



# Synthesis and Pharmacological Evaluation of a Novel Peptide Based on *Anemonia sulcata* BDS-I Toxin as a New $K_V3.4$ Inhibitor Exerting a Neuroprotective Effect Against Amyloid- $\beta$ Peptide

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There is increasing evidence that the fast-inactivating potassium current  $I_A$ , encoded by  $K_V3.4$  channels, plays an important role in Alzheimer's Disease (AD), since the neurotoxic  $\beta$ -amyloid peptide $_{1-42}$  ( $A\beta_{1-42}$ ) increases the  $I_A$  current triggering apoptotic processes. The specific inhibition of  $K_V3.4$  by the marine toxin extracted from *Anemonia sulcata*, named blood depressing substance-I (BDS-I), reverts the  $A\beta$  peptide-induced cell death. The aim of the present study was to identify the smallest fragments of BDS-I, obtained by peptide synthesis, able to inhibit  $K_V3.4$  currents. For this purpose, whole-cell patch clamp technique was used to evaluate the effects of BDS-I fragments on  $K_V3.4$  currents in CHO cells heterologously expressing  $K_V3.4$ . We found that BDS-I[1-8] fragment, containing the N-terminal octapeptide sequence of full length BDS-I, was able to inhibit  $K_V3.4$  currents in a concentration dependent manner, whereas the scrambled sequence of BDS-I[1-8] and all the other fragments obtained from BDS-I full length were ineffective. As we demonstrated in a previous study, BDS-I full length is able to counteract  $A\beta_{1-42}$ -induced enhancement of  $K_V3.4$  activity, preventing  $A\beta_{1-42}$ -induced caspase-3 activation and the abnormal nuclear morphology in NGF-differentiated PC-12 cells. Similarly to BDS-I, we found that BDS-I[1-8] blocking  $K_V3.4$  currents prevented  $A\beta_{1-42}$ -induced caspase-3 activation and apoptotic processes. Collectively, these results suggest that BDS-I[1-8] could represent a lead compound to be developed as a new drug targeting  $K_V3.4$  channels.

**Keywords:** BDS-I, voltage gated potassium channel,  $K_V3.4$ , Alzheimer's disease,  $A\beta$  peptide

## INTRODUCTION

Voltage-gated potassium ( $K_V$ ) channels are transmembrane proteins with strong selectivity for  $K^+$  ions and sensitivity to voltage changes. In neuronal cells,  $K_V$  channels are key regulators of membrane excitability, resting membrane potentials and spontaneous firing rate, action potential waveform and duration, neurotransmitter release, and apoptosis (Rudy et al., 1999).

Several neuronal populations in the central nervous system (CNS) are required to generate action potentials with high frequency (Coetzee et al., 1999). K<sub>v</sub> channels of the K<sub>v</sub>3 family are prominently expressed in these neurons and have a central role in facilitating sustained and/or repetitive high frequency firing (Moreno et al., 2001). The subunits of the K<sub>v</sub>3 *Shaw*-related subfamily (K<sub>v</sub>3.1-K<sub>v</sub>3.4), assembled into homotetramers with identical subunits or heterotetramers with different subunits, display unique biophysical properties such as high thresholds of activation, rapid activation and deactivation kinetics and relatively large conductance (Coetzee et al., 1999; Rudy et al., 1999). In rodents, three of four known K<sub>v</sub>3 genes (K<sub>v</sub>3.1-K<sub>v</sub>3.3) are conspicuously expressed in the CNS, whereas K<sub>v</sub>3.4 transcripts are abundant in skeletal muscle, sympathetic neurons, and weakly expressed in a few neuronal types in the brain in physiological conditions (Heinemann et al., 1996; Rudy et al., 1999). On the other hand, many evidence have been provided revealing the increase of K<sub>v</sub>3.4 protein expression in several pathological conditions such as hypoxia (Kaab et al., 2005), oxidative stress (Song et al., 2017), and neurodegeneration (Baranauskas et al., 2003; Angulo et al., 2004; Boscia et al., 2017).

Moreover, among the K<sub>v</sub>3 subfamily, only the K<sub>v</sub>3.4 channel carries the fast inactivating potassium currents I<sub>A</sub> and exhibits peculiar pharmacological characteristics. In fact, similarly to the other K<sub>v</sub>3 channels, K<sub>v</sub>3.4-mediated currents are highly sensitive to external tetraethylammonium or 4-aminopyridine and, in addition, a selective responsiveness of K<sub>v</sub>3.4 to some toxins have been reported. In this regard, Diochot et al. reported that two 43-amino acid peptides from the sea anemone *Anemonia sulcata*, the blood depressing substance (BDS)-I and BDS-II, show a specific blocking activity against the I<sub>A</sub> mediated by this channel (Diochot et al., 1998). These toxins directly inhibit K<sub>v</sub>3.4 channels in a reversible manner, with a similar IC<sub>50</sub> (Diochot et al., 1998), despite a difference in the time-dependence of their effect has been suggested (Diochot et al., 1998; Song et al., 2017). Another aspect that deserves a mention is that the drug sensitivity of K<sub>v</sub>3.4 pore-forming  $\alpha$  subunit, as well as its gating kinetics, conductance and ion selectivity, is modulated by the interaction with its accessory  $\beta$  subunit, the MinK-related peptide 2 (MiRP2) (Abbott et al., 2001; McCrossan and Abbott, 2004).

The I<sub>A</sub> currents mediated by K<sub>v</sub>3.4 channel subunits have been recognized as a relevant player in Alzheimer's disease (AD) and are now emerging as a new target candidate for AD (Angulo et al., 2004; Pannaccione et al., 2005, 2007; Boda et al., 2012; Boscia et al., 2017). In previous studies, we demonstrated that  $\beta$ -amyloid peptide 1-42 (A $\beta$ <sub>1-42</sub>) alters the properties of K<sup>+</sup> currents in primary hippocampal neurons (Pannaccione et al., 2005) through the selective up-regulation of K<sub>v</sub>3.4 channels mediated by the activation of the transcriptional factor NF- $\kappa$ B (Pannaccione et al., 2007). Moreover, we also reported an increase in cell death caused by the reduction of cytoplasmic K<sup>+</sup> concentrations, because of the enhanced expression and function of this K<sup>+</sup> channel upon A $\beta$ <sub>1-42</sub> exposure (Pannaccione et al., 2007). In addition, it has been demonstrated the correlation between altered intracellular K<sup>+</sup> concentrations and apoptotic processes (Yu, 2003). Interestingly, the concept that K<sub>v</sub>3.4 channels are involved in the A $\beta$ <sub>1-42</sub>

neurotoxic effects (Abbott et al., 2001) was further supported by our results showing that BDS-I treatment, by blocking the K<sub>v</sub>3.4 channel, prevents the apoptotic cascade triggered by A $\beta$ <sub>1-42</sub> fragment in hippocampal neurons, thus exerting a potent neuroprotective action (Pannaccione et al., 2007). Importantly, these previous results identified the K<sub>v</sub>3.4 channel as a new molecular target, thus revealing a possible new strategy in the scenario of pharmacological therapies against AD progression.

For these reasons, the main goal of the present study has been to identify the smallest BDS-I amino acid sequence able to exert a neuroprotective effect by blocking K<sub>v</sub>3.4 activity and to provide a new opportunity in the development of potential drugs for AD treatment.

