



# Vdac1 Downregulation Causes Mitochondrial Disintegration Leading to Hippocampal Neurodegeneration in Scopolamine-Induced Amnesic Mice

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## Abstract

Our previous report on hippocampal proteome analysis suggested the involvement of voltage-dependent anion channel (Vdac) 1 in scopolamine-induced amnesia. Further silencing of Vdac1 in young mice reduced the recognition memory. Vdac1 is a porin protein present abundantly on outer mitochondrial membrane. It acts as a transporter of energy metabolites ATP/ADP and Ca<sup>2+</sup> ions and helps in communication between mitochondrial matrix and cytosol. As Vdac1-associated energy metabolism may be affected during amnesia, we determined the downstream function of Vdac1 in the present study. The expression of Vdac1 and total ATP level was decreased in the hippocampus of scopolamine-induced amnesic mice. Also, the mitochondrial membrane potential, cristae organization, and morphology were disrupted leading to increased ROS generation and reduced SOD and catalase activity. On the other hand, there was increase in the expression of pro-apoptotic marker proteins (Bax, Bad, Casp 3), leading to rising degenerated neuronal cells in the dentate gyrus and Cornu ammonis 3 and 1 subregions of the hippocampus during amnesia. Further, to check whether Vdac1 downregulation is associated with neurodegeneration, we infused Vdac1 siRNA stereotaxically in the hippocampus of normal young mice. As compared to control, Vdac1 silencing decreased ATP level and mitochondrial membrane potential leading to increase in the number of degenerated neuronal cells in subregions of the hippocampus. Taken together, our study shows that downregulation of Vdac1 causes neurodegeneration through mitochondrial disintegration in the hippocampus of scopolamine-induced amnesic mice.

**Keywords** Scopolamine · Vdac1 · Mitochondria · Apoptosis · Neurodegeneration · Hippocampus

## Introduction

Scopolamine, a nonselective antagonist of muscarinic cholinergic receptor, causes memory loss by altering synaptic plasticity associated gene expression [1–3]. It is used widely to generate amnesia in animal models. Using scopolamine-induced amnesic mice, we reported earlier alteration in the protein profile of hippocampus. Among 18 altered proteins,

Vdac1 expression was maximally downregulated, showing common interaction with other altered proteins and its silencing in the hippocampus of normal mice impaired recognition memory [4]. Weeber et al. [5] have also shown that Vdac1 deficiency leads to impairment in fear conditioning and spatial memory with disruption in long- and short-term synaptic plasticity in the hippocampus of mice. Vdac1 protein forms a channel in the outer mitochondrial membrane and allows the communication between mitochondrial matrix and cytoplasm, and thus regulates cell metabolism, apoptosis, and reactive oxygen species (ROS) through transportation of ATP/ADP [6]. Vdac1 deficiency reduces the cellular production of ATP in Vdac1 knockout mice [7]. Such failure to produce energy might cause impairment in neural transmission leading to neurodegeneration [8, 9]. Intracerebroventricular injection of amyloid  $\beta$ 1–42 peptide causes impairment in learning and memory through depletion of ATP and increase of ROS level leading to Alzheimer's disease (AD) like symptoms in cerebral cortex and hippocampus of mice [10]. Also, exposure to lead

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(Pb) and dopamine causes toxicity through downregulation of Vdac1 expression, ATP depletion, ROS elevation, and disruption in mitochondrial membrane potential (MMP) leading to apoptosis during in vitro and in vivo conditions [11, 12]. Also, cannabinoid intoxication induces amnesia through alteration in cellular respiration and energy depletion [13–15]. These studies suggest the involvement of Vdac1 in energy metabolism, cellular function, synaptic transmission, and learning and memory. As Vdac1 was maximally downregulated in the hippocampus of scopolamine-induced amnesic mice, this might be a key regulator of amnesia. Therefore, in the present study, we have explored the downstream function of Vdac1 in scopolamine-induced amnesic mice and reported that its downregulation causes failure of energy production leading to mitochondrial disintegration-mediated hippocampal neurodegeneration.

## Experimental Details

### Materials

Scopolamine hydrobromide was purchased from Sigma–Aldrich (USA); ENLITEN ATP Assay kit from Promega (USA); carboxy-H2DCFDA from Invitrogen (USA); Rhodamine 123 from Thermo Fisher Scientific (USA), primary antibodies; anti-Vdac1 (ab14734) from Abcam (UK), anti-cytochrome c (sc-13561), anti-Bcl-2 (sc-8392), and anti-caspase 3 (sc-7148) from Santa Cruz Biotechnology (USA); anti-Bax (no. 2772) and Bad (no. 9292) from Cell Signaling Technology (USA); Vdac1 siRNA and negative control siRNA (cat. no. 4390771, 4390843) from Ambion (USA). All other biochemicals were purchased from Merck (Germany).

