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Dendrimers reduce toxicity of Aß 1-28 peptide during aggregation and accelerate fibril

formation

Barbara Klajnert, Prof.*1, Tomasz Wasiak, M.Sc.1, Maksim Ionov, Ph.D.1, Marcos Fernandez-

Villamarin, M.Sc.², Ana Sousa, M.Sc.², Juan Correa, Ph.D.², Ricardo Riguera, Prof.², and

Eduardo Fernandez-Megia, Prof.*2

¹Department of General Biophysics, University of Lodz, 141/143 Pomorska St., Lodz, Poland,

e-mail: aklajn@biol.uni.lodz.pl, phone: +48 42 635 44 29, fax: +48 42 635 44 74;

²Department of Organic Chemistry and Center for Research in Biological Chemistry and

Molecular Materials (CIQUS), University of Santiago de Compostela, Jenaro de la Fuente

s/n, 15782 Santiago de Compostela, Spain, e-mail: ef.megia@usc.es, phone: +34 881 815

727, fax: +34 881 815 704

* Corresponding authors

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Abstract

The influence of a GATG (gallic acid-triethylene glycol) dendrimer decorated with 27 terminal morpholine groups ([G3]-Mor) on the aggregation process of Alzheimer's peptide has been investigated. Amyloid fibrils were formed from the A\Beta 1-28 peptide and the process was monitored by a ThT assay, changes in CD spectra, and transmission electron microscopy. In the presence of [G3]-Mor, more fibrils were built and the process significantly accelerated compared to a control. The cytotoxicity of (1) Aβ and (2) the system [G3]-Mor/Aβ was monitored at different stages of the aggregation process. Prefibrillar species were more toxic

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than mature fibrils. [G3]-Mor significantly reduced the toxicity of $A\beta$, probably due to lowering the amount of prefibrillar forms in the system by speeding up the process of fibril formation.

Alzheimer's disease (AD) is the major cause of disability and death in the elderly. Amyloid deposits surrounding dying neurons, neurofibrillar degeneration with tangles, and cerebrovascular angiopathies are three main characteristics of AD [1]. One of the main pathological hallmarks of AD is the formation of amyloid plaques in the affected brain as a result of protein misfolding [2]. The major component of amyloid deposits is a 40-42 residue peptide called β-amyloid peptide (Aβ), produced by endoproteolytic cleavage of an amyloid precursor protein (APP) [3]. The structure of Aβ 1-40 has recently been published [4]. It adopts a compact, partially folded structure (with a central hydrophobic region forming a 3₁₀ helix from H13 to D23) predicted to be crucial in-pathway intermediates in fibrilogenesis that leads to creation of oligomers, protofibrils, and fibrils in equilibrium. Amyloid aggregates are formed when $A\beta$ is excessively produced, cleared too slowly, or in contact with proaggregating factors. Various aggregated forms of Aβ act as potent and direct neurotoxic agents, eliciting a cascade of events leading to neuronal pathology and clinical manifestation of AD. Thus, clearance of Aβ from the brain is first choice in therapies for AD patients [5]. A complementary and actively pursued therapeutic strategy for treating AD is preventing aggregation of Aβ. For instance, efforts to synthesize small molecules that target Aβ aggregates and at the same time complex metal-ions (that are known to facilitate Aß aggregation and produce reactive oxygen species associated to oxidative stress and neuropathology of AD) have been made [6, 7]. Inorganic nanoparticles strongly inhibit Aβ fibrillation by preferentially binding to oligomers [8]. Alternatively, dendrimers - globular, highly branched polymers - either inhibit or accelerate the production of fibrils [9, 10]. A breakthrough in the use of dendrimers as therapeutic agents for AD has been the discovery that polyamidoamine (PAMAM) dendrimers can purge scrapie forms of prion protein (PrPSc) in vitro from infected brain extracts [11, 12]. Several mechanisms have been proposed to explain this dendrimer activity; interaction with peptide monomers, blocking of fibril ends, and disruption of existing fibrils [13].

One of the most important aspects of AD is that neurotoxicity relates to A β aggregates, not to newly solubilized peptides [14]. Mechanisms has been suggested to explain the cytotoxicity induced by A β aggregates, such as activation of false signal transduction pathways that lead to apoptosis [15, 16], the generation of free radicals that result in oxidative stress and mitochondrial dysfunction [17], and the opening of ion-channels in membranes that leads to their depolarization [18].