## MATERIALS AND METHODS

### Materials

Fmoc-protected amino acids, Rink amide resin, and coupling reagents for SPPS compatibility, all were purchased from GL Biochem (Shanghai, China). Common solvents for SPPS and HPLC purification and characterization were obtained from VWR (Milano, Italy), whereas trifluoroacetic acid (TFA) was purchased from Iris Biotech GmbH (Marktredwitz, Germany). Other unmentioned materials were from Sigma Chemicals (St. Louis, MO, USA). Rabbit polyclonal anti-K<sub>v</sub>3.4, rabbit polyclonal anti-MiRP2, NGF 2.5S, TTX, nimodipine were from Alomone Labs (Jerusalem, Israel). Rabbit polyclonal anti-cleaved caspase-3 and rabbit monoclonal anti-A $\beta$  were purchased from Cell Signaling (Massachusetts, USA). DMEM, FBS, non-essential amino acids, penicillin, streptomycin, and PBS were from Gibco-BRL (Grand Island, NY, USA). Lipofectamine 2000 was from Invitrogen Corp (Carlsbad, CA, USA). The A $\beta$ <sub>1-42</sub> peptide was synthesized by INBIOS (Pozzuoli, Naples, Italy). The BDS-I

**TABLE 1** | Peptide library on BDS-I sequence. BDS-I[1-8], BDS-I[7-14], BDS-I[13-20], BDS-I[19-26], BDS-I[25-32], BDS-I[31-38], BDS-I[37-43], and BDS-I[1-8scr] peptides were designed following the mimotopic strategy.

Code	Sequence
BDS-I[1-43]	AAPCFCSGKPGRGDLWILRGTCPPGGYGYTSNCKYKWPNICCYPH
*BDS-I[1-43]	AAPAFASGKPGRGDLWILRGTPGGYGYTSNAYKWPNIAYPH
BDS-I[1-8]	H-Ala-Ala-Pro-Ala-Phe-Ala-Ser-Gly-OH
BDS-I[7-14]	H-Ser-Gly-Lys-Pro-Gly-Arg-Gly-Asp-OH
BDS-I[13-20]	H-Gly-Asp-Leu-Trp-Ile-Leu-Arg-Gly-OH
BDS-I[19-26]	H-Arg-Gly-Thr-Ala-Pro-Gly-Gly-Tyr-OH
BDS-I[25-32]	H-Gly-Tyr-Gly-Tyr-Thr-Ser-Asn-Ala-OH
BDS-I[31-38]	H-Asn-Ala-Tyr-Lys-Trp-Pro-Asn-Ile-OH
BDS-I[37-43]	H-Asn-Ile-Ala-Ala-Tyr-Pro-His-OH
BDS-I[1-8scr]	H-Gly-Ser-Ala-Phe-Ala-Pro-Ala-Ala-OH

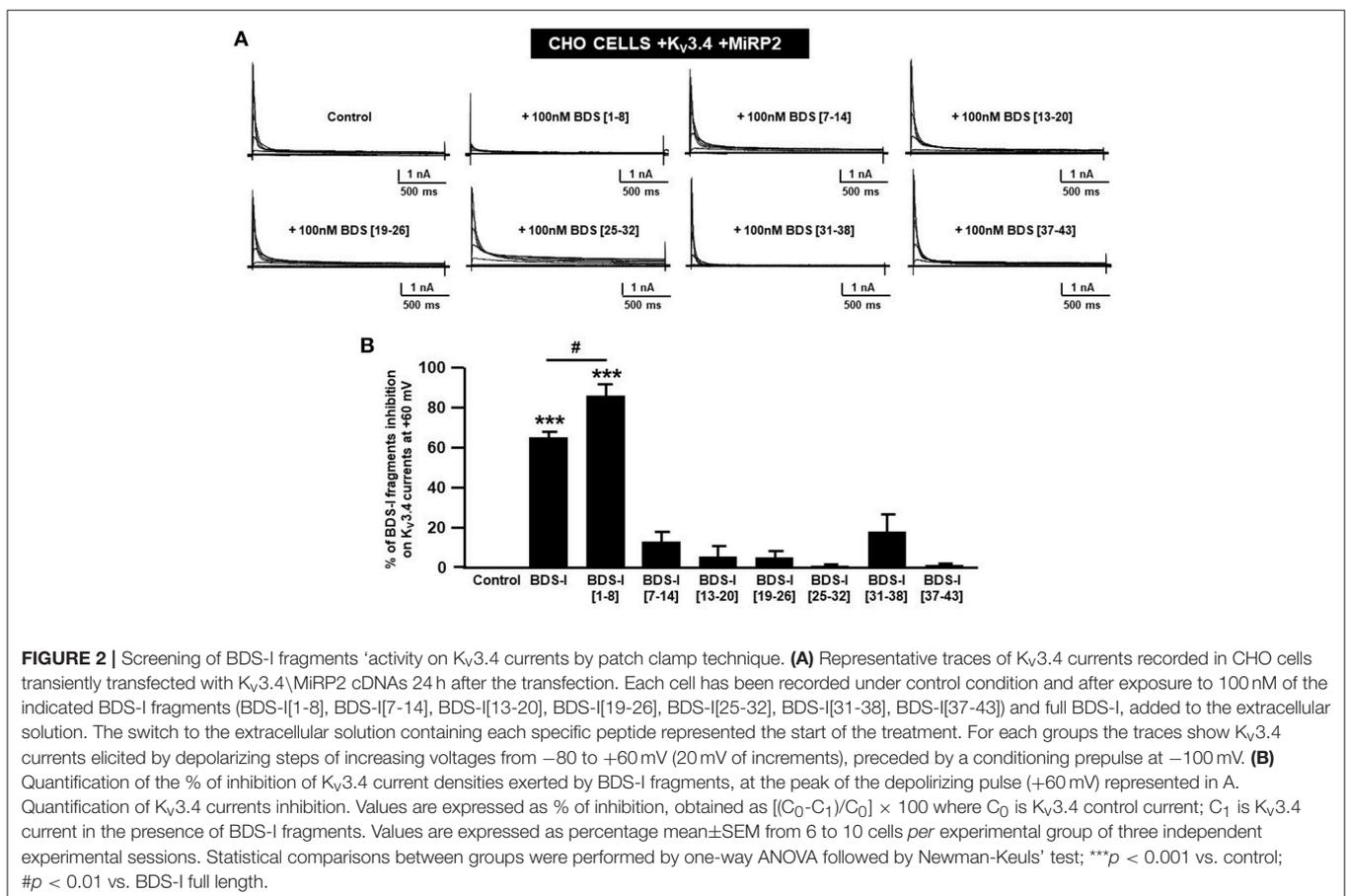
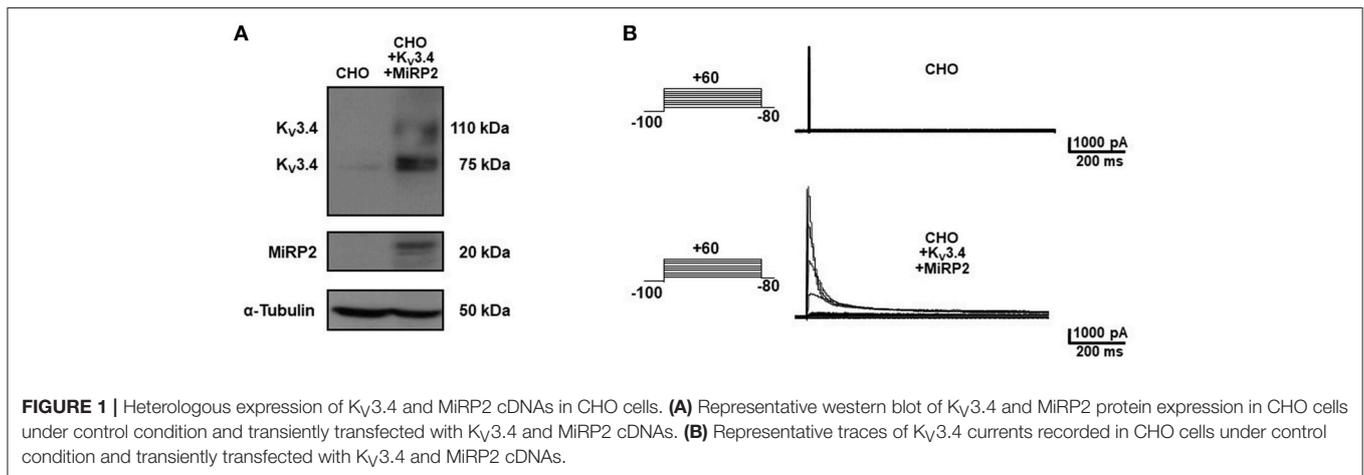
The synthesis was based on BDS-I full length sequence replacing the cysteine in position 4, 6, 22, 32, 39, and 40 by alanine order to prevent the formation of disulphide bonds and to obtain unfolded peptides. We obtained a small library characterized by overlapping peptides of equal length, containing 8 amino acids, except for the sequence BDS-I[37-43]. \*BDS-I[1-43] with cysteine  $\rightarrow$  lanine substitution.

