell-transducing agents, including cell penetrating peptides (CPP). A number of reviews on CPP and their therapeutic application have been reported.<sup>62–65</sup> A triphenyl-phosphonium-tagged PNA targeted to HIV-1 TAR was found to be efficiently taken up by cells and to block HIV-1 production in cell culture.<sup>66</sup> The major drawback in the PNA biodelivery system based on cell penetrating peptides is the endosomal entrapment of PNA in the cytosol which limits its target efficacy.<sup>67</sup>

The most efficient cellular uptake of anti-HIV-1 PNA was noted with the use of cell-penetrating peptides (CPP), including penetratin, transportan and HIV-1 Tat-derived peptide as vehicle to deliver anti-HIV-1 PNA into cells.<sup>68–70</sup> CPP vehicle specially the cationic CPP such as Tat and oligo-Arg carrying PNA cargoes has been shown to enter the cell via endosomal pathway.<sup>65,71</sup> Another major deterrent is that peptide carrier are unsuitable for oral biodelivery as they are prone to cleavage by gut proteases prior to their entry into the cells.

Methods for conjugation of PNA with aminoglycosides have also been reported.<sup>52,72–76</sup> In one unique study in the antisense approach, we have conjugated the neamine core **2** (Figure 1) corresponding to rings I and II of neomycin B **1** with the 16-mer PNA **3** targeting HIV-1 TAR RNA (anti-TAR PNA).<sup>75</sup> In the resulting conjugate **4** (Figure 1), the presence of the neamine core not only imparts greater solubility and enhanced cellular uptake of PNA, it also confers a unique metal-ion-independent target specific RNA cleavage property to the conjugate.<sup>75,76</sup> We also noted that unlike PNA-CPP conjugate, PNA-neamine is compatible with oral administration in mice and efficiently absorbed in the gut (unpublished data).

Probably, the presence of positive charges is necessary for the penetration of the conjugate **4** in human cells through interaction with the membrane phospholipids and disruption of the membrane integrity. In order to decrease the number of amine functions in conjugate **4** that may be protonated under physiological conditions and could be source of toxicity and non specific interaction with the nontarget nucleic acid, our strategy was to conjugate the anti-TAR PNA sequence **3** to the small aminoglycosidic moiety 6-amino-6-deoxyglucosamine (neosamine) which corresponds to ring II of neamine (and neomycin).

*N*-acetyl-D-glucosamine is the major constituent of the peptidoglycan polymer (murein) forming the bacterial cell wall and of the chitin polymer forming the exo-skeleton of arthropods such as crustaceans (e.g. crabs). The chitosan polymers obtained by hydrolysis of chitin under basic conditions have been intensively used for preparing gene delivery vectors.<sup>77</sup> Aminosugars such as D-glucosamine have been also found to be able to weakly induce single-strand breakage in DNA that was increased in the presence of copper (II) ions.<sup>78–82</sup> In the search for antibacterial agents, 6-aminoglucosamine derivatives incorporating at the 1-position acyclic deoxystreptamine mimetics (ring I of neomycin B)

have been synthesized but none of them shows a high potency comparable to that of the natural aminoglycosides.<sup>83</sup> Here, we report on the synthesis of a 6-amino-6-deoxyglucosamine derivative and its conjugation to sequence specific PNA targeted to TAR element on HIV-1 genome. The resulting conjugate **5a** (Figure 1) is rapidly taken up by the cells and exhibits potent antiviral activity and thus suggesting a strong therapeutic potential for this class of molecules.

The prepared conjugate **5a** and the corresponding fluorescently tagged conjugate **5b** (Figure 1) were designed to carry only two positive charges under physiological conditions whereas in the neamine conjugate **4** four positive charges can be present. In **5a** and **5b**, the PNA and the linking chain were attached at the 4-oxygen atom of the neosamine moiety for a maximal distance of both protonated amine functions in regard to the possible crucial role of these functions for an efficient cellular uptake. A methyl group was introduced at the anomeric position of the neosamine moiety (1-methylneosamine) in order to stabilize the sugar ring and to minimize the steric hindrance at this position as well as the lipophilicity for clearly defining a charged polar head in the conjugate.

## SYNTHESIS

To link the 6-amino-D-glucosamine ring to the anti-TAR PNA sequence, inexpensive *N*-acetyl-D-glucosamine was used as a starting compound. The target glucosamine derivative **15** used for further coupling to the protected PNAs **3a** and **3b** was prepared in 14 steps (Scheme 1). In comparison to the glucosamine moiety, a second amine function should be added at the 6-position and a methyl group at the 1-hemicetal function as well as a linking arm at the 4-position for attachment to the PNA. This arm should carry a terminal carboxylate group for coupling through an amide junction to the *N*-terminus of the protected anti-TAR PNAs.

The first steps in the synthesis were performed according to the Literature for obtaining the intermediate **6**<sup>84</sup> through (i) methylation of the anomeric hemiacetal function, (ii) protection of the 4- and 6-hydroxyl functions by formation of the benzilidene derivative and (iii) deacetylation to release the 2-amine function.

In order to protect the 3-hydroxyl function allowing the selective introduction of the linking chain from the free 4-hydroxyl group, the 2-amine function was protected in two steps as an oxazolidinone spanning the *N*-2 and *O*-3 atoms and bearing a benzyl group on the nitrogen atom.<sup>85</sup> Treatment of the resulting compound **7** with triethylsilane and BF<sub>3</sub> diethyl etherate allowed the selective deprotection of the 4-hydroxyl function to lead to **8** which was alkylated with ethyl 6-bromohexanoate in order to introduce the protected linking arm. From the prepared ester **9**, the 6-*O*-benzyl group was removed by catalytic hydrogenation to release the 6-hydroxyl group.

A 6-azido group precursor of the additional 6-amine function was introduced through tosylation and then substitution of the tosyl group by reaction with sodium azide to lead to the derivative **12**. Treatment of **12** with aqueous NaOH opened the oxazolidinone ring releasing the 3-hydroxyl function and the benzylated 2-amino function as well as a carboxylate function at the end of the linking chain in compound **13**. In the presence of palladium hydroxide and ammonium formate, it was possible to reduce concomitantly the 6-azido group and to debenzylate the 2-amino function of **13** to lead to the methyl-2, 6-diaminoglucoopyranoside **14** bearing the linking chain with a terminal carboxylate function.

In order to (i) link the methyl-2,6-diaminoglucoopyranoside ring to the anti-TAR PNA sequence, (ii) deprotect the obtained conjugate under the conditions used to remove the

benzhydryloxy carbonyl (Bhoc) groups employed in the PNA synthesis for protection of the amine functions of the bases and (iii) cleave the conjugate from the solid support, trityl acidic-labile protecting groups were introduced to protect the 2- and 6-amino functions of **14** and afforded compound **15** in two steps. It was not possible to introduce a 4-methoxybenzyl group to protect the remaining free 3-hydroxyl group of compound **13** probably due to a strong steric hindrance generated by the O-4-linking arm and the N-2-trityl group. Thus **15** was used without protection of the 3-hydroxyl function for the coupling to the protected PNAs after exchange of the sodium counterion for a triethylammonium cation.

This salt was coupled at the *N*-terminus of the protected 16 mer anti-TAR PNA **3a** attached to its solid support of synthesis using EDC/HOBt as coupling agents. Concomitant deprotection of the neosamine moiety, of the PNA bases and cleavage from the solid support were performed in one step using TFA/anisole (1:1). The resulting conjugate **5a** was purified by HPLC on C18 reversed phase (50% coupling yield) and its structure was confirmed by mass spectrometry (MALDI: calcd 4574.88, found 4576.00).

In order to study the cellular uptake of the anti-TAR PNA-neosamine conjugate, a second anti-TAR conjugate incorporating at its C-terminus extremity a fluorescent probe was prepared from a protected lysine residue functionalized with fluorescein isothiocyanate (FITC) and attached to a solid support. For this the neosamine derivative **15** was coupled to the protected anti-TAR PNA **3b** bearing fluorescein at the C-terminus. The resulting conjugate was deprotected and cleaved from the solid support with TFA/anisole. It was purified by HPLC to give rise to the fluorescent conjugate **5b**; its structure was confirmed by mass spectrometry (MALDI: calcd 5206.09, found 5207.28).

## STABILITY OF THE PNA-GLUCOSAMINE CONJUGATE (5a) UNDER ACIDIC CONDITION

The DNA oligonucleotides based antisenses are susceptible to apurination under acidic condition and hence not suitable for oral administration. Although naked PNAs are highly stable molecule under acidic condition, we expected that its glucosamine conjugate should also be stable under similar condition. To examine this, we incubated **5a** with 0.1 M aqueous HCl at 37 °C for 3 days and then compared its HPLC elution profile with untreated sample. We found no difference in the retention time of treated and untreated samples and thus confirming its stability at high acidic condition (data not shown).

## CELLULAR UPTAKE, LOCALIZATION AND ANTIVIRAL ACTIVITY

### Cellular uptake

Since in our earlier studies we have shown that fluorescein labeled PNA-neamine when incubated with the cells at a 2  $\mu$ M concentration, approximately 33% cells became fluorescence positive after overnight incubation.<sup>76</sup> We therefore examined if conjugate **5b** carrying one 6-aminoglucosamine ring is a better candidate for biodelivery. The CEM cells were incubated with increasing concentration of **5b** at 37 °C. Cells were harvested at different time points (30 min, 3 h and 14 h) and measured for fluorescence by FACScan. We noted that **5b** is efficiently taken up by the cells in a time and concentration dependent manner, (Figure 2 A, B). At 0.2  $\mu$ M concentration, 10% of the cells were fluorescence positive within 30 min which reached to 85% within 14 h (Figure 2B). By increasing the concentration to 0.3  $\mu$ M, nearly 100% cells were positive within 14 h of incubation (Figure 2B). We further noted that the cellular uptake of **5b** was independent of temperature suggesting that its uptake is not via endocytosis which has been shown to be temperature dependent and occurs at physiological temperature.<sup>86</sup> We observed that the extent of cellular



uptake of **5b** remained unchanged at both 4 °C and 37 °C (results not shown). We also noted that PNA-aminoglucosamine is not cytotoxic since greater than 95% cells remained viable at all the concentrations (0.2 – 1 μM) tested. Thus, the cellular uptake of **5b** which carries only ring II of the neamine core was found to be improved several fold as compared to that observed with PNA-neamine conjugate **4**.<sup>76</sup> As against conjugate **4** which displayed only 30% uptake at 2 μM concentration after overnight incubation, **5b** displayed 100% cellular uptake at 0.3 μM concentration within 14 h of incubation.

