



Polycationic peptide R⁷-G-Aβ₂₅₋₃₅ selectively induces cell death in leukemia Jurkat T cells through speedy mitochondrial depolarization, and CASPASE-3 -independent mechanism

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ABSTRACT

Background: Acute lymphoblastic leukemia (ALL) is still incurable hematologic neoplasia in an important percentage of patients. Therefore, new therapeutic approaches need to be developed.

Methods: To evaluate the cellular effect of cell-penetrating peptides (C-PP) on leukemia cells, Jurkat cells -a model of ALL were exposed to increasing concentration (50–500 μM) Aβ₂₅₋₃₅, R⁷-G-Aβ₂₅₋₃₅ and Aβ₂₅₋₃₅-G-R⁷ peptide for 24 h. 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay, flow cytometry (FC), and fluorescent microscopy (FM) analysis were used to assess metabolic viability, cell cycle and proliferation, mitochondria functionality, oxidative stress, and cell death markers.

Results: We report for the first time that the R⁷-G-Aβ₂₅₋₃₅, but not Aβ₂₅₋₃₅ peptide, induced selective cell death in Jurkat cells more efficiently than the Aβ₂₅₋₃₅-G-R⁷ peptide. Indeed, R⁷-G-Aβ₂₅₋₃₅ (200 μM) altered the metabolic activity (–25%), arrested the cell cycle in the G2/M-phase (15%), and induced a significant reduction of cellular proliferation (i.e., –74% reduction of Ki-67 nuclei reactivity). Moreover, R⁷-G-Aβ₂₅₋₃₅ induced the dissipation of mitochondrial membrane potential (ΔΨ_m, 51%) and produced an important amount of reactive oxygen species (ROS, 75% at 8 h) in Jurkat cells. The exposure of cells to antioxidant/cytoprotectant N-acetylcysteine (NAC) did not prevent R⁷-G-Aβ₂₅₋₃₅ from a loss of ΔΨ_m in Jurkat cells. The peptide was also unable to activate the executor CASPASE-3, thereby preserving the integrity of the cellular DNA corroborated by the fact that the caspase-3 inhibitor NSCI was unable to protect cells from R⁷-G-Aβ₂₅₋₃₅ -induced cell damage. Further analysis showed that the R⁷-G-Aβ₂₅₋₃₅ peptide is specifically localized at the outer mitochondria membrane (OMM) according to colocalization with the protein translocase TOMM20. Additionally, the cytotoxic effect of the poly-R⁷ peptide resembles the toxic action of the uncoupler FCCP, mitocan oligomycin, and rotenone in Jurkat cells. Importantly, the R⁷-G-Aβ₂₅₋₃₅ peptide was innocuous to menstrual mesenchymal stromal cells (MenSC) –normal non-leukemia proliferative cells.

Conclusion: Our findings demonstrated that the cationic Aβ peptide possesses specific anti-leukemia activity against Jurkat cells through oxidative stress (OS)- and CASPASE-3-independent mechanism but fast mitochondria depolarization.

1. Introduction

Acute lymphoblastic leukemia (ALL) is a malignant transformation and proliferation of either B-cell or T-cell lymphoid progenitor cells in the bone marrow, blood, and extramedullary sites affecting children, young adolescents, and adults [1] around the world [2]. According to the American Cancer Society, there are approximately 5690 new cases of

ALL, and about 1580 deaths in 2021. While childhood ALL studies have shown improved 5-year overall survival (OS) rates exceeding 90% [3], only 30–40% of adult patients with ALL will achieve long-term remission [4]. Although conventional cytotoxic chemotherapy [5,6] or in combination with monoclonal antibodies (e.g., Ref. [7]) used to treat ALL results in high cure rates, treatment is suboptimal in an important percentage of pediatric and adult patients with relapsed and refractory

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ALL. Therefore, new treatment options are needed for the treatment of resistance ALL. Indeed, leukemia cells avoid apoptosis -a regulated form of cell death [8] contributing to their pathological features. Since apoptosis can be initiated through the intrinsic pathway that depends on mitochondrial outer membrane permeabilization or the extrinsic pathway that depends on external signals via cell receptors [9], a reasonable strategy would be to trigger apoptosis by agents that directly or indirectly target mitochondria or by external signals to boost cell death for anti-leukemic therapy.

Polycationic peptides are a large class of short amino acid sequences (5–30 residues) composed mainly of multiple lysines (K) and/or arginine (R) residues usually used as anti-microbial agents (e.g., Refs. [10–14]). It is the cationic properties that promote the preferential binding of peptides to the negatively charged bacterial cytoplasmic membrane instead of the zwitterionic membrane of mammalian cells [15]. Interestingly, not only they can impair microbial cell walls and plasma membrane phospholipids but also mitochondria in free-cell assays [16,17]. Currently, polycationic peptides have appeared as specifically anti-cancer therapeutic agents [18], targeting mitochondria acting as class 6 mitocan agents [19,20]. Since R oligomers have been shown to facilitate cellular uptake of cargo proteins [21] probably through membrane multilamellar and subsequently entrance via formation of a fusion pore [22], thereby enhancing their capacity to directly permeabilize mitochondria (e.g., Ref. [23]). Indeed, synthetic cell-penetrating peptides (C-PP) poly-Rⁿ (n > 6) have currently been used as the delivery vector [24,25]. For instance, the peptide r7-kla (D-hepta-arginine linked to D-forms of KLA) induced apoptosis (48%) in leukemia Jurkat cells -a T-cell ALL line [23]. Moreover, it has been shown that cationic peptides such as P7-4 (R⁷-GG-IYLATALAKWALKQGF) and P7-5 (IYLATALAKWALKQGF-GG-R⁷) significantly induced demise (29% and 31%, respectively) of leukemia Jurkat cells [17]. Taken together these data suggest that some poly-Rⁿ-spacer-cargo peptides provoke apoptosis in leukemia cells mainly through targeting the mitochondrial membrane.

The A β _{25–35} with sequence GSNKGAIIGLM is an eleven-residue peptide produced by proteolytic conversion of racemized β -amyloid ([d-Ser²⁶] A β _{1–40}) in the brains of aged Alzheimer's disease patients [26]. Since the A β _{25–35} peptide is amphiphilic and tends to aggregate, it has been shown that the peptide displays neurotoxic properties [27] probably through disruption of the structure of the plasma membrane causing uncontrollable permeation of Ca²⁺ ions [28]. However, no data are available to establish whether A β _{25–35} peptide might be cytotoxic to other non-neuronal cells. Based on the above information, we used A β _{25–35} as a paradigm peptide. Indeed, the C-PP vector R⁷ was fused to A β _{25–35} at their N- or C-termini through one unstructured glycine residue used as a spacer to obtain R⁷-G-A β _{25–35} and A β _{25–35}-G-R⁷, respectively. We hypothesized that A β _{25–35} peptide induces apoptosis in leukemia cells. We also theorized that either R⁷-G-A β _{25–35} or A β _{25–35}-G-R⁷ increase several folds apoptosis when compared to A β _{25–35} peptide. To test these hypotheses, we used Jurkat T cell to elucidate the molecular mechanism of cell death induced by either cationic A β _{25–35} or non-cationic A β _{25–35} peptides. To test the selectivity of peptides to induce cell death, menstrual-derived mesenchymal stromal cells (MenSCs) were used as control. We report for the first time that the R⁷-G-A β _{25–35}, but not A β _{25–35} peptide, induced selective cell death in Jurkat cells more efficiently than the A β _{25–35}-G-R⁷ peptide. The R⁷-G-A β _{25–35} peptide altered several metabolic parameters in the leukemia cell line such as metabolic activity, cell cycle, cellular proliferation, $\Delta\Psi_m$ leading to cell death by a mechanism independent of oxidative stress (OS) and CASPASE-3 activation but fast mitochondrial depolarization. Our findings suggest the cationic A β peptide possesses anti-leukemia activity against ALL.