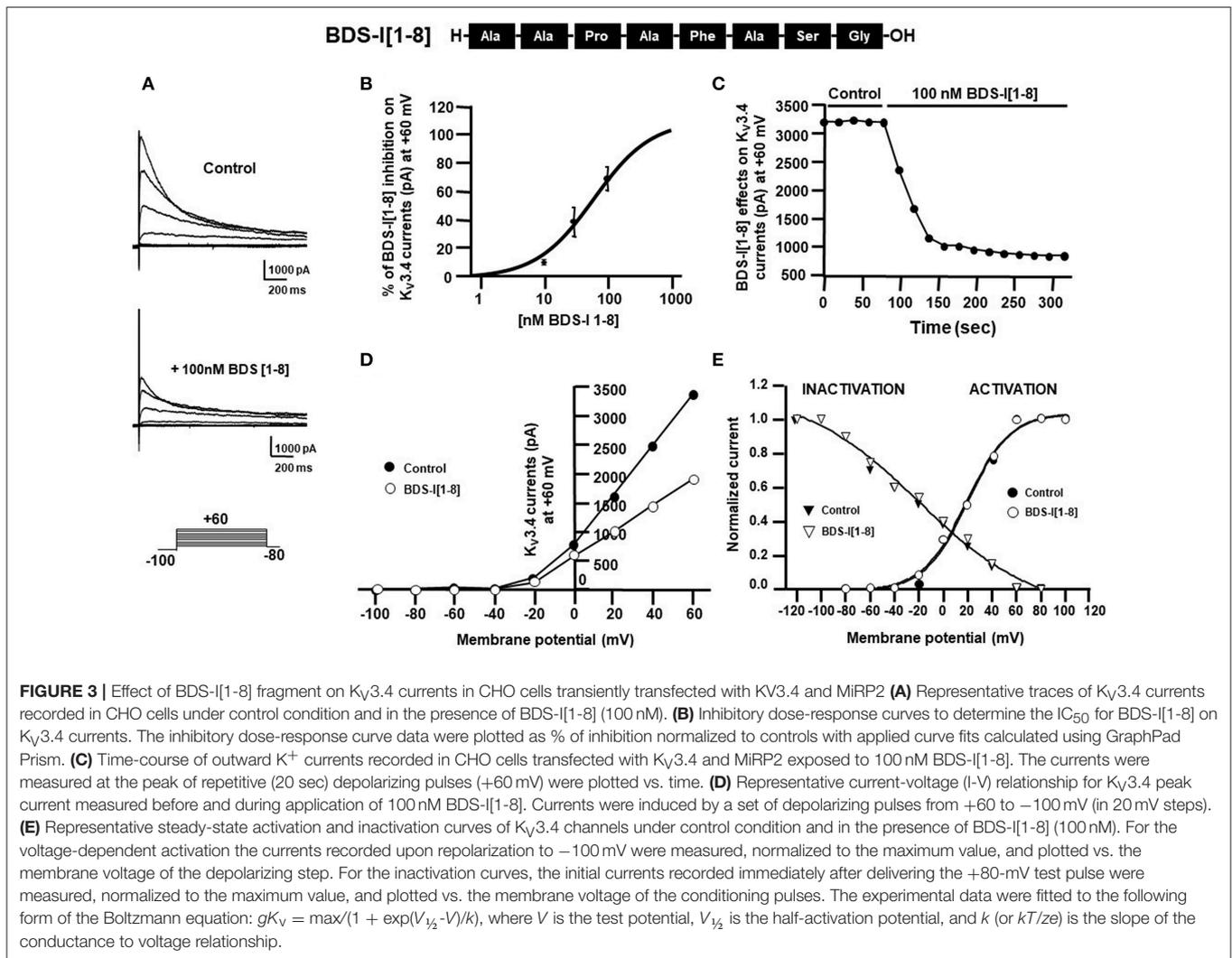
toxin and BDS-I fragments, BDS-I[1-8], BDS-I[1-8scr], BDS-I[7-14], BDS-I[13-20], BDS-I[19-26], BDS-I[25-32], BDS-I[31-38], BDS-I[37-44], were synthesized by Prof Paolo Grieco from Dept. of Pharmacy, University of Naples "Federico II," Naples, Italy. Non-fat dry milk and precast gels 4-20% were from Bio-Rad Laboratories (Milan, Italy). Protease Inhibitor Cocktail II was from Roche Diagnostic (Monza, Italy). Glass coverslips were from Carolina Biological Supply Company

(Burlington, NC, USA). The nitrocellulose membranes and ECL Western Detection Kit were from Amersham Bioscience (Buckinghamshire, UK).

## Heterologous Expression of K<sub>v</sub>3.4 and MiRP2 cDNAs

K<sub>v</sub>3.4 and MiRP2 channel subunits were expressed in Chinese hamster ovary (CHO) cells by transient transfection. CHO





cells were grown in 100 mm plastic Petri dishes in Dulbecco's Modified Eagle Medium containing 10% fetal bovine serum (FBS), non-essential amino acids (0.1 mM), penicillin (50 U/ml), and streptomycin (50  $\mu$ g/ml) in a humidified atmosphere at 37°C with 5%  $CO_2$  (Miceli et al., 2015).  $K_v3.4$  and  $MiRP2$  cDNAs were kindly provided by the Professor Goldstein of the Department of Pediatrics and Institute for Molecular Pediatric Sciences, University of Chicago, USA. For electrophysiological experiments, cells were seeded on glass cover-slips (Glaswarenfabrik Karl Hecht KG, Sondheim, Germany) coated with poly-L-lysine (50  $\mu$ g/ml) and transfected the next day with the appropriate cDNAs using Lipofectamine 2000 according to guidelines for transfection. We performed electrophysiological experiments 48 h after transfection.