To further confirm the cellular uptake of **5b** in CEM cells, we captured the images of cells treated with FITC labeled naked PNA and FITC labeled PNA-aminoglucosamine (**5b**) using confocal microscope. The cells were incubated with the indicated amount of FITC labeled naked PNA or FITC labeled PNA-aminoglucosamine **5b**. After 12 h, cells were harvested and visualized by confocal microscope (Figure 2C). The cells incubated in the presence of 0.5 μM and 1 μM of **5b**, showed approximately 90% and 99% fluorescence positive cells. In contrast FITC labeled naked PNA was unable to enter in the cell even after prolonged incubation.

It has been recognized that cellular delivery of PNAs conjugated with cell penetrating peptides occurs mainly through endocytotic pathways and thus retaining a major fraction of internalized PNA in the endosome-lysosome compartments.<sup>87</sup> Endosome disrupting agent such as chloroquine have shown to improve the release of the endosomal entrapped PNA into the cytosol. Confocal microscopy studies carried out by Turner et al.,<sup>67</sup> have demonstrated that fluorescently labeled PNA-peptide conjugates sequestered in endosomes or membrane-bound compartments in HeLa cells are significantly reduced when co-treated with chloroquine and thus resulting in improved distribution of the conjugates in the cytosol.<sup>67</sup>

We therefore resorted to examine if there is any difference in the cellular distribution of **5b** upon co-treatment with chloroquine. We used Huh7.5 cells which have much larger cytoplasmic space as compared to CEM cells and incubated these cells with 1 μM of the conjugate **5b** in the absence and presence of chloroquine as described before.<sup>67</sup> After 10 h of incubation, cells were washed by PBS and fixed with 4% paraformaldehyde. The cellular nucleus and the membrane glycoproteins were stained by DAPI and wheat germ agglutinin conjugated with rhodamine (WGA), respectively, and examined by confocal microscopy. The result shown in Figure 2D indicates that cellular distribution of **5b** remained unaffected upon co-treatment with chloroquine suggesting the absence of endosomal entrapment of **5b** in the cell.

This result is exciting since most of the biodelivery system so far developed for PNA have limitations due to their endosomal entrapment in the cytosol.<sup>67</sup> Our results with **5b** indicate that the problem associated due to endosomal compartmentalization could be circumvented by conjugating target specific PNA to the 6-aminoglucosamine ring. The localization and distribution of **5b** in both cytosol and nucleus indicate that **5a** could efficiently block Tat mediated transactivation of HIV-1 transcription in the nucleus and abort reverse transcription process of viral RNA in the cytosol.

## TARGET SPECIFICITY

Since cellular uptake of PNA conjugated with 6-aminoglucosamine was significantly enhanced, we examined if this improvement in uptake is not compromised by target specificity. We then examined the binding specificity of the PNA-aminoglucosamine conjugate (**5a**) to the TAR RNA by gel mobility shift assay using the <sup>32</sup>P-labeled 82 bp long TAR RNA transcript. We also used mutated TAR RNA carrying a deletion in the stem

(TAR-BS) and a 98 base long RNA fragment from hepatitis C virus 3' non translated region (HCV-3'NTR) as the non-target RNA and scrambled PNA-glucosamine as the negative controls. As shown in Figure 3, a distinct shift in the mobility of TAR RNA was observed due to the formation of PNA-RNA complex. The extent of gel shift of the target RNA by **5a** was stoichiometric. At 0.5:1 molar ratio of **5a** to TAR RNA, 50% gel shift was seen while complete shift was noted at 1:1 molar ratio (lane 5). We have earlier demonstrated that PNA-neamine conjugate **4** exhibits metal independent RNA nuclease activity which specifically cleaves the target RNA at the binding site upon prolonged incubation.<sup>75</sup> However, we could not demonstrate similar target specific RNA cleavage activity with conjugate **5a** either in the absence or presence of different metals ions (results not shown).

## INHIBITION OF TAT MEDIATED TRANSACTIVATION OF HIV-1-LTR

Since the PNA-glucosamine conjugate **5a** targeted to HIV-1 TAR is efficiently taken up by the cells, we examine if **5a** is able to block Tat-mediated transactivation of transcription of HIV-1 long terminal repeat (HIV-1 LTR) when supplemented in the culture medium. The significance of Tat-TAR interaction in regulating HIV-1 gene expression is well documented.<sup>88,89</sup> Activation of transcriptional elongation occurs following the recruitment of Tat to the transcription machinery via a specific interaction with TAR, a 59-residue RNA leader sequence in the HIV-1 LTR.<sup>89</sup> The main advantage of targeting the TAR element is that it is conserved and folds into a stable stem-loop structure. It was expected that binding of **5a** to sequences on the stem-loop region of TAR may destabilize the stem-loop structure of TAR and abolish Tat-TAR interaction. If this mode of action is the case then **5a** should be able to negatively impact Tat/LTR driven reporter while should have no effect in the absence of Tat.

To examine this, we used a reporter plasmid construct (pHIV-1 LTR-Luc) expressing firefly luciferase under the control of the HIV-1 LTR and pRL-SV40 expressing Renilla luciferase under the control of SV40 as an internal control. The reporter plasmids were co-transfected in CEM cells with and without pCMV-Tat. The transfected cells were grown in RPMI medium supplemented with or without different concentrations of **5a** or scrambled PNA-glucosamine conjugate. After 24 h, cells were harvested, lysed and assayed for the determination of relative expression of firefly and Renilla luciferase activity (Fig. 4). Expression of Renilla luciferase driven by the CMV promoter did not change in the absence or presence of the Tat expression clone or in the presence or absence of **5a** and was therefore used as a control to normalize for transfection efficiency. On the other hand, expression of the firefly luciferase driven by the HIV-1 LTR directly correlated with the concentration of **5a**, thus indicating its promoter specificity. These results are presented in Figure 4, as the percent inhibition of Tat-mediated transactivation of the HIV-1 LTR at indicated concentrations of **5a** or scrambled PNA-glucosamine conjugate calculated from the respective ratios of the firefly and Renilla luciferase activities.

As shown in the figure, relative luciferase activity was barely detectable in the absence of Tat (lane 2) as compared to that observed in the presence of Tat (lane 1). The expression of firefly luciferase was strongly inhibited when transfected cells were grown in the presence of **5a** (lanes 7–10). Approximately 75% inhibition of Tat-mediated transactivation of firefly luciferase was noted at 1  $\mu$ M concentration of **5a**. In contrast, only a basal level of LTR driven firefly luciferase expression was observed in the absence of Tat (lane 2) which remained unaffected by **5a** (lanes 3–6).

The naked PNA (lanes 11, 12) or scrambled PNA-glucosamine conjugate (lanes 13, 14) had no effect on the expression of firefly luciferase when supplemented in the culture medium.

These results clearly indicated that **5a** effectively blocks Tat-mediated transactivation of the HIV-1 LTR.

## ANTIVIRAL ACTIVITY AND CYTOTOXICITY

Since PNA-aminoglucosamine conjugate efficiently internalized in the cell and binds the target sequence in HIV-1 TAR RNA, we examined if the conjugate is able to block TAR mediated transactivation of HIV-1 transcription in HIV-1 infected lymphocytes. To evaluate this, we used pseudotyped HIV-1 particles carrying luciferase reporter gene to infect CEM T-lymphocyte cells. The infected cells were grown at 37 °C for 48 h and then supplemented with increasing concentration of **5a**. After 76 h, cells were lysed and assayed for firefly luciferase activity. The results shown in Figure 5 indicate that **5a** strongly inhibited HIV-1 transcription by blocking the TAR as evident by low level of expression of luciferase activity. Approximately 50% reduction in luciferase activity was observed at 800 nM concentration of **5a**. We further confirmed that dose-dependent reduction in luciferase expression is not due to cellular cytotoxicity of **5a**. As shown in Figure 5B, we did not find any difference in the [<sup>3</sup>H]thymidine incorporation when infected CEM cells were grown in the absence or presence of indicated concentration of PNA-aminoglucosamine. The cells grown in the presence of 0.1 mM and 0.5 mM PNA-aminoglucosamine (**5a**) showed similar extent of [<sup>3</sup>H]thymidine incorporation, suggesting that the cellular proliferation was not affected in the presence of **5a**.

## CONCLUSION

In conclusion, the PNA-aminoglucosamine conjugate **5a** carrying reduced number of amino groups as compared to the PNA-neamine conjugate **4**, displayed several-fold enhanced cellular uptake and target efficacy at subnanomolar concentration. It efficiently enters the cells, strongly binds to its target sequence on HIV-1 genome, blocks Tat-mediated transactivation of HIV-1 LTR and inhibits virus production in HIV-1 infected cells. The most exciting results was that, unlike peptide based PNA biodelivery system, PNA conjugated with glucosamine is uniformly localized and distributed in the cytosol and nucleus and not subjected to endosomal compartment in the cytosol. Using this novel class of small molecule to conjugate sequence specific PNA for sequestering critical regulatory elements in the viral RNA genome may have great potential in PNA based therapeutics. A strategy to enhance oral biodelivery of PNAs will certainly improve their therapeutic since PNA-aminoglucosamine conjugates being highly stable under acidic conditions corresponding to the stomachal environment (results not shown) may also be expected to be compatible with oral administration.

## EXPERIMENTAL SECTION

### General

All reagents were used as purchased from suppliers without further purification. The protected 16-mer PNA oligomers were purchased from Eurogentec. DMF was distilled in the presence of CaH<sub>2</sub>, and stored under argon atmosphere prior to use. Thin layer chromatographies were performed on silica gel (Alugram Sil G/UV<sub>254</sub>) or Alumina gel (Alugram Alox N/UV<sub>254</sub>) from Macherey-Nagel and spots were detected either by UV absorption or by charring with ninhydrin. HPLC purifications were carried out on a C<sub>18</sub> reversed-phase column (Macherey-Nagel, 10.0 × 25.0 mm) with a Dionex Ultimate 3000 apparatus. Melting points were determined with a BUCHI 510 apparatus and are reported uncorrected. <sup>1</sup>H NMR (400 MHz) and <sup>13</sup>C NMR (100 MHz) spectra were recorded with a BRUKER ADVANCE 400 spectrometer using the residual solvent signal as internal standard. HRMS were obtained from the Mass Spectrometry Service, CRMPO, at the



University of Rennes I, France, using a MICROMASS ZABSPEC-TOF spectrometer and a VARIAN MAT311 spectrometer or with a 4700 reflector spectrometer. The purities of the conjugates **5a** and **5b** determined by C<sub>18</sub> reversed phase HPLC were found to be 96.2 and 98.5%, respectively (see Supporting Information).