Herein we report on the influence of GATG (gallic acid-triethylene glycol) dendrimers on the aggregation of an amyloid model peptide AB 1-28, which contains the core region responsible for Aβ aggregation. GATG has recently emerged as an interesting family of dendrimers that benefit from an easy structural modification, and an adequate aqueous solubility and biocompatibility due to their ethylene glycol side-chains [19-21]. The presence of peripheral azides in GATG dendrimers has been exploited for efficient functionalization by means of Cu(I)-catalyzed azide-alkyne cycloaddition (CuAAC) [22, 23]. The resulting activated dendrimers have emerged as interesting agents in the study of the multivalent carbohydrate-receptor interaction [24], the dynamics of dendrimers [25], and the preparation of polyion complex micelles and dendriplexes for gene therapy [26,27]. GATG-based contrast agents for MRI have recently been reported [28]. Encouraged by the promising properties of GATG dendrimers as inhibitors of the dimerization of the capsid protein (CA) of HIV-1 [29], we have investigated their potential to influence the formation of amyloid fibrils and the toxicity of A\beta 1-28. With this aim, a novel GATG dendrimer of generation 3, functionalized with 27 terminal morpholinoethyl groups ([G3]-Mor, Figure 1), has been prepared with CuAAC. N-alkylated morpholines are characterized by physiological pH pKa values, which makes them attractive ligands in dendrimer-amyloid peptide interactions. The formation of fibrils in the presence of [G3]-Mor has been confirmed using the thioflavin T (ThT) assay, TEM observations and CD experiments. The cytotoxicity of Aβ species was monitored by the MTT assay during the aggregation process in the absence and presence of the dendrimer. The cytotoxicity of [G3]-Mor was also assessed.

2. Material and methods

2.1. Materials

A synthetic peptide A β 1-28 [DAEFRHDSGYEVHHQKLVFFAEDVGSNK] (purity >90%) was purchased from JPT Peptide Technology GmbH (Germany). A peptide stock solution was kept in 10 mM HEPES buffer at pH 7.4. Thioflavin T (ThT), heparin-sodium salt from porcine intestinal mucosa (H4784) and 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) were purchased from Sigma Chemical Company. All other chemicals were of analytical grade. Double-distilled water was used to prepare the other solutions. [G3]-Mor: [G3]-N₃ (27 mg, 3.44 μ mol, prepared according to Fernandez-Megia et al. [20]) was dissolved in *t*-BuOH (0.46 ml) and H₂O (0.30 ml). 4-(3-butyn-1-yl)morpholine (26 mg, 0.186 mmol, prepared according to Smaill el at. [30]) and freshly prepared aqueous solutions of CuSO₄ (46 μ L, 4.64 μ mol) and sodium ascorbate (116 μ L, 23.2 μ mol), were added. The solution was stirred at room temperature for 72 h. Aq HCl (0.1 ml, 0.1 M) was added and the mixture purified by ultrafiltration (Amicon YM1, 5 x 30 ml H₂O) before being freeze-dried to

yield the hydrochloride salt of [G3]-Mor (38.8 mg, 90%) as a pale yellow foam. 1 H-NMR (500 MHz, D₂O) δ : 7.90 (br s, 27H), 7.23-6.99 (m, 26H), 4.56 (br s, 54H), 4.21-3.48 (m, 498H), 3.26 (br s, 2H), 3.19-2.71 (m, 216H), 1.56 (br s, 2H), 0.90 (br s, 3H). 13 C NMR (100 MHz, D₂O) δ : 169.6, 169.5, 152.9, 152.7, 144.6, 140.7, 130.3, 125.0, 107.2, 73.2, 71.0, 70.9, 70.7, 70.4, 70.0, 69.7, 69.4, 65.6, 57.4, 52.9, 50.9, 40.7, 21.4, 11.8. IR (KBr, cm⁻¹) 3288, 2928, 2870, 1651, 1118. [G3]-Mor was dissolved in 10 mM HEPES buffer at pH 7.4.

