Cytotoxicity assessment, inflammatory properties, and cellular uptake of Neutraplex lipid-based nanoparticles in THP-1 monocyte-derived macrophages

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Abstract
Current antiretroviral drugs used to prevent or treat human immunodeficiency virus type 1 (HIV-1) infection are not able to eliminate the virus within tissues or cells where HIV establishes reservoirs. Hence, there is an urgent need to develop targeted delivery systems to enhance drug concentrations in these viral sanctuary sites. Macrophages are key players in HIV infection and contribute significantly to the cellular reservoirs of HIV because the virus can survive for prolonged periods in these cells. In the present work, we investigated the potential of the lipid-based Neutraplex nanosystem to deliver anti-HIV therapeutics in human macrophages using the human monocyte/macrophage cell line THP-1. Neutraplex nanoparticles as well as cationic and anionic Neutraplex nanolipoplexes (Neutraplex/small interfering RNA) were prepared and characterized by dynamic light scattering. Neutraplex nanoparticles showed low cytotoxicity in CellTiter-Blue reduction and lactate dehydrogenase release assays and were not found to have pro-inflammatory effects. In addition, confocal studies showed that the Neutraplex nanoparticles and nanolipoplexes are rapidly internalized into THP-1 macrophages and that they can escape the late endosome/lysosome compartment allowing the delivery of small interfering RNAs in the cytoplasm. Furthermore, HIV replication was inhibited in the in vitro TZM-bl infectivity assay when small interfering RNAs targeting CXCR4 co-receptor was delivered by Neutraplex nanoparticles compared to a random small interfering RNA sequence. This study demonstrates that the Neutraplex nanosystem has potential for further development as a delivery strategy to efficiently and safely enhance the transport of therapeutic molecules into human monocyte-derived macrophages in the aim of targeting HIV-1 in this cellular reservoir.

Keywords
Nanomedicine, drug delivery nanosystem, siRNA, HIV, THP-1

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Introduction

Since the discovery of HIV-1, the spread of the virus has grown from an epidemic to one of the world’s most serious health challenges with approximately 37 million people living with the virus at the end of year 2016.\textsuperscript{1} Highly active antiretroviral therapy (ART) allows the control of viral replication, delay or prevention of the progression to AIDS, increase of survival, and improvement of the quality of life of HIV-infected individuals. However, ART does not eliminate the virus. Hence, HIV is able to persist in the infected host even during prolonged ART.\textsuperscript{2,3} Indeed, residual viremia is still detected in patients on ART when very sensitive methods are used\textsuperscript{4} and HIV-1 reverts to measurable plasma level in less than 2 weeks when ART is interrupted.\textsuperscript{5,6} It has been shown that HIV is able to establish reservoirs in anatomically sequestered sites such as the brain, gut, liver, and secondary lymphoid tissue where drug penetration is suboptimal, or within cells where the virus is latent, thus allowing reinfection and development of drug resistance.\textsuperscript{7–12}

Although resting CD4\textsuperscript{+} T cells are known to be the main cellular reservoir for latent HIV-1 infection, cells belonging to the monocyte/macrophage lineage have been described as the most important reservoirs outside the bloodstream.\textsuperscript{13–15} Findings suggest that in the presence of antiretroviral drugs, macrophages are likely the main source of plasma viremia.\textsuperscript{16,17} In contrast to T cells, HIV infection in monocyte/macrophages is less cytotoxic and renders them more resistant to apoptosis, thus extending their lifespan and making these cells a persistent reservoir of HIV regardless of the presence of ART.\textsuperscript{13,18,19} Moreover, high expression of efflux transporters in monocyte/macrophages has been shown to contribute to the resistance of these cells to antiretroviral treatment.\textsuperscript{20,21} Altogether HIV infection in macrophages may result in continued viral replication and the formation of HIV reservoirs. Therefore, the development of new strategies for eliminating or preventing viral replication in monocyte/macrophages is urgently needed to achieve viral eradication in HIV-1-infected patients.

One promising approach to enhance drug delivery and target HIV in reservoir sites is the use of nanomaterials.\textsuperscript{22–25} We and others have shown the efficiency and benefits of using nanoformulations to deliver HIV therapeutics and target HIV in infected cells and reservoir sites including macrophages.\textsuperscript{26–31} The aim of this study was to evaluate the potential of the Neutraplex (Nx) nanosystem to deliver therapeutic molecules in HIV reservoir sites using the THP-1 monocyte-derived macrophages (MDMs) as the in vitro cellular model. Cytotoxicity of Nx nanoparticles (NPs) was assessed on cell viability and membrane integrity of THP-1 MDMs. In addition, their inflammatory profile was determined by evaluating their effects on polymorphonuclear neutrophils (PMNs) apoptosis and on pro-inflammatory cytokine secretion in exposed THP-1 MDMs. Furthermore, their efficiency to deliver anti-HIV therapeutics in human macrophages was evaluated by confocal microscopy studies in THP-1 MDMs using small interfering RNAs (siRNAs) complexed to Nx NPs (nanolipoplexes) as a proof-of-concept strategy. The unique properties of the Nx nanosystem are that it allows the production of nanolipoplexes NPs of opposite surface charges with similar size and great stability without changing the chemical composition, therefore the possibility to study the impact of surface charge on cellular uptake, internalization, and drug delivery. Here, two nanolipoplexes formulations with opposing surface net charge but similar size, Nx+12/siRNA (cationic) and Nx-40/siRNA (anionic), were prepared and compared for their capacity to be internalized into human macrophages and to deliver active siRNAs to block HIV replication using the in vitro TZM-bl infectivity assay.

