Electrostatic complex of neurotrophin 4 with dendrimer nanoparticles: controlled release of protein in vitro and in vivo

Maria Dąbkowska1
Dorota Rogińska2
Patrycja Kłos1
Anna Sobus2
Małgorzata Adamczak3
Zofia Litwińska2
Anna Machalińska4
Bogusław Machaliński2

1Department of Medical Chemistry, Pomeranian Medical University, Szczecin 70-204, Poland; 2Department of General Pathology, Pomeranian Medical University, Szczecin 70-204, Poland; 3Department of Pharmacy, School of Pharmacy, University of Oslo, Blindern, Oslo 0316, Norway; 4First Department of Ophthalmology, Pomeranian Medical University, Szczecin 70-204, Poland

Background: NT4 has been regarded as a promising therapeutic protein for treatment of damaged retinal pigment epithelium cells.

Purpose: Here, we studied physicochemical parameters of an NT4–polyamidoamine (PAMAM) electrostatic complex, which can provide a sustained concentration of protein in intraocular space over an extended period after delivery. Adsorption/desorption of NT4 molecules to/from positively charged PAMAM dendrimers were precisely determined to control the concentration of bounded/unbounded protein molecules, diffusion coefficient, and size of a protein-laden dendrimer structure. We determined kinetics of NT4 desorption in PBS, vitreous, and damaged retina.

Methods: Initially, adsorption of NT4 molecules on PAMAM dendrimers was studied in PBS using dynamic light scattering, electrophoresis, solution depletion, ELISA, and atomic force microscopy. This allowed us precisely to determine desorption of NT4 from nanoparticles under in situ conditions. The maximum coverage of irreversibly adsorbed NT4 determined by ELISA allowed us to devise a robust procedure for preparing stable and well-controlled coverage of NT4 on PAMAM nanoparticles. Thereafter, we studied diffusion of nanospheres containing NT4 molecules by injecting them into vitreous cavities of mice exposed to intravenous injections of sodium iodate and evaluated their intraocular desorption kinetics from drug carriers in vivo.

Results: Our measurements revealed NT4–dendrimer nanoparticles can be used for continuous neurotrophic factor delivery, enhancing its distribution into mouse vitreous, as well as damaged retina over 28 days of postinjury observation.

Conclusion: Understanding of polyvalent neurotrophin interactions with dendrimer nanoparticles might be useful to obtain well-ordered protein layers, targeting future development of drug-delivery systems, especially for neuroprotection of damaged retinal neurons.

Keywords: neurotrophin 4, NT4, retinal pigment epithelium, therapeutic protein, dendrimer functionalization, controlled release of protein, NT4–PAMAM electrostatic complex

Introduction

NT4 is a growth factor that belongs to a family of neurotrophic factors/neurotrophins that act via different classes of receptors,1,2 leading to the subsequent activation of various signaling pathways in target cells. Two classes of receptor have been identified: p75, which is a member of the TNF-receptor family, and Trk, belonging to the integral tyrosine-kinase family. In our previous work3 we studied NT4–TrkB signaling in mesenchymal stem cells in retina and subsequent Akt- and MAPK-related pathway activation. These pathways may modulate the development and...
maintenance of the vertebrate nervous system via promoting survival, migration, proliferation, differentiation, and death of neurons. Synthesis, secreted concentration, and half-life of NT4 in the human body are limited. NT4 is not available for all neurons; therefore, its delivery from cells to tissue results in concentration gradients. Decreased neurotrophin concentration has been associated with several neurodegenerative diseases and their symptoms. Accordingly, improved administration of exogenous NT4 and consequent neuroprotection has been considered a potentially novel treatment for neurodegenerative diseases, including retinal degeneration. The survival-promoting activity of neuroprotective agents in diverse neuronal systems has been studied recently to determine their use in retinal disorders. Neurotrophins have been used for the rescue of photoreceptors from retinal damage, as described by La Vail et al., Paskowitz et al, and Birch et al.

NT4 delivery can be achieved more easily in the eye than other organs, because the treated tissue surface is small inside the ocular globe, very little diffusion of proteinaceous compound is expected, due to ocular barriers, and thus proteins with molecular weight of 55 kDa can easily reach the retina through diffusion from the vitreous. However, carrier-free administration of NT4 is relatively unstable, because of rapid degradation in biological media as a result of very short in vivo half-life and low bioavailability. On the other hand, biodistribution of nanosize carrier systems after intravitreous injection is highly influenced by the surface of nanoparticles.