2.2. Formation of amyloid fibrils – ThT assay

Aggregation was monitored using the dye, ThT, whose fluorescence depends on the aggregate formation. A stock solution of A β 1-28 peptide (1 mM) in HEPES, pH 7.4 (kept at 4 °C and used within few days) was diluted to 50 μ M. ThT was then added and its concentration in the sample set to match 35 μ M and a pH of 5.5 with aliquots of aq HCl. Aggregation was triggered by the addition of 0.041 mg/ml heparin. Fluorescence was measured with a Perkin-Elmer LS-50B spectrofluorimeter. Experiments were performed at 37°C with continuous stirring of the sample. The kinetics of aggregation was monitored by the changes in fluorescence intensity during the course of the experiment, with excitation and emission wavelengths of 450 and 490 nm, respectively. The excitation and emission slit widths were set at 5 nm.

2.3. Formation of amyloid fibrils – transmission electron microscopy observations

Fifteen µl of a sample were removed from a fluorimetric cuvette after completion of aggregation (Section 2.2) and placed on a copper grid with carbon surface for 10 min and dried with a filter paper. The sample was stained with 2% (m/v) uranyl acetate for 2 min and dried. Transmission electron microscopy images were taken with a Hitachi H-7000 (75 kV) microscope.

2.4. Formation of amyloid fibrils – Circular dichroism experiments

CD in the far UV region measurements made with a J-815 CD spectrometer (Jasco). A β 1-28 samples was prepared analogously as for a ThT assay except by adding ThT. CD spectra were recorded between 260 and 190 nm using 0.05 cm path-length quartz sandwich cell (Helma). The recording parameters were bandwidth – 1.0 nm, slit width – auto, response – 1 s, scan speed – 50 nm/min, and step resolution – 0.2 nm. The number of scans varied between 3 and 5 for each sample. Measurements were made at different time intervals during aggregation (0, 5, 60, and 120 min from the beginning of the process). CD spectra were corrected by subtracting CD spectra obtained for [G3]-Mor dissolved in a buffer without the peptide. The mean residue ellipticity, θ , expressed as a value deg cm²dmol⁻¹, was calculated.

2.5. Cell culture

Immortalized Chinese hamster fibroblast (B14, ATCC no. CCL-14) were used to measure the cytotoxicity of [G3]-Mor, as also the monomeric and aggregated forms of A β 1-28. Cells were grown in Dulbecco's Modified Eagle's Medium Gluta-MAX (DMEM) (Gibco, UK) supplemented with 10% inactivated fetal bovine serum (Sigma Chemical Company) at 37°C in a humidified incubator with 5% CO₂ in air. Medium was replaced every 3 to 4 days and cells were split for subcultures 1:10 every 4 to 5 days.

2.6. Cytotoxicity of [G3]-Mor – MTT assay

Five \times 10³ cells were seeded per well in 96-well plates with growth media (100 µl). Cells were washed with phosphate-buffer saline (PBS: 2 mM KH₂PO₄, 10 mM Na₂HPO₄, 150 mM NaCl, 2.7 mM KCl, 1 mM CaCl₂, 0.5 mM MgCl₂, pH 7.4). Fresh medium and [G3]-Mor (0.05-1.5 µM) were added 24 h after seeding. After 2 h, the medium with [G3]-Mor was removed, cells were washed twice with PBS, and fresh medium added. After 72 h, 50 µl MTT (0.5 mg/ml in PBS) was added to each well for 4 h, and the supernatant (containing unreacted dye) was replaced with dimethyl sulfoxide (DMSO) (100 µl/well). Plates were shaken and absorbance measured at 540 nm with a reference at 720 nm at a Cary 50 BIO UV-Visible Spectrophotometer. Yellow MTT formed purple formazan by mitochondrial dehydrogenases in the living cells, while no activity of mitochondrial dehydrogenases is detected in dead and damage cells [31]. Cell viability was calculated according to equation:

cell viability[%] =
$$(x/x_c) \times 100$$

where x is the absorbance of cells treated with [G3]-Mor and x_C the absorbance of untreated (control) cells. An IC₅₀ value of cell viability was obtained graphically.

2.7. Cytotoxicity of AB 1-28 forms

Two and a half × 10^3 cells were seeded per well in a 96-well plate with growth media (50 µl). A β 1-28 peptide samples (collected during the aggregation process) were added to cells at $10 \, \mu M$. A β was removed after 2 h later, the cells washed twice with PBS, and fresh medium added. The MTT assay was used as before (section 3.4). The same procedure was applied when A β 1-28 aggregated in the presence of [G3]-Mor.