## Synthesis

**Glucosamine derivative 7**—To a solution of compound **6**<sup>84</sup> (3.30 g, 10.7 mmol) in dry DMF (15 mL) were added successively K<sub>2</sub>CO<sub>3</sub> (7.4 g, 54 mmol, 5 equiv) and benzyl bromide (3.18 mL, 26.7 mmol, 2.5 equiv). The solution was stirred at room temperature for 10 h under argon atmosphere. A saturated NH<sub>4</sub>Cl solution was added and the mixture was extracted with ethyl acetate (3 × 20 mL). The combined organic layers were dried over MgSO<sub>4</sub>, filtered and concentrated. The residue obtained was purified over silica gel column (EtOAc/cyclohexane, 1/3) to afford **7** (3.57 g, 84%) as a white solid; mp 186–188 °C; <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) δ 3.19 (s, 3H, OMe), 3.35 (dd, J = 12.0 Hz, 1H, H-2), 3.78–3.85 (m, 2H, H-5, H-6'), 3.98 (q, J = 8.0, 12.0 Hz, 1H, H-4), 4.21–4.24 (m, 1H, H-6), 4.45 (s, 2H, NCH<sub>2</sub>Ph), 4.58 (d, J = 4.0 Hz, 1H, H-1), 4.69 (q, J = 8.0, 12.0 Hz, 1H, H-3), 5.57 (s, 1H, CHPh), 7.33–7.38 (m, 8H, aromatic), 7.47–7.49 (m, 2H, aromatic); <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>) δ 49.0 (NCH<sub>2</sub>Ph), 55.8 (OMe), 62.3 (C-2), 65.6 (C-5), 68.7 (C-6), 73.2 (C-3), 80.4 (C-4), 97.4 (C-1), 101.6 (CHPh), 126.3, 128.5, 128.9, 129.0, 129.2, 129.4, 135.3, 136.7, 158.9 (CO); HRMS (ESI) Calcd. for C<sub>22</sub>H<sub>23</sub>NO<sub>6</sub>Na [M+Na]<sup>+</sup>: 420.14231, found 420.1411, Calcd. for C<sub>22</sub>H<sub>23</sub>NO<sub>6</sub>K [M+K]<sup>+</sup>: 436.1162, found 436.1160.

**Glucosamine derivative 8**—To a stirred solution of compound **7** (2.20 g, 5.54 mmol) in dry dichloromethane (10 mL), under argon atmosphere, was added Et<sub>3</sub>SiH (5.47 mL, 66.5 mmol, 12 equiv) and the mixture was cooled to 0 °C. To this mixture, BF<sub>3</sub>·OEt<sub>2</sub> (1.39 mL, 11.08 mmol, 2 equiv) was added dropwise then reaction mixture was stirred at 0 °C for 2 h. A saturated NaHCO<sub>3</sub> solution was added and the resulting aqueous mixture was extracted with dichloromethane (3 × 20 mL). The combined organic layers were dried over MgSO<sub>4</sub>, filtered, and concentrated. The residue was purified over silica gel column (EtOAc/cyclohexane, 1/1) to afford **8** (1.79 g, 81%) as a yellow foam; <sup>1</sup>H NMR (400 MHz, CD<sub>3</sub>OD) δ 3.13 (s, 3H, OMe), 3.34 (dd, J = 4.0, 8.0 Hz, 1H, H-2), 3.57–3.61 (m, 1H, H-5), 3.72–3.73 (m, 2H, H-6, H-6'), 3.89 (t, J = 8.0 Hz, 1H, H-4), 4.29 (d, J = 12.0 Hz, 1H, NCH<sub>2</sub>Ph), 4.37–4.57 (m, 4H, H-3, OCH<sub>2</sub>Ph, NCH<sub>2</sub>Ph), 4.61 (d, J = 4.0 Hz, 1H, H-1), 7.31–7.36 (m, 10H, aromatic); <sup>13</sup>C NMR (100 MHz, CD<sub>3</sub>OD) δ 49.5 (NCH<sub>2</sub>Ph), 55.9 (OMe), 62.8 (C-2), 69.7 (C-4), 69.9 (C-6), 74.5 (OCH<sub>2</sub>Ph), 75.7 (C-5), 78.4 (C-3), 97.7 (C-1), 128.8, 128.9, 129.1, 129.5, 129.7, 129.8, 137.1, 139.6, 161.4 (CO); HRMS (ESI) Calcd. for C<sub>22</sub>H<sub>25</sub>NO<sub>6</sub>Na [M+Na]<sup>+</sup>: 422.1580, found, 422.1587. calcd. for C<sub>22</sub>H<sub>25</sub>NO<sub>6</sub>K [M+K]<sup>+</sup>: 438.1319, found, 438.1299.

**Glucosamine derivative 9**—To a solution of compound **8** (2.80 g, 7.01 mmol) in DMF (10 mL) was added NaH (0.56 g, 14.02 mmol, 2 equiv, 60% suspension) and the mixture was stirred at room temperature for 10 min under argon atmosphere. Ethyl 6-bromo hexanoate (2.61 mL, 14.02 mmol, 2 equiv) was added and the mixture was stirred at room temperature for 8 h. A saturated NH<sub>4</sub>Cl solution was added and the resulting aqueous mixture was extracted with ethyl acetate (3 × 20 mL). The combined organic layers were dried over anhydrous MgSO<sub>4</sub>, filtered and concentrated. The residue was purified over silica gel column (EtOAc/cyclohexane, 1/3) to afford **9** (3.49 g, 92%) as a yellow foam; <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) δ 1.26 (t, J = 8.0 Hz, 3H, Me), 1.27–1.31 (m, 2H, H-3'), 1.45–1.49 (m, 2H, H-2'), 1.59–1.62 (m, 2H, H-4'), 2.27 (t, J = 8.0 Hz, 2H, CH<sub>2</sub>CO), 3.15 (s, 3H, OMe), 3.27 (dd, J = 4.0, 12.0 Hz, 1H, H-2), 3.36–3.37 (m, 1H, OCH<sub>2</sub>), 3.58–3.61 (m, 2H, H-5, H-6'), 3.68–3.71 (m, 2H, H-4, H-6), 3.78 (m, 1H, OCH<sub>2</sub>), 4.13 (q, J = 8.0, 16.0 Hz, 2H, CH<sub>2</sub>-ethyl ester), 4.41 (d, 2H, NCH<sub>2</sub>Ph), 4.45–4.52 (m, 2H, H-3, OCH<sub>2</sub>Ph), 4.55 (d, J = 4.0

Hz, 1H, H-1), 4.63 (d,  $J = 12.0$  Hz, 1H, OCH<sub>2</sub>Ph), 7.29–7.37 (m, 10H, aromatic); <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>)  $\delta$  14.4 (Me), 24.9 (C-4'), 25.7 (C-3'), 29.7 (C-2'), 34.4 (CH<sub>2</sub>CO), 48.9 (NCH<sub>2</sub>Ph), 55.6 (OMe), 60.4 (CH<sub>2</sub>-ethyl ester), 61.3 (C-2), 67.9 (C-6), 71.3 (OCH<sub>2</sub>), 72.7 (C-5), 73.7 (OCH<sub>2</sub>Ph), 75.5 (C-4), 77.4 (C-3), 96.4 (C-1), 127.9, 128.0, 128.4, 128.6, 128.9, 135.4, 138.0, 159.3 (CO-amide), 173.8 (CO-ester); HRMS (ESI) Calcd. for C<sub>30</sub>H<sub>39</sub>NO<sub>8</sub>Na [M+Na]<sup>+</sup>: 564.2573, found 564.2574, Calcd. for C<sub>30</sub>H<sub>39</sub>NO<sub>8</sub>K [M+K]<sup>+</sup>: 580.2313, found 580.2345, Calcd. for C<sub>29</sub>H<sub>39</sub>NO<sub>6</sub>Na [M-CO<sub>2</sub>+Na]<sup>+</sup>: 520.2675, found 520.2688.

**Glucosamine derivative 10**—To a solution of compound **9** (3.20 g, 5.91 mmol) in a MeOH/H<sub>2</sub>O (9/1 mL) mixture were added Pd(OH)<sub>2</sub>/C (20%) (0.17 g, 1.18 mmol, 0.2 equiv) and HCO<sub>2</sub>NH<sub>4</sub> (1.86 g, 29.5 mmol, 5 equiv). The resulting mixture was refluxed for 1.5 h, filtered through a pad of celite and concentrated. The residue was purified over silica gel column (EtOAc/pentane, 2/1) to afford **10** (2.61 g, 98 %) as a yellow foam; <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>)  $\delta$  1.25 (t,  $J = 8.0$  Hz, 3H, Me), 1.36–1.38 (m, 2H, H-4'), 1.55–1.65 (m, 4H, H-2', H-3'), 2.29 (t,  $J = 8.0$  Hz, 2H, CH<sub>2</sub>CO), 3.15 (s, 3H, OMe), 3.21 (dd,  $J = 12$  Hz, 1H, H-2), 3.49–3.53 (m, 2H, H-5, OCH<sub>2</sub>), 3.67 (t,  $J = 8.0$  Hz, 1H, H-4), 3.76 (d, 2H, H-6, H-6'), 3.83–3.87 (m, 1H, OCH<sub>2</sub>), 4.12 (q,  $J = 8.0$ , 16.0 Hz, 2H, CH<sub>2</sub>-ethyl ester), 4.41 (d, 2H, NCH<sub>2</sub>Ph), 4.50–4.55 (m, 2H, H-1, H-3), 7.30–7.36 (m, 5H, aromatic); <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>)  $\delta$  14.2 (Me), 24.6 (C-4'), 25.5 (C-3'), 29.4 (C-2'), 34.2 (CH<sub>2</sub>CO), 48.6 (NCH<sub>2</sub>Ph), 55.4 (OMe), 60.2 (CH<sub>2</sub>-ethyl ester), 61.0 (C-6), 61.2 (C-2), 71.0 (OCH<sub>2</sub>), 73.3 (C-5), 75.4 (C-4), 77.1 (C-3), 96.2 (C-1), 128.2, 128.6, 128.7, 135.2, 159.1 (CO, amide), 173.8 (CO, ester); HRMS (ESI) Calcd. for C<sub>23</sub>H<sub>33</sub>NO<sub>8</sub>Na [M+Na]<sup>+</sup>: 474.2104, found 474.2117. Calcd. for C<sub>23</sub>H<sub>34</sub>NO<sub>8</sub> [M+H]<sup>+</sup>: 452.2284, found, 452.2284.