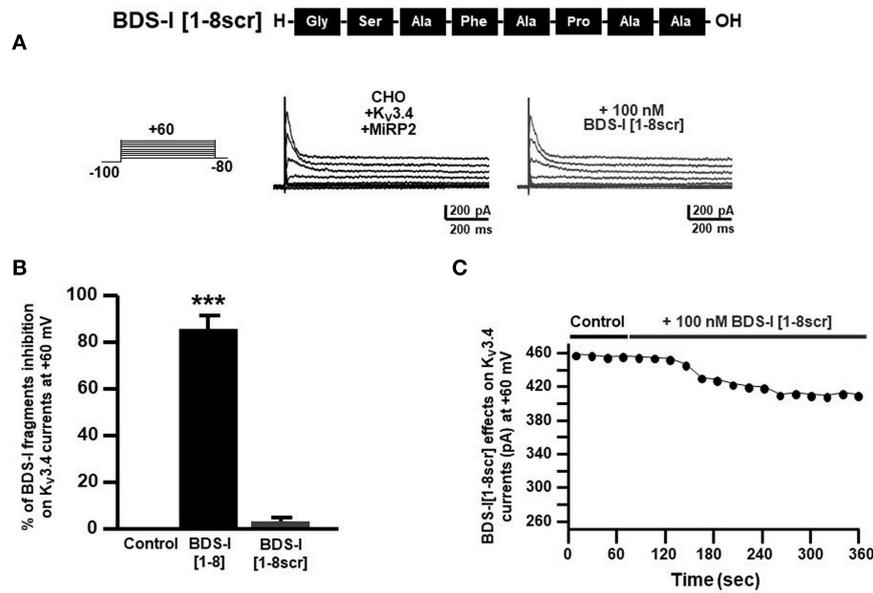
## Cell Culture

Rat pheochromocytoma cells (PC-12 cells) were grown as previously described (Pannaccione et al., 2005, 2007). For all the experiments, cells were seeded at low density on glass cover-slips coated with poly-L-lysine (50  $\mu$ g/ml). Differentiation of PC-12 cells was achieved by treatment with neuronal growth factor

(NGF) 2.5 S (50 ng/ml) for 7–9 days (Greene and Tischler, 1976). For electrophysiological experiments, cells were seeded on glass cover-slips and used after 7–9 days of NGF.

## Whole-Cell Electrophysiology

Currents from CHO cells were recorded at room temperature 1–2 days after transfection. The extracellular solution contained the following (in mM): 138 NaCl, 5.4 KCl, 2  $CaCl_2$ , 1  $MgCl_2$ , 10 glucose and 10 HEPES, adjusted pH 7.4 with NaOH. The pipette solution contained (in mM) the following: 140 KCl, 2  $MgCl_2$ , 10 EGTA, 10 HEPES, 5 Mg-ATP, adjusted pH 7.4 with KOH (Miceli et al., 2015).  $K_v3.4$  currents were measured by applying voltage steps from -100 mV to +60 mV from a holding potential of -80 mV (Abbott et al., 2001). Moreover, the total outward  $K^+$  currents were recorded in NGF-differentiated PC-12 cells using the extracellular solution contained the following (in mM): 150 NaCl, 5.4 KCl, 3  $CaCl_2$ , 1  $MgCl_2$ , 10 HEPES, adjusted pH 7.4 with NaOH. The pipette solution contained the following (in mM): 140 KCl, 2  $MgCl_2$ , 10 HEPES, 10 glucose, 10 EGTA, and 1 Mg-ATP adjusted at pH 7.4 with KOH (Diocot et al., 1998). All experiments were performed in the presence of 50 nM tetrodotoxin (TTX) and 10  $\mu$  M nimodipine in the extracellular



**FIGURE 4 |** Effect of BDS-I[1-8scr] fragment on Kv3.4 currents in CHO cells transfected with Kv3.4 and MiRP2. **(A)** Representative traces of Kv3.4 currents recorded in CHO cells under control condition and in the presence of BDS-I[1-8scr] 100 nM. Each cell has been recorded under control condition and after exposure to 100 nM of BDS-I[1-8scr] added to the extracellular solution. The switch to the extracellular solution containing each specific peptide represented the start of the treatment. For each groups the traces show Kv3.4 currents elicited by depolarizing steps of increasing voltages from  $-80$  to  $+60$  mV (20 mV of increments), preceded by a conditioning prepulse at  $-100$  mV. **(B)** Quantification of the % of inhibition of Kv3.4 current densities exerted by BDS-I fragments, at  $+60$  mV represented in A. Quantification of Kv3.4 current inhibition. Values are expressed as % of inhibition, obtained as  $[(C_0 - C_1)/C_0] \times 100$  where  $C_0$  is Kv3.4 control current;  $C_1$  is Kv3.4 current in the presence of BDS-I[1-8scr]. Values are expressed as percentage mean  $\pm$  SEM from 6 to 10 cells per experimental group of 3 independent experimental sessions. Statistical comparisons between groups were performed by one-way ANOVA followed by Newman-Keuls' test;  $***p < 0.001$  vs. control. **(C)** Time-course of outward K<sup>+</sup> currents recorded in CHO cells transfected with Kv3.4 and MiRP2 exposed to 100 nM BDS-I[1-8scr]. The currents were measured at the peak of repetitive (20 s) depolarizing pulses ( $+60$  mV) were plotted vs. time.

solution. To discriminate K<sup>+</sup> current components appropriate electrophysiological protocols were used (Pannaccione et al., 2007). Steady-state activation and inactivation properties were measured by using appropriate protocols (Pannaccione et al., 2005). Capacitive currents were elicited by 5-mV depolarizing pulses from  $-80$  mV and acquired at a sampling rate of 50 kHz. The capacitance of the membrane was calculated according to the following equation:  $C_m = \tau_c \cdot I_0 / \Delta E_m (1 - I_\infty / I_0)$ , where  $C_m$  is membrane capacitance,  $\tau_c$  is the time constant of the membrane capacitance,  $I_0$  is the maximum capacitance current value,  $\Delta E_m$  is the amplitude of the voltage step, and  $I_\infty$  is the amplitude of the steady-state current. K<sup>+</sup> currents were recorded by patch-clamp technique in CHO and NGF-differentiated PC-12 cells using a commercially available amplifier (Axopatch 200B, Molecular Devices) and Digidata 1322A interface (Molecular Devices, USA). The whole-cell configuration of the patch-clamp technique was adopted using glass micropipettes with a resistance of 2.5–5.0 M $\Omega$ . Data were acquired and analyzed using the pClamp software (version 9, Molecular Devices).