Materials and methods

Materials

Dioctadecylamidoglycylspermine (DOGS) was purchased from PolyPeptide Laboratories (Strasbourg, France), dioleoylphosphatidylethanolamine (DOPE), fluorescein-labeled DOPE, and cardiolipin from Sigma-Aldrich (Oakville, Ontario, Canada). SiRNAs and RNase-free water were purchased from Applied Biosystems (Foster City, California, USA). THP-1 monocyte/macrophage cell line was obtained from the American Type Culture Collection.\textsuperscript{32,33} The human epithelial cancer cervical cell line TZM-bl was obtained through the US National Institutes of Health AIDS Research and Reference Reagent Program from Dr John C Kappes, Dr Xiaoyun Wu, and Tranzyme Inc (Durham, North Carolina, USA).\textsuperscript{34,35} Roswell Park Memorial Institute (RPMI)-1640 and Dulbecco’s minimum essential medium (DMEM) cell culture medium, phosphate-buffered saline (PBS), Hank’s balanced salt solution (HBSS), phorbol myristate acetate (PMA), trypsin–ethylene diamine tetraacetic acid (EDTA) solution, dimethyl sulfoxide (DMSO), LysoTracker\textsuperscript{®} Blue DND-2, and Texas Red\textsuperscript{®} wheat germ agglutinin (WGA) were all purchased from Life Technologies (Burlington, Ontario, Canada). CellTracker Blue\textsuperscript{®} (CTB) Cell Viability, CytoTox-96\textsuperscript{®} (lactate dehydrogenase (LDH)), and Beta-Glo\textsuperscript{®} assay kits were obtained from Promega (Madison, Wisconsin, USA). Capture beads were purchased from Lumix (Austin, Texas, USA). Cytokine antibody pairs and standards were obtained from R & D Systems Inc. (Minneapolis, Minnesota, USA) (interleukines IL-1\textbeta, IL-8) and interleukins IL-6, IL-10, tumor necrosis factor alpha TNF-\textalpha, and streptavadin-phycocerythrin (streptavadin-PE) was obtained from Thermo Fisher Scientific, Waltham, Massachusetts, USA. Milli-Q and deionized water were produced by a Millipore water purification system (EMD Millipore, Etobicoke, Ontario, Canada). Analytical grade solvents and reagents were purchased from Sigma-Aldrich unless otherwise specified.
SiRNA

Synthetic double-stranded siRNA sequences were dissolved in sterile RNase-free water at a concentration of 50 μM and stored as indicated by the manufacturer. SiRNAs used in this study include siCCR5 targeting C–C chemokine receptor type 5 (CCR5) labeled at the 3’ end with Cy3 (sense strand: 5’-GGGCUCUAAUUAAUAGGCUCdTdT-3’), siCXCR4 targeting chemokine C–X–C motif receptor 4 (CXCR4; sense strand: 5’-GGAGCUGUGGCUGAAAdtdt-3’), and a scrambled siRNA that bears no homology to the human genome which was used as a non-targeting random control siRNA (sense strand: 5’-AGUACUGCUUACGAUCGdtdt-3’).

Preparation of lipidic NPs and nanolipoplexes

Nx lipid-based NPs were prepared as previously described. Briefly, an ethanol solution of a lipid mixture containing 1-mg DOGS, 1-mg of DOPE, and 0.5-mg cardiolipin was injected in an aqueous solution to form small unilamellar vesicles (SUVs) with a final concentration of 6.25-mg/mL lipids. This preparation can be kept for over 9 months at 4°C. To prepare fluorescein-labeled Nx (fluorescein isothiocyanate (FITC)-Nx) NPs, FITC-DOPE was slightly mixed and stored at 4°C until their use. Nanolipoplexes (Nx/siRNA) were prepared as previously described by mixing the SUV suspension with siRNA in sterile RNase-free water at two different lipids: nucleic acid (NA) weight ratios to obtain cationic (Nx+) and anionic (Nx-40/siRNA) nanolipoplexes depending on the NA load charge in the formulation. After incubating for 30 min at room temperature, the mixture was slightly mixed and stored at 4°C until their use. Nanolipoplexes were used within 2 weeks from preparation.

Particle size and surface charge

Particle size and zeta potential were measured by dynamic light scattering (DLS) using a Zetasizer Nano ZS (Malvern Instruments Limited, Worcestershire, UK) with 633-nm laser wavelength and a measurement angle of 173° (backscatter detection) at 25°C in water (pH 7.0). Zeta (ζ) potential was employed to evaluate surface charge density of the nanoformulations. The width of the DLS hydrodynamic diameter distributions is indicated by the polydispersity index. Nanoformulations were vortexed before analysis and each sample was measured in triplicate.

Cell culture

The TZM-bl indicator cell line was generated from a clone of HeLa cells (JC.53, clone 13). TZM-bl cells express human CD4 and the human chemokine receptors CXCR4 and CCR5 and are highly sensitive to infection with diverse isolates of HIV-1. TZM-bl cells express β-galactosidase and luciferase under the control of HIV-1 promoter, which is transactivated by the Trans-Activator of Transcription (HIV-1 Tat) protein in relation to the level of virus replication. THP-1 cells were maintained in RPMI 1640 and TZM-bl cells in DMEM, supplemented with 10% heat-inactivated fetal bovine serum, 50-μg/mL gentamicin, and 2-mM L-glutamine at 37°C, 5% carbon dioxide (CO2). THP-1 MDMs were obtained by adding PMA to the culture medium at a final concentration of 10 ng/mL. THP-1 cells were incubated for 3–5 days to allow the monocytes to differentiate into macrophages. MDMs will be referred to in the text and figures hereafter as macrophages.