**Glucosamine derivative 11**—To a solution of compound **10** (1.60 g, 3.54 mmol) in dry pyridine (10 mL) was added TsCl (1.69 g, 8.85 mmol, 2.5 equiv) and the mixture was stirred at ambient temperature under argon atmosphere. After completion of the reaction (TLC), saturated NaHCO<sub>3</sub> solution was added and the resulting aqueous mixture was extracted with ethyl acetate (3 × 10 mL). The combined organic layers were dried over anhydrous MgSO<sub>4</sub>, filtered and concentrated. The residue was purified over silica gel column (EtOAc/pentane, 1/2) to afford the desired compound **11** (2.06 g, 96%) as a yellow oil; <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>)  $\delta$  1.25 (t,  $J = 8.0$  Hz, 3H, Me), 1.27–1.30 (m, 2H, H-3'), 1.41–1.46 (m, 2H, H-2'), 1.57–1.61 (m, 2H, H-4'), 2.26 (t,  $J = 8.0$  Hz, 2H, CH<sub>2</sub>CO), 2.43 (s, 3H, Me), 3.05 (s, 3H, OMe), 3.17 (dd,  $J = 4.0$ , 12.0 Hz, 1H, H-2), 3.31–3.33 (m, 1H, OCH<sub>2</sub>), 3.54–3.57 (m, 2H, H-4, H-5), 3.74–3.77 (m, 1H, OCH<sub>2</sub>), 4.12 (q,  $J = 8.0$ , 16.0 Hz, 2H, CH<sub>2</sub>-ethyl ester), 4.16 (d,  $J = 4.0$  Hz, 1H, H-6'), 4.24 (dd,  $J = 4.0$ , 8.0 Hz, 1H, H-6), 4.30 (d,  $J = 16.0$  Hz, 1H, NCH<sub>2</sub>Ph), 4.40–4.47 (m, 3H, H-1, H-3, NCH<sub>2</sub>), 7.27–7.34 (m, 7H, aromatic), 7.75 (d,  $J = 8.0$  Hz, 2H, aromatic); <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>)  $\delta$  14.4 (Me), 21.8 (Me), 24.8 (H-4'), 25.6 (H-3'), 29.6 (H-2'), 34.3 (CH<sub>2</sub>CO), 48.9 (NCH<sub>2</sub>Ph), 55.7 (OMe), 60.3 (CH<sub>2</sub>-ethyl ester), 61.2 (C-2), 68.1 (C-6), 70.9 (C-5), 71.2 (OCH<sub>2</sub>), 75.2 (C-4), 76.8 (C-3), 96.3 (C-1), 128.1, 128.4, 128.6, 128.8, 128.9, 130.0, 133.0, 135.2, 145.1, 158.9 (CO-amide), 173.7 (CO-ester); HRMS (ESI) Calcd. for C<sub>30</sub>H<sub>39</sub>NO<sub>10</sub>NaS [M+Na]<sup>+</sup>: 628.2192, found 628.2187. Calcd. for C<sub>30</sub>H<sub>39</sub>NO<sub>10</sub>SK [M+K]<sup>+</sup>: 644.1932, found, 644.1927.

**Glucosamine derivative 12**—To a solution of compound **11** (1.90 g, 3.14 mmol) in DMF (10 mL) was added NaN<sub>3</sub> (2.04 g, 31.4 mmol, 10 equiv), then the solution was heated at 80 °C for 3 h under argon atmosphere. A saturated NH<sub>4</sub>Cl solution was added and the resulting aqueous mixture was extracted with ethyl acetate (3 × 10 mL). The combined organic layers were dried over anhydrous MgSO<sub>4</sub>, filtered and concentrated under reduced pressure. The residue was purified over silica gel column (EtOAc/pentane, 1/3) to afford **12** (1.36 g, 91%) as a yellow gummy liquid; <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>)  $\delta$  1.24 (t,  $J = 8.0$  Hz, 3H, Me), 1.32–1.36 (m, 2H, H-3'), 1.53–1.56 (m, 2H, H-2'), 1.58–1.64 (m, 2H, H-4'), 2.28

(t,  $J = 8.0$  Hz, 2H, CH<sub>2</sub>CO), 3.13 (s, 3H, OMe), 3.24 (dd,  $J = 4.0, 12.0$  Hz, 1H, H-2), 3.43–3.47 (m, 3H, H-6, H-6', OCH<sub>2</sub>), 3.55 (t,  $J = 8.0, 12.0$  Hz, 1H, H-4), 3.62–3.65 (m, 1H, H-5), 3.81–3.87 (m, 1H, OCH<sub>2</sub>), 4.11 (q,  $J = 8.0, 16.0$  Hz, 2H, CH<sub>2</sub>-ethyl ester), 4.34 (d,  $J = 16.0$  Hz, 1H, NCH<sub>2</sub>Ph), 4.45–4.51 (m, 3H, H-1, H-3, NCH<sub>2</sub>), 7.30–7.36 (m, 5H, aromatic); <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>)  $\delta$  14.4 (Me), 24.8 (H-4'), 25.7 (H-3'), 29.6 (H-2'), 34.3 (CH<sub>2</sub>CO), 48.9 (NCH<sub>2</sub>Ph), 51.0 (C-6), 55.7 (OMe), 60.4 (CH<sub>2</sub>-ethyl ester), 61.5 (C-2), 71.2 (OCH<sub>2</sub>), 72.5 (C-5), 76.5 (C-4), 76.9 (C-3), 96.3 (C-1), 128.4, 128.8, 128.9, 135.3, 159.0 (CO-amide), 173.7 (CO-ester); HRMS (ESI) Calcd. for C<sub>23</sub>H<sub>32</sub>N<sub>4</sub>O<sub>7</sub>Na [M+Na]<sup>+</sup>: 499.2169, found 499.2183. Calcd. for C<sub>23</sub>H<sub>32</sub>N<sub>4</sub>O<sub>7</sub>K [M+K]<sup>+</sup>: 515.1908, found, 515.1916.

**Glucosamine derivative 13**—A solution of compound **12** (1.10 g, 2.31 mmol) in a mixture of 1M NaOH/1,4-dioxan (5/5 mL, v/v) was heated at 80 °C for 20 h. The mixture was diluted with ethyl acetate and washed with brine. The separated aqueous layer was repeatedly washed with ethyl acetate. The combined organic layers were dried over anhydrous MgSO<sub>4</sub>, filtered and concentrated. The residue obtained was purified over silica gel column (CH<sub>2</sub>Cl<sub>2</sub>/MeOH, 9/1) to afford **13** (0.72 g, 74%) as a yellow gummy liquid; <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>)  $\delta$  1.37–1.43 (m, 2H, H-3'), 1.53–1.64 (m, 4H, H-2', H-4'), 2.30 (t,  $J = 4.0, 8.0$  Hz, 2H, CH<sub>2</sub>CO), 2.70 (dd,  $J = 4.0, 12.0$  Hz, 1H, H-2), 3.13 (t,  $J = 8.0$  Hz, 1H, H-4), 3.31 (s, 3H, OMe), 3.40 (dd,  $J = 8.0, 12.0$  Hz, 1H, H-6'), 3.48 (dd,  $J = 12.0$  Hz, 1H, H-6), 3.51–3.55 (m, 1H, OCH<sub>2</sub>), 3.62–3.66 (m, 1H, H-5), 3.76 (t,  $J = 8.0, 12.0$  Hz, 1H, H-3), 3.88–3.94 (m, 1H, OCH<sub>2</sub>), 3.96 (s, 2H, NCH<sub>2</sub>Ph), 4.59 (d,  $J = 4.0$  Hz, 1H, H-1), 6.63 (bs, NH), 7.32–7.37 (m, 5H, aromatic); <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>)  $\delta$  24.9 (C-4'), 25.6 (C-3'), 30.0 (C-2'), 35.2 (CH<sub>2</sub>CO), 51.6 (NCH<sub>2</sub>Ph), 51.7 (C-6), 55.5 (OMe), 61.6 (C-2), 70.2 (C-5), 72.2 (C-3), 72.3 (OCH<sub>2</sub>), 78.9 (C-4), 97.1 (C-1), 128.2, 128.8, 129.0, 137.6, 179.5 (CO); HRMS (ESI) calcd. for C<sub>20</sub>H<sub>30</sub>N<sub>4</sub>O<sub>6</sub>Na [M+Na]<sup>+</sup>: 445.2069, found 445.2063, calcd. for C<sub>20</sub>H<sub>31</sub>N<sub>4</sub>O<sub>6</sub> [M+H]<sup>+</sup>: 423.2239, found, 423.2244. Calcd. for C<sub>19</sub>H<sub>27</sub>N<sub>4</sub>O<sub>5</sub> [M-CH<sub>3</sub>OH +H]<sup>+</sup>: 391.1981, found, 391.1972.

**Glucosamine derivative 14**—To a solution of compound **13** (0.64 g, 1.44 mmol) in a of MeOH/H<sub>2</sub>O (9/1 mL) mixture were added Pd(OH)<sub>2</sub>/C (20%) (0.04 g, 0.29 mmol, 0.2 equiv) and HCO<sub>2</sub>NH<sub>4</sub> (0.45 g, 7.20 mmol, 5 equiv). The mixture was refluxed for 1.5 h, filtered through a pad of celite and concentrated to afford **14** (0.45 g, 95%) as a white crystalline solid; mp 87–89 °C; <sup>1</sup>H NMR (400 MHz, D<sub>2</sub>O)  $\delta$  1.30–1.35 (m, 2H, H-4'), 1.51–1.60 (m, 4H, H-2', H-3'), 2.16 (t,  $J = 8.0$  Hz, 2H, CH<sub>2</sub>CO), 2.93 (dd,  $J = 4.0, 12.0$  Hz, 1H, H-2), 3.11–3.22 (m, 2H, H-4, H-6'), 3.37 (dd,  $J = 4.0, 12.0$  Hz, 1H, H-6), 3.40 (s, 3H, OMe), 3.61–3.70 (m, 2H, H-3, OCH<sub>2</sub>), 3.81–3.87 (m, 2H, H-5, OCH<sub>2</sub>), 4.85 (s, 1H, H-1); <sup>13</sup>C NMR (100 MHz, D<sub>2</sub>O)  $\delta$  25.0 (C-4'), 25.4 (C-3'), 28.9 (C-2'), 37.3 (CH<sub>2</sub>CO), 40.3 (C-6), 54.4 (C-2), 55.4 (OMe), 67.1 (C-5), 72.0 (C-3), 73.1 (OCH<sub>2</sub>), 79.7 (C-4), 98.5 (C-1), 183.6 (CO); HRMS (ESI) calcd. for C<sub>13</sub>H<sub>26</sub>N<sub>2</sub>O<sub>6</sub>Na [M+Na]<sup>+</sup>: 329.1689, found 329.1690, calcd. for C<sub>13</sub>H<sub>27</sub>N<sub>2</sub>O<sub>6</sub> [M+H]<sup>+</sup>: 307.1869, found, 307.1864. Calcd. for C<sub>13</sub>H<sub>25</sub>N<sub>2</sub>O<sub>6</sub>Na<sub>2</sub> [M-H +2Na]<sup>+</sup>: 351.1508, found, 351.1516.