2.8. Statistics

Data were expressed as means \pm SD, and their statistical significance was analyzed by Tucey's t-test using StatSoft, Inc. (2000) STATISTICA software for Windows.

3. Results

Amyloid fibrils of A β 1-28 were formed *in vitro* and the aggregation process was monitored over time by fluorescence in the ThT assay (Figure 2), having previously checked that ThT did not bind to [G3]-Mor. ThT is a dye with a fluorescence intensity that depends strictly on the presence of fibrils in the system - the more fibrils formed, the higher the ThT fluorescence intensity. A curve characteristic for a nucleation-dependent process was taken as a control

(the system without dendrimer). The lowest concentration of [G3]-Mor added (0.01 μ M) had almost no effect on aggregation, whereas a 10 x higher concentration caused a slight increase in fibril production. The most pronounced effect was obtained with a 1 μ M dimer, which led to the production of far more fibrils and a significant acceleration of aggregation. A plateau was reached after 120 min, compared with ~210 min.

The effect of [G3]-Mor on the morphology of A β 1-28 amyloid aggregates was followed by TEM. For control samples (Figure 3A) and samples containing 1 μ M of [G3]-Mor (Figure 3B), characteristic long fibrils were seen. However, the amount of fibrils formed in 1 μ M [G3]-Mor during aggregation was significantly higher.

The amount of fibrils produced, as judged by TEM and changes in ThT fluorescence intensity were compared with alterations in CD spectra. CD experiments were carried out for the control (without [G3]-Mor) and for the system when [G3]-Mor was present at 1 μ M. CD spectra were recorded in the region of 190-260 nm (Figure 4A). The CD signal here reflects the basic secondary structural features of peptides. The spectrum obtained for non-aggregated A β 1-28 had a broad minimum between 195 and 210 nm that decreased as the aggregation developed. After addition of [G3]-Mor to A β 1-28, the process was faster, as illustrated by the changes in residue mean ellipticity for λ =201 nm during aggregation (Figure 4B).

Different states of A β peptides that are formed during the aggregation process are believed to be associated with neurotoxicity in AD. The effect of [G3]-Mor on the toxicity of A β 1-28 was studied during the progress of amyloid fibril formation by the MTT test (Figure 5). The cytotoxicity of samples taken every 30 min was determined for a control and for the experiments with [G3]-Mor at 0.2 and 0.02 μ M (molar ratios A β /[G3]-Mor equaled to 50 and 500 that correspond to [G3]-Mor concentrations of 1 and 0.1 μ M in the ThT assay). Before the aggregation process started, the A β was almost non-toxic. Toxicity increased significantly after 30 min and reached a maximum after 90 min in the control. Toxicity decreased and after 180 min, cell viability reached the same level as at 30 min. For samples in which [G3]-Mor was present, no toxicity was seen during the first 30 min of aggregation. Nevertheless, [G3]-Mor significantly reduced the cytotoxicity of samples taken at 60, 90, and 120 min (times for which the highest toxic effect was observed in control samples) in a concentration-dependent fashion.

The cytotoxicity of [G3]-Mor was also checked to confirm it was innocuous at 0.02 and 0.2 μ M. Indeed, cell viability started decreasing when the concentration of [G3]-Mor reached 0.5 μ M (Figure 6).

4. Discussion

Dendrimers have been intensively explored as anti-amyloid agents in the last few years [13, 32, 33]. The formation of insoluble, fibrillar protein aggregates is a common of many neurodegenerative disorders, such as AD, Parkinson's disease and prion disorders. Dendrimers have a high affinity for proteins [34, 35], which has been used to explain their interference with amyloidogenesis through several mechanisms (see Introduction). Interestingly, most of the previously tested dendrimers block fibril formation at high concentrations, whereas they accelerate fibril production at lower concentrations [9, 13, 36]. This is typically the pattern for fibril disrupting agents. Inhibitors administered in low doses cause a moderate level of fibril breakage in amyloidogenic processes, and may be responsible for speeding up the formation of aggregates by creating new free-ends that serve as sites of replication. Conversely, the very same inhibitors can be effective at higher doses when breakage is very fast.