2. Materials and methods

2.1. Peptide synthesis

Peptide A β _{25–35}-G-R⁷ was synthesized by Dr. F Guzman (Núcleo Biotecnología Curauma, Valparaíso, Chile). Peptide synthesis was performed in a Liberty Blue™ automated microwave peptide synthesizer (CEM

Corp., Matthews, NC, USA) following a standard fluorenylmethoxycarbonyl protecting group (Fmoc)/tert-butyl-L-tyrosine (tBu) protocol as previously described [29]. Briefly, a Rink Amide AM resin (loading 0.6 mmol/g) was used as the solid support. Standard couplings of amino acids were carried out at 0.2 M in dimethylformamide (DMF) using diisopropylcarbodiimide (DIC)/OxymaPure® activation (the synthesis method optimization according to Liberty Blue™ recommended operation (CEM Corp., Matthews, NC, USA)), and the corresponding amino acid. Fmoc removal was done with three different reagents: 20% v/v 4-methylpyrazole (4 MP) in DMF; 20% v/v. Deprotection and coupling were performed as described [29]. For arginine, an additional coupling step was performed. Once synthesis was complete, peptides were cleaved manually from the resin with trifluoroacetic acid (TFA) under gentle agitation for 4 h at room temperature (r.t.) in the presence of scavengers (standard cleavage solution: TFA/tri-isopropylsilane (TIS)/Water 95:2.5:2.5). After filtration, the crude peptide was precipitated by cold diethyl ether, centrifuged, washed with cold Et₂O five times, dried, dissolved in ultrapure water, frozen, and lyophilized. Peptide R⁷-G-A β _{25–35} was synthesized by Dr. S Estrada-Gomez and CF Salinas-Restrepo (Ofidism/Esorpionism Program, Universidad de Antioquia). The peptide was chemically synthesized through the solid phase using Fmoc/tBu as the orthogonal protection strategy in the Rink AM resin using the simultaneous synthesis strategy described in Ref. [30]. The (Benzotriazole-1-yl) tetramethyluronium hexafluorophosphate (HBTU), (Benzotriazolyl) tetramethyluronium tetrafluoroborate (TBTU), DIC, and O-(6-Chloro-1-hydroxybenzotriazol-1-yl)-1,1,3,3-tetramethyluronium tetrafluoroborate (TCTU) were used as activators, and the deprotection was carried away with piperidine 20% in DMF. For the cleavage a 92.5% TFA, 2.5% TIS, 2.5% 2-mercaptoethanol and 2.5% water solution was used as described in Ref. [29]. The crude and folded peptide was solubilized in 200 μ L of solution A (0.1% TFA in water) and centrifuged at 1.000 g for 3 min. The supernatant was applied to a reverse-phase CNW Athena C18-WP (CNW Technologies, Düsseldorf, Germany) column (4.6 \times 100 mm, 5 μ m, 100 Å), and separated by RP-HPLC on a Shimadzu Prominence chromatograph (Kyoto, Japan). The crude peptide was precipitated by cold diethyl ether, centrifuged, washed with cold Et₂O three times, dried, dissolved in ultrapure water, frozen, and lyophilized. All reactants were from Merck Millipore (Merck Millipore, Billerica, MA, USA). Peptide A β _{25–35} was acquired from Sigma-Aldrich (Cat#A4559). For culture experiments, peptides were resuspended in Roswell Park Memorial Institute (RPMI)-1640 culture medium with glucose (11 mM; Gibco/Invitrogen, Grand Island, New York, USA).

2.2. Cell line and reagents

Jurkat clone E6-1 cells (Catalog no. TIB-152; American Type Culture Collection (ATCC), Manassas, Virginia, USA) were cultured according to the supplier's indications. The cell suspension (1 \times 10⁶ cells/well in 1 mL final volume) was exposed to increasing concentrations of synthetic peptides (0–500 μ M) freshly prepared in RPMI-1640 medium with glucose (11 mM; Gibco/Invitrogen, Grand Island, New York, USA) in the absence or presence of different products of interest (e.g., antioxidant, inhibitors) for 24 h at 37 °C. All other reagents were from Sigma-Aldrich (St Louis, Missouri, USA).

2.3. Isolation of mesenchymal stromal cells (MSCs)

Isolation of MenSCs was performed according to previous reports [31]. Briefly, a menstrual blood sample (tissue bank code (TBC # 69308) was delivered to the laboratory and mixed with an equal volume of phosphate buffer saline (PBS) containing 1 mM ethylenediamine tetra-acetic acid (EDTA), with 100 U/mL penicillin/streptomycin 0.25 mg/mL amphotericin B, and standard Ficoll procedures. After centrifugation, the cells were suspended in a buffy coat and were transferred into a new tube, washed in PBS twice, and resuspended in a growth medium (low-glucose Dulbecco's modified Eagle's medium (DMEM) medium supplemented with 10% fetal bovine serum (FBS, Gibco, USA),

100 U/mL penicillin/streptomycin, and 0.25 mg/mL amphotericin B), and seeded into 25-cm² plastic cell culture flasks at 37 °C with 5% humidified CO₂.

2.4. MTT assay

The proliferation of Jurkat cells was evaluated using the 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay by determining the conversion of the water-soluble MTT to an insoluble formazan [32]. Briefly, cells were pelleted and were incubated under increasing (0–500 μM) concentrations of synthetic peptides for 24 h. Thereafter, the medium was removed and replaced with fresh medium without phenol red. Cells were incubated with 1.2 mM MTT at 37 °C for 3 h and after incubated with dimethyl sulfoxide (DMSO). Finally, samples were mixed, and the absorbance was read at 570 nm using a Stat Fax 3200 microplate spectrophotometer (Awareness Technology, Palm City, Florida, USA). MTT absorbance was assessed 3 times in independent experiments.

2.5. Ki-67 immunofluorescence

To evaluate the effect of synthetic peptides on Jurkat cells proliferation, we evaluated the percentage of Ki-67 positive nuclei according to Ref. [33]. Briefly, the treated cells were fixed with cold ethanol (–20 °C) for 20 min followed by 10% bovine serum albumin (BSA) blockage. Thereafter, cells were incubated overnight with anti-Ki-67 monoclonal mouse antibody conjugated with fluorescein isothiocyanate isomer 1 (FITC, Cat #F0788, Dako) 1:200. The nuclei were stained with 1 μM Hoechst 33342 (life technologies). Ki-67 reactivity was quantified in a Zeiss AxioStart 100 Fluorescence Microscope equipped with a Zeiss AxioCam Cm1 (Zeiss Wöhlk-Contact-Linsen, Gmb Schöckirchen, Germany) by assessing the labeling percentage from the ratio of the number of nuclei-stained Ki-67 to the total number of nuclei counted per section. At least 10 different randomly selected areas were counted.

2.6. Determination of DNA fragmentation and cell cycle by flow cytometry

DNA fragmentation and cell cycle were determined by using a hypotonic solution of propidium iodide (PI, [34]). After treatments, cells (1x10⁵) were washed with PBS (pH 7.2) and stored in 95% ethanol overnight at –20 °C. Thereafter, cells were washed and incubated in a 400 μL solution containing PI (50 μg/ml), RNase A (100 μg/mL), EDTA (50 mM), triton X-100 (0.2%) for 30 min at 37 °C. The cell suspension was analyzed for PI fluorescence by using a BD LSRFortessa II flow cytometer (BD Biosciences). Quantitative data and figures were obtained using FlowJo 7.6.2 Data Analysis Software. Cells entering the sub-G1 phase were used as a marker of apoptosis (DNA fragmentation). For cell cycle analysis, the sub-G1 population was subtracted from the total acquired events, and the Dean Jett Fox analysis was applied (root mean square (RMS) < 10). The experiment was conducted three times, and 10,000 events were acquired for analysis.

