



ORIGINAL ARTICLE

An *in vitro* study on probable inhibition of cerebrovascular disease by salidroside as a potent small molecule against induction of protein amyloid fibrils and cytotoxicity



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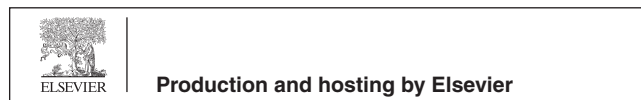
Abstract Protein aggregation and associated amyloid formation is linked with several harmful human pathophysiologies including Alzheimer's, Parkinson's, and cerebrovascular diseases. A potential approach for modulating and exploring amyloid fibrillization is the control of protein self-assembly. Herein, anti-aggregation effects of salidroside, its influence on the kinetics of amyloid fibrillization of $A\beta_{1-42}$ peptide and its cytotoxicity against cerebrovascular endothelial cells (bEnd.3) were assessed by using a wide range of spectroscopic and cellular techniques. The present outcome of Thioflavin T (ThT) and 8-anilino-1-naphthalenesulfonic acid (ANS) fluorescence, Congo red (CR), and circular dichroism (CD) analyses indicated that salidroside potentially inhibits protein fibril formation. The cellular studies inferred that salidroside protects bEnd.3 cells against $A\beta_{1-42}$ oligomers -triggered cytotoxicity through modulation of oxidative stress [reactive oxygen species (ROS), superoxide dismutase (SOD) and catalase (CAT) activities] and apoptosis (caspase-3 activity). Therefore, the data signifies the role of salidroside as a promising small molecule in inhibiting $A\beta_{1-42}$ aggregation and associated cerebrovascular endothelial cell toxicity. Hence, salidroside can serve as a potential inhibitor in the therapeutic advancement to combat cerebrovascular diseases.

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1. Introduction

Cerebrovascular dysfunction is an important contributor in the protein aggregation-stimulated neurodegenerative disorders (Kamat et al., 2015; Bos et al., 2017). During this process, the blood vessels function in the cerebral cells is downregulated (Smith and Greenberg, 2009). Indeed, several reports have indicated that amyloid plaque formation results in hypoxia-triggered inappropriate blood supply causing improper function in vascular activity (Peers et al., 2009; Zlokovic, 2011). The vascular damage influences the endothelial cells in the vascular system, and amyloid β ($A\beta$) peptide leads to cerebrovascular endothelial cytotoxicity (Thomas et al., 1997). $A\beta$ -triggered adverse effects cause downregulation of the relaxing mediators in the endothelium (Xi et al., 2012). Therefore, $A\beta$ peptide-stimulated endothelial dysfunction is heavily associated with the vasoconstriction (Smith et al., 2004). In addition, several reports have indicated that different vascular factors are associated with neurodegeneration in neurodegenerative diseases (Iadecola, 2010; Sweeney et al., 2018; Cheng et al., 2020). Structural and molecular alterations in endothelium like the formation of $A\beta$ are known as the key parameters influencing cerebrovascular dysfunction in neurodegenerative diseases (Han et al., 2008).

Structural changes of $A\beta$ with a molecular weight of around 4 kDa can cause its aggregation into amyloid fibrils which become neurotoxic (Maltsev et al., 2011). $A\beta$ peptides are one of the natural products of metabolism, which is composed of 36–43 amino acids, where $A\beta_{1-40}$ is more common than others. (Maltsev et al., 2011) $A\beta$ peptides are formed by proteolysis of the amyloid precursor protein (APP) by the activity of the secretase family (Maltsev et al., 2011). Most of the processes related to the neurotoxic influences of $A\beta$ are related to type $A\beta_{1-42}$ (Marshall et al., 2016), which causes damage to synaptic activity. APP is a membrane protein located at the end of presynaptic axons (Portelius et al., 2011). This protein has a large extracellular amine portion and a small intracellular carboxylic portion (Portelius et al., 2011; Vaillant-Beuchot et al., 2021). This part of the precursor protein is a stimulus factor for neuronal survival (Portelius et al., 2011; Vaillant-Beuchot et al., 2021).