### Animals and Drug Administration

Swiss albino young ( $10 \pm 2$  weeks) male mice were inbred and maintained at  $24 \pm 2$  °C under 12 h light and dark cycle with ad libitum access to standard mice feed and drinking water in the animal house of Department of Zoology, Banaras Hindu University, India. They were handled as per guidelines of the Animal Ethical Committee, Banaras Hindu University. Scopolamine hydrobromide (SC) (3 mg/kg body weight) was administered intraperitoneally to experimental mice and saline (SA) (0.9% NaCl) as a vehicle to control mice between 9 and 10 am daily for 7 days. After treatment, control ( $n = 20$ ) and treated ( $n = 20$ ) mouse groups were sacrificed by cervical dislocation, brain was dissected out from the skull, and both cerebral hemispheres were separated in saline. Then, the hippocampi were removed for further experiments [16].

## Biochemical Analysis

### Analysis of Vdac1 and ATP Level

Protein extract was prepared by homogenizing hippocampus (10%) in RIPA buffer (50 mM TrisCl, pH 7.4, 150 mM NaCl, 0.1% Triton X-100, 0.1% SDS, 2 mM EDTA, 10 mM sodium deoxycholate, and 0.1 mM PMSF) followed by centrifugation at  $12,000 \times g$  for 20 min at 4 °C. The supernatant was collected in a microfuge tube and protein concentration was determined [17]. Equal amount of protein was separated by 10% SDS-PAGE (15% for cytochrome c), blotted to polyvinylidene difluoride (PVDF) membrane (Millipore) and blocked with 5% (w/v) nonfat skimmed milk for 2–3 h. After blocking, the blots were probed with primary antibody anti-Vdac1. After incubation, the blot was washed in phosphate-buffered saline (PBS) thrice and incubated with HRP-conjugated secondary antibody (1:2000) (Bangalore Genei, India) for 3 h. Then, the blot was washed in PBS and signal was detected using ECL reagents on X-ray film. The same blot was probed with anti- $\beta$ -actin HRP conjugated antibody (1:20,000) for 4–6 h and used for normalization.

To measure the energy level, ATP was extracted from the tissue using trichloroacetic acid (TCA) as previously described [18] and protocol supplied by the manufacturer with slight modification. The isolated hippocampus was immediately homogenized in 500  $\mu$ l ice-cold homogenization buffer (0.25 M sucrose and 10 mM HEPES–NaOH, pH 7.4) for three cycles of 30 s each followed by centrifugation at  $1000 \times g$  for 10 min at 4 °C. The supernatant (300  $\mu$ l) was quickly added to an equal volume of ice-cold 10% TCA, shaken for 20 s, and centrifuged at  $10,000 \times g$  for 10 min at 4 °C. Further, 400  $\mu$ l of supernatant was transferred to a fresh tube and 200  $\mu$ l 1 M Tris-acetate buffer (pH 7.75) was added for neutralization. The sample was diluted 30 times with ATP free water and ATP level was measured by luciferase assay through detection of bioluminescence emission. The tissue extract (10  $\mu$ l) was mixed with 90  $\mu$ l of luciferase reagent and luminescence was recorded immediately using multimode reader with bioluminescence detector (BioTek SYNERGY H1, USA). For normalization, readings were divided by tissue weight for respective control and experimental groups. Luciferase enzyme utilizes ATP and catalyzes the reaction to release energy in the form of bioluminescence. If high amount of ATP is available for luciferase reaction, it catalyzes more reaction and releases more bioluminescence ( $\text{ATP} + \text{Luciferin} + \text{O}_2 \rightarrow \text{Oxyluciferin} + \text{AMP} + \text{PPi} + \text{CO}_2 + \text{Light } 560 \text{ nm}$ ).

### Analysis of Mitochondrial Membrane Potential, Ultrastructure, and Calcium Level

To measure the mitochondrial membrane potential, we used Rhodamine 123 (Rh 123) which is a cell-permeant, green fluorescent, cationic dye and accumulates in active

mitochondria. The active mitochondria having normal membrane potential sequesters more amount of Rh 123 and proportionally gives more fluorescent intensity. Mitochondria were isolated from the hippocampus as described earlier with slight changes [19]. The tissue was minced on ice and homogenized in isolation buffer (225 mM mannitol, 75 mM sucrose, 5 mM HEPES, 1 mM EGTA, 1 mg/ml BSA, pH 7.4). The homogenate was centrifuged at  $600\times g$  for 5 min at 4 °C. The supernatant was collected and centrifuged at  $9000\times g$  for 8 min at 4 °C. The pellet obtained was resuspended in isolation buffer containing 0.02% heparin and centrifuged at  $9000\times g$  for 8 min at 4 °C. Finally, the brownish pellet was resuspended in isolation buffer and centrifuged at  $9000\times g$  for 10 min at 4 °C. The pellet was again suspended in the isolation buffer containing heparin and protein was estimated. Equal amount of mitochondria was taken and incubated with Rh 123 (0.3  $\mu$ M) dye for 20 min at 37 °C followed by centrifugation at  $9000\times g$  for 5 min to remove extra dye. Then, fluorescence was recorded in the multimode reader with fluorescence detector at 535 nm (excitation) and 580 nm (emission) wavelength. The fluorescence intensity value was used to reflect the mitochondrial membrane potential.