2.7. Evaluation of intracellular hydrogen peroxide (H₂O₂) by flow cytometry

H₂O₂ was determined with 2',7'-dichlorofluorescein diacetate (0.5 μM, DCFH₂-DA) according to Ref. [35]. Briefly, after cell treatment with compounds of interest, cells (1 × 10⁵) were incubated with DCFH₂-DA reagent for 30 min at 37 °C in the dark. Cells were washed and dichlorofluorescein (DCF) fluorescence was determined using a BD LSRFortessa II flow cytometer (BD Biosciences). The experiment was conducted three times, and 10,000 events were acquired for analysis. Quantitative data and figures were obtained using FlowJo 7.6.2 Data Analysis Software.

2.8. Analysis of mitochondrial membrane potential (ΔΨ_m) by flow cytometry

Assessment of the ΔΨ_m was performed according to Ref. [36]. We incubated cells (1x10⁵) for 20 min at r. t. in the dark with a deep red mitotracker (20 nM final concentration) compound (Thermo Scientific, cat# M22426). Cells were analyzed by using a BD LSRFortessa II flow cytometer (BD Biosciences). The experiment was conducted three times, and 10,000 events were acquired for analysis. Quantitative data and figures were obtained using FlowJo 7.6.2 Data Analysis Software.

2.9. Assessment of cell death by fluorescent microscopy

MenSC and Jurkat cells were incubated under increasing (0–500 μM) concentrations of synthetic peptides for 24 h. Nuclei were stained for 20 min at 37 °C in the dark with Hoechst 33342 (0.5 μM), deep red mitotracker (20 nM), and DCFH₂-DA, (0.5 μM). Thereafter, cells were deposited in a microscope slide and covered with a coverslip. Fluorescence microscopy analysis was performed with a Zeiss AxioStart 100 fluorescence microscope equipped with a Zeiss AxioCam Cm1 (Zeiss Wöhlk-Contact-Linsen, Gmb Schöckirchen, Germany). The adjustment of the images obtained was performed with the software provided by the manufacturer (ZEN 2 Core).

2.10. Antioxidant and pharmacological inhibitor protection experiments

Jurkat cell suspension (1x10⁶/well in 1 ml final volume) was left untreated or treated with R⁷-G-Aβ₂₅₋₃₅ synthetic peptide (200 μM) alone or in combination with either antioxidant N-acetyl-L-cysteine (NAC, 1 mM) or the caspase-3 inhibitor 1-(4-Methoxybenzyl)-5-[2-(pyridin-3-ylloxymethyl)pyrrolidine-1-sulfonyl]-1H-indole-2,3-dione (NSCI, 10 μM, Sigma-Aldrich, catalog N1413) at 37 °C for 24 h. The cells were evaluated for ΔΨ_m by flow cytometry. The assessment was repeated three times in independent experiments.

2.11. Flow cytometry analysis for CASPASE-3

Flow cytometry acquisition was used to determine the percentage of Caspase-3 positive cells according to Ref. [37]. Jurkat cells were treated according to the above-mentioned procedures. The fixated Jurkat cells were washed and incubated with rabbit anti-Caspase-3 (Millipore, cat #AB3623) primary antibody (1:200) at 4 °C overnight. Cell suspensions were washed and incubated with DyLight 488 donkey anti-rabbit antibody (1:500). Finally, cells were washed and re-suspended in PBS for analysis on a BD LSRFortessa II flow cytometer (BD Biosciences). Twenty thousand events were acquired, and the acquisition analysis was performed using FlowJo 7.6.2 Data Analysis Software.

2.12. Analysis of mitochondrial membrane potential by fluorescent microscopy

The ΔΨ_m was evaluated according to Ref. [38]. Briefly, we incubated cells (1 × 10⁵) for 20 min at 37 °C in the dark with cationic and lipophilic 3,3'-dihexyloxycarbocyanine iodide (DiOC₆(3)), 20 nM final concentration compound (Calbiochem, Darmstadt, Germany; cat # D-273). Cells were incubated with mitochondrial-targeted drugs (i.e., carbonyl cyanide 4-(trifluoromethoxy)phenylhydrazone, FCCP, 10 μM), oligomycin (5 μg/ml), rotenone (100 μM) or R⁷-G-Aβ₂₅₋₃₅ peptide (200 μM) for 0, 30, 60, 120 and 240 s. Fluorescence microscopy analysis was performed with a Zeiss AxioStart 50 Fluorescence Microscope equipped with a Zeiss AxioCam Cm1 (Zeiss Wöhlk-Contact-Linsen, Gmb Schöckirchen, Germany). The adjustment of the images and the spectral images were obtained with the software provided by the manufacturer (ZEN 2 Core). The experiment was conducted three times.