Plant extracts and their bioactive compounds regulate the production and accumulation of $A\beta$ peptides associated with neurodegenerative diseases in the laboratory (Kwon et al., 2011). It has been shown that bioactive compounds of plant extracts can prevent the formation of amyloid structures in $A\beta$ peptides (Kumar et al., 2012; Doig and Derreumaux, 2015). The bioactive natural products have long been proven to be powerful anti-amyloid agents. Indeed, many researchers are exploring small molecules as potential inhibitors against protein aggregation *in vitro* as a starting point and the next step is to formulate these molecules in order to investigate their ability to interfere with the formation of amyloid plaque *in vivo* (Velander et al., 2017). For this reason, the bioactive compounds should cross the blood brain barrier and prevent the aggregation of $A\beta$ peptides in the brain, especially to interfere with cerebrovascular/neurodegenerative diseases-related plaques (Brahmachari et al., 2017). Together, the researchers should assess the new molecule's ability to inhibit the potential toxicity of amyloid plaques containing proteins.

Salidroside as one of the major phenolic glycosides is known as the major bioactive compound extracted from *Rhodiola crenulata*, has been demonstrated to provide promising influences on inhibiting oxidative stress and cell mortality (Cao et al., 2005; Zuo et al., 2007; Zhang et al., 2009; Ye et al., 2011; Zhu et al., 2011) and has benefit for several disorders including ischemic cardiomyocytes (Zhong et al., 2010), lung damage (Guan et al., 2012) and diabetes (Li et al., 2011). It has been also indicated that salidroside can prevent H_2O_2 or $A\beta$ -triggered oxidative damage and subsequent apoptosis in neuron-like cells (Cai et al., 2008; Jang et al., 2003) and neuroblastoma cells (Zhang et al., 2010) and hippocampal neurons (Chen et al., 2009). *In vivo* data have also shown the ability of salidroside to mitigate cerebral ischemia–reperfusion damage by its potential reactive oxygen species (ROS)-scavenging characteristics (Zou et al., 2009; Shi et al., 2012).

Although, it has been shown that salidroside can prevent the $A\beta$ -stimulated neurotoxicity, its anti-amyloid properties against $A\beta$ aggregation and associated cytotoxicity against cerebrovascular endothelial cell line (bEnd.3) have not been well explored in detail. Therefore, in this study as the main novelty of this work we aimed to evaluate the inhibitory characteristics of salidroside against $A\beta$ aggregation and relevant cytotoxicity.

2. Experiments

2.1. Sample preparation

The $A\beta_{1-42}$ was prepared as explained in literature (Wang et al., 2011; Du et al., 2015 Jan 23) For formation of $A\beta_{1-42}$ aggregation, 70 μ M of protein sample was incubated at 37 °C for 50 h under constant stirring (100 rpm) with and without 35 and 70 μ M of salidroside prepared in DMSO based on the previous study (Wang et al., 2011; Du et al., 2015; Alam et al., 2016).

2.2. ThT fluorescence spectroscopy assay

ThT fluorescence analysis was done using Shimadzu fluorescence spectrophotometer (RF-6000). ThT powder was dissolved in double distilled water and filtered. Then, the $A\beta_{1-42}$ samples co-incubated with or without salidroside, from each sample were taken out at different time intervals and well-mixed with ThT to reach final $A\beta_{1-42}$ and ThT concentration of 20 μ M. The samples were then incubated in dark for 15 min at room temperature followed by excitation at 440 nm and reading the emission spectra at 480 nm with both excitation and emission slit widths of 10 nm. The protein samples were diluted with sodium phosphate buffer (20 mM, pH 7.4) and resulting data were subtracted from associated blanks and were fitted as explained previously (Chaturvedi et al., 2015).