Dendrimer is a monodisperse therapeutic scaffold that allows NT4 delivery to damaged retinal pigment epithelium (RPE) cells, enhancing its local concentration and protein stability against enzymatic degradation. Moreover, distribution of well-controlled, multiple reactive surface groups that have great structural adaptability qualifies dendrimers to improve cargo of NT4 as a functional molecule. Physicochemical properties of this densely branched polyelectrolyte with well-defined spherical geometry enhances stability and surface functionality of the NT4-delivery system. With the use of dendrimer, NT4 can be effective only where it is needed. The size of polyamidoamine (PAMAM) dendrimer 6 generation enables its efficient diffusion efficiently across the vitreous body. It can be internalized into cells and removed rapidly by the kidneys. Dendrimers present strong ability to escape uptake by the specific reticuloendothelial system and consequently remain for plenty of time in the blood circulation, increasing their biological potency in specific tissue sites in therapeutic approaches. Using dendrimers as a rapidly removable platform for delivery of protein helps to avoid long-term toxicity. In a localized manner, NT4–PAMAM nanoparticles can be administered as an ocular drug to treat ophthalmic disease. Due to their high degree of structural control (monodispersity and tunable chemical structure), dendrimers are widely utilized in vivo as biological functional nanocarriers for drugs, biomacromolecules, gene delivery, imaging agents, and diagnostic products. The polyvalent interaction of dendrimers with protein resembles a common type of interaction between biological entities, such as receptors and ligands or viruses and cell surfaces. Although NT4 can be a promising therapeutic agent, adsorption of this protein molecule at dendrimer-nanoparticle surfaces remains not fully elucidated.

Therefore, the present study was designed to elucidate the mechanism of NT4 adsorption/desorption at well-characterized PAMAM dendrimer nanoparticles and investigate diffusion of NT4–PAMAM after administration into eyes in real time up to 28 days. In this paper, we formulate reliable clues as to the adsorption/desorption mechanism of NT4 to/from PAMAM nanoparticles, which is generally governed by electrostatic interactions between the protein and polymer surface, van der Waals forces, and structural adaptation of the protein. The main goal of our study was to determine the physicochemical parameters of PAMAM, NT4, and NT4–PAMAM drug-delivery system to assess the usefulness of NT4–PAMAM in an in vitro model. This NT4–PAMAM electrostatic complex allows controlled release of NT4; therefore, different complementary techniques were performed to confirm adsorption/desorption of NT4 protein to/from the PAMAM surface. ELISA measurements were performed to monitor the progress of protein desorption in situ in PBS, as well as in vitro in vitreous humour, retinas, and whole mouse eyes. The reversibility of adsorbed/desorbed protein was determined precisely via laser Doppler velocimetry (LDV) and atomic force microscopy (AFM) measurements and compared with the ELISA results, thus providing reliable data for bulk NT4 concentration as low as 0.1 mg·L\(^{-1}\). In this way, one can unequivocally prove the irreversibility of NT4 adsorption for a low-concentration protein range. It should be mentioned that this method has been efficiently used before to determine adsorption mechanisms of lysozymes, human serum albumin, and fibrinogen on polystyrene microparticles.
As such, here we studied NT4 as a low-molecular-weight therapeutic agent in an ocular drug-delivery system. Sustained controlled release of NT4 for 14 days and up to 28 days from the proposed PAMAM drug-delivery system could promote damaged RPE regeneration and functional recovery. Understanding the adsorption/desorption process of NT4 to/from PAMAM dendrimers could constrain the use of polyelectrolytes or surfactant-mediated systems for eye application.

Methods

NT4, PAMAM, and NT4–PAMAM conjugates

Filtered (Centrifree ultrafiltration device; Merck, Darmstadt, Germany) stock solutions of carrier-free recombinant human NT4 (268-N4-025/CF; R&D Systems) of known concentrations (typically 250 mg·L⁻¹) in PBS (pH 7.4 ±0.2; Biomed, Lublin, Poland) were prepared to remove aggregates and provide a constant, free-form protein-molecule concentration in the solvent. To minimize errors in concentration measurements, two complementary spectro-photometric techniques were used: a BCA protein-quantification kit for low concentration (Abcam), and ultraviolet absorbance at 280 nm measured with microplate spectrophotometry (BioTek Epoch). Prior to each measurement, the stock solution was diluted to a desired bulk concentration, typically 10⁻⁵–50 mg·L⁻¹. The exact concentration of these solutions after membrane filtration was determined by commercially available ELISA kits (DY992, DY990, DY994, DY999, DY995, WA126, DY006, and DY268; R&D Systems). The temperature of experiments was kept at a constant 298±0.1 K.

A suspension of PAMAM G6 ethylenediamine-core sixth-generation dendrimers, (536717; Sigma-Aldrich, St Louis, MO, USA) to remove low-molecular-weight impurities. Using ultraviolet-visible spectrophotometry, we determined the exact concentration of G6 PAMAM dendrimers after purification using the ninhydrin assay. This procedure is based on the interaction of the primary amino group of PAMAM dendrimers with ninhydrin (Sigma-Aldrich) to form a blue product. According to this protocol, we were able to determine the concentration of dendrimer molecules in the range of 25–200 mg/ml. Afterward, the stock suspension was diluted prior to each adsorption experiment to a desired mass concentration of 50 mg·L⁻¹ and filtered to remove aggregates with the lowest binding membrane for protein solutions (0.1 μm, VDF Durapore; Merck Millipore).