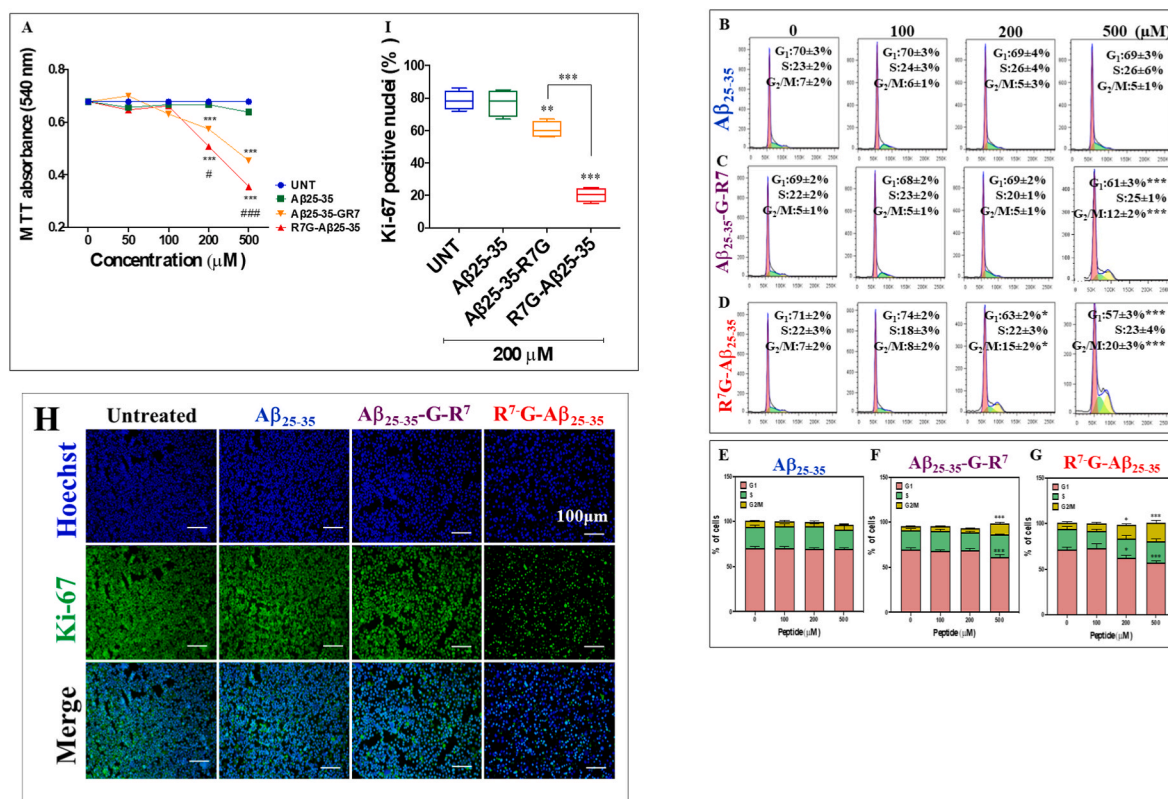


Fig. 1. Effect of Aβ₂₅₋₃₅, Aβ₂₅₋₃₅-G-R⁷, and R⁷-G-Aβ₂₅₋₃₅ synthetic peptides on the metabolic viability, cell cycle, and proliferation in Jurkat cells. (A) Cells were left untreated or treated with Aβ₂₅₋₃₅, Aβ₂₅₋₃₅-G-R⁷, and R⁷-G-Aβ₂₅₋₃₅ (0–500 μM; 24 h), and metabolic viability was measured by MTT assay. (B–D) Representative histograms show cell cycle phase in Jurkat cells treated with 0, 100, 200, and 500 μM of Aβ₂₅₋₃₅ (B), Aβ₂₅₋₃₅-G-R⁷ (C), and R⁷-G-Aβ₂₅₋₃₅ (D) synthetic peptides for 24 h. (E–G) Quantitative data show the cell cycle phase mean percentage of Jurkat cells treated with 0, 100, 200, and 500 μM of Aβ₂₅₋₃₅ (E), Aβ₂₅₋₃₅-G-R⁷ (F), and R⁷-G-Aβ₂₅₋₃₅ (G) synthetic peptides. (H) Representative immunofluorescence images show the Ki-67 and nuclei staining in cells untreated or treated with Aβ₂₅₋₃₅, Aβ₂₅₋₃₅-G-R⁷, and R⁷-G-Aβ₂₅₋₃₅ (200 μM) synthetic peptides. (I) Quantitative data show the mean percentage of Ki-67 positive nuclei in Jurkat cells untreated or treated with Aβ₂₅₋₃₅, Aβ₂₅₋₃₅-G-R⁷, and R⁷-G-Aβ₂₅₋₃₅ (200 μM) synthetic peptides. Figures represent 1 out of 3 independent experiments. The numbers represent the mean percentage of three independent experiments. **p* < 0.05; ***p* < 0.01; ****p* < 0.001. Image magnification 20x.

2.13. Immunofluorescence colocalization analysis of the Aβ peptide and cluster of differentiation (CD)45 or translocase of the outer mitochondrial membrane (TOMM)20

The immunofluorescent staining procedure was according to the standard procedure [39]. Briefly, untreated, and treated cells (0 or 200 μM, respectively) with synthetic peptides (Aβ₂₅₋₃₅, R⁷-G-Aβ₂₅₋₃₅) were plated in a positively charged slide and air-dried. Cells were fixed using 4% formaldehyde for 20 min. After permeabilization, cells were incubated overnight at 4 °C with primary anti-amyloid (25–35) rabbit polyclonal antibody (GenScript, cat# A00687-40), in combination with CD45 mouse monoclonal antibody (BD Pharmingen, cat# 555482) or TOMM20 mouse monoclonal antibody (ab115746). All primary antibodies were prepared at 1:200. After several washes, cells were incubated with Alexa Fluor488 and 594 donkey anti-rabbit and anti-mouse IgG secondary antibodies, respectively (Life Technologies), according to the supplier's protocol (Life Technologies, Eugene, Oregon, USA).

2.14. Photomicrography and image analysis

The fluorescent microscopy photographs were taken using a Zeiss Axiostart 100 Fluorescence Microscope equipped with a Zeiss AxioCam Cm1 (Zeiss Wöhlk-Contact-Linsen, Gmb Schöckirchen, Germany). Images were analyzed by ImageJ software [40]. The figures were transformed into 8-bit images and the background was subtracted. The cellular measurement regions of interest (ROI) were drawn over cell structures (i.e., membrane or mitochondria) and the fluorescence

intensity was subsequently determined by applying the same threshold for controls and treatments.

2.15. Statistical analysis

Statistical analyses were performed using the GraphPad Prism 6 scientific software (GraphPad, Software, Inc. La Jolla, CA, USA). Data are expressed as the mean ± S.D. of a minimum of three independent experiments. One-way ANOVA with a Tukey post hoc test was used to compare the differences between the experimental groups. A P-value < 0.05 (*), < 0.01 (**), and < 0.001 (***) were statistically significant.

3. Results

3.1. R⁷-G-Aβ₂₅₋₃₅ and Aβ₂₅₋₃₅-G-R⁷, but not Aβ₂₅₋₃₅ peptide, induce cell cycle arrest, reduce cellular proliferation, and diminish cellular metabolic activity in Jurkat cells

We first wanted to evaluate the effect of Aβ₂₅₋₃₅, R⁷-G-Aβ₂₅₋₃₅, and Aβ₂₅₋₃₅-G-R⁷ peptides on cellular metabolic activity, cell proliferation, and cell cycle in Jurkat cells. To this aim, cells were left untreated or treated with an increasing concentration of the peptides (0–500 μM) for 24 h. As shown in Fig. 1A, Jurkat cells exposed to Aβ₂₅₋₃₅ peptide remain unaffected according to cellular metabolic activity assay when compared to untreated cells. However, leukemia cells showed a progressive and significant concentration-dependent viability loss by both R⁷-G-Aβ₂₅₋₃₅ (e.g., –25, –48% reduction at 200–500 μM, respectively)

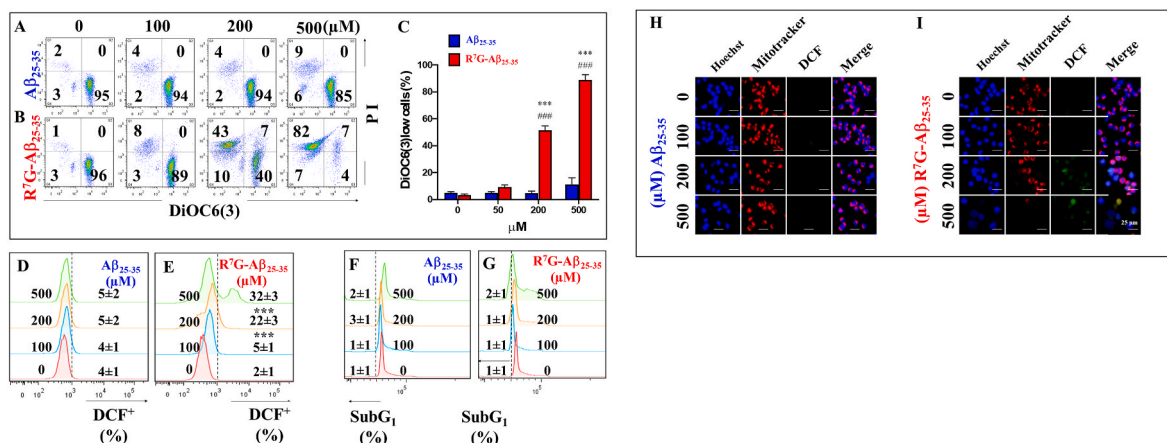


Fig. 2. Synthetic peptide R⁷-G-Aβ₂₅₋₃₅ induces dissipation of mitochondrial membrane potential ($\Delta\Psi_m$), and production of reactive oxygen species (ROS), but no damage to nuclei in Jurkat cells.