## BDS-I Fragments Synthesis

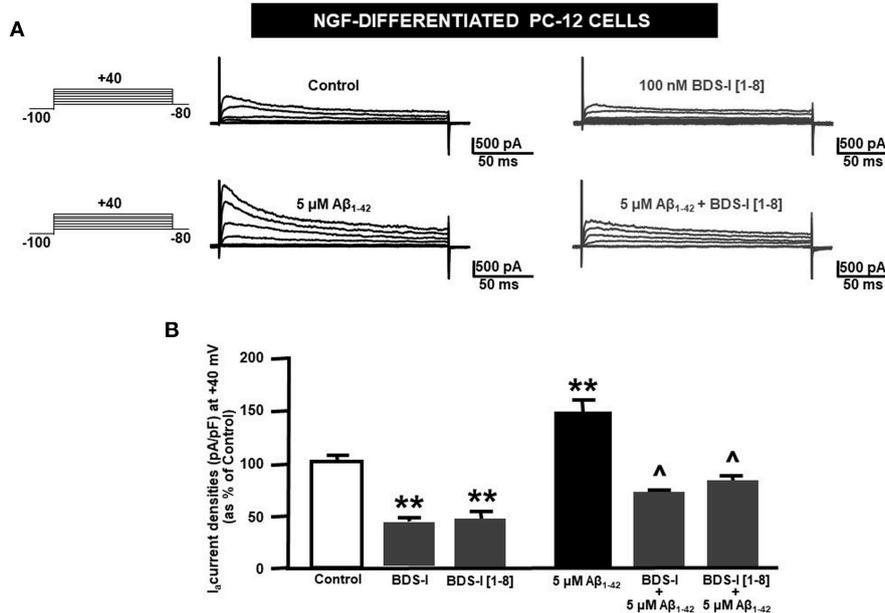
Peptides BDS-I[1-8], BDS-I[13-20], BDS-I[19-26], BDS-I[25-32], BDS-I[31-38], BDS-I[37-43], and BDS-I[1-8scr] were designed following the mimotopic strategy (Table 1). All compounds were synthesized manually by adopting a standard solid-phase

peptide synthesis (SPPS) using the Fmoc/tBu orthogonal strategy (Merrifield, 1963; Carpino and Han, 1970; Merlino et al., 2016). The use of the Wang resin 0.7 mmol/g (GL Biochem, Shanghai, China) as solid support in the entire synthesis resulted in amidated C-terminal peptides after TFA-mediated cleavage (TFA/TIS/H<sub>2</sub>O, 95:2.5:2.5, 3 h). The crude peptides were purified by reverse-phase HPLC (RP-HPLC, Shimadzu Nexera) on a preparative column (PhenomenexKinetex, C18 column, 150  $\times$  21.2 mm, 5  $\mu$ m, 100  $\text{\AA}$ ) using a linear gradient of acetonitrile (0.1% TFA) in water (0.1% TFA), from 10 to 90% over 20 min, with a flow rate of 10.0 mL/min and UV detection at 220 nm. The purity (>98%) was ascertained by analytical RP-HPLC on a PhenomenexKinetex C18 column (150  $\times$  4.6 mm, 5  $\mu$ m, 100  $\text{\AA}$ ) and molecular weights of all peptides were confirmed by LC-ESI MS system (Agilent Technology).

BDS-I[1-8]: AAPAFASG-OH, (3.2 mg, yield 31%, purity >98%), observed mass 690.75 (M+H)<sup>1+</sup>, calculated mass 691.32.

BDS-I[7-14]: GKPRGDL-OH, (2.8 mg, yield 27%, purity >98%), observed mass 798.89 (M+H)<sup>1+</sup>, calculated mass 799.03.

BDS-I[13-20]: GDLWILRG-OH, (4.2 mg, yield 34%, purity >98%), observed mass 929.08 (M+H)<sup>1+</sup>, calculated mass 929.76.



**FIGURE 5 |** Effect of BDS-I[1-8] fragment on  $K_v3.4$  currents in NGF-differentiated PC-12 cells under control condition and exposed to  $A\beta_{1-42}$ . **(A)** Representative traces of  $K_v3.4$  currents recorded in NGF-differentiated PC-12 cells under control condition and exposed to 100 nM of BDS-I[1-8] (top); representative traces of  $K_v3.4$  currents recorded in NGF-differentiated PC-12 cells exposed to  $A\beta_{1-42}$  oligomers (5  $\mu$ M for 24 h) in the absence and in the presence of BDS-I[1-8] fragment (30 min of pre-treatment before adding  $A\beta_{1-42}$  oligomers) (bottom). For each group, traces show  $K_v3.4$  currents elicited by depolarizing steps of increasing voltages from  $-80$  to  $+40$  mV (20 mV of increments), preceded by a conditioning prepulse at  $-100$  mV. **(B)** Quantification of  $K_v3.4$  current densities, at the peak of the depolarizing pulse ( $+40$  mV) represented in A. Values are expressed as percentage mean  $\pm$  SEM from 6 to 10 cells *per* experimental group of 3 independent experimental sessions. Statistical comparisons between groups were performed by one-way ANOVA followed by Newman-Keuls' test; \*\* $p < 0.01$  vs. control;  $\wedge p < 0.01$  vs.  $A\beta_{1-42}$ .

BDS-I[19-26]: RGTAPGGY-OH, (3.4 mg, yield 25%, purity >98%), observed mass 777.83 (M+H) $^+1$ , calculated mass 778.21.

BDS-I[25-32]: GYGYSNAY-OH, (5.7 mg, yield 38%, purity >98%), observed mass 995.00 (M+H) $^+1$ , calculated mass 995.67.

BDS-I[31-38]: NAYKWPNI-OH, (4.6 mg, yield 30%, purity >98%), observed mass 1005.13 (M+H) $^+1$ , calculated mass 1005.90.

BDS-I[37-43]: NIAAYPH-OH, (5.1 mg, yield 41%, purity >98%), observed mass 784.86 (M+H) $^+1$ , calculated mass 784.97.

BDS-I[1-8scr]: GSAFAPAA-OH, (2.4 mg, yield 43%, purity >98%), observed mass 690.77 (M+H) $^+1$ , calculated mass 691.32.

## BDS-I Fragment Application

All BDS-I fragments, BDS-I[1-8], BDS-I[7-14], BDS-I[13-20], BDS-I[19-26], BDS-I[25-32], BDS-I[31-38], BDS-I[37-43] as well as the negative control peptide BDS-I[1-8scr] have been applied by acute treatment.

For the acute treatment, each peptide has been added in the extracellular solution. For each cell, we first recorded  $K^+$  currents as previously described, in the absence of peptides. After that, we switched from the extracellular solution without BDS-I peptides to the extracellular solution containing the BDS-I peptide. In these conditions, we

recorded  $K^+$  currents until the steady-state was reached. This protocol has been performed to test the effect on  $K_v3.4$  currents of each BDS-I fragment as well as of BDS-I[1-8scr].