NT4 adsorption at PAMAM dendrimers was performed: the reference electrophoretic mobility of bare PAMAM nanoparticles was measured, NT4 monolayers were formed by mixing equal volumes of solutions of bulk concentration (0–40 mg·L⁻¹) with nanoparticle suspensions of bulk concentrations of 100 mg·L⁻¹, and the electrophoretic mobility of NT4–PAMAM nanoparticles was measured and the corresponding ζ-potential calculated. Final experiments were performed at pH 7.4, ionic strength 0.15 M, temperature 20°C. This procedure enables direct in situ determination of hydrodynamic size, electrophoretic mobility and ζ-potential variations as a function of bulk concentration of NT4. The adsorption time of NT4 at PAMAM dendrimers was 1,800 seconds in the primary adsorption experiments. The relaxation time of the lysozyme (similar molecular mass to NT4 molecule) monolayer on microparticles was around 2 seconds for full monolayer. The time for formation of additional layers is much longer than monolayer formation. In all cases, NT4 should be irreversibly adsorbed onto the surface as a result of short-range electrostatic interactions between NT4 and PAMAM, since the adsorption process is mediated by electrostatic interactions.

The diffusion coefficient of NT4, PAMAM, and the NT4–PAMAM conjugate was determined by dynamic light scattering (DLS; Malvern Zetasizer Nano ZS). The electrophoretic mobility of NT4 molecules, bare PAMAM, and NT4–PAMAM conjugate was measured with LDV with the aid of the aforementioned Malvern device. Suspensions of nanoparticles and protein diluted in PBS to a suitable concentration were used for these analyses. Data analysis was performed in automatic mode at 25°C. Measured size was presented as the average value of 20 runs, with triplicate measurements within each run.

NT4 surface concentration

Ruby muscovite mica (Continental Trade, Poland) was used as a substrate for NT4, PAMAM, and NT4–PAMAM adsorption measurements. Fresh solid pieces of mica were cleaved into thin sheets prior to every experiment. AFM was used to obtain information about the binding strength.
of NT4 on PAMAM dendrimers. In parallel with ELISA measurements, the unbound (residual) NT4 surface concentration after adsorption was determined directly by AFM imaging. In the former case, the nanoparticle–NT4 mixture and supernatant acquired after the adsorption step were left to deposit on mica sheets placed in the diffusion cell over a controlled time, and then the substrate was removed and rinsed for half an hour in ultrapure water. Samples were left for air-drying until the next day. AFM measurements were carried out under ambient air conditions using a NanoWizard (JPK Instruments). Intermittent contact–mode images were obtained in the air using ultrasharp silicon cantilevers (NSC35/AIBS; MicroMash), and the cone angle of the tip was <20°. Images were recorded at a scan rate of 1 Hz for the six randomly chosen places. Images were flattened using an algorithm provided with the instrument. We captured all images in random areas within a scan size of 0.5×0.5 µm or 1×1 µm. NT4 surface dimension and concentration were determined using ImageJ software by gathering the number and coordinates of single protein molecules. Manual counting of NT4 molecules was based on comparing the original image and the same picture altered by digital image filters by cutting off the picture background. Once the number of protein molecules had been established, the residual bulk concentration of NT4 and consequently its coverage on PAMAM nanoparticles was calculated.

Quantification of NT4
Adsorption/desorption of NT4 molecules at/from nanoparticles was studied with ELISA according to the manufacturer’s protocol. Initially, the residual (unbound) NT4 concentration in the filtrate was determined by applying a sandwich ELISA technique to monitor simultaneously the maximum concentration of NT4 in the supernatant suspensions, acquired after adsorption at PAMAM nanoparticles. Standard and supernatant samples containing NT4 after ultrafiltration of PAMAM suspensions were added to the 96-well microplate precoated. Therefore, it was possible precisely to determine the concentration of unadsorbed NT4 molecules at PAMAM nanoparticle surfaces separated by ultrafiltration.

NT4–PAMAM–AF488 conjugates
For detection of NT4–PAMAM conjugates on mouse-eye slides by confocal microscopy, a detectable fluorescent molecule, eg, PAMAM–AF488 was needed. We used AF488-NHS ester as fluorescent label. PAMAM at 2.5 mg·ml⁻¹ in pH 8 sodium bicarbonate buffer reacted with 1 mg·ml⁻¹ amine-reactive dye dissolved in dimethylformamide as described in the manufacturer’s protocol to obtain fluorescently labeled PAMAM–AF488 conjugates. These conjugates were purified by extensive dialysis against pH 7.4 to remove unreacted labeling. After dialysis, we further purified conjugates through centrifugal filtration until filtrate absorbance at 490 nm had reached background levels. Protein concentrations were determined using a total-protein assay (Micro BCA; Pierce). Afterward, NT4 adsorption at PAMAM–AF488 molecules was performed according to the procedure described in the “NT4, PAMAM, and NT4–PAMAM conjugates” section.

Animals and experimental procedures
Pathogen-free 12-week-old mature female BALB/c mice (Center for Experimental Medicine, Medical University of Bialystok, Poland) weighing 25±2 g were used in the experiment. Animals were maintained under standard conditions with a 12-hour day (160 lux)–night cycle and ad libitum feeding. All animal procedures were performed according to the regulations in the Association for Research in Vision and Ophthalmology Statement for the Use of Animals in Ophthalmic and Vision Research and were approved by the local ethics committee for experiments on animals in Poznań, Poland (approval 27/2017).