2.3. ANS fluorescence measurements

$A\beta_{1-42}$ samples with a final concentration of 15 μ M were mixed with ANS (15 μ M) and incubated in dark for 30 min at room temperature. ANS fluorescence intensity was then read with excitation at 380 nm and emission between 440 and 600 nm. Both excitation and emission slit width were set at 10 nm.

2.4. CR binding analysis

CR was prepared in a phosphate buffer (20 mM, pH 7.4) supplemented with NaCl (50 mM) and filtered and the final concentration was calculated employing ϵ_M 45,000 $M^{-1} cm^{-1}$ at 495 nm. CR (15 μ M) and $A\beta_{1-42}$ (15 μ M) samples were mixed and incubated in the dark for 30 min at room temperature. The CR absorbance spectra (440–650 nm) were read using a UV–Visible spectrophotometer (Perkin Elmer Lambda 25).

2.5. Far-UV CD analysis

The CD study was performed on a J-815-JASCO spectropolarimeter, where the analysis was performed with $A\beta_{1-42}$. Signals were read in the range of 260–190 nm using a cuvette with 0.1 cm path length at scanning rate of 100 nm/min.

2.6. Cell culture

bEnd.3 endothelial cells (ATCC, VA, USA) were cultured in Dulbecco's Modified Eagle's Medium (DMEM, #30-2002 from ATCC, VA, USA) supplemented with 10 % FBS (Gibco, UK) and incubated at 37 °C with 5 % CO₂ in accordance with the supplier's instruction.

2.7. Cell viability assay

MTT assay was performed to assess the cell viability of bEnd.3. Sample solutions of A β ₁₋₄₂ with and without salidroside were added to the bEnd.3 cells for 24 h and MTT assay was done based on the literature (Telli et al., 1999) using a microplate absorbance reader (Bio-Rad). Cell viability was reported statistically significant as compared to control cells. The cells exposed to A β ₁₋₄₂ monomer (7 μ M) and salidroside (7 μ M) were also used to determine the cytotoxicity of native protein and salidroside at the highest studied concentration.

2.8. ROS assay

The level of ROS was quantified by using 2', 7'-dichlorodihydrofluorescein as a fluorescence probe as determined in the literature (Chen et al., 2010). The fluorescence intensity (λ_{ex} = 485 nm; λ_{em} = 535 nm) was read using the Perkin Elmer Multimode reader (Waltham, USA).

2.9. Superoxide dismutase (SOD), catalase (CAT), and caspase-3 activities assays

After preparation of protein extract and determination of protein concentration in the supernatant by Bradford assay (Jones et al., 1989), 25 μ g of protein was used to determine the SOD and CAT activity using the relative kits (Abcam, UK) based on the manufacturer's protocol.

2.10. Statistical analysis

Data were reported as mean \pm SD of three independent assays and analyzed by Student *t*-test to explore the significant differences between samples. *P* < 0.05 was reported as significant.

3. Results and discussion

3.1. Spectroscopic studies

The kinetic of A β ₁₋₄₂ aggregation was explored via ThT fluorescence analysis. The sigmoidal ThT fluorescence plot was detected for A β ₁₋₄₂ aggregation with a detectable lag phase followed by fibrillization step and finally a plateau step, indicating a probable nucleation dependent aggregation rate. To analyze the preventive influence of salidroside, A β ₁₋₄₂ aggregation was performed in the presence of different concentrations of salidroside. As depicted in Fig. 1a, in addition to an obvious influence on the lag phase, salidroside led to an apparent reduction in the steady-state fluorescence (F_{max}) in a concentration-dependent fashion. It can be indicated that F_{max} is directly proportional to the level of aggregation and the

reduction in the amount of F_{max} can be used to determine the inhibitory effect of salidroside (Sabaté et al., 2003). The obvious reduction in the F_{max} can propose that salidroside can inhibit the A β ₁₋₄₂ aggregation. As demonstrated in Fig. 1a, salidroside with various concentrations of 35 and 70 μ M potentially prevented A β ₁₋₄₂ aggregation as calculated by about 44 % and 63 % reduction in F_{max} , respectively. Moreover, the apparent rate growth constant (k_{app}) can be regarded as an important indicator of protein aggregation. It was found that the k_{app} values were 0.331 ± 0.029 h⁻¹, 0.213 ± 0.014 h⁻¹, 0.133 ± 0.011 h⁻¹ upon interaction of A β ₁₋₄₂ with various concentrations of salidroside; 0, 35 and 70 μ M, respectively.