Cytotoxicity assays

Exposure experiments. THP-1 cells were seeded in flat-bottomed 96-well plates (Corning) at a density of 2 × 105 cells per well and allowed to differentiate. Then macrophages were washed with PBS and 100 μL of fresh complete culture medium containing different concentrations of nanoformulations was added to the cells. After the indicated time of exposure (2–48 h), cytotoxicity assays were performed as described below.

Cell viability. Cell viability was determined using the CTB Assay as per the manufacturer instructions. Various control wells were added to each plate: wells with culture medium only (background controls), wells with culture medium containing Nx (interference controls), cells without Nx (untreated controls), and cells + 10% DMSO (positive controls for cytotoxicity). After exposure to Nx, 20 μL of CTB reagent was added to each well. Absorbance was measured by top reading at 560 nm using 590 nm as a reference wavelength using a BioTek Synergy Mx microplate spectrophotometer (Thermo Fisher Scientific, Nepean, Ontario, Canada). CTB reduction was calculated by absorbance at 2-h minus absorbance at 0 h to control for background absorbance. The percentage cell viability was obtained after normalizing the data by untreated controls. Untreated cells served as 100% cell viability.

LDH release. The enzymatic activity of cytosolic LDH released extracellularly in the cell culture supernatants was measured using the CytoTox 96 nonradioactive colorimetric assay per the manufacturer instructions. After exposure to Nx nanoformulations, supernatants were collected and frozen at −80°C until their analysis for LDH activity; 50 μL of supernatant of each sample was plated in 96-well plates and 50 μL of substrate from the LDH assay kit was added to each well. The plate was then incubated for 30 min at room temperature in the dark. Absorbance was read at 490 nm using a BioTek Synergy Mx spectrophotometer. Culture medium control wells were used to correct for phenol red and endogenous LDH activity in the serum and untreated cells served to measure LDH spontaneously released by the cells.
released from cells. Cells treated with 10% DMSO served as positive controls. Data are reported as percentage release of LDH compared to untreated control cells.

**Pro-inflammatory assays**

**Cytokine analysis.** THP-1 macrophages were exposed to Nx NPs for 48 h and cytokine secretion was measured in supernatants by a Luminex multiplex bead assay using an in-house developed panel for pro-inflammatory cytokines as described earlier. Capture antibodies were linked to carboxylated beads and used for detection of cytokines (IL-1β, IL-6, IL-8, IL-10, and TNF-α). Captured cytokines were measured using biotinylated-antibodies and streptavidin-PE. Beads were analyzed using a Luminex-100 instrument with data collection software version 1.7. Cytokine concentrations were determined from standard curves calculated using the curve fitting software MasterPlex QT v. 5.0.

**Neutrophil apoptosis assays.** Freshly isolated human PMNs from healthy donors were isolated by dextran sedimentations and used for detection of cytokines (IL-1β, IL-6, IL-8, IL-10, and TNF-α). Captured cytokines were measured using biotinylated-antibodies and streptavidin-PE. Beads were analyzed using a Luminex-100 instrument with data collection software version 1.7. Cytokine concentrations were determined from standard curves calculated using the curve fitting software MasterPlex QT v. 5.0.

**Confocal studies**

THP-1 cells were seeded in a glass flat-bottomed 96-well tissue culture plates for confocal microscopy (MatTek, Ashland, Massachusetts, USA) at a density of 2 × 10⁴ cells per well with culture medium containing PMA to allow differentiation. After 5 days of incubation, differentiated THP-1 cells were washed with PBS and fresh culture medium containing FITC-labeled Nx NPs at 4.4 µg/mL or nanolipoplexes (Cy3-labeled anti-CCR5 siRNA complexed to nonfluorescent Nx NPs at a final concentration of 50 nM, corresponding to 8.8 µg/mL NPs for N+12/Cy3-siCCR5 or 2.6 µg/mL NPs for Nx-40/Cy3-siCCR5) was added to the wells. In control wells, only culture medium was added to the cells. Cells were incubated at 37°C for 30 min to 24 h, then washed with PBS and fresh culture medium without NPs was added to each well. Cellular uptake and internalization of FITC-Nx NPs or Nx/Cy3-siCCR5 nanolipoplexes were visualized using confocal laser scanning microscopy (LSM700; Carl Zeiss Canada, Toronto, Ontario, Canada). For localization studies, cells were stained for late endosomes/lysosomes using the blue fluorescent dye LysoTracker and for cellular membrane using the red fluorescent dye WGA as per the manufacturer protocols as described earlier. Briefly, live cells were stained with 75 nM LysoTracker Blue DND-22 for 30 min prior to microscopy observation. For membrane staining, cells were fixed with cold methanol for 5 min at –20°C, then washed three times with 2-mL HBSS without phenol red and incubated with WGA conjugated with Texas Red for 10 min at room temperature.

**HIV inhibition assay**

TZM-bl cells were seeded at 6 × 10³ cells per well in a 96-well microtiter plate and allowed to adhere overnight in a humidified CO₂ incubator at 37°C. Then, supernatants were removed and replaced by fresh complete DMEM culture medium containing various concentrations of cationic or anionic nanolipoplexes (N+12/ or N–40/siCXCR4 or N+12/ or N–40/random siRNA sequence) and cells were incubated for 24 h at 37°C. Following incubation, cells were rinsed to remove the nanolipoplexes suspension and were infected with HIV-1/HIV at multiplicity of infection 0.1 for 2 h at 37°C. After infection, cells were rinsed to eliminate unbound viruses and fresh complete culture medium was added. Then, cells were incubated for an additional 24 h at 37°C. After that time, HIV replication was evaluated by measuring β-galactosidase activity using the Beta-Glo assay accordingly to the manufacturer.
instructions. Briefly, 100 μL of Beta-Glo reagent was added to each well and plates were incubated for 30 min at room temperature. Light intensity of each well was measured using a BioTek Synergy 2 multimode instrument (Thermo Fisher Scientific, Nepean, Ontario, Canada). Various control wells were added to each plate: wells without cells (background controls), uninfected cells without nanolipoplexes (negative controls), infected cells without nanolipoplexes (positive controls), and infected cells treated with nanolipoplexes containing a nontargeting scrambled siRNA sequence (random controls).