**Glucosamine derivative 15**—A solution of compound **14** (0.22 g, 0.67 mmol) in a DMF/triethylamine (7/1 mL) mixture, under argon atmosphere, was stirred at room temperature for 30 min, and then a solution of trityl chloride (0.93 g, 3.35 mmol, 5 equiv) in a DMF/triethylamine (5/1) mixture was added. The resulting solution was stirred at room temperature for 8 h. A saturated NH<sub>4</sub>Cl solution was added and the resulting aqueous mixture was extracted with ethyl acetate (3 × 10 mL). The combined organic layers were dried over anhydrous MgSO<sub>4</sub>, filtered and concentrated. The residue obtained was purified over silica gel column (EtOAc/cyclohexane, 1/8 containing triethylamine) to give the tritrylated compound (0.47 g, 68%) as a white amorphous solid; HRMS (ESI) Calcd. for C<sub>70</sub>H<sub>69</sub>N<sub>2</sub>O<sub>6</sub> [M+H]<sup>+</sup>: 1033.5156, found 1033.5163.

A solution of this compound (0.45 g, 0.43 mmol) in a mixture 1M NaOH/dioxane (4/4 mL, v/v) was heated at 80 °C for 6 h, then diluted with ethyl acetate and washed with a saturated solution of NaCl. The separated aqueous layer was repeatedly washed with ethyl acetate. The combined organic layers were dried over anhydrous MgSO<sub>4</sub>, filtered and concentrated. The obtained residue was purified over silica gel column de-activated with triethylamine (CH<sub>2</sub>Cl<sub>2</sub>/MeOH, 9/1) to give **15** (0.33 g, 85%) as a white crystalline solid; mp 107–109 °C; <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>) δ 1.21–1.27 (m, 2H, H-4'), 1.30 (t, J = 4.0, 8.0 Hz, 9H, Et<sub>3</sub>N), 1.41–1.44 (m, 2H, H-3'), 1.55–1.57 (m, 2H, H-2'), 1.97–1.98 (m, 1H, H-6'), 2.19 (t, J = 8.0 Hz, 2H, CH<sub>2</sub>CO), 2.50–2.53 (m, 1H, H-6), 2.77 (d, J = 4.0 Hz, 1H, H-1), 2.81 (dd, J = 4.0, 8.0 Hz, 1H, H-2), 2.92 (q, J = 4.0, 8.0 Hz, 1H, H-4), 2.99 (s, 3H, OMe), 3.05 (q, J = 8.0, 12.0 Hz, 6H, Et<sub>3</sub>N), 3.28–3.31 (m, 1H, OCH<sub>2</sub>), 3.64 (m, 1H, H-5), 3.76–3.83 (m, 2H, H-3, OCH<sub>2</sub>), 7.16–7.30 (m, 20H, aromatic), 7.40–7.42 (m, 5H, aromatic), 7.56–7.58 (m, 5H, aromatic); <sup>13</sup>C NMR (100 MHz, CDCl<sub>3</sub>) δ 8.9 (Me, Et<sub>3</sub>N), 25.6 (C-4'), 26.1 (C-3'), 30.3 (C-2'), 36.1 (CH<sub>2</sub>CO), 45.1 (C-6), 45.4 (CH<sub>2</sub>, Et<sub>3</sub>N), 54.2 (OMe), 58.1 (C-2), 66.4 (C trityl), 70.2 (C-5), 70.3 (C trityl), 71.9 (OCH<sub>2</sub>), 73.8 (C-3), 80.7 (C-4), 97.5 (C-1), 126.3, 126.7, 126.8, 127.8, 127.9, 128.0, 128.1, 128.3, 128.8, 129.0, 129.1, 146.2, 147.0, 148.7, 179.1 (CO); HRMS (ESI) calcd. for C<sub>51</sub>H<sub>54</sub>N<sub>2</sub>O<sub>6</sub>Na [M-Na]<sup>+</sup>: 813.3880, found 813.3889, calcd. for C<sub>51</sub>H<sub>53</sub>N<sub>2</sub>O<sub>6</sub>Na<sub>2</sub> [M-H+2Na]<sup>+</sup>: 835.3699, found, 835.3696. Calcd. for C<sub>51</sub>H<sub>52</sub>N<sub>2</sub>O<sub>6</sub>Na<sub>3</sub> [M-2H+3Na]<sup>+</sup>: 857.3519, found, 857.3553.

### Procedure for the synthesis of PNA-aminoglucosamine conjugates **5a** and **5b**

To a solution of the protected glucosamine derivative **15** (18 mg, 20 μmol) in dry DMF (1 mL), under argon, were added 1-[3-(dimethylaminopropyl)]-3-ethylcarbodiimide hydrochloride (5.2 mg, 30 μmol) and 1-hydroxybenzotriazole (HOBt) (4.0 mg, 30 μmol). The resulting solution was stirred for 15 min and then the protected PNA (**3a** or **3b**) on its solid support was added under argon (PNA synthesis at 2 μmol scale). The mixture was stirred at room temperature for 1 h and filtered. The neamine PNA conjugate was then cleaved from the solid support with concomitant deprotection by treatment with TFA/anisole (1/1) for 1 h. The mixture was filtered and, then, diethyl ether was added to the solution to precipitate the PNA. HPLC purification was carried out on a C<sub>18</sub> reversed-phase column (Macherey-Nagel, 10.0 × 250 mm) with a Dionex Ultimate 3000 (detection at 260 and 280 nm). Elution was performed at 60 °C at a flow rate of 2 mL/min by building up the following gradients: 0.1% TFA in acetonitrile/0.1% TFA in water (10/90 v/v) for 10 min, then 0.1% TFA in acetonitrile/0.1% TFA in water/methanol (10/85/5 v/v/v) for **5a** or 0.1% TFA in acetonitrile/0.1% TFA in water (10/90 to 40/60 v/v for 30 min) for **5b**.

Conjugate **5a**: MALDI-TOF MS, m/z found, 4576.0088, calcd. for C<sub>183</sub>H<sub>239</sub>N<sub>91</sub>O<sub>54</sub>, 4574.8753, (non coupled and free PNA: 4286.7068). Conjugate **5b**: MALDI-TOF MS, m/z found, 5207.2808, calcd. for C<sub>216</sub>H<sub>272</sub>N<sub>94</sub>O<sub>64</sub>, 5206.0919 (non coupled free PNA: 4917.9234).

### Biological assays

**Cellular uptake of Fluorescein labeled PNA-aminoglucosamine by flow cytometry**—CEM (T-cell lymphocytes) cells were used to evaluate the efficiency of cellular uptake of conjugate **5b**. The cells were grown at 37 °C in 5% CO<sub>2</sub> environment in complete RPMI1640 medium (GIBCO, Life Technology, USA) containing 10% fetal bovine serum and 100 units each of penicillin and streptomycin. The cells growing at log phase were harvested, washed with phosphate buffered saline and resuspended in RPMI media with 2% FBS. Cells were aliquoted in 12-well plate at 0.5 × 10<sup>6</sup> cells/mL and incubated for 30 min at 37 °C. Fluorescein labeled PNA-aminoglucosamine conjugate **5b** was added to cells at varying concentrations (50 nM – 500 nM) and incubated at 37 °C. At different time points, cells were harvested, washed twice with PBS and resuspended in 1 mL PBS for

FACScan analysis on Becton Dickinson flow cytometer. To exclude the uptake by dead cells, the scanning was done in the presence of propidium iodide (1  $\mu\text{g}/\text{mL}$ ). Cell Quest Pro software (Becton Dickinson) was used to acquire and analyze events detected by FL1 detector (for FITC), which excluded FL3 detector (Propidium iodide). Graphs were generated using GraphPad Prism 4.0. We also used Huh7.5 cells which are relatively larger and have more cytoplasmic space than CEM cells.

Cells were incubated with 1  $\mu\text{M}$  PNA-glucosamine conjugate **5b** in the absence and presence of an endosome disrupting agent, chloroquine.<sup>67</sup> After 10 h of incubation, cells were washed with PBS and stained with wheat germ agglutinin conjugated with rhodamine to label the oligosaccharides component of the membrane glycoprotein (red) and with DAPI to stain the nuclear DNA (blue). Uptake of FITC labeled PNA-glucosamine **5b** shows green fluorescence measured at 488 nm. The images were acquired on Nikon AIR confocal microscope.

**Preparation of  $^{32}\text{P}$ -labeled HIV-1 TAR RNA and HCV 3'NTR RNA fragment**—The plasmid pEM-7 encoding the HIV-1 TAR under the control of the T7 promoter was linearized using HindIII and transcribed to generate 82 base long run-off transcript of HIV-1 TAR RNA using T7 RNA polymerase transcription kit (Roche Diagnostics). For preparing hepatitis C virus RNA fragment corresponding to 3x' tail of HCV-3'NTR, we amplified this region by PCR and then transcribing the PCR product as described before.<sup>90</sup> The RNA transcript was internally labeled using  $\alpha$ - $^{32}\text{P}$ -CTP (3000 Ci/mmol; Perkin-Elmer Life and Analytical Sciences). The run-off transcript was purified by phenol:chloroform:isoamyl alcohol extraction followed by ethanol precipitation. The RNA was air-dried and resuspended in water pretreated with diethyl pyrocarbonate (DEPC) and finally stored at  $-70$  °C.

**Gel mobility shift assay**—The Binding affinity of the PNA-aminoglucosamine for the TAR RNA was examined by gel mobility shift assay. An aliquot of  $^{32}\text{P}$ -labeled 82 bp long TAR RNA transcript (10 nM) was incubated with varying concentrations of conjugate **5a** in a binding buffer containing 50 mM Tris-HCl (pH 7.8), 60 mM KCl, 5 mM  $\text{MgCl}_2$  in a final volume of 10  $\mu\text{L}$ . After incubation at room temperature for 1 h, 10  $\mu\text{L}$  of gel loading dye (0.27% bromophenol blue and 20% glycerol) was added to the samples and were run (100 volts) on a native 8% polyacrylamide gel using Tris-borate-EDTA buffer at room temperature for 4 h. The gel was subjected to PhosphorImager analysis (Molecular Dynamics).

**Tat-mediated transactivation of HIV-1 LTR**—CEM cells ( $0.5 \times 10^6$  cells per well in 12 well plate) were transfected with the plasmid cocktail containing pHIV-1 LTR Luc (2  $\mu\text{g}$ ) and pRL-SV40 (500 ng) in the presence and absence of pCMV-Tat (2  $\mu\text{g}$ ) using XtremeGENE 9 DNA transfection reagent (Roche). The transfection complexes were prepared in Opti-MEM medium using 6  $\mu\text{L}$  transfection reagent. The transfected cells were further grown in normal growth medium (RPMI medium containing 10% fetal bovine serum, 4 mM L-glutamine, 100 U of penicillin and 100  $\mu\text{g}$  of streptomycin per ml) at 37 °C in 5%  $\text{CO}_2$  environment. After 2 h, PNA-conjugates were supplemented in the medium and cells were further grown. After 24 h, cells were harvested and washed twice with PBS. The firefly and Renilla luciferase activities were measured using dual-luciferase reporter assay system (Dual Luciferase Assay kit). In brief, the cells were lysed using 100  $\mu\text{L}$  passive lysis buffer on a rocking shaker for 20 min. The lysate were centrifuged at 15000 rpm for 1 min and supernatant were collected in fresh tubes. Luciferase assay was performed by mixing 10  $\mu\text{L}$  cell lysate in 100  $\mu\text{L}$  Luciferase assay reagent in a 96 well plate (Fluotrac). The light emission was measured using Peckard Top count luminescence counter. Firefly luciferase



activities were quenched by the addition of 100  $\mu$ L Stop and Glo reagent to measure the Renilla luciferase activity. Transfection efficiencies were normalized by the expression of the Renilla luciferase reporter gene construct co-transfected along with the experimental plasmid.