(A-I) Representative dot plot figures show DiOC₆(3)/PI double staining in Jurkat cells treated with 0, 100, 200, and 500 μM (A) Aβ₂₅₋₃₅, and (B) R⁷-G-Aβ₂₅₋₃₅ synthetic peptides for 24 h. (C) Quantitative data show the DiOC₆(3)^{low} mean percentage (Q1 + Q4) of Jurkat cells treated with 0, 100, 200, and 500 μM Aβ₂₅₋₃₅ and R⁷-G-Aβ₂₅₋₃₅ synthetic peptides. Representative histograms show the DCF⁺ mean percentage in Jurkat cells treated with 0, 100, 200, and 500 μM (D) Aβ₂₅₋₃₅ and (E) R⁷-G-Aβ₂₅₋₃₅ peptides. Representative histograms show the SubG₁ mean percentage in Jurkat cells treated with 0, 100, 200, and 500 μM (F) Aβ₂₅₋₃₅ and (G) R⁷-G-Aβ₂₅₋₃₅ peptides. Representative fluorescence images show the Hoechst/Mitotracker/DCF triple staining in cells treated with 0, 100, 200, and 500 μM (H) Aβ₂₅₋₃₅ and (I) R⁷-G-Aβ₂₅₋₃₅ peptide. Figures represent 1 out of 3 independent experiments. The numbers represent the mean percentage of three independent experiments. **p* < 0.05; ***p* < 0.01; ****p* < 0.001. Image magnification 20x.

and Aβ₂₅₋₃₅-G-R⁷ (e.g., -15 -33%) peptides. Importantly, R⁷-G-Aβ₂₅₋₃₅ significantly reduced cellular metabolic activity compared to the Aβ₂₅₋₃₅-G-R⁷ peptide. Cell cycle analysis revealed that Aβ₂₅₋₃₅ did not affect the Jurkat cell cycle (Fig. 1B and E), whereas Aβ₂₅₋₃₅-G-R⁷ (Fig. 1C and F) and R⁷-G-Aβ₂₅₋₃₅ (Fig. 1D and G) induced a significant G₂/M-phase cell arrest in a concentration-dependent manner, albeit R⁷-G-Aβ₂₅₋₃₅ (200 μM) was more effective altering the Jurkat cell cycle than the Aβ₂₅₋₃₅-G-R⁷ peptide. Similarly, we found that Aβ₂₅₋₃₅ peptide (e.g., 200 μM) did not affect proliferative-associated Ki-67 protein expression (Fig. 1H and I), whereas R⁷-G-Aβ₂₅₋₃₅ peptide (e.g., 200 μM) and Aβ₂₅₋₃₅-G-R⁷ induced a significant reduction (-74% and -23%) of Ki-67 nuclei reactivity, respectively (Fig. 1I). Because the R⁷-G-Aβ₂₅₋₃₅ peptide altered cellular metabolic activity, cell proliferation, and cell cycle more efficiently than Aβ₂₅₋₃₅-G-R⁷ in Jurkat cells, the former peptide was selected for further experiments. The Aβ₂₅₋₃₅ peptide was included for comparative purposes.

3.2. R⁷-G-Aβ₂₅₋₃₅, but not Aβ₂₅₋₃₅ peptide, induces dissipation of mitochondrial membrane potential ($\Delta\Psi_m$) and produced reactive oxygen species (ROS) preserving cellular DNA integrity in Jurkat cells

Next, we wanted to determine whether the synthetic peptides induce $\Delta\Psi_m$, ROS production, and DNA fragmentation in leukemic cells. Therefore, Jurkat cells were left untreated or treated with increasing concentrations (0–500 μM) of Aβ₂₅₋₃₅, or R⁷-G-Aβ₂₅₋₃₅ for 24 h at 37 °C. As shown in Fig. 2, Aβ₂₅₋₃₅ neither alter the $\Delta\Psi_m$ (Fig. 2A and C) nor influenced ROS production (Fig. 2D) nor induced DNA fragmentation (Fig. 1F), whereas R⁷-G-Aβ₂₅₋₃₅ peptide not only significantly induced a concentration-dependent loss of $\Delta\Psi_m$ (Fig. 2B and C) but induced ROS production (Fig. 2E) according to flow cytometry analysis. These observations were confirmed by fluorescent microscopy (Fig. 2H and I). Analysis of the content of cellular DNA indicated that SubG₁ remained unaffected when exposed to either peptide (Fig. 2F and G).

3.3. The toxic effect of the R⁷-G-Aβ₂₅₋₃₅ peptide is independent of ROS production and CASPASE-3 activation in Jurkat cells

Since ROS plays an important role as a signaling molecule [41], we

evaluated whether the generation of ROS by R⁷-G-Aβ₂₅₋₃₅ peptide-induced activation of CASPASE-3 over time. As shown in Fig. 3, R⁷-G-Aβ₂₅₋₃₅ induced a dramatic increment of ROS production up to 8 h (i.e., ~73% increase) followed by a significant reduction (e.g., ~-71% reduction) after 24 h according to DCFH₂-DA oxidation into DCF fluorescence assay (Fig. 3A). CASPASE-3 activity remained unaffected during observational time points (Fig. 3B). To further corroborate these observations, we left Jurkat cells untreated or treated with R⁷-G-Aβ₂₅₋₃₅ peptide alone or in combination with NAC – a well-known cytoprotective and antioxidant agent [42,43], or NSCI – a caspase-3 inhibitor. Fig. 3 shows that neither NAC (5 mM) nor NSCI (10 μM) protected Jurkat cells against R⁷-G-Aβ₂₅₋₃₅ peptide-induced loss of $\Delta\Psi_m$ (Fig. 3C and D).

3.4. The R⁷-G-Aβ₂₅₋₃₅ peptide dramatically reduces $\Delta\Psi_m$ by its mitochondrial accumulation in Jurkat cells

To further characterized the effect of the R⁷-G-Aβ₂₅₋₃₅ peptide on mitochondria, we wanted to specifically localize the cellular site of action of the peptide. To this aim, we left Jurkat cells untreated or treated with Aβ₂₅₋₃₅ or R⁷-G-Aβ₂₅₋₃₅ peptides for 24 h. Thereafter, we used the plasma membrane marker CD45, or outer mitochondrial membrane receptor TOMM20 together with antibody Aβ₂₅₋₃₅ for co-localization purposes. As shown in Fig. 4, Aβ₂₅₋₃₅ localizes neither at the plasma membrane (Fig. 4B, i.e., positive CD45 /negative Aβ₂₅₋₃₅ fluorescence) nor at mitochondria (Fig. 4E, i.e., positive TOMM20/negative Aβ₂₅₋₃₅ fluorescence) compared to untreated cells (Fig. 4A and D), whereas R⁷-G-Aβ₂₅₋₃₅ peptide was positively detected at mitochondria (Fig. 4C, 4F–H).

3.5. The R⁷-G-Aβ₂₅₋₃₅ peptide induces a rapid loss of $\Delta\Psi_m$ by affecting mitochondria oxidative phosphorylation and uncoupling mitochondrial potential in Jurkat cells

The above observations prompted us to evaluate the effect R⁷-G-Aβ₂₅₋₃₅ peptide on the mitochondria in Jurkat cells over time. Flow cytometry analysis indicates that R⁷-G-Aβ₂₅₋₃₅ peptide significantly reduced the $\Delta\Psi_m$ as early as 4 h in Jurkat cells (Fig 5A, 50% = 37% early (green circle, Q4) plus 13% late (orange circle, Q1) mitochondrial

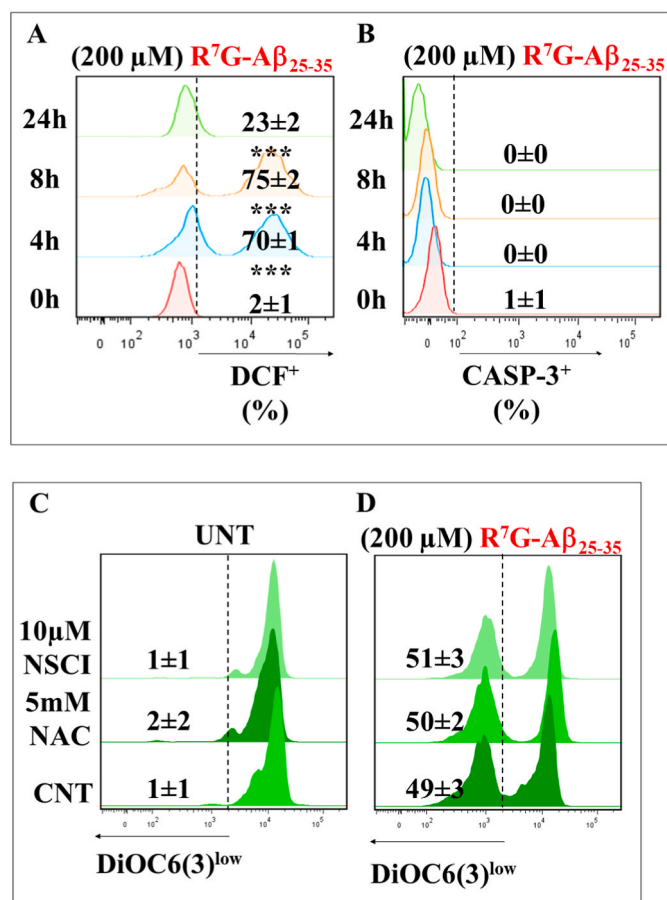


Fig. 3. Synthetic peptide R^7 -G- $A\beta_{25-35}$ induces cell death in a CASPASE-3 (CASP-3)- and reactive oxygen species-independent mechanism in Jurkat cells