Statistics
Statistical analysis was performed using SigmaPlot (version 12.5). A p value of less than 0.05 was considered as statistically significant.

Results

Physicochemical characteristics of the nanoformulations

Particle size analysis and surface charge of Nx NPs and Nx/random siRNA nanolipoplexes were determined by DLS in water and summarized in Table 1. Small particles with a diameter under 100 nm were formed. The size of both the cationic Nx+12/random siRNA (92.2 ± 5.6 nm) and the anionic Nx-40/random siRNA (93.5 ± 17.0 nm) nanolipoplexes was slightly higher compared to Nx only NPs (83.4 ± 12.5 nm), indicating the presence of the siRNAs in the nanolipoplex formulations. Homogenous lipid NPs and nanolipoplexes were obtained with polydispersity varying from 0.192 to 0.327 depending on the nanofomulations (Table 1). Zeta potential varied depending on the ratio lipid: NA as expected being less positive when higher amount of NA was used in the preparation of the nanolipoplexes. Nx NPs containing no NA formed the highest positively charged particles (+50.7 ± 2 mV) while Nx+12 nanolipoplexes containing NA at a ratio of lipid: NA of 13 formed less positively charged particles (+37 ± 5.5 mV) and the Nx-40 nanolipoplexes containing the highest amount of NA with a lipid: NA ratio of 3.9 formed highly negatively charged particles (−55.9 ± 3.3 mV) compared to Nx NPs only. The DLS results indicated a narrow size distribution and a good colloidal stability of the NPs (Online Supplementary Figure S1). The results were reproducible in multiple batches.

Cytotoxicity assessment

First we evaluated the effects of the Nx NPs on THP-1 macrophages cell viability using the CTB assay and on cell integrity by measuring the amount of LDH released from the cells after 48 h of exposure to increased concentration of NPs from μg/mL 2.19 to 35 μg/mL. As illustrated in Figure 1(a), Nx NPs did not affect macrophages cell viability or cytoplasmic membrane integrity up to 17.5 μg/mL. When the cells were treated with 35 μg/mL, cell viability decreased to 84.5% and release of LDH increased by 18% when compared to untreated control cells (p < 0.001). Next, we wanted to investigate the effects of NPs on THP-1 macrophages cell viability and integrity over time (Figure 1(b)) looking at times earlier than 48 h to confirm we have not missed any possible transient cytotoxic effect that might have been observed within 24 h of exposure. Since we found that exposure to 35 μg/mL of NPs for 48 h did not potently affect the cells (Figure 1(a)), for this next series of experiments, we only used the two highest concentrations tested in the previous assay (17.5 and 35 μg/mL) and added a higher concentration (70 μg/mL) to further evaluate the cytotoxicity of the NPs (Figure 1(b)). We observed that the maximal effect on macrophages cell viability was already obtained after 4 h of incubation, 73 and 79% for 70 and 35μg/mL, respectively, but was not found statistically different from 48 h (for the 35 μg/mL concentration; 84.5%, Figure 1(a)). The difference in cell viability over time was only found to be statistically significant at 35 μg/mL between 4-h and 24-h post-incubation (p < 0.05). The rate of CTB reduction within each time of incubation was not found to be concentration-dependent between 2 h and 24 h, although it was previously found statistically significant at 48 h (17.5 vs. 35 μg/mL, p < 0.05; Figure 1(a)). To note, NPs did not interfere with the CTB reduction assay as shown in Online Supplementary Figure S2. Maximum release of LDH (122%) was observed at 16 h in cells treated with 70 μg/mL (Figure 1(c)) but was only found significantly higher when compared with cells incubated for 2 h (2 vs. 16h, p < 0.01) and was not found to be statistically different from LDH release at 48 h (119%; data not shown). At 35 μg/mL, LDH release was not found significantly higher at any time point between 2 h and 24 h than at

Table 1. Physicochemical characterization of the lipoplexes used in this study.a

<table>
<thead>
<tr>
<th>Lipoplexe</th>
<th>Ratio lipid–NA</th>
<th>Size (nm)</th>
<th>Polydispersity</th>
<th>Zeta potential (mV)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Nx only</td>
<td></td>
<td>83.4 ± 12.5</td>
<td>0.327 ± 0.035</td>
<td>+50.7 ± 2.0</td>
</tr>
<tr>
<td>Nx+12</td>
<td>13</td>
<td>92.2 ± 5.6</td>
<td>0.192 ± 0.016</td>
<td>+37.0 ± 5.5</td>
</tr>
<tr>
<td>Nx-40</td>
<td>3.9</td>
<td>93.5 ± 17.0</td>
<td>0.229 ± 0.043</td>
<td>−55.9 ± 3.3</td>
</tr>
<tr>
<td>Nx-40 3.9</td>
<td></td>
<td>12.5</td>
<td>0.043</td>
<td>−55.9 ± 3.3</td>
</tr>
</tbody>
</table>

Nl: Neutraplex; Nl+: cationic Nl lipoplexes; Nl−: anionic Nl lipoplexes; NA: nucleic acid.

*aParticles mean diameter was measured by dynamic light scattering and surface charge by zeta potential. Nx lipoplexes were prepared as described in the Materials and methods using the control scrambled siRNA sequence. Values represent mean ± standard deviation of four different experiments (n = 4).
48 h (118%; Figure 1(a)). A concentration-dependent effect on LDH release was only observed at 16-h post-incubation (Figure 1(c)) between 35 μg/mL and 70 μg/mL (p < 0.05).