To study the penetration of encapsulated NT4 protein in vivo after intravitreal injection of PAMAM–NT4, we monitored vitreal and retinal levels of NT4 over 28 days. Prior to NT4 administration, mice were anesthetized (n=40), and sodium iodate (NaIO₃; Sigma-Aldrich) in PBS at a dose of 30 mg/kg body weight was injected intravenously into the retroorbital sinus. Sodium iodate treatment is a well-characterized chemical model of acute retinal injury to study adjuvant or stem cell–based ocular therapy.¹⁻⁴,⁶ Subsequently, 24 hours after the retinal lesions were induced, 1 µL NT4 (20 µg/mL)–PAMAM (50 µg/mL) or PAMAM (50 µg/mL) alone (both solutions prepared in PBS) were injected into the vitreal cavity using a 32 G needle (Hamilton, Reno, NV, USA) to the left and right eyes, respectively (n=20 mice). Moreover, to track nanoparticles within the eye by confocal microscopy, some animals (n=10) were intravitreally injected with 1 µL NT4–PAMAM conjugated with AF88. Tissue samples were then collected on days 7 and 28 days after NaIO₃ and nanoparticle administration. Animals with no treatment (n=5) served as a control.
Tissue collection
Mice were euthanized by cervical dislocation at selected time points: before and on days 7, 14, and 28 after NaIO₃ alone, NaIO₃-PAMAM, NaIO₃/NT4-PAMAM, or NaIO₃-NT4–PAMAM–AF488 administration (n=10 eyes/group/time point). Eyes were enucleated and subjected into different analyses. To study tissue levels of NT4 protein, cornneas and lenses were removed from eyeballs by cutting along orae serratae with small scissors. Then, the vitreous was dissected and the remaining retina/RPE and vitreous were used separately to extract total protein for subsequent ELISA testing. For fluorescence imaging, enucleated eyes were fixed for 24 hours in Davidson’s fixative, dehydrated by immersion in alcohol solutions of increasing concentrations (50%, 70%, 80%, 96%, and 100%) for 5 minutes each and embedded in paraffin.

Protein extraction
To obtain total naïve protein, tissue samples were homogenized in 350 µL ice-cold cell-disruption buffer (mirVana PARIS kit; Thermo Fisher Scientific), supplemented with a Phosphatase Inhibitor Cocktail Set II (Merck). After centrifugation, the supernatant was collected and stored at −80°C until needed. The calculated concentration of NT4 protein on ELISA was normalized to the total protein concentration of the individual samples measured by Bradford assay (Sigma-Aldrich).

Confocal imaging
Paraffin blocks were cut into 5 µm–thick sections and mounted on microscopic slides for subsequent experiments. After 2-hour incubation in 58°C, paraffin sections were subjected to deparaffinization in xylene, using two changes of 15 minutes each. Sections were then gradually hydrated through graded alcohols — 100%, 96%, 80%, 70%, and 50%, 5 minutes in each — and subsequently washed in deionized H₂O and PBS pH 7.4, for 2 and 10 minutes, respectively. Next, nuclei were stained with DAPI (Thermo Fisher Scientific), for 5 minutes at room temperature in the dark, and specimens were washed three times in PBS. Tissue sections were mounted with Vectashield mounting medium (Vector Laboratories, Burlingame, CA, USA) and covered-slipped. Confocal imaging was performed by means of laser-scanning confocal microscopy (Axiolimger Z2; Carl Zeiss, Oberkochen, Germany) in 63× oil-immersion objective with excitation/emission wavelengths of λex=490 nm/λem=525 nm for AF488 and λex=358 nm/λem=461 nm for DAPI.

Statistical analysis
Statistical analysis within each study group was performed using the Kruskal–Wallis test. Two-way ANOVA was used for analysis between experimental groups. P<0.05 was considered statistically significant. Data are presented as mean ± SD.

Results and discussion
NT4 and PAMAM physicochemical characteristics
It was determined by DLS that for a PBS concentration of 0.15 M at pH 7.4, diffusion coefficients of NT4 and PAMAM molecules were 10.8×10⁻⁷±0.1 cm²·s⁻¹ and 6.3×10⁻⁶±0.1 cm²·s⁻¹ (at 298 K), respectively, which is in agreement with our previous results. Based on the known diffusion coefficient, the hydrodynamic diameters of NT4 and PAMAM were calculated using the Stokes–Einstein relationship:

\[ d_H = \frac{kT}{6\pi\eta D} \]

where \( d_H \) is the hydrodynamic diameter, \( k \) Boltzmann constant, \( T \) absolute temperature, \( \eta \) dynamic viscosity of water, and \( D \) diffusion coefficient of protein or nanoparticles. The hydrodynamic diameter obtained for NT4 was 4.0±0.5 nm and for PAMAM 7.6±0.1 nm. This corresponded to an NT4 molecule cross-section area of 14.2 nm². Therefore, the “hydrodynamic” size of NT4 can be treated as a good estimate of its geometric size, given the structural stability of the protein and its compact shape.

In parallel, AFM was used to determine the geometric size of NT4 and PAMAM adsorbed under diffusion-controlled transport conditions on mica surfaces at pH 7.4 and ionic strength 0.15 M PBS. The average NT4 molecule occupied an equivalent sphere area, with diameter of 4.5±1.5 nm, which was in good agreement with DLS bulk measurements. This is demonstrated in Figure 1A, where the presence of a larger number of NT4 oligomers at the mica surface in pH 7.4 and ionic strength 0.15 M PBS was not observed after ultrafiltration. As can be seen in Figure 1B, the average PAMAM molecule occupied an equivalent sphere area, with diameter of 3.5±1.5 nm. According to these results, the size measured by AFM was in agreement with DLS bulk measurements, which are presented in Figure 1.