(A) Representative histograms show the DCF⁺ mean percentage in Jurkat cells treated with R^7 -G- $A\beta_{25-35}$ peptide (200 μ M) for 0, 4, 8, and 24 h. (B) Representative histograms show the CASP-3⁺ mean percentage of Jurkat cells treated with R^7 -G- $A\beta_{25-35}$ (200 μ M) peptide for 0, 4, 8, and 24 h. (C) Representative histograms show the DiOC₆(3)^{low} mean percentage of Jurkat cells untreated, or with NSCI (10 μ M) or NAC (5 mM) only for 24 h. (D) Representative histograms show the DiOC₆(3)^{low} mean percentage of Jurkat cells treated with R^7 -G- $A\beta_{25-35}$ peptide (200 μ M) only or with NSCI (10 μ M) or with NAC (5 mM) for 24 h. The numbers represent the mean percentage of three independent experiments. * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$.

damage), and this effect was stable over time up to 24 h (~50%, Fig. 5B) followed by a stepwise progressive cell membrane permeabilization according to propidium iodide-stained cells (Fig. 5A). We wanted to determine whether the R^7 -G- $A\beta_{25-35}$ peptide was linked to the impairment of the mitochondria oxidative phosphorylation constituents. To this aim, we compared the effect of R^7 -G- $A\beta_{25-35}$ peptide with other well-known mitochondrial-targeted drugs such as mitochondria Complex I (NADH: ubiquinone oxidoreductase) inhibitor rotenone (class 5 mitocan), complex V (ATP synthase) inhibitor oligomycin (class 5 mitocan), and classical uncoupler of oxidative phosphorylation FCCP (carbonyl cyanide 4-(trifluoromethoxy) phenylhydrazone) reflected as $\Delta\Psi_m$ changes on Jurkat cells. As expected, in about 240 s, FCCP (Fig. 5C), oligomycin (Fig. 5D), and rotenone (Fig. 5E) induced a rapid and progressive loss of $\Delta\Psi_m$ according to DiOC₆(3) fluorescence change analysis (Fig. 5G). Strikingly, R^7 -G- $A\beta_{25-35}$ peptide -a class 6 mitocan had a similar effect on the $\Delta\Psi_m$ when compared to mitochondrial-targeted drugs (Fig. 5F and G).

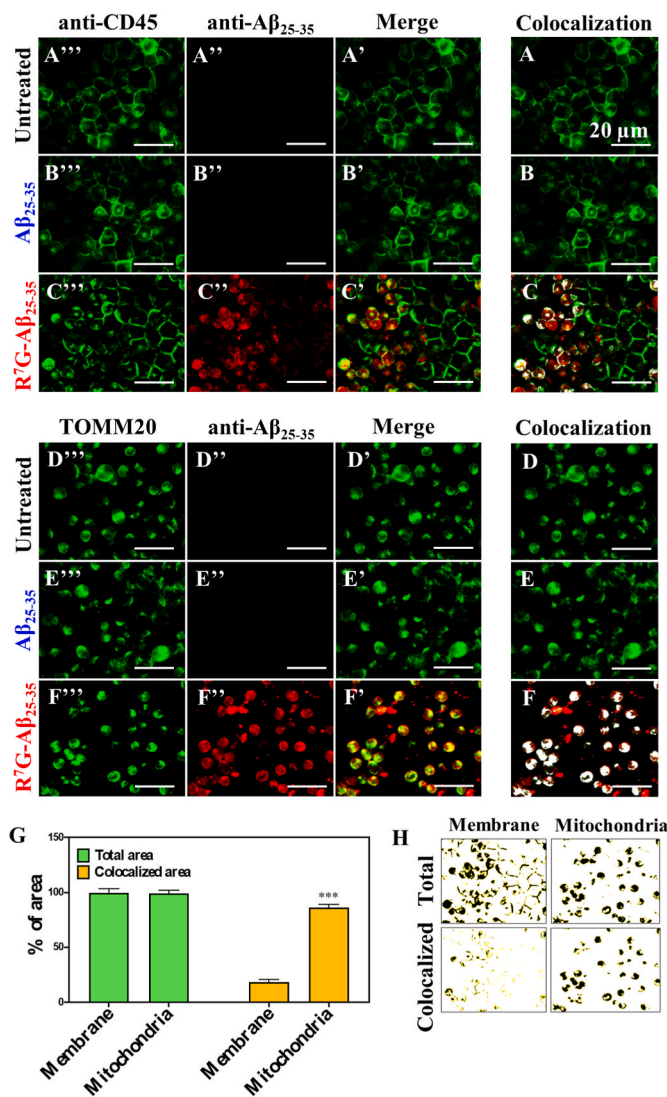


Fig. 4. Synthetic peptide R^7 -G- $A\beta_{25-35}$ co-localizes with mitochondria but not with the cell membrane in Jurkat cells.

(A-C) Representative colocalization images (A-C) and merge images (A'-C') show $A\beta_{25-35}$ epitope sequence (red; A'-C') and CD45 (green; A''-C'') of untreated (A), treated with $A\beta_{25-35}$ (B) or with R^7 -G- $A\beta_{25-35}$ peptide (C) in Jurkat cells for 24 h. (D-F) Representative colocalization images (D-F) and merge images (D'-F') show $A\beta_{25-35}$ epitope sequence (red; D'-F') and TOMM20 (green; D''-F'') of untreated (D), treated with $A\beta_{25-35}$ (E) or with R^7 -G- $A\beta_{25-35}$ peptide (F) in Jurkat cells for 24 h. (G) Quantitative data show the mean percentage of membrane or mitochondria area co-localized with β_{25-35} epitope sequence in Jurkat cells treated with R^7 -G- $A\beta_{25-35}$ (200 μ M) peptide. (H) Representative images show the measurement of total and co-localized area. The numbers represent the mean percentage of three independent experiments. * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$. Image magnification 100x. (For interpretation of the references to colour in this figure legend, the reader is referred to the Web version of this article.)