It has been detected that structural alterations of biomolecules occurring upon protein aggregation can be determined by formation of hydrophobic moieties on the protein surface (Sirangelo et al., 1998). The increase in ANS fluorescence examination as a hydrophobic probe proved the appearance of obvious hydrophobic patches on the surface of the A β ₁₋₄₂ aggregated species (Fig. 1b). However, the presence of salidroside mitigated the ANS fluorescence intensity accompanied by an apparent red shift, suggesting that salidroside prevented the induction of hydrophobic patches in A β ₁₋₄₂ structure. This data in agreement with ThT fluorescence analysis suggested that salidroside potentially inhibited A β ₁₋₄₂ aggregation through inhibiting the formation of hydrophobic moieties on the protein.

To further study the inhibitory influence of salidroside against A β ₁₋₄₂ aggregation, CR adsorption analysis was carried out. Actually, the influence of salidroside on the A β ₁₋₄₂ aggregation was assessed through detecting maximal CR optical density. It was found that salidroside was outstandingly potent in preventing A β ₁₋₄₂ aggregation and this protective effect was directly related with the concentration of salidroside, as shown by the obvious decrease and blue shift detected in CR absorbance of A β ₁₋₄₂ samples incubated with salidroside (Fig. 1c).

To explore whether secondary structural changes of A β ₁₋₄₂ could be prevented by salidroside, far-UV CD technique was employed. As depicted in Fig. 1d, the ellipticity changes of A β ₁₋₄₂ after aggregation show an apparent transition from 195 nm to 218 nm, indicating the predominant formation of β -sheet structures (Böhm et al., 1992). Nevertheless, salidroside demonstrated a promising inhibition on secondary structural changes of A β ₁₋₄₂ and this aggregation inhibition potency was more pronounced in the case of higher concentration of this small molecule than lower concentrations. Also, it was realized that salidroside does not trigger any conformational changes on the random coil structure of A β ₁₋₄₂ monomers (data not shown).

This data also proved that salidroside protects A β ₁₋₄₂ against secondary structural changes induced by an aggregation buffer.

Since, A β ₁₋₄₂ oligomeric species are cytotoxic and the inhibition of protein aggregation by small molecules like salidroside can mitigate this associated cytotoxicity (Xu et al., 2019), cellular assays were performed to further support the spectroscopic data.

For the cellular assay, A β ₁₋₄₂ samples diluted to a final concentration of 7 μ M with corresponding salidroside concentrations of 3.5 μ M and 7 μ M.