At 48 h, we have also observed a significant difference between 17.5 μg/mL and 35 μg/mL (p < 0.05; Figure 1(a)). THP-1 macrophages treated with 10% DMSO served as positive controls in these experiments and the results are shown in Online Supplementary Figure S3. At this concentration of DMSO, macrophages cell viability was already dramatically reduced at 2 h, while LDH release increased over time to be maximal at 24 h. These results validate the CTB and LDH assays performed.

**Cellular uptake and intracellular localization of NPs**

Next, we wanted to evaluate the ability of human macrophages to internalize the Nx NPs. First, we performed a time course study by exposing THP-1 macrophage cells to 4.4 μg/mL of FITC-labeled Nx NPs for 30 min to 24 h. Uptake and localization of the FITC-Nx NPs were visualized using confocal microscopy and by staining the cells for late endosomes/lysosomes (LysoTracker Blue) and cellular membrane (WGA; Red) markers. Results of the time course study are shown in Figure 3. After 30 min exposure, NPs were already found inside the cells (Figure 2(b)) and cell uptake increased over time significantly up to 24 h (Figure 2(b) and (d) to (f)). NPs seemed to be endocytosed by the macrophages since they were found to co-localize with the late endosome/lysosome marker at 30 min and 1.5 h (Figure 2(b) and (d), respectively). As early as 1.5-h post-exposure, fluorescent NPs showed a diffuse pattern in the cytoplasm of some cells indicating that they were able to escape the late endosomes/lysosomes compartment. After longer time of exposure (3 and 24 h), NPs were found mainly free in the cytoplasm of live cells (Figure 2(e) and (f), respectively). At the low concentration tested (4.4 μg/mL of NPs), after 24-h post-exposure, fluorescence in live cells was very bright and diffuse throughout the cytoplasm (Figure 2(f)). Next, we wanted to investigate whether the NPs could deliver siRNAs into macrophages. For these experiments, 50 nM Cy3-labeled siCCR5 (red fluorescence) was complexed with Nx NPs at two different lipids: NA ratios as described earlier and cellular uptake of nanolipoplexes was visualized 24 h after exposure in THP-1 macrophages stained for late endosomes/lysosomes (blue fluorescence). Results are shown in Figure 3. In

**Figure 1.** Cytotoxicity assays: Concentration-dependent effects and time-course of Nx nanoparticles. Macrophages were exposed to nanoparticles for A) 48h or B-C) the indicated time, then reduction of CellTiter-Blue and LDH release in culture supernatant were measured using the respective cytotoxicity assay as described in Materials and Methods. Data represent the means ± SEM of A) three separate experiments done in triplicates or B-C) at least two independent experiments done in triplicates. B-C) Dark grey column: 17.5μg/mL; Light grey column: 35μg/mL; Black column: 70μg/mL. Statistically significant based on a two-way ANOVA analysis A) Concentration, Assays or B-C) Concentration, Time. *p ≤ 0.05; **p ≤ 0.01; ***p ≤ 0.001.
control cells, no red fluorescence was observed (Figure 3(a) to (d)). In macrophages exposed to cationic nanolipoplexes \( \text{Nx}^{+12}/\text{Cy3-siCCR5} \) (Figure 3(e) to (h)), red fluorescence showed a punctate pattern and was found mainly inside the cells and not co-localized with late endosomes/lysosomes marker. Anionic nanolipoplexes \( \text{Nx-40/Cy3-siCCR5} \) were also efficiently taken up by macrophages (Figure 3(i) to (l)). At 24-h post-exposure, the red fluorescence showed also a punctate pattern but was not as bright as in the cells exposed to the cationic \( \text{Nx}^{+12}/\text{Cy3-siCCR5} \) nanolipoplexes (Figure 3(k) and (l)) and was found mainly co-localized with the late endosomes/lysosomes marker (Figure 3(l)). However, when incubation was prolonged to 48 h, \( \text{Cy3-siCCR5} \) uptake and delivery inside the cells was similar for both nanolipoplex formulations with very bright fluorescence mainly distributed throughout the cytoplasm and still showing a punctate pattern (Online Supplementary Figure S4). To note, cell morphology was not found different in control cultures than in cells exposed to either NPs alone or nanolipoplexes. Finally, Z-stacks analysis confirmed that the NPs were indeed internalized and not just adhering to the surface of the cells (data not shown).

**Pro-inflammatory effects of NPs**

The pro-inflammatory effects of \( \text{Nx} \) NPs were investigated by first evaluating whether the NPs could alter neutrophil apoptosis rates measured by cytology and their ability to spread onto glass. As shown in Figure 4(a), after 24 h of culture, control neutrophils were approximately 40% apoptotic as measured by cytology. As expected, GM-CSF delayed apoptosis (22.5%), whereas ATO induced neutrophil apoptosis (70%). In this assay, \( \text{Nx} \) NPs were not found to modulate neutrophil apoptosis (46%) at the concentration tested (100 \( \mu \)g/mL). To note, due to the limited number of donors (2–3 donors), these responses were not found to be statistically significant. In addition, because it was previously found that apoptotic neutrophils lose their ability to spread onto glass,42,43 we investigated whether \( \text{Nx} \) NPs could regulate this response. The results shown in Figure 4(b) are in agreement with the cytology data showing that \( \text{Nx} \) NPs did not alter the neutrophil apoptotic rate. After 12 h of culture, the number of spread cells remained stable after exposure to 100- \( \mu \)g/mL \( \text{Nx} \) NPs (25%) and was similar to the number of spread cells found in cells treated with water (17%). In contrast, cultures treated with GM-CSF, fMLP, or \( \text{TiO}_2 \) significantly increased the number of spread cells showing their ability to modify apoptotic rate in human neutrophils as we have previously reported.44,45 Furthermore, \( \text{Nx} \) NPs were tested for their capacity to alter the production of pro-inflammatory cytokines by human macrophages using the THP-1 cell model. Since in the confocal studies we found that with only 4.4 \( \mu \)g/mL, THP-1 cells were fully exposed to the \( \text{Nx} \) NPs, for this series of experiments we reasoned that exposure to the highest concentrations (35 and 70 \( \mu \)g/mL) will not be necessary. We evaluated three concentrations (2.2, 8.8, and 17.5 \( \mu \)g/mL). As illustrated in Figure 4(c), \( \text{Nx} \) NPs did not significantly induce pro-inflammatory cytokine secretion when compared to control cultures up to the highest concentration tested (17.5 \( \mu \)g/mL).