**Production and purification of pseudotyped HIV-1 virions**—Pseudotyped HIV-1 virions were produced in 293T cells as described before<sup>66</sup> by co-transfection of pNL4-3.Luc.R.E with the pVSV-G retroviral vector, encoding the vesicular stomatitis virus protein G under the control of the CMV immediate-early promoter (BD Biosciences Clontech.) using the calcium phosphate transfection system (Life Technologies). The transfected cells were grown in complete Dulbecco's modified Eagle's medium and the culture supernatant was saved at 24, 48 and 72 h post transfection, pooled together and analyzed for p24 antigen using the ELISA p24 antigen kit (Abbott Laboratories). The pseudotyped HIV-1 virions were isolated from the culture supernatant by filtration through a 0.45  $\mu$ m pore size PVDF membrane (Millipore) followed by ultracentrifugation at 70,000 g for 45 min. The viral pellet was resuspended in complete Dulbecco's medium and stored at  $-80$  °C.

**Antiviral efficacy of conjugate 5a**—The CEM CD4<sup>+</sup> lymphocytes 12D7 (13) were grown in RPMI-1640 medium supplemented with 10% fetal calf serum, 4 mM L-glutamine at 37 °C in 5% CO<sub>2</sub> containing humidified air. Early-mid log phase cells were harvested and washed with an equal volume of PBS w/o Ca<sup>2+</sup> and Mg<sup>2+</sup>. Approximately 1 $\times$ 10<sup>6</sup> cells in 1 mL of RPMI-1640 medium were incubated with pseudovirions (14 ng; equiv p24) on a rocker for 2 h. The cells were centrifuged and after washing with 1  $\times$  PBS and resuspended in fresh complete RPMI media in a 12-well plate. After 48 h post infection, increasing amount of conjugate 5a was supplemented in the culture medium. After 76 h, the cells were harvested, washed with 1 $\times$  PBS and lysed in 1 $\times$  passive lysis buffer (Promega). The cell lysates were centrifuged at 15,000 rpm for 15 min and an aliquot of the supernatant was assayed to determine inhibition of HIV-1 transcription by measuring reporter luciferase activity. The IC<sub>50</sub> value was calculated using CalcuSyn (Biosoft).

**Effect of conjugate 5a on [<sup>3</sup>H]thymidine incorporation into cellular DNA in CEM cells**—Cellular proliferation of CEM cells in the presence of conjugate 5a was determined by estimating the levels of [<sup>3</sup>H] thymidine incorporated in their nuclei. Briefly, freshly split cells were grown in the absence or presence of conjugate 5a (10  $\mu$ M) and supplemented with 10  $\mu$ Ci of [methyl-3H]thymidine/mL (83.7 Ci/mmol). The cells were harvested at 3, 12, 24, 36, and 48 h, washed with PBS, and resuspended in 200  $\mu$ L of lysis buffer containing 1% NP-40 in PBS. The nucleic acids were precipitated by adding cold 10% trichloroacetic acid (TCA). Precipitates were collected on GF/C glass fiber filters (Whatman, Inc., Maidstone, Kent, England) and washed extensively with ice-cold 10% TCA and once in 70% ethanol. Filters were dried and placed in scintillation vials, and radioactivity was counted in the scintillation counter. Cells treated with equal amounts of DMSO were kept as control. All tests were performed in triplicate.

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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## ABBREVIATIONS USED

<b>CPP</b>	cell-penetrating peptides
<b>DAPI</b>	4',6-diamidino-2-phenylindole
<b>FACS</b>	fluorescence activated cell sorter
<b>FITC</b>	fluorescein isothiocyanate
<b>ODN</b>	oligo-2'-deoxyribonucleotides
<b>PNA</b>	peptide (polyamide) nucleic acid
<b>TAR</b>	Trans-Activation Responsive region
<b>Tat</b>	Trans-Activator of Transcription.

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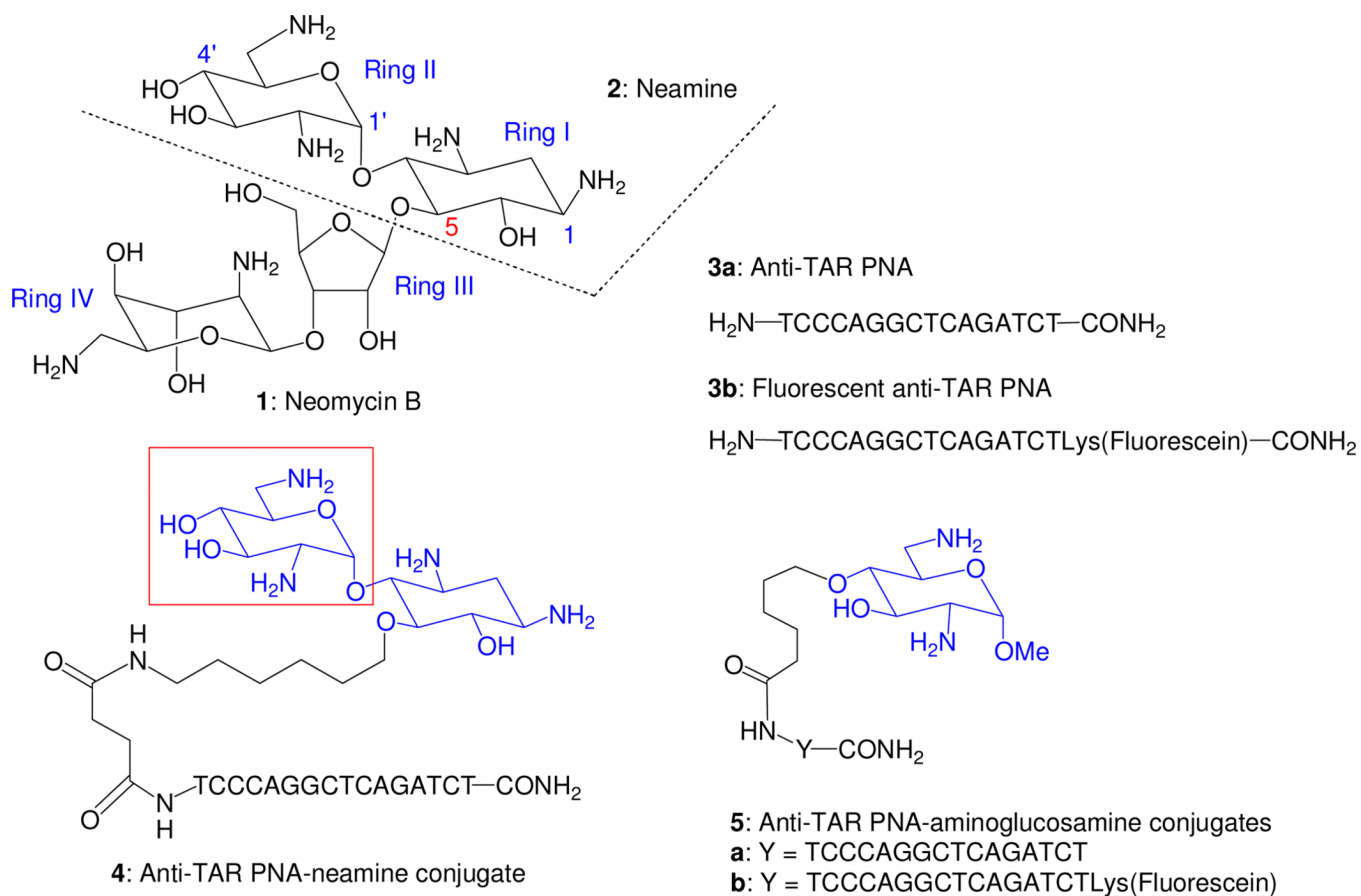
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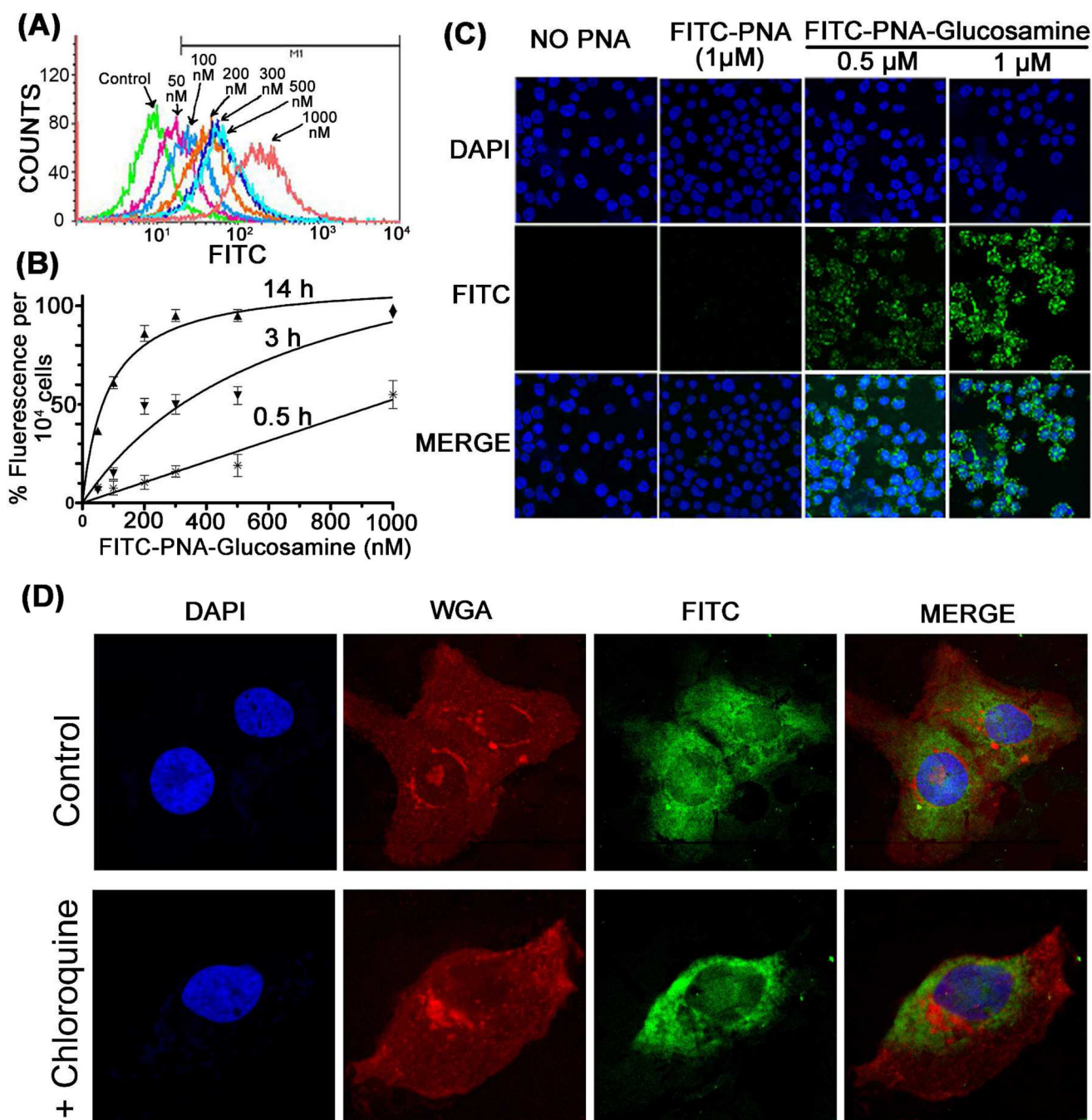
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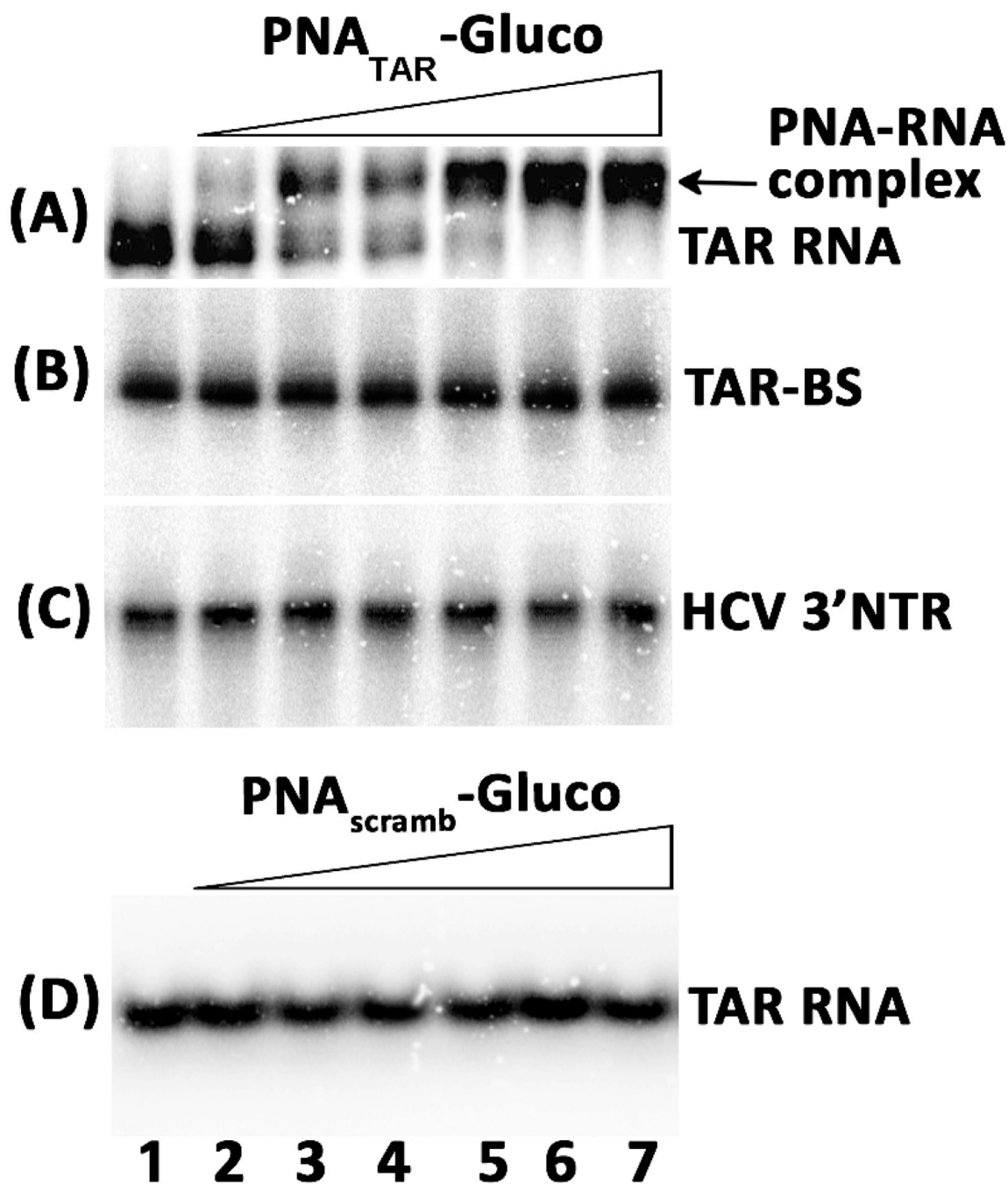
**Figure 1.**