3.6. The R^7 -G- $A\beta_{25-35}$ peptide is innocuous to non-leukemic proliferative stromal cells

To test the selective effect of R^7 -G- $A\beta_{25-35}$ peptide on normal non-leukemic proliferative cells, we left mesenchymal stromal cells (MSC) untreated or incubated with $A\beta_{25-35}$ or R^7 -G- $A\beta_{25-35}$ peptides (200 μ M) for 24 h, and evaluated the $\Delta\Psi_m$, ROS production and cell cycle analysis by flow cytometry and fluorescent microscopy (FM). Flow cytometry and FM analysis reveal that, like untreated cells (Fig. 6A and D, 6E''), neither $A\beta_{25-35}$ (Fig. 6B, D, 6F'') nor R^7 -G- $A\beta_{25-35}$ (Fig. 6C, D, 6G'')

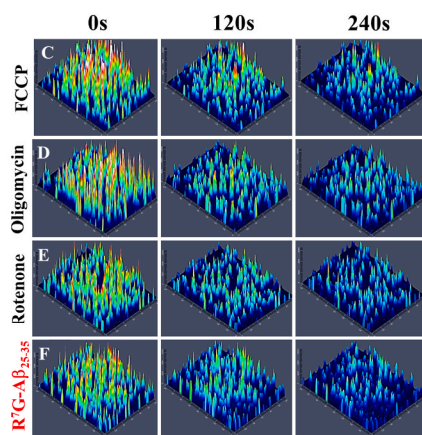
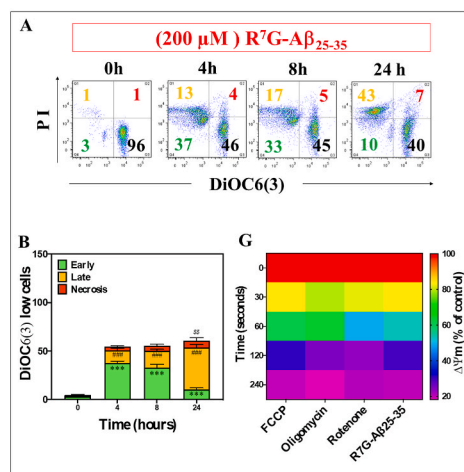


Fig. 5. Fast depolarization of mitochondrial membrane potential ($\Delta\Psi_m$) by synthetic peptide R^7 -G- $A\beta_{25-35}$ in Jurkat cells.

(A) Representative dot plot show $DiOC_6(3)/PI$ double staining in Jurkat cells treated with R^7 -G- $A\beta_{25-35}$ peptide (200 μ M) for 0, 4, 8, and 24 h. (B) Quantitative data show the mean percentage of early ($DiOC_6(3)^{low}/PI^-$; Q4) and late ($DiOC_6(3)^{low}/PI^+$; Q1) apoptotic cells, or the necrotic ($DiOC_6(3)^{high}/PI^+$; Q2) population. (C–F) Representative spectral images show the fluorescence intensity of Jurkat cells treated with (C) FCCP (10 μ M), (D) Oligomycin (5 μ g/ml), (E) rotenone (100 μ M), and (F) R^7 -G- $A\beta_{25-35}$ peptide (200 μ M) during 0, 120 and 240 s. (G) Heat map shows the mean fluorescence intensity (% of control at 0 s) of $DiOC_6(3)$ in Jurkat cells treated with FCCP (10 μ M), oligomycin (5 μ g/ml), rotenone (100 μ M), or R^7 -G- $A\beta_{25-35}$ peptide (200 μ M) during 0, 30, 60, 120 and 240 s. The numbers represent the mean percentage of three independent experiments. * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$. Image magnification 20x.

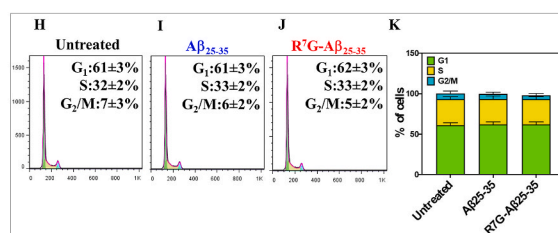
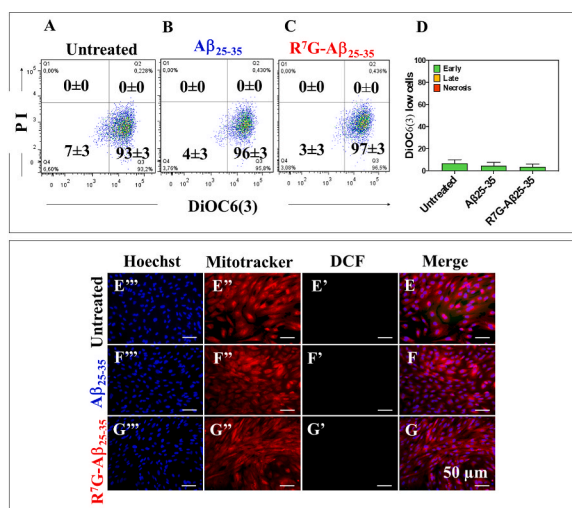


Fig. 6. Synthetic peptide R^7 -G- $A\beta_{25-35}$ induces no mitochondrial membrane potential ($\Delta\Psi_m$) dissipation, reactive oxygen species (ROS) production or cell cycle arrest in normal proliferating cells mesenchymal stromal cells (MSCs).

(A) Representative dot plot show $DiOC_6(3)/PI$ double staining in MSCs untreated or treated with $A\beta_{25-35}$ (200 μ M) or R^7 -G- $A\beta_{25-35}$ (200 μ M) peptides for 24 h. (B) Quantitative data show the $DiOC_6(3)^{low}$ mean percentage (Q1 + Q4) of MSCs untreated or treated with $A\beta_{25-35}$ and R^7 -G- $A\beta_{25-35}$ peptides. (C–E) Representative merge fluorescence images show DCF (C'–E'), Mitotracker (C''–E'') and Hoechst (C'''–E''') staining in MSCs untreated (C), treated with $A\beta_{25-35}$ (D) or R^7 -G- $A\beta_{25-35}$ (E) peptides for 24 h. (F) Representative histograms show cell cycle phase in MSCs untreated or treated with $A\beta_{25-35}$ or R^7 -G- $A\beta_{25-35}$ (200 μ M) synthetic peptides for 24 h. (G) Quantitative data show the cell cycle phase mean percentage of MSCs treated as described above. Figures represent 1 out of 3 independent experiments. The numbers represent the mean percentage of three independent experiments. * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$. Image magnification 20x.

peptides affected $\Delta\Psi_m$ (Fig. 6A–G). Further, the peptides neither induced ROS production (Fig. 6F' and 6G') nor affected the cell cycle progression in MSC (Fig. 6I–K) compared to untreated cells (Fig. 6H, K, and 6E').

4. Discussion

Cell-penetrating peptides (C-PP) are promising anti-cancer agents [18,25]. Here, we report for the first time that the synthetic cationic peptide R^7 -G- $A\beta_{25-35}$ and $A\beta_{25-35}$ -G- R^7 , but not $A\beta_{25-35}$ peptide, induced cell death in Jurkat cells, albeit with different strengths. Effectively, the R^7 -G- $A\beta_{25-35}$ reduced cellular metabolic activity (e.g., -25 to -48% reduction), induced G2/M-phase cell arrest (15–20%), and reduced cell proliferation (-74% reduction). Why is R^7 -G- $A\beta_{25-35}$ more cytotoxic than $A\beta_{25-35}$ -G- R^7 ? One possible explanation is that the poly- R^7 located at the amino-terminal of the $A\beta_{25-35}$ might provide a better

cell-penetrating capability of the peptide through a passive translocation mechanism [22] than when the poly- R^7 is located at the carboxyl-terminal position of the $A\beta_{25-35}$. Interestingly, we found that the $A\beta_{25-35}$ peptide was innocuous to Jurkat cells. This observation might be the result of significant differences in the net charge (at pH 7.0) between the $A\beta_{25-35}$ fragment (+1.00) and R^7 -G- $A\beta_{25-35}$ (+8.00). Indeed, the high net charge of the poly-R-peptide might favor its faster interaction with the highly negative charge of the cellular plasma membrane (~ -60 mV), thereby facilitating its cell entrance. Together these observations suggest that the poly- R^7 attached to the amino position is critical for efficient cell-penetrating peptide. However, further chemical structural studies are needed (e.g., circular dichroism spectroscopy) to fully understand the biochemical behavior of the R^7 -G- $A\beta_{25-35}$ ($A\beta_{25-35}$ -G- R^7) peptide. Whatever the mechanism of interaction between poly- R^7 peptides and plasma membrane, we found that once inside the cell, the R^7 -G- $A\beta_{25-35}$ targets mitochondria.