For mitochondrial ultrastructure, control and treated mouse groups were sacrificed and hippocampus was isolated. It was cut into 2-mm<sup>2</sup> sizes and fixed overnight at 4 °C with Karnovsky fixative containing 2.5% glutaraldehyde and 4% paraformaldehyde in 0.1 M phosphate buffer (pH 7.4). The tissue pieces were washed thrice for 15 min each with 0.1 M phosphate buffer at 4 °C and cut into 1-mm<sup>2</sup> size followed by post-fixation with 1% osmium tetroxide for 2 h at 4 °C. The tissue pieces were dehydrated in ice-cold acetone at 4 °C, cleared with toluene, and embedded in Araldite CY 212. Ultrathin sections (~60–90 nm) were cut by ultramicrotome, collected on copper grids, and stained with uranyl acetate and lead citrate. The grids were washed with distilled water and viewed using a transmission electron microscope (FEI Tecnai S Twin at All India Institute of Medical Sciences, New Delhi) at  $\times 5000$  magnification for visualization of mitochondrial ultrastructure in the hippocampus as described earlier with slight modification [20].

To measure calcium level, first, mitochondria were isolated and atomic absorption flame spectroscopy (AAS) was performed as described earlier [21]. Hippocampus (60 mg) was cut into small pieces in ice-cold medium (250 mM sucrose, 2 mM HEPES, and 3  $\mu$ M ruthenium red, pH 7.4). Once rinsed in buffer, tissue was homogenized in 2 ml buffer using homogenizer at speed of 5000 rpm with 5–7 strokes up and down. The homogenate was differentially centrifuged at  $1000\times g$  for 10 min, and then supernatant was further centrifuged at  $10,000\times g$  for 14 min to settle down the mitochondria. The pellet was washed once with buffer (equal volume to supernatant) and centrifuged at  $10,000\times g$  for 14 min. Finally, mitochondrial pellet was resuspended in 1.5 ml buffer and Janus Green staining was used to check the mitochondria.

Protein was estimated using Bradford method and equal amount of protein (0.15 mg) was taken to determine the calcium level by using AAS-containing cathode lamp. Samples were prepared through base digestion method with 1 N NaOH and heated at 75 °C in water bath for 15 min twice. After digestion, samples were passed through sterile 0.2- $\mu$ M syringe filter and used for analysis. The calcium standard curve was developed using different concentrations (1, 4.5, 10 mg/l) by AAS (Perkin Elmer) in the Department of Botany, Banaras Hindu University, as described earlier [22]. Samples were analyzed and readings were noted to calculate the calcium level in micromolars per milligram of protein.

### Analysis of ROS and Antioxidant Enzyme Activity

To determine total ROS level, 10% hippocampal homogenate was prepared in PBS followed by centrifugation at  $10,000\times g$  at 4 °C as described previously [23]. The supernatant was used for protein estimation; equal amount of protein was taken from control and treated samples, and incubated with 2 mM 2', 7'-dichlorofluorescein diacetate (H2DCFDA) in PBS at 37 °C for 30 min. The available ROS converted nonfluorescent H2DCFDA to highly fluorescent 2', 7'-dichlorofluorescein (DCF) which was recorded at 488 nm (excitation) and 525 nm (emission) using a multimode reader. The level of ROS in each sample was determined by calculating fluorescence intensity value per milligram of protein.

The antioxidant enzymes catalase and superoxide dismutase (SOD) play a role in defense mechanism of the cell. So, their activity was measured in mouse hippocampus as described earlier with slight modification [24]. Catalase activity was analyzed by native polyacrylamide gel electrophoresis followed by ferricyanide staining method. The homogenate (10%) was prepared in 0.2 mM Tris-HCl (pH 7.4) and centrifuged at  $10,000\times g$  for 10 min at 4 °C. The supernatant was used for protein estimation and an equal amount of protein was resolved by 8% native PAGE at 4 °C. After completion of electrophoresis, the gel was washed with distilled water and soaked in H<sub>2</sub>O<sub>2</sub> (0.003%) for 5–10 min followed by rinsing with distilled water to remove excess H<sub>2</sub>O<sub>2</sub>. The gel was then stained in a solution containing 1% ferric chloride and 1% potassium ferricyanide for 4–5 min. Thereafter, the achromatic bands appeared and the image was captured by Alpha Innotech gel documentation system for densitometric analysis. For analysis of SOD activity, an equal amount of protein was separated by 10% native PAGE at 4 °C and the gel was soaked in 1.23 mM nitro blue tetrazolium chloride (NBT) solution for 20 min in dark. Then, the gel was washed with distilled water and incubated in 100 mM phosphate buffer (pH 7.0) containing 28 mM TEMED and 0.28 mM riboflavin for 15–20 min in dark. Finally, the gel was exposed to light until achromatic bands appeared with the purple-blue

background; the image was captured and analyzed by Alpha Innotech gel documentation system.