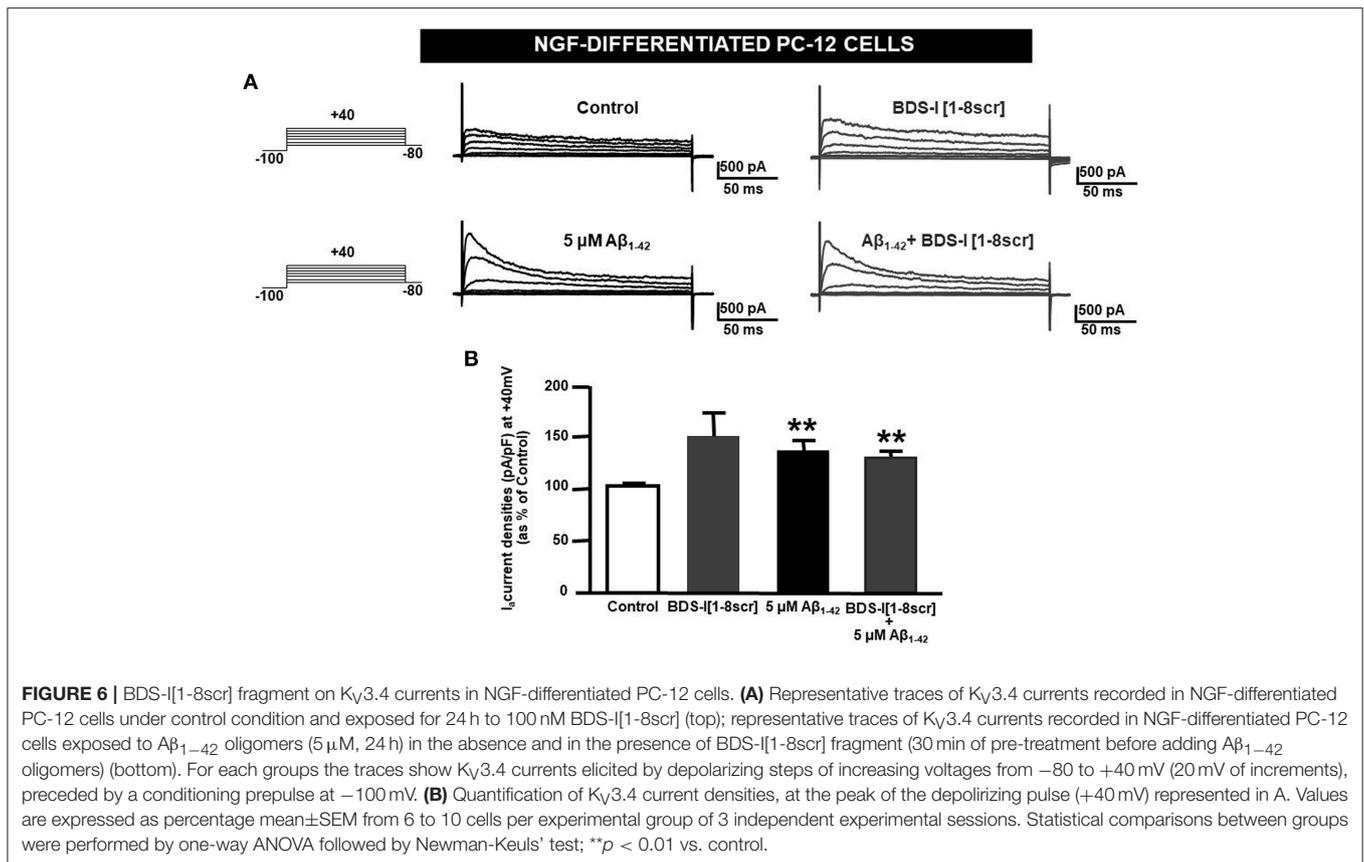
To evaluate the effect of the active fragments in the presence of  $A\beta_{1-42}$  oligomers, we pre-treated cells by adding the fragment in the culture medium at the final concentration of 100 nM, 30 min before exposing cells to  $A\beta_{1-42}$  oligomers and lasted for 24 h.

## Assessment of Nuclear Morphology

Nuclear morphology was evaluated by using the fluorescent DNA-binding dye Hoechst-33258 already described (Pannaccione et al., 2007).

## Western-Blot Analysis

NGF-differentiated PC-12 and CHO cells were washed in phosphate buffered saline (PBS) and collected by gentle scraping in ice-cold RIPA buffer containing in mM: 50 Tris pH 7.4, 100 NaCl, 1 EGTA, 1 PMSF, 1 sodium orthovanadate, 1 NaF, 0.5% NP-40, and 0.2% SDS supplemented with protease inhibitor cocktail II. After sonication and incubation for 1 h on ice, we centrifuged at 12,000 rpm at 4°C for 30 min and collected the supernatants. The protein content of resulting supernatant was determined using the Bradford reagent. Hundred microgram of proteins were mixed with a Laemmli sample buffer; then, they are applied and resolved on SDS-PAGE polyacrylamide gels.



Following transfer onto nitrocellulose membranes, non-specific binding sites were blocked by incubation for 2 h at  $4^{\circ}\text{C}$  with 5% non-fat dry milk in TBS-T buffer; subsequently, incubated with primary antibodies overnight at  $4^{\circ}\text{C}$ . After three washes with TBS-T, the membranes were incubated 1 h with the appropriate secondary antibody. Excessive antibodies were then washed away three times (10 min) with TBS-T. Proteins were visualized with peroxidase-conjugated secondary antibodies, using the enhanced chemiluminescence system. The software Image J (NIH, Bethesda, MA, USA) was used for densitometric analysis. Primary antibodies were: rabbit polyclonal anti-K<sub>v</sub>3.4 (1:1,000), rabbit polyclonal anti-MiRP2 (1:1,000), rabbit polyclonal anti-cleaved caspase-3 (1:1,000) and mouse monoclonal anti- $\alpha$ -Tubulin (1:3,000).