Figure 1B depicts monodisperse distribution of dendrimer size, where 85% of adsorbed PAMAM molecules at the mica surface had similar diameter. Figure 1B and DLS
measurements (polydispersity index = 1.03, in good accordance with Mullen et al48) proved that it was more likely for NT4 oligomers to appear at a solid surface than for PAMAM (Figure 1A). We observed some adsorbed NT4 aggregates with diameters 11.4±2 nm and 22.4±4 nm at the mica surface.

To understand the release mechanism from encapsulation-free NT4–PAMAM electrostatic complexes, we determined the distribution of positive and negative charges on nanoparticles and protein surfaces, which is a prerequisite for occurrence of attractive electrostatic interaction between them. The electrophoretic mobility ($\mu_e$) of NT4 and PAMAM molecules was measured for 0.15 M PBS concentrations at pH 7.4 using the LDV method and was 1.16 and $-0.47 \mu\text{m}\cdot\text{cm/(Vs)}^{-1}$, respectively. These values corresponded to $\zeta$-potentials of 22.2 mV and $-8.2$ mV, respectively, calculated using the Henry model. Knowing the electrophoretic
mobility, it was possible to calculate the electrokinetic charge per NT4 molecule, $Q_c$, from the Lorenz–Stokes relationship:\textsuperscript{49–51}

$$Q_c = 3\pi d_H \mu_e$$ (2)

which is expressed in Coulombs, with $\eta$ the dynamic viscosity of the solvent (water), $d_H$ expressed in nm, and $\mu_e$ expressed in $\mu$m·cm/V·s.

Equation 2 is valid for molecules of arbitrary shape and absolute $\zeta$-potential value $<50$ mV, though has some limitations for higher ionic strengths, where double-layer thickness becomes less than the protein dimension. Therefore, while calculating the electrokinetic charge $Q_c$ for ionic strength of 0.15 M, we applied a previously used equation:\textsuperscript{37,52,53}

$$Q_c = 2\pi \eta d_H \frac{1 + k d_H}{f_H(kd_H)} \mu_e = 2\pi \eta d_H \left(1 + k d_H\right)$$ (3)

The effective charge of PAMAM was a useful bulk characteristic for interpretation of electrostatic adsorption and desorption of NT4 molecules from these nanospheres. This approach does not require a chemical reaction, i.e., is used routinely as an electron-dense marker.\textsuperscript{54–56} As discussed in Mullen et al.\textsuperscript{48} dendrimers have a uniform surface-charge density. This can be obtained from electrokinetic mobility and the hydrodynamic diameter of a nanoparticle, and one can calculate the average number of charges per molecule from the Lorentz–Stokes relationship. The results of this calculation are collected in Table 1, according to Equations 1 and 2 at pH 7.4. NT4 molecules acquired a slightly negative net charge, whereas the charge of PAMAM nanoparticles was positive.

**Table 1: Electrophoretic mobility, $\zeta$-potential, and charge density of PAMAM nanoparticles and NT4 molecules**

<table>
<thead>
<tr>
<th>PBS concentration (M)</th>
<th>PAMAM (G6)</th>
<th>NT4</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>$\mu_e$ ($\mu$m·cm·[Vs]$^{-1}$)</td>
<td>$\zeta$ (mV)</td>
</tr>
<tr>
<td>0.15, pH 7.4</td>
<td>1.16</td>
<td>22.2</td>
</tr>
</tbody>
</table>

Notes: *Calculated from Lorenz model; **calculated from $2\pi \eta d_H f_H(kd_H) \mu_e$ (Henry model).

Abbreviation: PAMAM, polyamidoamine.

Reservoir-based NT4-delivery system

Adsorption of NT4 on PAMAM dendrimers in PBS

Electrokinetic studies offer a distinct possibility of direct in situ determination of electric properties of proteins and dendrimers, and can be conveniently interpreted by lateral electrostatic interactions between adsorbed molecules. In accordance with this approach, the bulk transport of proteins is described by the phenomenological continuity equation.\textsuperscript{57} Selecting a system of well-established physicochemical properties allowed us to combine dynamic aspects of interfacial interactions between NT4 and the PAMAM surface. Experimental data for the hydrodynamic diameter of the protein–dendrimer delivery system obtained with DLS after adsorption of NT4 (20 mg·L$^{-1}$) at PAMAM (50 mg·L$^{-1}$) nanoparticles showed a mean diameter of 13.5±5 nm for the electrostatic complex.

Additionally, we determined the geometric size of NT4–PAMAM conjugates adsorbed on mica surfaces at pH 7.4, 0.15 M PBS using AFM (Figure 2). Considering that NT4 has a molecular weight of 14 kDa and is 5 nm in length, the hydrodynamic diameter of a single PAMAM 6.0 molecule of molecular weight 58 kDa was 7.8 nm. As can be seen in Figure 2, the diameter of NT4–PAMAM conjugates was 18±5 nm after ultrafiltration.