**SiRNA activity of nanolipoplexes**

Next, we wanted to evaluate whether siRNAs complexed to \( \text{Nx} \) NPs could conserve their activity and be developed as anti-HIV therapeutics. To assess this, we used the TZM-bl
reporter cell line which expresses CCR5, CXCR4, and CD4 molecules allowing the cells to be infected by HIV and previously validated anti-CXCR4 siRNAs for their RNAi activity.\(^2^8\) First, we verified by confocal microscopy, the ability of the nanolipoplexes to be internalized by TZM-bl cells. We found that both nanolipoplexes were efficiently taken up by the cells and were able to escape the endosomes after 3 h of exposure (data not shown). Then, we performed the HIV inhibition assays. Data are shown in Figure 5 in terms of percent reduction of β-galactosidase activity compared to the non targeting control siRNA sequence. In this assay, the highest level of inhibition observed was around 30% when using Nx-40/siCXCR4 nanolipoplexes at 12.5 and 25 nM when compared to Nx-40/random siRNA controls. However, although the results showed a concentration-dependent effect, it was not found statistically significant for both nanolipoplex formulations at concentrations higher than 12.5 nM (\(p \leq 0.05\)). Anionic Nx-40 nanolipoplexes showed higher inhibitory effect than cationic Nx+12 nanolipoplexes at all concentrations tested except at 100 nM, but again the differences were not found statistically significant. We also evaluated the effects of the Nx NPs alone (no siRNAs) on the replication of HIV-1 (Online Supplementary Figure S5). We found that at concentrations of NPs of 8.8 \(\mu\)g/mL and higher, β-galactosidase activity was reduced when compared to untreated control cultures (\(p < 0.001\)) indicating that the Nx NPs themselves can interfere with HIV-1 replication in this infectivity assay. In addition, the effect of Nx NPs alone on the viability of TZM-bl cells was evaluated using the CTB assay and no cytotoxicity was observed up to 35 \(\mu\)g/mL, the highest concentration tested (Online Supplementary Figure S6). Finally, Nx/siRNA nanolipoplexes did not cause cytotoxicity in TZM-bl cells.

Figure 3. Uptake of nanolipoplexes in live macrophages. A-D) Untreated control cells incubated for 24h without nanolipoplexes, E-L) Cells exposed to nanolipoplexes (red fluorescence) at a final concentration of 50 nM for 24h: E-H) Nx+12/Cy3-siCCR5 (corresponding to 8.8 \(\mu\)g/mL of NPs) or I-L) Nx-40/Cy3-siCCR5 (corresponding to 2.6 \(\mu\)g/mL of NPs). After incubation cells were stained for late endosomes/lysosomes using Lysotracker (blue fluorescence), washed, and visualized by confocal microscopy. Multiple fields were analyzed and images are representative of two experiments.
up to the highest concentration tested: 111.05% ± 3.82% and 94.98% ± 5.01% of untreated control cells with 200 nM of \( \text{Nx}^{+12}/\text{siRNA} \) and 400 nM of \( \text{Nx}-40/\text{siRNA} \), respectively (data not shown).

**Discussion**

Macrophages are key players in HIV infection and contribute significantly to the cellular reservoirs of HIV. Hence, the virus can survive for prolonged periods in these cells despite ART (ART has been removed). Nanotechnology provides a means to overcome cellular and anatomical barriers to drug delivery and shows promises for its application in the area of HIV eradication and viral reservoir targeting.\textsuperscript{24,31} In a previous study, Nx nanolipoplex formulations were compared to the commercially available cationic Lipofectamine\textsuperscript{®} RNAiMAX reagent (Thermo Fisher Scientific), for their cytotoxicity and capacity to deliver active siRNAs in HeLa-derived cells.\textsuperscript{28} We have found that Nx NPs showed advantages over the commercial cationic RNAiMAX reagent in terms of stability, cytotoxicity, and cellular delivery. In addition, we reported for the first time the influence of surface charge on cytotoxicity and delivery properties of nanolipoplexes using the Nx nanosystem.\textsuperscript{28,46} Here, we wanted to investigate the potential of the lipidic Nx nanosystem as a delivery strategy to target HIV reservoirs. Nx NPs and two nanolipoplex formulations (\( \text{Nx}^{+12}/\text{siRNA} \) and \( \text{Nx}-40/\text{siRNA} \)) were prepared and characterized in terms of their size and surface charge. Physicochemical results obtained were similar to the findings of the previous
study,28 with stable and homogenous nanoformulations with sizes smaller than 100 nm and surface charges corresponding to the lipid: NA ratio. In addition, the Nx nanoformulations and NPs were stable at 4°C up to at least 1 and 9 months, respectively, in solution which represents a great advantage for future use in clinical application. Importantly, the nanoformulations did not aggregate in the presence of serum as per our microscopic observations (data not shown) allowing to perform experiments in biological conditions.