### Analysis of Apoptotic Marker Protein Level

To check apoptosis, we have analyzed the level of apoptotic marker proteins in the hippocampus by western blotting as mentioned in the previous section. Equal amount of protein was resolved by 10% SDS-PAGE and blotted to PVDF membrane. For cytochrome c, we isolated the cytosolic and mitochondrial fraction as described earlier [25] and equal amount of cytosolic and mitochondrial fraction protein was separated by 15% SDS-PAGE and blotted to PVDF membrane. Blots were blocked for 2–3 h and probed with primary antibodies anti-cytochrome c (1:500), anti-Bax (1:1000), anti-Bad (1:1000), caspase 3 (1:1000), and anti-Bcl-2 (1:500) diluted in blocking buffer overnight at 4 °C. After incubation, the blots were washed in PBS thrice and incubated with HRP-conjugated secondary antibodies (1:2000) for 3 h. Then, the blots were washed in PBS and signal was detected on X-ray film. The same blot was probed with anti- $\beta$ -actin HRP-conjugated antibody for 4–6 h and used for normalization.

### Analysis of Neurodegeneration

For detection of degenerated neurons in the hippocampus of control and treated mice, the brain was processed as described earlier. The tissue was first fixed in 4% paraformaldehyde (PFA) overnight, then immersed in 30% sucrose and finally embedded in optimum cutting temperature (OCT) solution. Transverse 8- $\mu$ m thick sections were cut by a cryostat (Leica Biosystem, Germany), mounted on poly-L-lysine-coated slides and stored at –80 °C for FJC staining. FJC dye is a polyanionic fluorescein-derived molecule which selectively binds to degenerated neurons and is used to detect neurodegeneration [26]. To visualize degenerated neuronal cells, the brain sections were immersed in a solution of 1% NaOH for 5 min, followed by rinsing with 70% ethanol and distilled water for 2 min each [27]. Then, sections were incubated in 0.06% potassium permanganate solution for 10 min and rinsed in distilled water for 2 min. Further, slides were incubated in FJC staining solution (stock solution 0.01% in water; diluted in 0.1% acetic acid) for 25 min. Then, slides were washed for 1 min thrice and air-dried at 50 °C for 30 min. After clearing in xylene, sections were mounted with DPX solution and a coverslip was applied. Slides were observed and hippocampal subregions dentate gyrus (DG) and Cornu ammonis (CA) 3 and CA1 were focused under the fluorescence microscope (Leica, Germany) with FITC filter. The images

were captured at  $\times 40$  resolution and FJC-positive cells were counted using ImageJ software (NIH, USA).

### Intrahippocampal Vdac1 siRNA Infusion

To silence Vdac1 expression, Vdac1 siRNA and nontargeting sequence as a negative control (NC) were infused bilaterally into the hippocampus stereotaxically as described earlier [4]. The mice were anesthetized by administration of sodium pentobarbital (40 mg/kg bw i.p.). Hair was removed from the head and fixed on stereotaxic stage. Initially, coordinates (2 mm frontal to lambda, 2 mm lateral to midline, and 1.4 mm depth from the skull surface) were set to hippocampus and slowly a hole was made by additional needle. The Vdac1 siRNA sequence (sense 5'-GUCAGAGAAUGGAUUGGAAtt-3', antisense 5'-UCCCAAUCCAUUCUCUGACtt-3') was validated through siRNA-Check online bioinformatics tool (<http://projects.insilico.us/SpliceCenter/siRNACheck>). Vdac1 siRNA (3  $\mu$ g/5  $\mu$ l) and NC was infused at the rate of 1  $\mu$ l/min. After infusion, the animal was removed from the apparatus and incision was stitched and nebasulf powder was employed for normal recovery. All the animals were kept in separate cages and exposed to table lamp for warming. After 5 days of the recovery, mice were sacrificed and the hippocampus was separated. The expression of Vdac1 protein was analyzed by western blotting, ATP level by luciferase assay, MMP by Rh 123 dye, and neurodegeneration by FJC staining as described in previous sections.

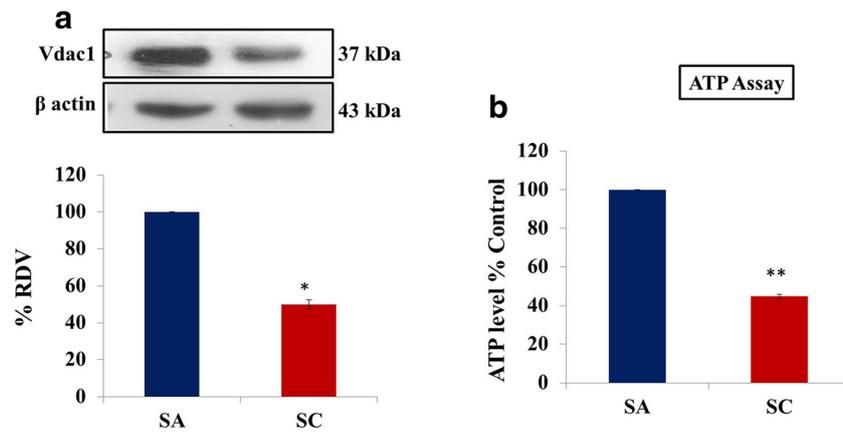
### Statistical Analysis

The data of control and experimental groups obtained from western blotting, ATP assay, ROS level, the density value of achromatic band of antioxidant enzymes activity, the fluorescence intensity of Rh123, and FJC-positive cell number were analyzed and values were presented as the mean  $\pm$  SEM. Student's *t* test was used to determine statistical significance by SPSS 16.0 (Statistical Products and Service Solutions, IBM Corporation, Armonk, NY, USA). The *p* value < 0.05 was considered as statistically significant.