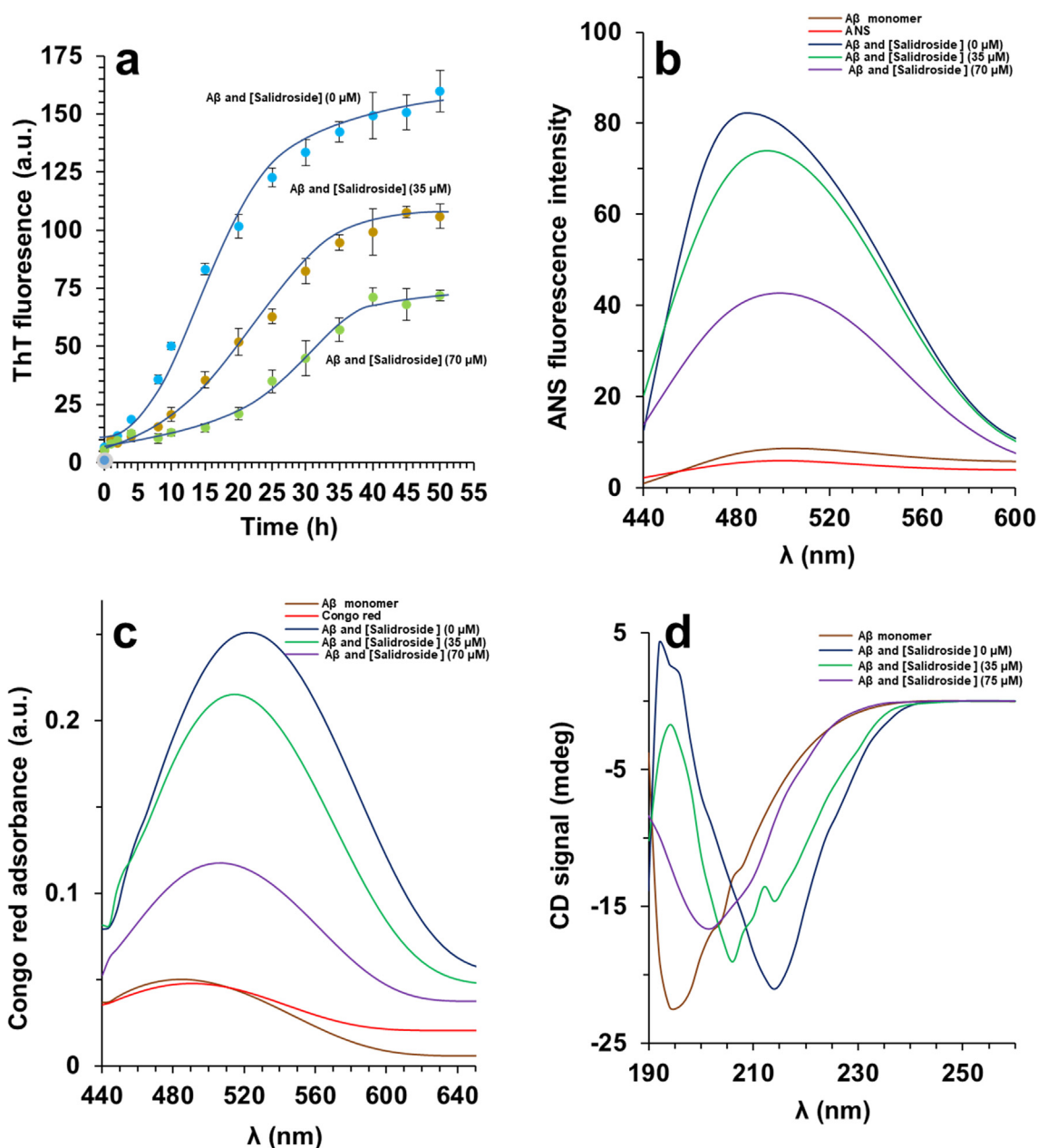


Fig. 1 (a) ThT fluorescence assay of A β_{1-42} aggregation with or without salidroside. (b) ANS fluorescence assay of A β_{1-42} aggregated samples with or without salidroside. (c) Congo red absorbance assay of A β_{1-42} aggregated samples with or without salidroside. (d) CD analysis of A β_{1-42} aggregated samples with or without salidroside.

3.2. MTT assay

The cytotoxicity of different A β_{1-42} samples against bEnd.3 cells for 24 h was explored via MTT assay (Fig. 2). It was seen that not only salidroside does not induce a significant cytotoxicity, but also the toxicity of A β_{1-42} oligomeric species aged for 25 h with salidroside against bEnd.3 cells was remarkably lower than that of A β_{1-42} aggregated species alone, and this protective effect was more pronounced in the case of higher concentration of this small molecule than lower concentration. This outcome indicated that the salidroside as a biocompatible compound can prevent the A β_{1-42} aggregation and associated cytotoxicity.