With [G3]-Mor, it was soon revealed that it affected the process of amyloidogenesis in a strikingly different way to that of most other dendrimers. Indeed, when [G3]-Mor was used at high concentrations, fibril formation was not inhibited, but aggregation was faster and increased amounts of fibrils were formed, a seemingly undesirable effect. Thus, although for many years the central role of aggregated forms of Aβ in AD pathogenesis has stimulated the development of therapeutic approaches designed to prevent fibril formation (fibrillar forms of Aβ being necessary for neurotoxicity) [37], recent studies show that alternative Aβ assemblies, including small oligomers and fibril intermediates, can also be neurotoxic [38]. For instance, correlation of neurological symptoms of AD with insoluble amyloid load seems to be much worse than with oligomers [39], which have been found in brains of AD patients [40]. In another study, young APP transgenic mice underwent synaptic electrophysiological and behavioral changes before amyloid plaques were formed [41,42]. It is also known that soluble oligomeric Aβ blocks synaptic plasticity is required for memory formation, particularly in the hippocampus where neuronal loss occurs as the disease progresses [43]. In addition, others have shown that pre-fibrillar assemblies interact with synthetic phospholipid bilayers and cell membranes, possibly destabilizing them and impairing specific membrane-bound proteins [44,45]. Disclosure of oligomer toxicity has an immediate practical aspect: therapeutic strategies aimed at the destabilization of Aβ fibrils might prove counterproductive and lead to an increase in the level of oligomers. It seems more important to check the toxicity of these systems and how this changes during aggregation than trying to stop fibril formation.

In our studies, cytotoxicity of was checked by the MTT assay. A small quantity of a sample was taken every 30 minutes from a cuvette where the aggregation progressed (control). Before the aggregation started, no toxic effect was found (time zero). Regarding the

cytotoxicity of non-aggregated and aggregated forms of A β 1-28, this is consistent with the monomeric α -helical conformation being neurotrophic and becoming neurotoxic only after transformation to an oligomeric β -sheet conformation [46]. The toxic effect progressed for up to 90 min, with minimal cell viability. Then toxicity decreased, and after 180 min reached a similar level to in the initial 30 min. Mature amyloid fibrils were substantially less harmful to cells than their precursors. When similar experiments involved A β /[G3]-Mor mixtures prepared at a molar ratio equal to 50, toxicity was substantially decreased. This effect was even more pronounced when [G3]-Mor was 10x higher.

The most likely role played by [G3]-Mor in the reduction of toxicity relates to the faster and higher production of fibrils, with concomitant reduction of the concentration of prefibrillar forms, which in the control experiment are seen as ultimately responsible for its toxicity. Moreover, good correlation is observed between the times for minimal cell viability and ~50% of maximal fluorescence intensity in the ThT assay (90 min for the control and 30 min for the A β /[G3]-Mor system at ratio of 50). It means that agents that accelerate the fibril formation might have a beneficiary effect in terms of reducing A β toxicity. This represents an alternative approach to the search for inhibitors of the formation of both fibrillar and oligomeric morphologies, such as trehalose. Trehalose inhibits formation of fibrils and oligomers for A β 1-40 that results in reduced toxicity against SH-SY5Y cells. However, when trehalose was co-incubated with A β 1-42, it only inhibited formation of the fibrillar morphology, and no protective effect on SH-SY5Y cells was seen [47]. The behavior of the tested dendrimer proves that toxicity can be reduced not only by stopping aggregation, but also by accelerating the formation of mature fibrils that are much less harmful to cells than prefibrillar forms.

To conclude, a GATG dendrimer decorated with 27 terminal morpholine groups ([G3]-Mor) accelerates at non-toxic concentrations A β fibril formation at A β /dendrimer ratios 50 and 500. Even though the final concentration of fibrils was higher in the presence of [G3]-Mor, the A β toxicity was significantly lower. This encourages the monitoring of whether the same phenomenon is observed ith A β 1-40 and A β 1-42. If this is the case, it may have a practical significance in nanomedicine.

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Figure legends:

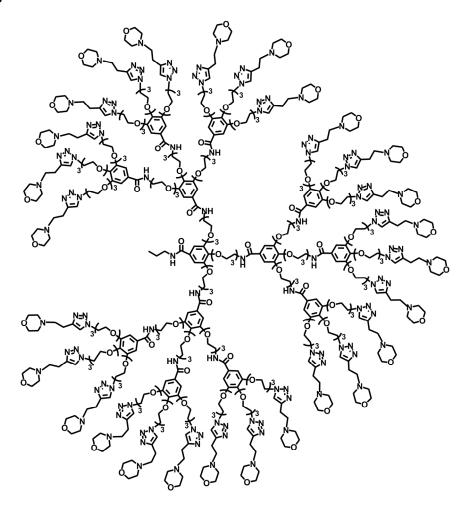


Figure 1. Structure of [G3]-Mor.