## Results

### Effect of Scopolamine on Vdac1 and ATP Level

The level of Vdac1 protein was downregulated (45%) and ATP level was reduced (55%) in the hippocampus of scopolamine-induced amnesic mice as compared to control (Fig. 1). Such depletion in ATP level might occur through Vdac1 downregulation.



**Fig. 1** Effect of scopolamine on Vdac1 and ATP level. **a** Western blot analysis of Vdac1 expression. Histogram represents relative density value (RDV) (Vdac1/ $\beta$  actin). **b** Luciferase assay analysis to measure ATP level. Histogram shows bioluminescence intensity-based % ATP level.

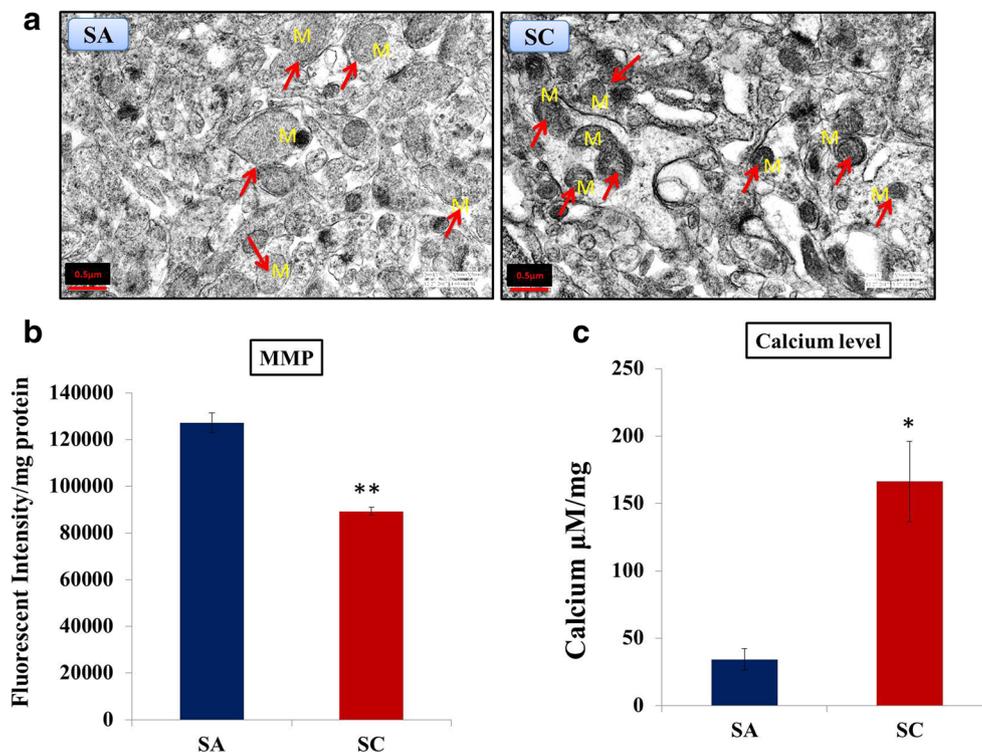
Data are represented as mean  $\pm$  SEM of three animals in each group. Statistically significant difference is represented as one ( $p < 0.01$ ) and two ( $p < 0.001$ ) asterisks in scopolamine (SC) group as compared to saline control (SA)

### Effect of Scopolamine on Mitochondrial Membrane Potential, Ultrastructure, and ATP Level

As reduced ATP level affects mitochondrial integrity, we checked it and analyzed mitochondrial membrane potential using Rh 123 dye. Both mitochondrial membrane potential

and accumulation of Rh 123 in mitochondria declined significantly, suggesting disruption of mitochondrial membrane potential in the hippocampus of scopolamine-induced amnesic mice as compared to control (Fig. 2a).

In electron micrograph, we observed that mitochondria were small in size, more spherical in shape, and have damaged



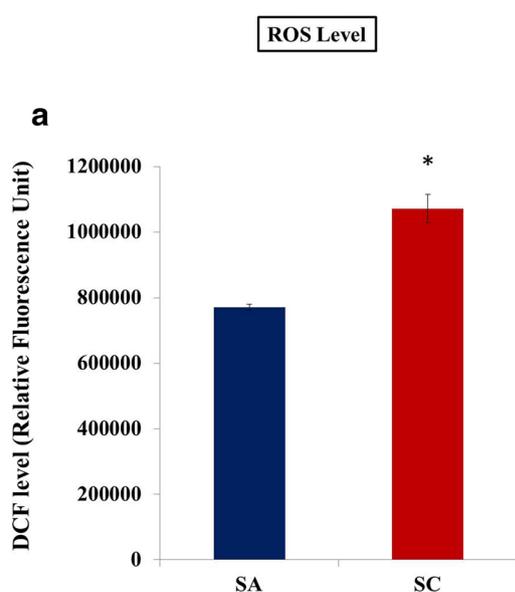
**Fig. 2** Effect of scopolamine on mitochondrial ultrastructure, membrane potential, and calcium level. **a** Electron micrograph revealed small size, more spherical shape, and damaged cristae in mitochondria of scopolamine-induced amnesic mice. Mitochondria appeared blackish in amnesic mice. M, mitochondria. Arrows indicate mitochondria. Scale bar, 0.5  $\mu$ m. **b** Analysis of mitochondrial membrane potential: histogram