3.3. Oxidative stress assay

The effect of different A β_{1-42} samples alone or with salidroside on generation of intracellular ROS and SOD and CAT activity was assessed to examine the protective effect of salidroside against A β_{1-42} oligomeric samples-induced oxidative stress in bEnd.3 cells after 24 h (Fig. 3). It was determined that A β_{1-42} oligomers stimulated a significant enhancement in the generation of intracellular ROS (Fig. 3a), and reduction in the SOD activity (Fig. 3b) and CAT activity (Fig. 3c) in bEnd.3 cells. Nevertheless, treatment of cells with A β_{1-42} oligomers cocubated with salidroside reduced the generation of ROS and recovered the SOD and CAT activity. Therefore, it was

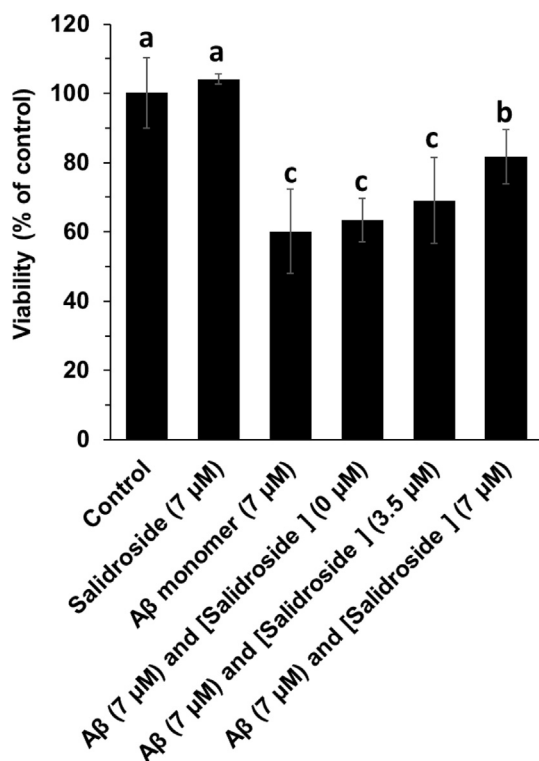


Fig. 2 MTT assay of bEnd. 3 cells after incubation with different A β_{1-42} samples (aged for 25 h) for 24 h. a, b, and c letters show the significance ($P < 0.05$).

deduced that the level of oxidative stress induced by A β_{1-42} oligomers can be mitigated upon treatment of cells with salidroside.

3.4. Caspase-3 activity

The effect of different A β_{1-42} oligomers alone or with salidroside on caspase-3 activity was explored to assess the probable inhibitory effect of salidroside against A β_{1-42} samples-triggered apoptosis in bEnd.3 cells after 24 h (Fig. 4). It was observed that A β_{1-42} oligomers induced a significant increase in caspase-3 activity in bEnd. 3 cells, whereas treatment of cells with A β_{1-42} samples co-incubated with salidroside mitigated the caspase-3 activity. Therefore, it was determined that the probable apoptosis stimulated by A β_{1-42} oligomers can be reduced after treatment of cells with salidroside.

4. Discussion

In recent years, several emerging diseases have been identified in humans directly related to abnormal and expanded aggregation of protein (Chiti and Dobson, 2017). The ability to form amyloid structures is one of the general characteristics of proteins. Therefore, the development of amyloid structures in model proteins and the investigation of the protective effect of small molecules on the formation of these structures can be useful in designing potentially effective drug compounds for disease-related amyloid (Alam et al., 2017). The present study was designed to investigate the effect of salidroside as a flavonoid small molecule on the formation of amyloid structure in A β_{1-42} as a model protein and its associated cytotoxicity in bEnd.3 endothelial cells as a marker of cerebrovascular diseases.