Structure of neomycin B, neamine, the anti-TAR PNA and the corresponding neamine and 6-aminoglucosamine conjugates **4**, **5a** and **5b**.

**Figure 2.**

(A) Uptake of FITC labeled PNA-aminoglucoamine (**5b**) in CEM cells: cells were incubated with varying concentration of **5b** (50 nM to 1  $\mu$ M) in complete RPMI medium at 37  $^{\circ}$ C for 14 h. Cells were washed and resuspended in PBS containing 2% FBS and fluorescence positive cells were measured by FACSscan. (B) The graphical presentation of percent uptake of **5b** as a function of time and concentration. (C) Cellular localization of FITC labeled PNA-glucosamine **5b** in CEM cells: Cells were incubated in the absence of PNA or in the presence of 1  $\mu$ M FITC labeled naked PNA or 0.5  $\mu$ M and 1  $\mu$ M of **5b**. After 12 h, cells were washed and images were acquired. DAPI was used to stain (blue) cellular

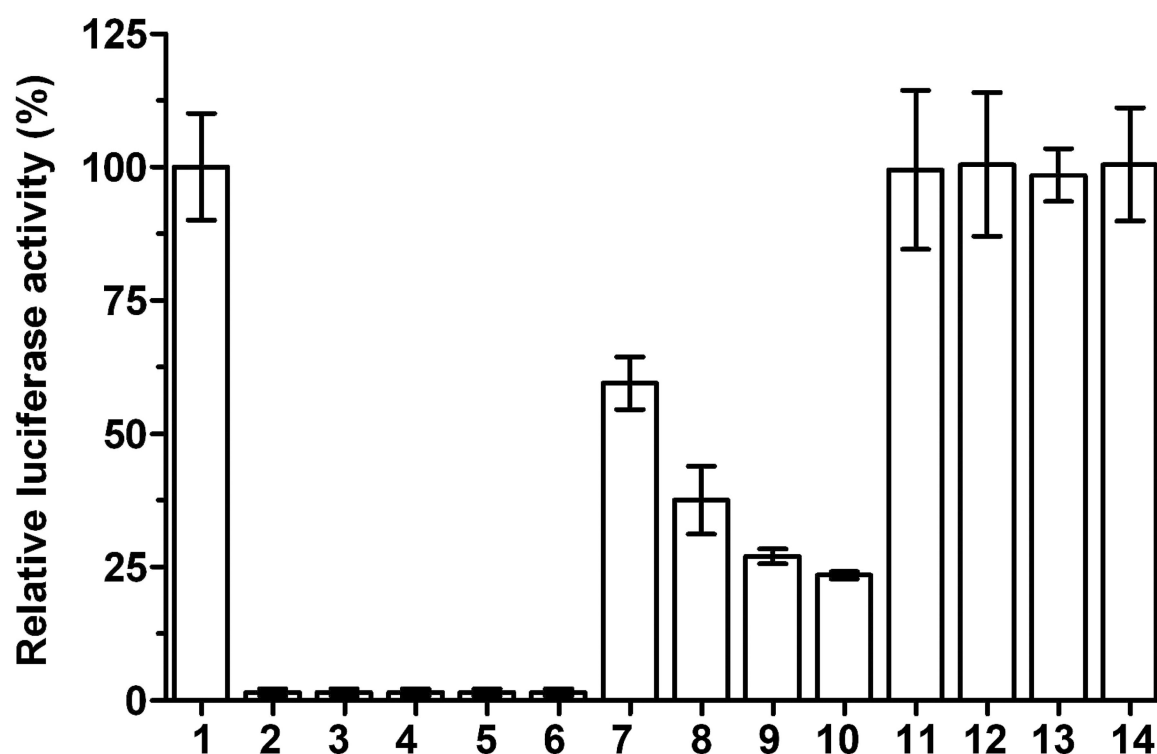
nuclei; FITC indicates the fluorescence of FITC labeled PNA or its glucosamine conjugate **5b**, measured at 488 nm. MERGE shows the distribution of **5b** in the cell. (D) Cellular localization of **5b** in Huh7.5 cells: Cells were incubated with 1  $\mu\text{M}$  of **5b** in the absence (control) and presence of chloroquine as described by Turner et al.<sup>67</sup> Images were acquired after 10 h of incubation, washing and staining with WGA-rhodamine and DAPI to label the membrane glycoproteins (red) and nuclear DNA (blue) respectively. The FITC labelled PNA-glucosamine **5b** is shown in green fluorescence. The merge indicate superimposition of blue, red and green.



**Figure 3.**

(A) Binding specificity of PNATAR-aminoglucoamine (conjugate **5a**) to its target sequence. The affinity of the conjugate **5a** (A) for its target sequence on the TAR RNA was assessed by gel mobility shift analysis. (B) TAR mutant RNA carrying a deletion at the top of the stem (TAR BS). (C) The 98 mer HCV 3'NTR RNA fragment as the non target RNA. (D) TAR RNA incubated with scrambled PNA-glucoamine conjugate **5a** as a negative control. Lane 1 through 7 represents molar ratios of PNA<sub>TAR</sub>-glucoamine or scrambled PNA-aminoglucoamine to TAR RNA of 0.0, 0.2, 0.5, 0.8, 1.0, 2.0 and 5.0, respectively.