The electrophoretic mobility of NT4–PAMAM conjugates was studied by the microelectrophoretic method using the Henry equation to calculate $\zeta$-potential, as described elsewhere.\textsuperscript{40–42} Data obtained in the primary adsorption experiments for 0.15 M NaCl, pH 7.4 PBS are shown in Table 1. The $\zeta$-potential of the NT4 (20 mg·L$^{-1}$)–PAMAM (50 mg·L$^{-1}$) conjugate was 12±8 mV for pH 7.4 and 0.15 M PBS. As expected, adsorption of protein with slightly negative charge led to a reduction in the surface charge of nanoparticles, which gained less positive charge. We also determined the dependence of the $\zeta$-potential of microparticles on the initial concentration of NT4 in the PAMAM suspension (after mixing), denoted by $c_p$ (Figure 3). The $\zeta$-potential abruptly decreased with NT4 concentration and approached $c_p>0.5$ mg·L$^{-1}$ at plateau values of −5 mV, for pH 7.4 0.15 M PBS.

Adsorption of the NT4-molecule monolayer on PAMAM nanoparticles was connected with significant changes in their apparent $\zeta$-potential, which can be efficiently monitored by LDV method, studied multiple times before for fibrinogen, recombinanthuman serum albumin, plasmid protein, and albumin. As depicted in Figure 3, $\zeta$-potential abruptly decreased with increasing NT4 concentration and approached plateau
values of \(-5.8\, \text{mV}\), which was slightly below the \(\zeta\)-potential of NT4 in the bulk \((-8.2\, \text{mV} \text{ at } 0.15\, \text{M ionic strength})\). The electrophoretic mobility of the NT4–PAMAM complex was 0.7 the electrophoretic mobility of bulk protein, which corresponded to the formation of an NT monolayer on PAMAM from 0.5 mg\(\text{L}^{-1}\) NT4 concentration. The use of higher NT4 concentration and prolonged adsorption up to 24 hours possibly resulted in multilayer formation (Figure 4). Moreover, in additional experiments, we confirmed that NT4 adsorption at the first layer/molecule was irreversible. We did not observe changes in electrophoretic mobility (\(\zeta\)-potential) over time up to 360 hours, which suggests that the \(\zeta\)-potential of PAMAM after NT4 adsorption (for NT4 bulk concentration up to 0.6 mg\(\text{L}^{-1}\); Figure 3) was only dependent only on NT coverage. Based on the data shown in Figure 3, we concluded that the densely packed monolayer of NT4 on nanoparticles started from 0.6 mg\(\text{L}^{-1}\). NT4 adsorption onto PAMAM nanoparticles was found to be in agreement with previous results observed by Serra et al\(^{58}\) for polyclonal IgG, Bratek-Skicke et al\(^{42}\) for human serum fibrinogen, and Dąbkowska et al\(^{40}\) for lysozymes, where results were interpreted in terms of the electrokinetic model previously applied in simple and reliable LDV measurements. LDV measurement of the electrophoretic mobility of microparticles enables determination of the bulk concentration of protein, even for a range of 0.1–1 mg\(\text{L}^{-1}\), ie, the nanomolar concentration range,\(^{41}\) thus indicating that electrostatic interactions have a pivotal role in adsorption process, which we have already observed for lysozymes, another protein with low molecular weight, just as with NT4.

Our results are in accordance with maximum coverage of similar molecular weight to NT4 protein, and can be found in our recent work.\(^{40}\) Similar data were also obtained for protein-molecule adsorption at silica particles (20 nm in diameter),\(^{59}\) colloidal alumina particles (180 nm)\(^{60}\) and soda lime glass (74.5 nm).\(^{51}\) However, in other work, much larger values for maximum protein coverage were reported, which can most probably be attributed to adsorption of aggregates.

The small slope seen on the chart in Figure 3 means that estimation of maximum protein coverage directly from electrophoretic mobility measurements was less accurate for concentration >0.8 mg\(\text{L}^{-1}\) at higher ionic strength, eg, 0.15 M PBS. Therefore, we used ELISA to determine residual (free) NT4 concentration after adsorption at PAMAM nanoparticles (Figure 4).

As can be observed, for low initial NT4 concentration (1 \(\text{mg\text{L}^{-1}}\)) unbounded protein molecules in PBS were negligible, indicating that adsorption onto 50 mg\(\text{L}^{-1}\) PAMAM nanoparticles was almost complete. In this way, we conclude that NT4 molecules after adsorption at PAMAM (pH 7.4 in 0.15 M PBS) are likely to form a monolayer of single molecules or a multilayer of aggregated molecules, which correlates with adsorption time and protein concentration. Moreover, it can be noticed that at pH 7.4, close to the isoelectric point of NT4, double-layer forces were weak, due to lowering protein net charge (Table 1). Under these conditions, double-layer forces were remarkably weak, and thus interactions between NT4 molecules increased compared to the same

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**Figure 2** Adsorbed NT4–PAMAM (PBS, 0.15 M, pH 7.4) indicated by direct AFM enumeration obtained for low surface-molecule concentration.

**Notes:** Figure created by taking into account ten randomly chosen areas, where each micrograph of the NT4–PAMAM monolayer at the mica surface has a size of 1\(\times\)1 \(\mu\text{m}\).