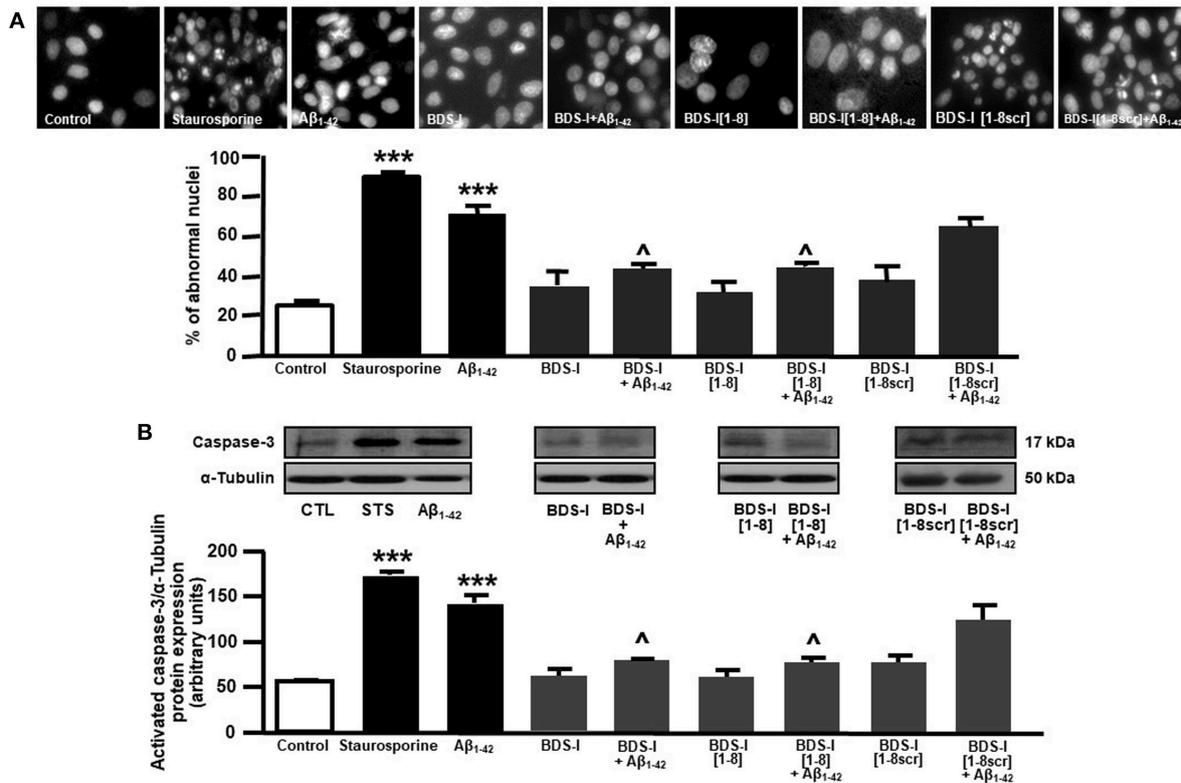
## A $\beta$ Treatments

The peptide used in our study was synthesized by INBIO (Pozzuoli, Naples, Italy) using the A $\beta_{1-42}$  sequence of human APP [UniProtKB-P05067 (A4\_HUMAN)]. The purity of the peptide was assessed by high-performance liquid chromatography ( $>95\%$  pure) and the amino acid composition was verified by mass spectrometry. The A $\beta$  oligomers was resuspended in 1,1,1,3,3,3-hexafluoro-2-propanol (HFIP; Sigma, Milan, Italy), at a final concentration of 1 mM. Then, the preparation was aliquoted, microcentrifuged, dried by

SpeedVac and stored at  $-20^{\circ}\text{C}$ . Before their use, A $\beta$  oligomer aliquots were first resuspended in dimethyl sulfoxide (DMSO) at a final concentration of 5 mM, and then diluted in ice-cold cell culture medium (phenol red-free Ham's F-12, Sigma, Milan, Italy), to the final concentration of 100  $\mu\text{M}$ . Next, after vortexing for 30 sec, the resuspension was incubated for 24 h at  $4^{\circ}\text{C}$ . Finally, the solution was centrifuged at 14,000 rpm at  $4^{\circ}\text{C}$  for 10 min, and the supernatant, containing A $\beta_{1-42}$  oligomers, aliquoted and stored at  $-20^{\circ}\text{C}$  (Stine et al., 2003). A $\beta_{1-42}$  were added to the culture medium at the final concentration of 5  $\mu\text{M}$  for 24 h. The pre-aggregated preparation of A $\beta$  oligomers was analyzed in SDS-PAGE using the rabbit monoclonal anti-A $\beta$  on precast gels 4-20%. Oligomers between 4 and 15 kDa were the major species of the A $\beta$  oligomers in the preparation.

## Data and Statistical Analysis

GraphPad Prism 6.02 was used for statistical analyses (GraphPad Software, La Jolla, CA). The data are expressed as the mean  $\pm$  S.E.M. of the values obtained from individual experiments. Statistical comparisons between groups were performed by Student's  $t$ -test or one-way analysis of variance (ANOVA) followed by Newman-Keuls' test;  $p < 0.05$  was considered significant.



**FIGURE 7** | Effect of BDS-I[1-8] on A $\beta_{1-42}$ -induced abnormal nuclear morphology and caspase-3 activation in NGF-differentiated PC-12 cells. **(A)** Assessment of nuclear morphology with Hoechst-33258 in NGF-differentiated PC-12 cells under control condition and exposed to A $\beta_{1-42}$  oligomers (5  $\mu$ M, 24 h) in the absence and in the presence of BDS-I[1-8] fragment, BDS-I[1-8scr] fragment, BDS-I full length (3 h of pre-treatment before adding A $\beta_{1-42}$  oligomers) (top). Quantification of the percentage of abnormal nuclei in NGF-differentiated PC-12 cells under control condition and exposed to A $\beta_{1-42}$  oligomers (5  $\mu$ M, 24 h) in the absence and in the presence of BDS-I[1-8] fragment, BDS-I[1-8scr] fragment, BDS-I full length (30 min of pre-treatment before adding A $\beta_{1-42}$  oligomers) obtained in 3 separate experiments in which at least 10 microscopic fields were analyzed (~1,000 cells per group) (Bottom). Statistical comparisons between groups were performed by one-way ANOVA followed by Newman-Keuls' test; \*\*\* $p$  < 0.001 vs. control;  $\Delta$   $p$  < 0.01 vs. A $\beta_{1-42}$ . **(B)** Representative western blot (top) and densitometric quantification (bottom) of caspase-3 protein expression in NGF-differentiated PC-12 cells under control condition and exposed to A $\beta_{1-42}$  oligomers (5  $\mu$ M, 24 h) in the absence and in the presence of BDS-I[1-8] fragment, BDS-I[1-8scr] fragment, BDS-I full length (30 min of pre-treatment before adding A $\beta_{1-42}$  oligomers). Values are expressed as mean  $\pm$  SEM of 3 independent experimental sessions. Statistical comparisons between groups were performed by one-way ANOVA followed by Newman-Keuls' test; \*\*\* $p$  < 0.001 vs. control;  $\Delta$   $p$  < 0.01 vs. A $\beta_{1-42}$ .

## RESULTS

### Synthesis of BDS-I Fragments

To identify the key amino acidic residues of BDS-I essential for its inhibitory action on  $K_v3.4$  channels, we performed an overlapping peptide library on BDS-I sequence (Table 1). This small library is characterized by overlapping peptides of equal length, containing 8 amino acids, except for the sequence BDS-I[37-43]. Moreover, the cysteine in position 4, 6, 22, 32, 39, and 40 in the sequence of BDS-I[1-43], presumably involved in the tertiary structure of BDS-I, have been replaced by alanine in order to prevent the formation of disulphide bonds and to obtain unfolded peptides. This approach allowed us to focus on the importance of the amino acid sequence rather than the tertiary structure of BDS-I fragments.

### Screening of BDS-I Fragments 'Activity on $K_v3.4$ Currents by Patch Clamp Technique

To identify the smallest BDS-I amino acid sequence able to block  $K_v3.4$  activity, we evaluated the effect of the newly synthesized BDS-I fragments in CHO cells transiently transfected with  $K_v3.4$ /MiRP2, by patch clamp technique in whole cell configuration.  $K_v3.4$  expression and activity in transfected CHO cells were monitored by western blot analysis and electrophysiological experiments, respectively (Figures 1A,B). After acute application of the peptides, we found that only BDS-I[1-8] fragment (100 nM), containing the N-terminal octapeptide sequence of full length BDS-I, similarly to the full length, was able to inhibit  $K_v3.4$  currents (Figures 2A,B), whereas the other BDS-I fragments, BDS-I[7-14], BDS-I[13-20], BDS-I[19-26], BDS-I[25-32], BDS-I[31-38], and BDS-I[37-44] (100 nM) did not

(Figures 2A,B, 3A). The percentage of inhibition of  $K_V3.4$  currents exerted by BDS-I[1-8] fragment (100 nM) was more than 80%. Interestingly, BDS-I, displaying a percentage of inhibition of ~60%, appeared less effective than BDS-I[1-8] fragment (Figure 2B). Moreover, we found that BDS-I[1-8] reduced  $K_V3.4$  currents in a concentration-dependent manner, with an  $IC_{50}$  value of 75 nM (Figure 3B). The time-course of the effect of BDS-I[1-8] on  $K_V3.4$  currents showed that the steady-state of inhibition was reached at 150 s of peptide exposure (Figure 3C). Inhibition of the  $K_V3.4$  current induced by BDS-I[1-8] was not voltage-dependent as evidenced by I-V relationship for  $K_V3.4$  peak current measured before and during application of 100 nM BDS-I[1-8] (Figure 3D). Furthermore, BDS-I[1-8] treatment did not modify the steady-state parameters for the activation and inactivation of  $K_V3.4$  currents (Figure 3E).