shows comparative fluorescent intensity values. **c** Measurement of calcium level in mitochondria: histogram shows calcium level in isolated mitochondria. Values are represented as mean  $\pm$  SEM of three animals in each group. Statistically significant difference is represented as one ( $p < 0.05$ ) and two ( $p < 0.01$ ) asterisks in SC group as compared to SA

cristae in the hippocampus of scopolamine-induced amnesic mice. Mitochondria appeared blackish in the hippocampus of amnesic mice, but they were small and elongated with clearly visible larger cristae in control. The blackish appearance of mitochondria could be due to the deposition of calcium ions during apoptosis (Fig. 2b). To confirm this, we analyzed the calcium level in mitochondria of control and scopolamine-induced amnesic hippocampus. Interestingly, we observed that calcium level was very high in mitochondria of scopolamine-induced amnesic hippocampus as compared to control (Fig. 2c). This indicates that the blackish appearance of mitochondria might be due to accumulation of calcium.

### Effect of Scopolamine on ROS Level and Antioxidant Enzyme Activity

As reduced ATP level in the hippocampus of amnesic mice might affect the normal mitochondrial function and disrupt the cell homeostasis, we checked ROS level and the activity of its scavenging enzymes (SOD and catalase). ROS level (fluorescent intensity) was increased (Fig. 3a), and the activity of antioxidant enzymes SOD (35.47%) and catalase (42.47%) decreased in the hippocampus of scopolamine-induced amnesic mice as compared to control (Fig. 3b).



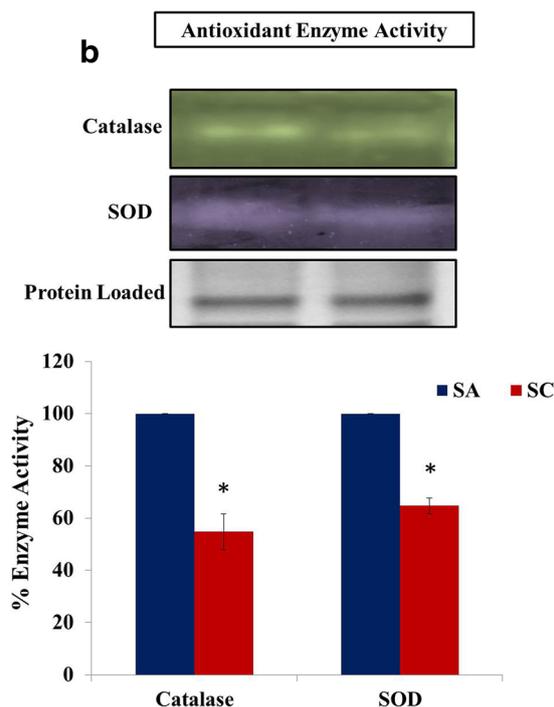
**Fig. 3** Effect of scopolamine on ROS and antioxidant enzyme activity. **a** Measurement of ROS level: histogram shows DCF fluorescence intensity values. **b** Activity of antioxidant enzymes (SOD and catalase) was analyzed in the hippocampus of amnesic mice and density values are

### Effect of Scopolamine on Apoptotic Marker Protein Level

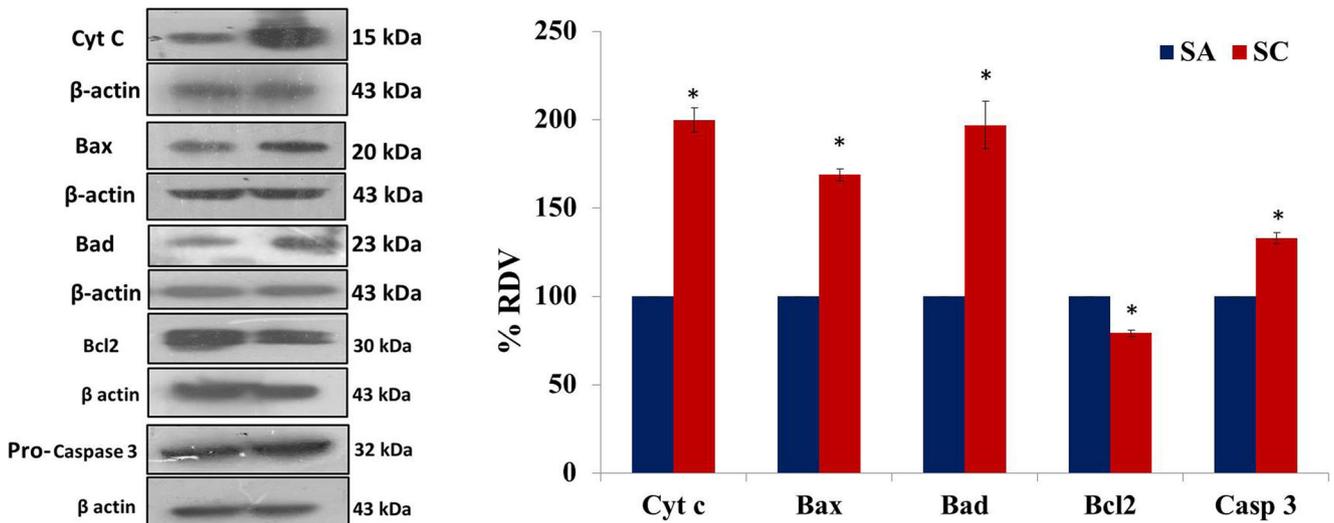
As alteration in the mitochondrial integrity during amnesia might trigger the intrinsic pathway of apoptosis, we analyzed the level of cytosolic cytochrome c, Bax, Bad, Bcl-2, and caspase 3 protein. The level of cytosolic cytochrome c and pro-apoptotic proteins (Bax, Bad, Pro-Caspase 3) increased but the level of the cyt c in mitochondrial fraction (Supplementary Fig. 2) and anti-apoptotic protein (Bcl-2) decreased in the hippocampus of scopolamine-induced amnesic mice as compared to control (Fig. 4).