There is ample evidence that any defect in the control and clearance mechanisms of misfolded proteins may lead to protein amyloid formation and the development of neurodegenerative diseases (Kylkilahti et al., 2021). On the other hand, the aggregation of proteins stimulates the cellular stress and

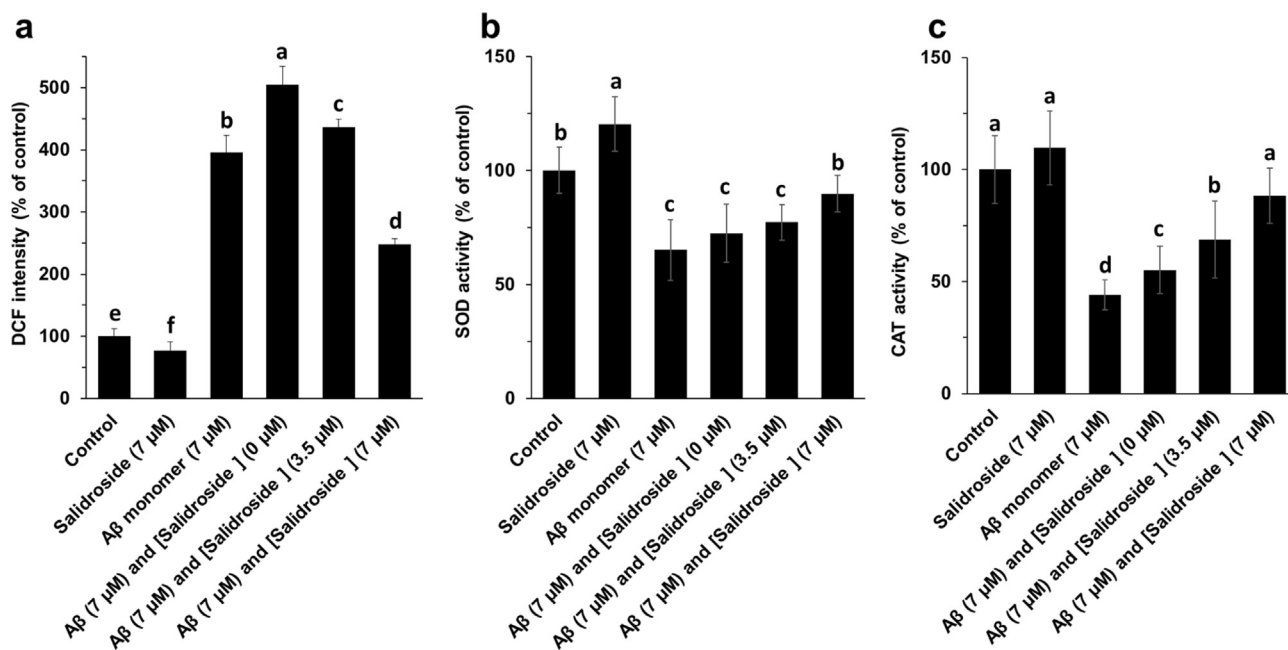


Fig. 3 ROS (a), SOD activity (b), CAT activity (c) assays of bEnd. 3 cells after incubation with different A β_{1-42} samples (aged for 25 h) for 24 h. a, b, c, and d letters show the significance ($P < 0.05$).