Macrophages obtained by differentiation of THP-1 monocytes were chosen as the in vitro cellular model to assess cytotoxicity and pro-inflammatory effects of the Nx NPs, because it has been successfully used to investigate the toxicity of NPs on macrophages and, in addition, they secrete macrophage-specific cytokines. First, cytotoxicity was assessed by two different bioassays (CTB reduction and LDH release), and time point experiments were performed to investigate time-dependent influence on the cytotoxic results. Here, we found that both cytotoxicity assays provided similar results in terms of the magnitude of the effect on the cells. Hence, Nx NPs did not seriously affect cell viability or membrane integrity of human macrophages since maximal effects observed were 27% decrease in cell viability and 22% increase in LDH release with the highest concentration evaluated (70 μg/mL). Considering that it has been suggested that a decrease in cell viability by more than 30% is to be considered as a cytotoxic effect and that high level of cellular uptake was observed with only 4.4 μg/mL of Nx NPs, we can conclude that Nx NPs have a low cytotoxic profile for human macrophages. Furthermore, our time point experiments revealed that the effect on enzymatic activity (CTB reduction) was observed more rapidly than the impact on cell integrity (LDH release) on cells exposed to the highest concentration (70 μg/mL) of NPs, at 4 h compared to 16 h, respectively. This suggests that cellular enzymatic activity is more susceptible to the effect of Nx NPs, and impact on intracellular pathways might consequently lead to the cell membrane integrity damage observed later. We and others have previously reported a difference in sensitivity between cell viability and LDH release assays with nanomaterials. For instance, we found that the LDH release assay was more sensitive than the CTB reduction assay when evaluating carbon nanotubes, standard reference materials: silica and TiO₂ microparticles, and with different size-fractionated particles collected in a small urban area in Canada. In contrast, others have reported less sensitivity with the LDH release assay compared to the cell viability MTT assay when assessing cytotoxicity of solid lipid NPs. Comparison between different nanotoxicity studies is difficult because they vary in terms of the choice of concentrations of nanomaterials, time of exposure, bioassays, and cell models. In addition to the bioassay, the nature and physicochemical characteristics of the nanomaterial itself influence the results of cytotoxicity assessment.

Furthermore, it is well known that the cytotoxicity potential of a given compound varies with the cell type. In the present study, THP-1 macrophages were found to be more sensitive to Nx NPs than HeLa-derived epithelial cells as no CTB reduction was observed in TZM-bl cells treated with 35 μg/mL of Nx NPs for 48 h. These findings are in line with previous studies which have also reported that macrophages were more responsive to nanomaterials than epithelial cells. Furthermore, Nx NPs did not affect cell viability of neuronal cells in CTB assays up to the highest concentration evaluated, 70 μg/mL (unpublished results). These findings indicate the potential of the Nx nanosystem for drug delivery in the central nervous system either to directly target HIV-1 in this sanctuary site including infected macrophages in the brain or via Nx-loaded monocytes that would migrate into the brain. Hence, we have previously shown the efficiency of the Nx nanosystem to deliver siRNAs in the brain of baboons showing their potential for in vivo application for neurodegenerative diseases and HIV-1 targeting in the central nervous system. Others have reported using a murine model of neuroAIDS that ex vivo loaded MDMs with indinavir-encapsulated lipid NPs injected intravenously were able to target the brain and reduce HIV-1 replication in HIV-1 encephalitis brain regions.