### Effect of Scopolamine on Neurodegeneration

As elevated expression of apoptotic proteins is a sign of apoptosis and cell death, we analyzed neurodegeneration during scopolamine-induced amnesia. Through microscopic visualization of FJC staining, we observed significantly high number of degenerated FJC positive neuronal cells in DG, CA3, and CA1 subregions of the hippocampus of scopolamine-induced amnesic mice as compared to control. This indicated relatively higher neurodegeneration in the hippocampus of amnesic mice (Fig. 5).



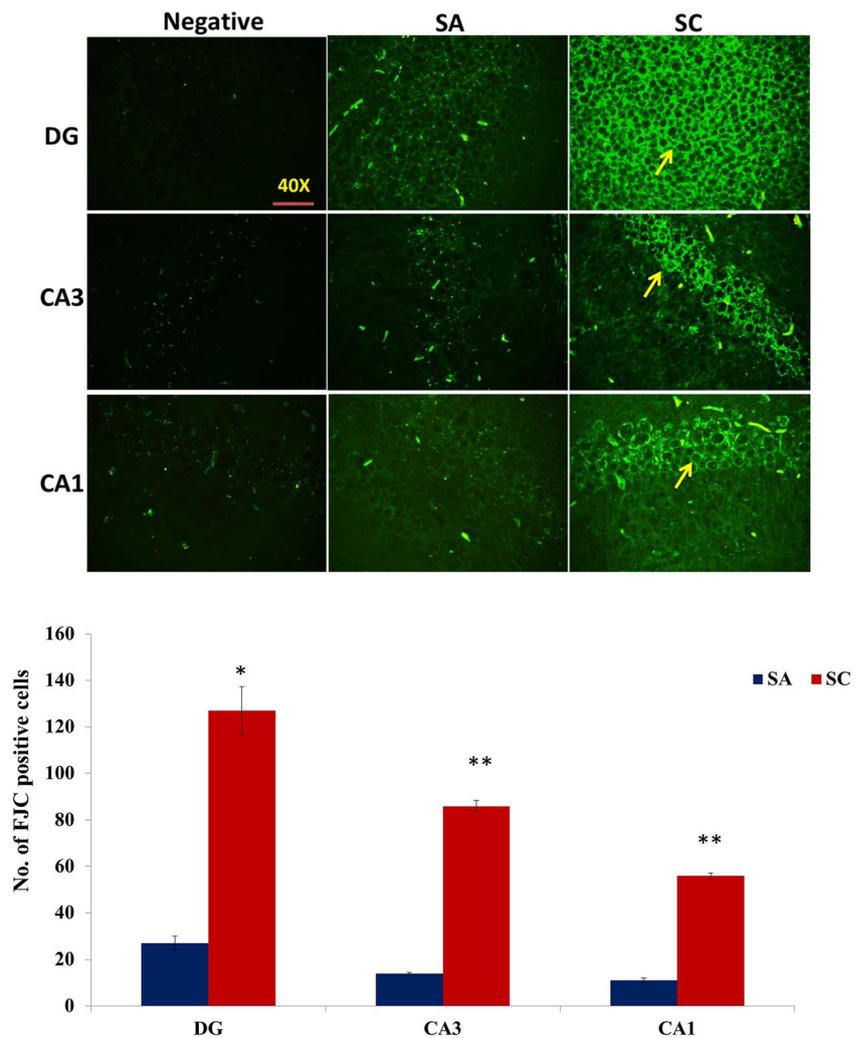
plotted. Data are represented as mean  $\pm$  SEM of three animals in each group. Statistically significant difference is represented as one asterisk ( $p < 0.05$ ) in SC group as compared to SA