**Abbreviations:** PAMAM, polyamidoamine; AFM, atomic force microscopy.

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**Figure 3** Dependence of \(\zeta\)-potential of PAMAM nanoparticles on bulk concentration of NT4 in the suspension after mixing \(c_b\).

**Notes:** Points show experimental results obtained by laser Doppler velocimetry for 0.15 M PBS (pH 7.4) and PAMAM concentration 50 mg\(\text{L}^{-1}\). Solid lines show linear fit of data. Dashed lines show \(\zeta\)-potential of bare microparticles.

**Abbreviation:** PAMAM, polyamidoamine.
electrostatic interaction at lower pH. Because of that, protein aggregation occurred, i.e., aggregates consisting of two or three NT4 molecules assembled on the dendrimer surface. The large number of NT4 aggregates in the bulk formed due to the decrease in the range of lateral electrostatic interactions (double-layer thickness of 0.79 nm) among adsorbed NT4 molecules at 0.15 M ionic strength. The concentration of unbounded protein molecules increased systematically with increasing initial protein concentration from 1 mg L\(^{-1}\) to 20 mg L\(^{-1}\) (Figure 4), which clearly indicated the formation of NT4 aggregates on the dendrimer surface.

More importantly, maximum concentration of adsorbed protein at the PAMAM surface was obtained at 0.6 µg/mL, representing a densely packed protein monolayer. The almost completely irreversible adsorption regime is found only for low protein concentration characterized by formation of an NT4 monolayer at the nanoparticle surface and hence there was no observable desorption within the 4 weeks. Increasing adsorption time and protein bulk concentration caused the creation of additional protein layers at PAMAM nanoparticles (seen as a decline in residual NT4 concentration in PBS). The decline in protein molecules from the bulk solution during adsorption finally led to a steady state, which indicated slow alteration in the adlayer structure.

Desorption of NT4 from PAMAM dendrimers in PBS

In order to determine the possibility of NT4 desorption from dendrimers, once adsorption reached equilibrium, we studied NT4 release in PBS. At this point, we used ELISA to evaluate more accurately concentrations of desorbed NT4 from PAMAM nanoparticles in electrolytes. The release profiles of NT4 molecules from nanoparticles at initial protein concentration of 20 mg L\(^{-1}\) is shown in Figure 4 (pH 7.4 and 0.15 M PBS).

Initial protein concentration decreased with desorption time, reaching final steady-state values after 500 minutes (Figure 5A). Therefore, proteins were released in a biphasic mode. The first phase was characterized by a fast release of NT4 molecules, which probably resulted from the solubilization of protein that usually exists near the surface. Spontaneous electrostatic interaction led to release of 20% of NT4 in the first 2 hours, but the remaining NT4 was delivered as an irreversibly adsorbed protein at dendrimer nanoparticles. The second phase was characterized by a slow/sustained NT4-desorption profile, which could be attributed to reversible adsorption of the protein layer at polymer nanoparticles, leading to diffusion of the entrapped protein. Our results are in agreement with profiles of protein release previously presented in the literature for nanosize systems. Abruptly desorbing fraction of reversibly bound NT4 molecules from dendrimer nanoparticles indicates the existence of adsorbed NT aggregates or...
Figure 5 Desorption characteristics of NT4 from PAMAM G6 dendrimers for PBS electrolytes. Notes: NT4 detection by ELISA over the first (A) 10 hours and (B) 32 days of incubation. Abbreviation: PAMAM, polyamidoamine.

Figure 6 Desorption characteristics of NT4 proteins in vitreous body after injection of control mice (before treatment), NaIO₃ injection, and NaIO₃–PAMAM or NaIO₃–NT4–PAMAM injection. Notes: (A) ELISA detection of NT4; (B) NT4 concentration normalized to total protein concentration of individual tissue samples. Results shown as means ± SD. *P<0.05; **P<0.01; ***P<0.001. Abbreviation: PAMAM, polyamidoamine.
various conformations of single NT4 molecules in the adsorbed state. The binding strength of asymmetric shapes of protein aggregates could be responsible for the reversibility of the adsorption process. A fraction of NT4, which formed the next layers on the dendrimer surface, was characterized by slow desorption from the carrier over 28 days.

Desorption of NT4 from PAMAM dendrimers in vitreous bodies and injured retinas in mouse model

We investigated the effect of NT4 adsorption on the in vivo transportation of PAMAM nanoparticles to the vitreous body as well as into retinal layers of mice exposed to intravenous injection of NaIO$_3$. Systemic administration of sodium iodate is known to selectively impair the RPE, resulting in patchy loss of RPE and subsequent degeneration of photoreceptors.$^{63}$ To assess the distribution of injected dendrimer-NT conjugates, we analyzed vitreous NT4 concentrations after injection of PAMAM and PAMAM–NT4 electrostatic complex with ELISA up to 28 days postinjection. Eyes exposed to retinal injury without subsequent intravitreal injection served as controls. The results are summarized in Figure 6.