Effects of nanomaterials on cells not only depend on the initial concentration at the time of exposure but also to the actual amount of NPs taken up by a single cell. Cellular uptake of a nanomaterial is mainly determined by its nanoscale size but also by its surface charge, shape, and its...
In the current study, using live-cell confocal microscopy, we have investigated cellular uptake of Nx NPs by THP-1 macrophages, and we found that Nx NPs are rapidly taken up by the cells. Even though the exact mechanism of entry was not part of this study, we can speculate that they entered through an endocytic process since once internalized the Nx NPs were found in endosomal and lysosomal compartments. A transition into the endosomal pathway might represent an advantage for a delivery system aiming to target HIV in macrophages since it has been shown that the virus assembles in the endosomal compartments of macrophages. Other entry mechanisms might also be involved such as temperature-independent uptake as we have previously reported in epithelial HeLa cells or phagocytosis. Further studies are needed to confirm the exact uptake process of Nx NPs in THP-1 macrophages. Here, we observed that Nx NPs were able to rapidly escape the endosomal pathway as the NPs were found into the cytoplasm as soon as 1.5 h after exposure. Although the exact mechanism and kinetics or release of siRNAs has not yet been investigated, we can speculate based on previous studies that the presence of cardiolipin in Nx NPs might help to transition from lamellar to hexagonal ultrastructure and thus enhance lipid membrane fusion with endosomes allowing their content to leak into the cytosol (unpublished results). Furthermore, siRNAs complexed to the NPs were found in the cytoplasm of macrophages in a nonhomogenous pattern outside the endosomal/lysosomal vesicles but never inside the nucleus demonstrating the efficiency of the Nx nanosystem to deliver siRNAs to their site of action. Indeed, we and others have previously reported and suggested that siRNAs localize in cytosol and to regions in close proximity to the nuclear membrane in a nonhomogenous pattern near RISC components creating a focal point for RNAi factories. Our data indicating that siRNAs delivered by Nx NPs can localize in cytosol are in consonance with the previous studies and demonstrate the protective effect of Nx. Furthermore, the fact that Nx NPs localized in cytosol in a homogenous diffuse pattern while siRNAs harbored a more punctate distribution supports the breakdown of the nanolipoplex and delivery of siRNA for RNAi. However, although silencing activity of the siRNAs delivered by Nx NPs has been confirmed in the TZM-bl reporter cell model, silencing activity still need to be confirmed in macrophages and further investigation is ongoing. Interestingly, anionic nanolipoplexes were also efficiently taken up by the cells and able to escape the endosomes/lysosomes compartment but they stayed longer in this compartment compared to the cationic Nx nanolipoplexes suggesting that anionic Nx nanofomulations might be favorable for endosomal targeting. In addition, delivery of siRNAs could also be achieved with high level of efficiency using anionic Nx nanolipoplexes and siRNAs were found in the cells for up to 96 h, the last time point examined (data not shown). Considering that anionic nanolipoplexes contain less lipids for the same amount of NAs than their cationic counterparts thus suggesting potentially less cytotoxicity, therefore, anionic nanoformulations represent an advantage for in vivo application. Furthermore, we and others have previously shown that anionic NPs achieved more widespread dispersal in the brains of rats and baboons than cationic NPs showing their potential for brain delivery. Also, the fact that the intracellular uptake of the Nx nanoformulations was not affected by serum make them suitable for therapeutic application. Experiments are ongoing to further examine the interaction of the anionic and cationic nanolipoplexes with biological fluids such as blood and serum and to better investigate the possibility of their distinct mechanisms of uptake. Finally, in contrast to several NPs which have been found to exert pro-inflammatory properties such as zinc oxide (ZnO) or TiO2, Nx NPs were not found to modulate neutrophil apoptosis or enhance the production of pro-inflammatory cytokines by THP-1 macrophage cells and were comparable to polylactide-co-glycolide (PLGA) NPs known for their low pro-inflammatory profile. These findings are promising but warrant further in vivo immunotoxicological studies to clearly establish the immunomodulatory effect of Nx NPs and confirm their low inflammatory profile. Clearly, THP-1 is a valid and useful model for peripheral blood mononuclear cells (PBMCs). However, it has also been shown that THP-1 cells produce less IL-6, IL-8, and IL-10 than PBMCs while showing a robust TNF-α production upon stimulation with lipopolysaccharides (LPS). Here, we report that THP-1 cells produced no detectable amount of TNF-α and IL-6 and did not increase IL-8 and IL-1β level in response to Nx NPs. It is interesting to note that Nx NPs do not induce IL-8 production in THP-1 cells, indicating the absence of endotoxins in the NP preparations as endotoxins are known to be potent inducers of IL-8. Furthermore, although not found to be statistically significant, Nx NPs slightly induced IL-10 production at the highest dose tested, which is anti-inflammatory. This contrasts with our previous results showing the pro-inflammatory profile of secreted cytokines in THP-1 cells in response to silica particles (personal communication) or silver NPs exposure. Altogether, these observations suggest that Nx NPs are not potent inducers of cytokine production in THP-1 cells. It should be interesting to verify the low inflammatory potential of Nx NPs using PBMCs from a panel of subjects, which should give insight into interindividual variability in NPs reactivity. Further studies are needed to further characterize the inflammatory profile of Nx NPs.

Our results demonstrate the potential of Nx nanosystem to deliver active silencing siRNA targeting CXCR4 co-receptor. In a previous study, we have shown that siCXCR4 when delivered with cationic or anionic Nx nanolipoplexes downregulated CXCR4 in HeLa-derived cells; however, anionic nanoformulations showed less potency despite their high cellular uptake. However, the impact of the down-regulation of CXCR4 on the ability of HIV to replicate was
allowing a multi-drug approach. Finally, further HIV latency. Furthermore, Nx nanosystem can be designed to varying the siRNA sequence used in the nanoformula-
tions equivalent and above 8.8 μg/mL. This anti-HIV activity might potentially be due to an interaction with Nx NPs with the virus particles inside the cells preventing them to complete their replication cycle. This anti-HIV property might be exploited as well in vivo to prevent HIV transmission and deserve further investigation. Also, by varying the siRNA sequence used in the nanoformulations, Nx nanolipoplexes could be designed to target any genes and diseases including those involved in viral latency. Furthermore, Nx nanosystem can be designed to deliver a combination of different siRNA sequences, allowing a multi-drug approach. Finally, further HIV inhibitory experiments using HIV-infected MDMs are ongoing to confirm the potential of the Nx nanosystem to target HIV in this cellular reservoir.

Conclusion

Nx NPs were prepared and evaluated for their potential to efficiently and safely deliver siRNA in cellular models. Cationic and anionic siRNA nanolipoplexes were proved to be efficient to protect and deliver NAs into the cytoplasm of macrophages with negligible toxicity and immunogeni-
city for the cells. Furthermore, both nanolipoplexes showed anti-HIV activity in a cell indicator assay. Although further investigation in vivo is required, altogether, these results demonstrate that the Nx nanosystem is a promising strategy to deliver therapeutic agents in human MDMs and could be used to target pathogens that proliferate inside macrophages including HIV-1, visceral leishmaniasis, malaria, and tuberculosis.

Author contributions

Carole Lavigne is the principal investigator of this study. She designed the study, analyzed and interpreted the data, and wrote the manuscript. Dalibor Breznan and Renaud Vincent contributed to the design of the cytotoxicity assessment and the analysis and interpretation of the data and critically revised the manuscript. Denis Girard and Sylvie Faucher contributed to the design, analysis, and interpretation of the data for the inflammatory experiments and critically reviewed the manuscript. Eric Berger, Sandra Stals, Viraj J Jasinghe, and David Gonçalves contributed to the acquisition and analysis of data and reviewed the manu-
script for its final approval. Alain R Thierry contributed to the conception and design of the study and reviewed the manuscript.

Declaration of Conflicting Interests

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Supplementary material

Supplementary material for this article is available online.

References