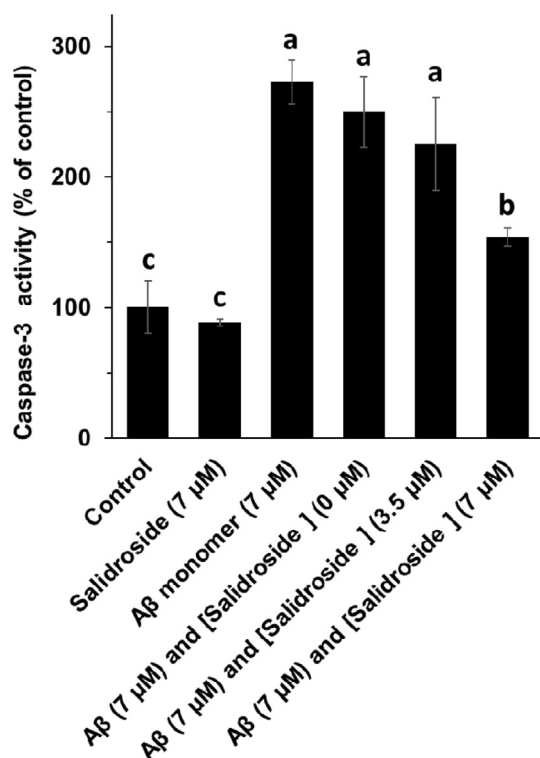


Fig. 4 Caspase-3 assay of bEnd. 3 cells after incubation with different A β_{1-42} samples (aged for 25 h) for 24 h. a, b, and c letters show the significance ($P < 0.05$).

appearance of neurodegenerative diseases (Zaman et al., 2019). Several human proteins that cause amyloid disease have been identified as capable of forming structures called amyloid fibrils in various tissues, including neurodegenerative diseases such as Huntington's, Alzheimer's and cerebrovascular (Tabner et al., 2001). Studies have shown that proteins with very different structures can advance to these particular clusters by changing their natural structure (Gadad et al., 2011).

Different proteins that are prone to aggregation do not have identical amino acid sequence, or three-dimensional structures, but despite their different origins, the amyloids formed have similar structures with a fold structure and a similar cytotoxicity mechanism (Gadad et al., 2011). Due to the presence of unprotected hydrophobic groups, the misfolded protein has the potential to form larger clusters, leading to toxicity, cell damage, and eventually programmed cell death (apoptosis) (Li et al., 1996).

Despite extensive research on the process of amyloid formation and related diseases, few treatments are currently available for these diseases. Studies in various animal cells and models indicate that the prevention of amyloid formation is useful in reducing the complications of amyloidosis (Fändrich et al., 2018). In addition, anti-amyloid activity of various compounds such as antibodies, synthetic peptides, proteins known as chaperones, and chemical compounds have been identified (Härd and Lendel, 2012). Also, a large number of small molecules with different concentrations ranging from nanomolar to millimolar have been found to provide high potential for inhibiting the formation of amyloid aggregates, especially A β . Medicinal plants and their bioactive compounds as natural molecules have been shown to potentially inhibit the

formation and accumulation of amyloid and related diseases (Doig and Derreumaux, 2015). Based on the outcomes from *in vitro* and *in vivo* assays, it was found that A β could induce oxidative stress and apoptosis in cerebrovascular endothelial cells, which is in agreement with previous reports (Park et al., 2011; Song et al., 2017; Chen et al., 2018).

It has been shown that phenolic compounds can mitigate the bEnd.3 cytotoxicity from A β peptide-stimulated oxidative stress (Xi et al., 2012; Liu et al., 2017). Salidroside as a phenolic compound has been shown to protect neuron-like cells against A β -triggered apoptosis by modulation of AKT signaling pathway (Liao et al., 2019).

It can be noted that cerebrovascular-targeted small molecules relieving oxidative stress may provide potential results for delayed progression of neurodegenerative diseases.

5. Conclusion

In conclusion, with the use of several biophysical and cellular techniques, we demonstrated that salidroside prevented A β_{1-42} peptide aggregation and mitigated against amyloid triggered cerebrovascular endothelial cytotoxicity. The fluorescence analysis indicated that salidroside prevents the A β_{1-42} peptide aggregation and this potential inhibitory effect was more pronounced in the presence of higher concentration of salidroside. The aggregated species that appeared in the presence of salidroside showed fewer β -sheet structures, as evidenced by the CD and CR spectra. Interestingly salidroside was shown to be safe against bEnd.3 cells and also mitigated the cells against oligomers-triggered cytotoxicity. Though much more studies *in vivo* investigation is demanded to support it as a potential and clinically viable drug. This study will further expand the pool of promising drugs for cerebrovascular disease from bioactive compounds.

Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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