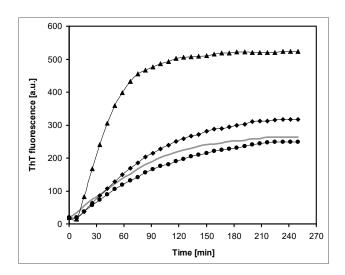


Figure 2. Variation of the fluorescence of ThT during the aggregation of A β 1-28 in the absence (grey line) and presence of [G3]-Mor at 0.01 μ M (circles), 0.1 μ M (rombs), 1 μ M (triangles).

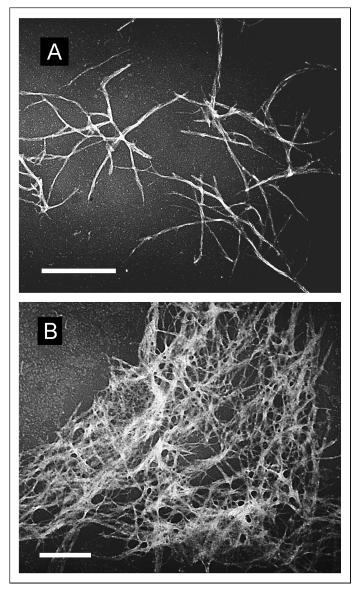


Figure 3. Electron micrographs of A β 1-28 samples at the end of aggregation without [G3]-Mor (A), and in the presence of 1 μ M [G3]-Mor (B). The length of the bar is 200 nm.

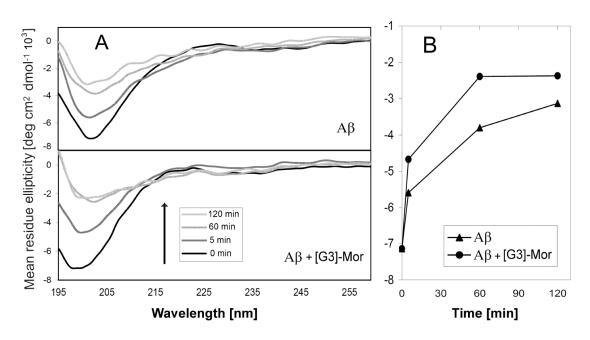


Figure 4. CD spectra of A β 1-28 during aggregation (A) in the absence of [G3]-Mor – top panel, and in the presence of 1 μ M [G3]-Mor – bottom panel. Changes in mean residue ellipticity (B).

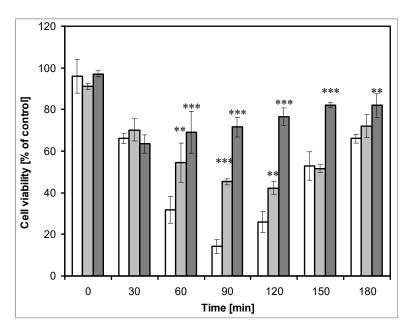


Figure 5. Changes in cell viability upon treatment with A β 1-28 in the absence (white bars) and presence of [G3]-Mor at 0.2 μ M (dark grey bars) and 0.02 μ M (light grey bars). Molar ratios A β /[G3]-Mor equaled to 50 and 500, respectively, that correspond to dendrimer concentrations of 1 and 0.1 μ M in the ThT assay. To check whether the presence of dendrimers affects the toxicity of A β 1-28 species (at different stages of aggregation), the statistical significance was compared between systems with [G3]-Mor and the control (** p<0.01, *** p<0.005).

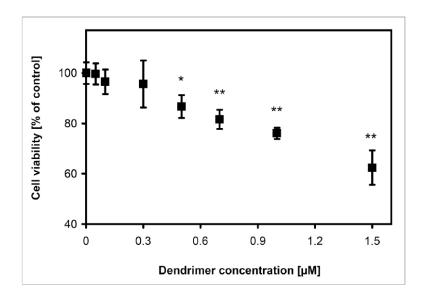


Figure 6. Changes in cell viability after treatment with [G3]-Mor at different concentrations (* p<0.05, ** p<0.01).