### Effect of BDS-I[1-8] Scrambled Peptide on $K_V3.4$ Currents

To confirm that the inhibitory effect of BDS-I[1-8] on  $K_V3.4$  currents was sequence-specific, we synthesized a peptide with the scrambled sequence of BDS-I[1-8], named BDS-I[1-8scr]. Electrophysiological experiments performed in CHO cells transiently transfected with  $K_V3.4$ /MiRP2 showed that BDS-I[1-8scr] at the same concentration of 100 nM was not able to modify  $K_V3.4$  activity (Figures 4A,B). The time-course of current inhibition during BDS-I[1-8scr] exposure is presented in Figure 4C.

### Effect of BDS-I[1-8] on $A\beta_{1-42}$ -Induced Upregulation of $K_V3.4$ Activity and Caspase-3 Activation in NGF-Differentiated PC-12 Cells

Analogously to the results obtained in CHO cells, BDS-I[1-8] fragment blocked  $K_V3.4$  activity in NGF-differentiated PC-12 cells (Figures 5A,B). Importantly, like BDS-I, BDS-I[1-8] fragment prevented the  $K_V3.4$  upregulation elicited by  $A\beta_{1-42}$  exposure (5  $\mu$ M for 24 h) (Figures 5A,B). By contrast, BDS-I[1-8scr] did not counteract  $A\beta_{1-42}$ -induced  $K_V3.4$  upregulation (Figures 6A,B). More interestingly, nuclear morphological studies by bisbenzimidazole Hoechst staining highlighted that, similar to the full length BDS-I, BDS-I[1-8], but not BDS-I[1-8scr], was able to significantly prevent the abnormal nuclear morphology induced by  $A\beta_{1-42}$  exposure in NGF-differentiated PC-12 cells (Figures 7A,B). These results were confirmed by western blot analysis of activated caspase-3. In fact, BDS-I[1-8] prevented  $A\beta_{1-42}$ -induced caspase-3 activation whereas BDS-I[1-8scr] did not (Figure 7C).

## DISCUSSION

In this study, a novel peptide obtained from the sequence of the sea anemone toxin BDS-I has been successfully identified as a new inhibitor of the  $K_V3.4$  channel subunits. In fact, BDS-I[1-8] fragment, containing the N-terminal octapeptide sequence of full length BDS-I, was able to inhibit  $K_V3.4$  currents

in a concentration dependent manner, with an  $IC_{50}$  value of 75 nM, in CHO cells transiently transfected with  $K_V3.4$ /MiRP2. Interestingly, the other BDS-I fragments, BDS-I[7-14], BDS-I[13-20], BDS-I[19-26], BDS-I[25-32], BDS-I[31-38], BDS-I[37-44], failed to modulate  $K_V3.4$  activity, thus confirming the key role played by the N-terminal octapeptide sequence of BDS-I in its pharmacological effect on  $K_V3.4$  currents. To confirm and validate the results obtained with BDS-I[1-8], we synthesized a scramble sequence of this peptide, as reported in Table 1. Importantly, the negative control peptide, BDS-I[1-8scr], did not display any pharmacological activity, thus highlighting that the sequence of BDS-I[1-8] is essential for the inhibitory effect of BDS-I on  $K_V3.4$  activity. The BDS-I[1-8], similarly to BDS-I, was able to selectively inhibit the fast inactivating  $K^+$  currents carried by the  $K_V3.4$  also in NGF-differentiated PC-12 cells, a well-known *in vitro* model mimicking the neuronal system, whereas BDS-I[1-8scr] did not exert any effect. More interestingly, BDS-I[1-8], but not BDS-I[1-8scr], fully counteracted the  $A\beta_{1-42}$ -induced enhancement of  $K_V3.4$  activity and prevented  $A\beta_{1-42}$ -induced caspase-3 activation and abnormal nuclear morphology. Indeed, we previously demonstrated that  $A\beta_{1-42}$  peptide induces a significant increase in  $K_V3.4$  expression and activity in NGF-differentiated PC-12 cells and hippocampal neurons and that BDS-I, by blocking  $K_V3.4$  channels, prevents the apoptotic cascade triggered by  $A\beta_{1-42}$  (Pannaccione et al., 2005, 2007). The present study clearly shows that, similarly to BDS-I, BDS-I[1-8] is able to exert a neuroprotective effect by counteracting the upregulation of  $K_V3.4$  channel induced by  $A\beta_{1-42}$ .

Importantly, it has been widely demonstrated that the dysregulation of  $K_V3.4$  channel subunits contributes to neuronal and glial alterations in AD. Previous evidence has shown that the overexpression of  $K_V3.4$  channel subunits, occurring both in the early and in the advanced stages of the disease, is crucially involved in the development of the disease (Angulo et al., 2004).  $K_V3.4$  upregulation has also been related to amyloid pathology, as high levels of these subunits have been detected in amyloid plaques in post-mortem human brain tissues (Angulo et al., 2004). Accordingly, we recently demonstrated that the expression and function of  $K_V3.4$  channel subunits are precociously upregulated in cultured astrocytes exposed to  $A\beta$  oligomers and in reactive astrocytes of AD Tg2576 mice (Boscia et al., 2017). Indeed, our results suggest that the upregulation of  $K_V3.4$  channel subunits in astrocytes may be an early event in the AD brain, since we observed astrocytes accumulating  $A\beta$  and overexpressing  $K_V3.4$  subunits in cortex, hippocampus, and cerebellum of 6 month old Tg2576 mice (Boscia et al., 2017). All together, these evidence identify  $K_V3.4$  channels as a crucial player and, hence, as a possible pharmacological target in AD. On the other hand,  $K_V3.4$  has been implicated in several diseases, including cancer and cardiovascular pathologies. In particular, it has been reported that  $K_V3.4$  channels play a key role in cancer cell migration and invasion as BDS, similarly to  $K_V3.4$  silencing, was able to reduce the number of invasive cells (Song et al., 2018). Moreover,  $K_V3.4$  currents have been implicated in the proliferation of vascular smooth muscle cells, a crucial

factor underlying the unwanted remodeling of arterial walls in pathological conditions (Miguel-Velado et al., 2010). In this scenario, the identification of the key residues of BDS-I essential for the inhibition of K<sub>v</sub>3.4 currents, namely BDS-I[1-8] peptide, provides a possible therapeutic opportunity for several diseases, despite subsequent approaches are needed to test BDS-I[1-8] stability and efficacy *in vivo*. Indeed, this small peptide could be a suitable lead compound for the development of new therapies targeting K<sub>v</sub>3.4 not only in neurodegeneration and neuroinflammation, but also in cancer and other pathological conditions.

## DATA AVAILABILITY

The raw data supporting the conclusions of this manuscript will be made available by the authors, without undue reservation, to any qualified researcher.

## AUTHOR CONTRIBUTIONS

AP conceived the study. AP and RC designed the experiments and performed data analysis. PG and FM synthesized BDS-I fragments. RC and IP performed

all the experiments. RC and IP wrote the manuscript. LA, AP, and PG reviewed and edited the final draft of the manuscript.

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