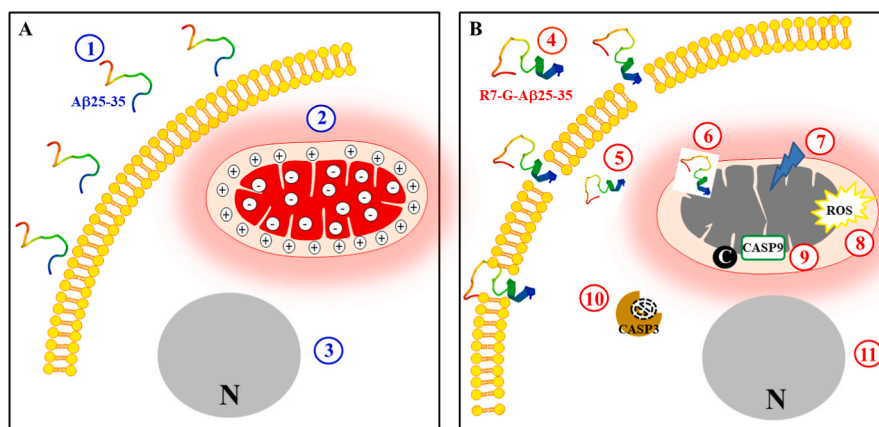


Fig. 7. Schematic representation of the cytotoxic effect of R⁷-G-Aβ₂₅₋₃₅ on Jurkat cells. (See text for explanation).

Mounting evidence has shown that mitochondria play a central role in the regulation of cellular metabolism, bioenergetics, and cell life/death decision during carcinogenesis [44]. Therefore, therapeutic interventions (e.g., mitocans) for the targeting of mitochondria exhibit enormous potential for future leukemia therapeutic strategies [20]. In line with this view, we show that R⁷-G-Aβ₂₅₋₃₅ induced a rapid depolarization of the $\Delta\Psi_m$ as early as 240s post-exposure in Jurkat cells. Remarkably, the R⁷-G-Aβ₂₅₋₃₅-induced loss of $\Delta\Psi_m$ resembled the rapid depolarization of the mitocan 5 rotenone, which specifically blocks mitochondrial complex I, and of the mitocan 5 oligomycin, which specifically blocks F₀ of the ATP synthase [19]. These observations suggest that R⁷-G-Aβ₂₅₋₃₅ might work as mitocan 5. However, whether R⁷-G-Aβ₂₅₋₃₅ directly interacts with mitochondrial complex I and/or ATP synthase, or if the mitochondrial depolarization is the consequence of the interaction of the poly-R⁷ peptide with the outer mitochondria membrane requires further investigation. However, since R⁷-G-Aβ₂₅₋₃₅-induced mitochondrial depolarization also resembled the effect of the uncoupler FCCP, our observations favor the view that loss of $\Delta\Psi_m$ might be the result of a complex interaction of the peptide with the inner mitochondrial membrane leading to uncoupling mitochondria [45], thereby inhibiting the complex I and ATP synthase, and generating ROS. This assumption is supported by 3 observations. First, R⁷-G-Aβ₂₅₋₃₅ accumulated at the outer mitochondria membrane according to colocalization with protein TOMM20. However, whether TOMM20 -a mitochondrial import receptor translocase [46] participates in the import of R⁷-G-Aβ₂₅₋₃₅ to the mitochondrial matrix needs further investigation. Second, R⁷-G-Aβ₂₅₋₃₅ induced an important production of ROS concomitant with mitochondrial depolarization, an effect that was not reversed by the antioxidant/cytoprotectant NAC [42,43]. Third, it is well established that a drastic loss of $\Delta\Psi_m$ generates ROS [47]. Taken together these observations suggest that R⁷-G-Aβ₂₅₋₃₅ induces a deleterious domino-like phenomenon in Jurkat cells, involving its accumulation at and ensuing depolarization of mitochondria through uncoupling mechanism, subsequent inhibition of Complex I, generation of ROS, and dysfunction of ATP synthase.

Unexpectedly, we found that the R⁷-G-Aβ₂₅₋₃₅ peptide did not induce CASPASE-3 activity, thus preserving nucleus integrity [48,49]. We speculate that this phenomenon might be due to the rapid depolarization of mitochondria, which in turn might trap effector cytochrome C, and pro-CASPASE-9 between the inner and outer mitochondria membrane space akin to “Venus’s flytrap” mechanism, and depletion of ATP content, thus disabling the apoptosome complex to process pro-CASPASE-3 into active CASPASE-3 [50]. Of note, the specific inhibitor caspase-3 NCSI was unable to protect Jurkat cells against R⁷-G-Aβ₂₅₋₃₅-induced toxicity. These results imply that R⁷-G-Aβ₂₅₋₃₅ can kill leukemia cells independently of CASPASE-3 [51]. In contrast to the

above observations, R⁷-G-Aβ₂₅₋₃₅ was innocuous to MenSC -normal non-leukemic proliferative cells. These data suggest that R⁷-G-Aβ₂₅₋₃₅ specifically deletes leukemia cells.

5. Conclusion

We demonstrate that R⁷-G-Aβ₂₅₋₃₅ (Aβ₂₅₋₃₅-G-R⁷) specifically induces cell death in Jurkat cells through a direct and fast disruption of the mitochondria membrane potential, and CASPASE-3 independent mechanism. In agreement with our observations, others have reported that C-PP (e.g., r7-kla; RRRRRR-GG-IYLATALAKWALKQGF) were able to kill cancer cell lines, including Jurkat cells through permeabilization of the mitochondrial inner membrane, and apoptosis [17,23]. These observations comply with the notion that some C-PP affect mitochondria membranes rather than other cellular membranes. Accordingly, the unmodified Aβ₂₅₋₃₅ (Fig. 7, step 1) is unable to pass the cellular plasma membrane. Therefore, functional mitochondria (s2) and intact nuclei are observed in Jurkat cells (s3). In contrast, when cells are exposed to R⁷-G-Aβ₂₅₋₃₅ peptide (s4), it translocates passively to the cell cytoplasm (s5). Due to its high net charge (+8.0), the peptide is attracted by the high negative mitochondrial potential (−140 to −180 mV [52]), leading to extensive accumulation of poly-R⁷ cation within mitochondria (s6) probably with the assistance of protein TOMM20. These actions can disrupt mitochondria membrane integrity, respiration, and ATP synthesis (s7), thereby generating a high amount of ROS (s8). As a result of those mitochondria alterations, depolarized mitochondria instantly trap pro-apoptogenic proteins pro-caspase-9 and cytochrome C (s9) avoiding activation of CASPASE-3 (s10). Therefore, despite mitochondria dysfunction, no nuclei damage is appreciably detected (s11). Taken together these observations suggest that the cationic Aβ peptide might be a potential anti-leukemia agent against ALL.

Ethical approval and consent to participate

Not applicable.

Consent for publication

Not applicable.

Availability of supporting data

All relevant data and materials are within the paper.

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Authors’ contributions

Conceptualization, C.V.-P., and M.J.-Del-R.; methodology, C.V.-P.; validation, M.M.-P.; formal analysis, M.M.-P., and C.V.-P.; investigation, M.M.-P.; resources, C.V.-P. and M.J.-Del-R.; data curation, C.V.-P.; writing—original draft preparation, C.V.-P., and M.J.-Del-R.; writing—review and editing, M.M.-P.; C.V.-P. and M.J.-Del-R.; supervision, C.V.-P., and M.J.-Del-R.; project administration, M.J.-Del-R.; funding acquisition, C.V.-P. All authors have read and agreed to the published version of the manuscript.

Declaration of competing interest

All the authors have no competing interests to declare. The funders of the study had no role in study design, data collection, data analysis, data interpretation, or writing of the report.

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