Our results (Figure 6) show unequivocally that injection of a negatively charged protein–nanoparticle complex had increased NT4 concentration in vitreous bodies of injured mouse eyes by day 7 postapplication. Moreover, injected PAMAM dendrimers alone did not influence NT4 concentration at all. NT4 vitreous level decreased gradually up to 28 days after administration, being in accordance with NT4-release profiles in PBS (Figure 5). This may indicate that irreversibly adsorbed NT4 on negatively

![Graph A](image1.png)

**Figure 7** Desorption characteristics of NT4 proteins in retina after injection of control mice (before any treatment), NaIO$_3$ injection, and NaIO$_3$–PAMAM or NaIO$_3$–PAMAM–NT4.

**Notes:** (A) ELISA detection of NT4; (B) NT4 concentration normalized to total protein concentration of individual tissue samples. Results shown as means ± SD. $^{*}$P<0.05; $^{**}$P<0.01; $^{***}$P<0.001.

**Abbreviation:** PAMAM, polyamidoamine.
charged dendrimers improved its tissue penetration and enabled its distribution into vitreous tissue. Furthermore, we determined desorption kinetics of NT4 from the protein–dendrimer conjugates within retinas at selected time points (days 7, 14, and 28) after intravitreal delivery (Figure 7A and B). This study was designed to quantitatively identify penetration of NT4 within damaged retinas. The results are summarized in Figure 7.

Administration of positively charged PAMAM dendrimers had increased NT4 concentration in impaired retinas compared to tissue without any carriers by day 7 postapplication. NT4 concentration decreased gradually over time. The initial concentration of NT4 in retinas without any modification on day 0 was maintained on day 28 after delivery (Figure 7). We speculate that the electrostatic or aspecific interaction of dendrimers with different types of proteins (protein corona) resulted in their removal by macrophage cells, hampering their integration with retinal layers. In fact, the role of the protein corona in the retina environment is vital for successful delivery of nanoparticles into retinal tissue.

Analysis of NT4 transport across mouse eyes revealed higher levels of protein concentration in the vitreous body than in damaged retina layers. The effective concentration of protein after administration of NT4–PAMAM in retinas was approximately 32 times lower than NT4 concentrations in vitreous body, which indicates that penetration of NT4 into retinal tissue was hampered. Intravitreally, distribution of NT4 was a transport- rather than elimination-limited process, as opposed to retinal tissue, where it was hampered, as illustrated in Figures 6 and 7. Penetration of the NT–dendrimer complex into retinal tissue was governed by electrostatic and hydrophobic interactions (connected with conformational transition of protein). Positively charged PAMAM nanoparticles and slightly negatively charged NT4–PAMAM complexes adsorbed onto negatively charged cell membranes. In this case, we suspect that the main force governing NT4 and NT4–PAMAM transport across the retinal tissue seen on 7 day after injection could be phagocytosis, causing NT4–PAMAM complex engulfment and sufficient removal from the bloodstream. Our results are in accordance with Ding and Ma.64 Additionally, confocal analysis of mouse-eye sections (Figure 8) revealed the presence of green fluorescent particles, which may correspond to clusters of the NT4–AF488–dendrimer nanoconstructs.

Particles were detected on day 7 after administration of PAMAM–NT4 complexes in vitreous bodies, but not in retinas. On day 28 after nanocomplex injection, the particles were

Figure 8 Confocal images showing NaIO₃-injured mouse eyes (vitreous body and retina) after administration of fluorescently-labeled NT4–PAMAM conjugates.

Notes: Top middle image reveals the presence of green fluorescent particles that may correspond to nanocomplexes in the vitreous body only 7 days after NT4–PAMAM injection. Particles were undetectable in the retina at all study time points (bottom row). Magnification 63×, oil immersion. Blue fluorescence, nuclei (DAPI); green background, tissue autofluorescence.

Abbreviation: PAMAM, polyamidoamine.
already undetectable in eyeballs, vitreous bodies, and retinas. Microscopy was consistent with the data obtained from ELISA, confirming the highest concentration of therapeutic complexes in vitreous bodies within 7 days of administration.

**Conclusion**

Our goal in this study was to quantitatively identify a transition from reversible to irreversible adsorption of NT4 to design a protein–dendrimer electrostatic complex for NT delivery to impaired retinal tissue. From our in situ measurements, we conclude that adsorption/desorption of NT4 from PAMAM nanoparticles in the PBS environment for the range of pH close to and below the isoelectric point is dependent on electrostatic interactions. We have pointed out that increasing adsorption time and protein bulk concentration caused the creation of additional protein layers at PAMAM nanoparticles (seen as a decline in residual NT4 concentration in PBS).

Our results obtained from desorption study in PBS by concentration-depletion methods are in agreement with data collected from the NT4-release profile in vitreous bodies. On the other hand, when the NT4–PAMAM conjugate interacted with cells in retinal tissue, we observed lower cellular uptake, indicating that tissue penetration of NT4 was an elimination–rather than transport–limited process. We established that efficacy of NT4–PAMAM electrostatic conjugate delivery across retinal tissue is likely dependent on electrostatic or aspecific interaction with the negatively charged cell membrane. Therefore, it may be concluded that understanding the adsorption/desorption transition of NTs to/from nanoparticles is essential for rational design of an NT-delivery system, which could overcome restricted retinal tissue penetration.

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**Disclosure**

The authors report no conflicts of interest in this work.

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