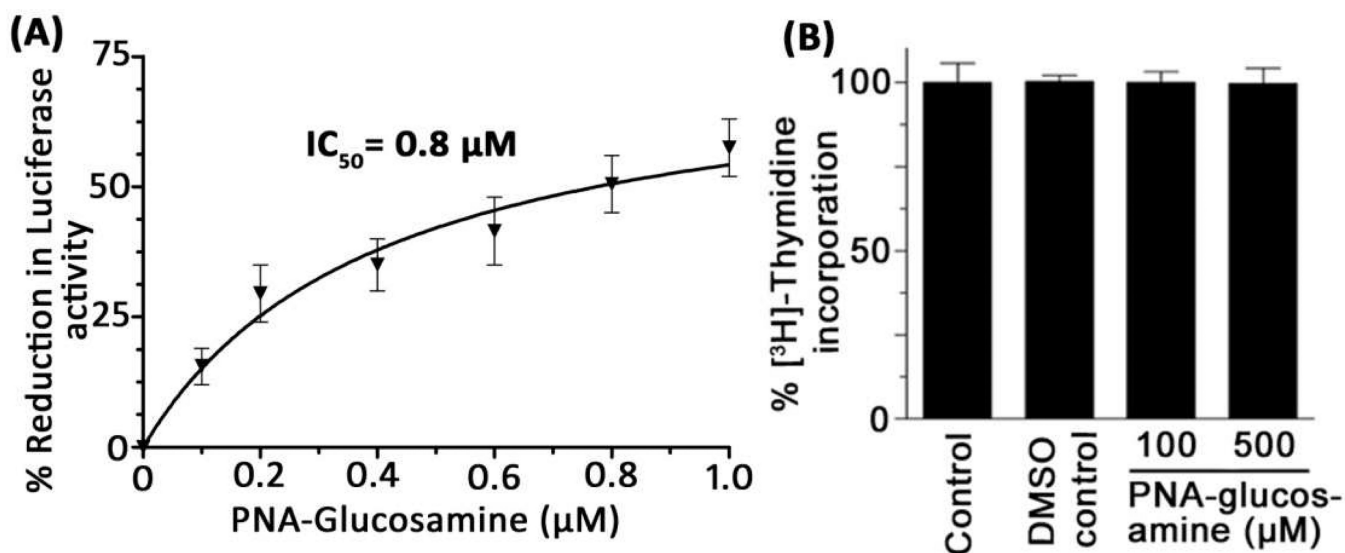




<b>pHIV-1 LTR-Luc</b>	+	+	+	+	+	+	+	+	+	+	+	+	+	+
<b>pRL-SV40</b>	+	+	+	+	+	+	+	+	+	+	+	+	+	+
<b>pCMV-Tat</b>	+	-	-	-	-	-	+	+	+	+	+	+	+	+
<b>PNA-Glucosamine</b>	-	-	0.1	0.5	1.0	2.0	0.1	0.5	1.0	2.0	-	-	-	-
<b>Naked PNA</b>	-	-	-	-	-	-	-	-	-	-	1.0	2.0	-	-
<b>Scrambled PNA</b>	-	-	-	-	-	-	-	-	-	-	-	-	1.0	2.0

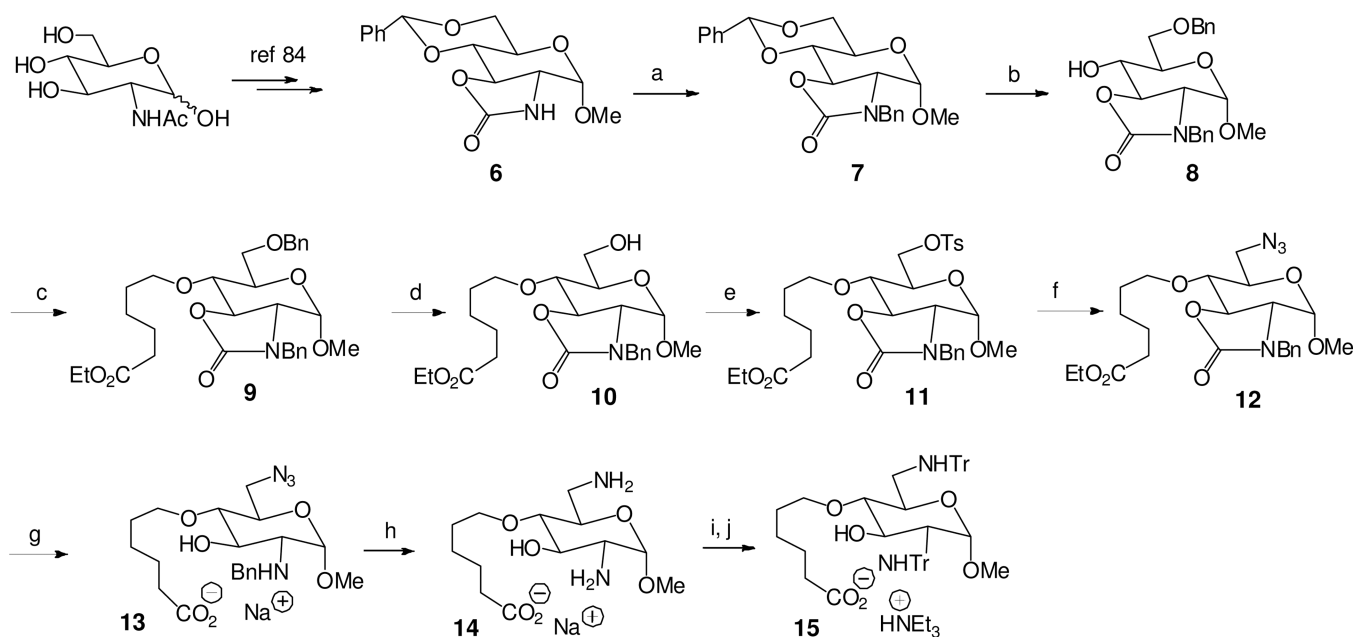
**Figure 4.**

Effect of the PNA-glucosamine conjugate **5a** on Tat-mediated transactivation: CEM cells ( $0.5 \times 10^6$  cells per well in 12 well plate) were transfected with the reporter plasmids pHIV-1 LTR-Luc ( $2 \mu\text{g}$ ) and pRL-SV40 ( $500 \text{ ng}$ ) in the presence and absence of Tat expression vector, pCMV-Tat ( $2 \mu\text{g}$ ). After 2 h of transfection, cells were supplemented with PNA conjugates in the concentration as indicated in  $\mu\text{M}$ . The transfected cells were harvested 24 h post-transfection and the luciferase activities were measured. Expression of Renilla luciferase was considered as a control for normalization of transfection efficiency. The results are presented as the relative ratio of firefly to Renilla luciferase activity in percentage by keeping the relative activity of untreated cells as 100%. Error bars represents the standard error of the mean of two independent experiments.



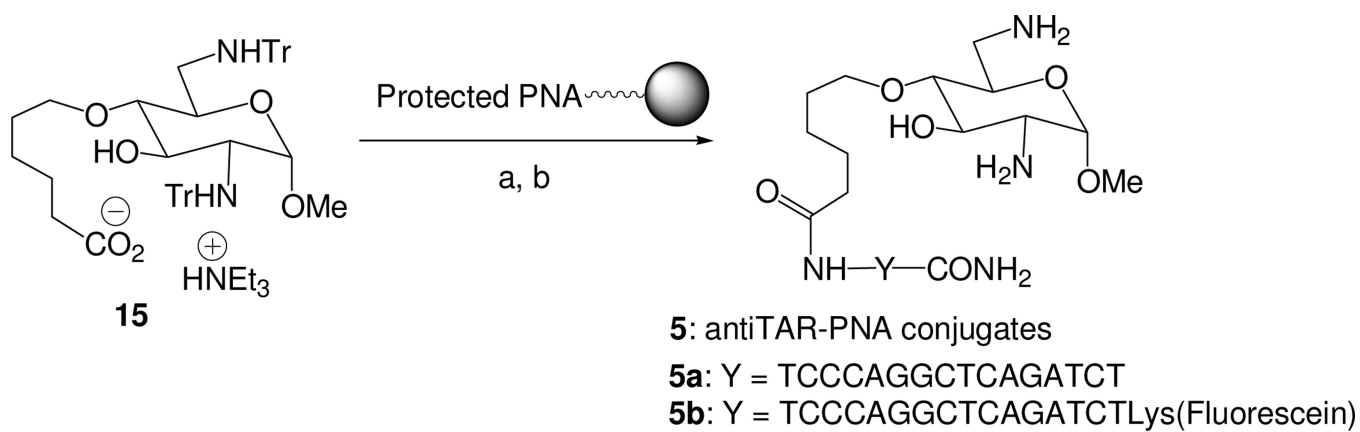
**Figure 5.**

(A) Inhibition of HIV-1 transcription by **5a**: the CEM cells were infected with pseudo-typed HIV-1 virus expressing the firefly luciferase reporter. After 48 h post-infection, indicated amount of PNA-aminoglucosamine was supplemented in the culture medium. After 76 h, cells were harvested and assayed for firefly luciferase activity in the cell extract. The results are expressed as mean value of two independent experiments with error bar. (B) Effect of **5a** on cellular proliferation: CEM cells were grown in the absence or presence of 0.1 mM and 0.5 mM of **5a** dissolved in DMSO. The final concentration of DMSO supplemented through **5a** in the medium was 5%. The culture medium was supplemented with 5 μCi of [methyl-<sup>3</sup>H]thymidine/mL. After 48 h, cells were harvested and [<sup>3</sup>H]-thymidine incorporated into the DNA was determined. The results expressed as percent of thymidine incorporation with respect to the control are the averages of three sets of independent experiments.

**Scheme 1.**

a Synthesis of the glucosamine derivative **15** for further coupling to the anti-TAR PNAs.

<sup>a</sup>Reagents and conditions: (a) BnBr, K<sub>2</sub>CO<sub>3</sub>, DMF, rt, 10 h, 84%; (b) Et<sub>3</sub>SiH, BF<sub>3</sub>·OEt<sub>2</sub>, CH<sub>2</sub>Cl<sub>2</sub>, 0 °C, 2 h, 81%; (c) 6-bromohexanoic acid ethyl ester, NaH, DMF, rt, 8 h, 92%; (d) H<sub>2</sub>, Pd/C (10%), EtOH, rt, 12 h, 98%; (e) TsCl, pyridine; rt, 10 h, 96%; (f) NaN<sub>3</sub>, DMF, 80 °C, 3 h, 91%; (g) 1M NaOH, dioxane/H<sub>2</sub>O (4/1), 100 °C, 14 h, 74%; (h) HCO<sub>2</sub>NH<sub>4</sub>, Pd(OH)<sub>2</sub>/C (20%), MeOH/H<sub>2</sub>O (9/1), reflux, 1.5 h, 95%; (i) TrCl, DMF, Et<sub>3</sub>N, rt, 8 h, 68%; (j) 1M NaOH/dioxane (1/1), 80 °C, 6 h, 85%.

**Scheme 2.**

a Synthesis of the PNA-aminoglucosamine conjugates **5a** and **5b**.

<sup>a</sup>Reagents and conditions: (a) EDC, HOBt, DMF, protected PNA **3a** or **3b**; (b) TFA/anisole (1/1), rt, 2 h.