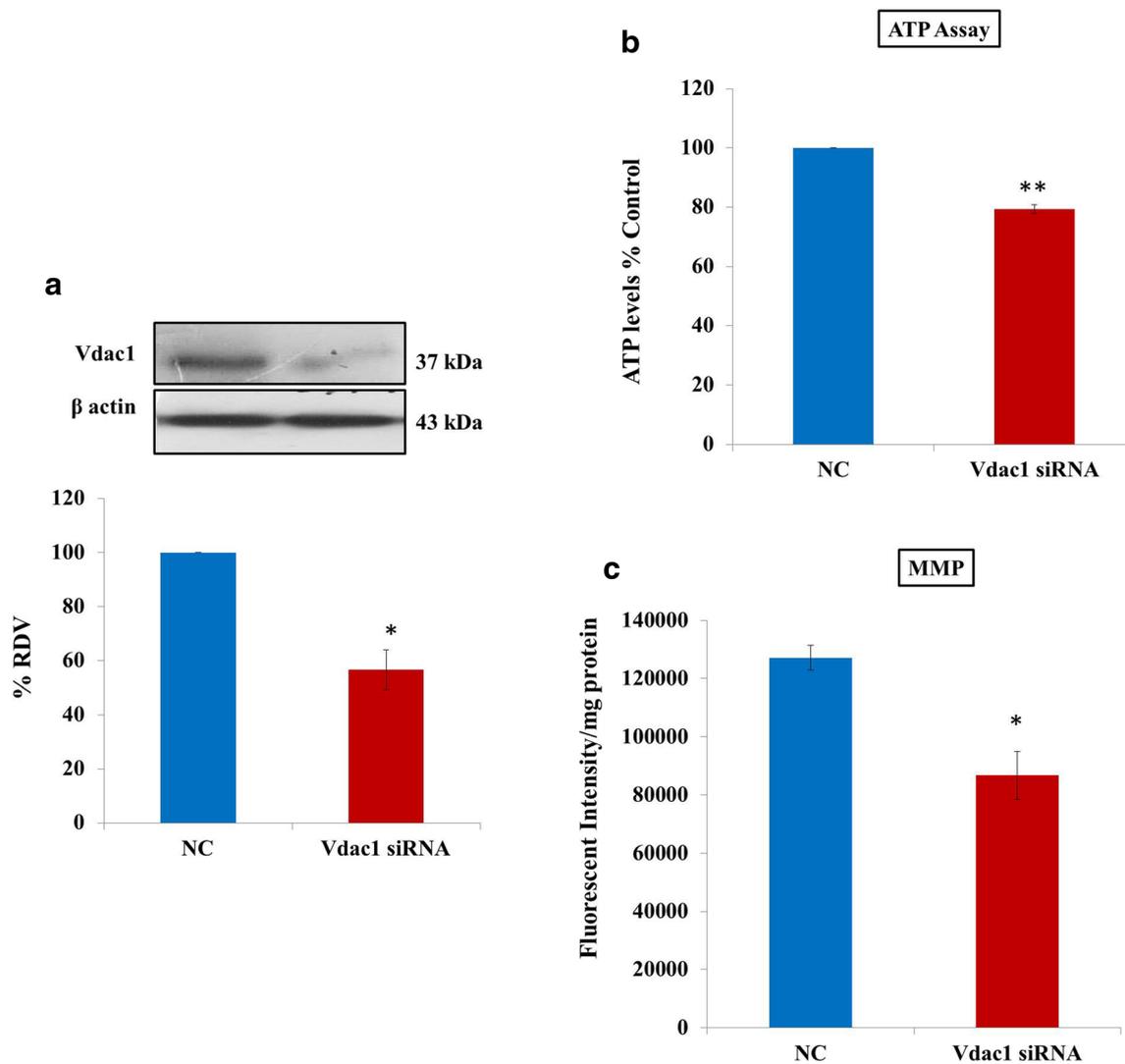


**Fig. 4** Effect of scopolamine on apoptotic marker protein level: Western blot analysis of pro-apoptotic and anti-apoptosis proteins. Histogram represents relative density value (RDV) (cyt c, Bax, Bad, Casp3,

Bcl1–2/β actin). Data are represented as mean ± SEM of three animals in each group. Statistically significant difference is represented as one ( $p < 0.05$ ) and two ( $p < 0.01$ ) asterisks in SC group as compared to SA

**Fig. 5** Effect of scopolamine on neurodegeneration: The panels show representative images of Fluro-Jade C staining in DG, CA3, and CA1 subregions of hippocampal neurons. Graph represents increased number of Fluro-Jade C-positive degenerated neuronal cells in the hippocampal subregions of scopolamine-induced amnesic mice. Arrows indicate FJC-positive cells. Data are represented as mean ± SEM of three animals in each group. Statistically significant difference is represented as one ( $p < 0.01$ ) and two ( $p < 0.001$ ) asterisks in SC group as compared to SA





**Fig. 6** Effect of Vdac1 silencing on Vdac1, ATP level, and mitochondrial membrane potential. **a** Western blot analysis of Vdac1 protein: histogram represents relative density value (RDV) (Vdac1/ $\beta$  actin). **b** Luciferase assay to measure ATP level: histogram shows bioluminescence intensity-based % ATP level. **c** Analysis of mitochondrial membrane

potential: histogram shows fluorescent intensity values. Data are represented as mean  $\pm$  SEM of three animals in each group. Statistically significant difference is represented as one ( $p < 0.01$ ) and two ( $p < 0.001$ ) asterisks in Vdac1 siRNA group as compared to negative control (NC)

### Effect of Vdac1 Silencing on Vdac1, ATP, MMP, and Neurodegeneration

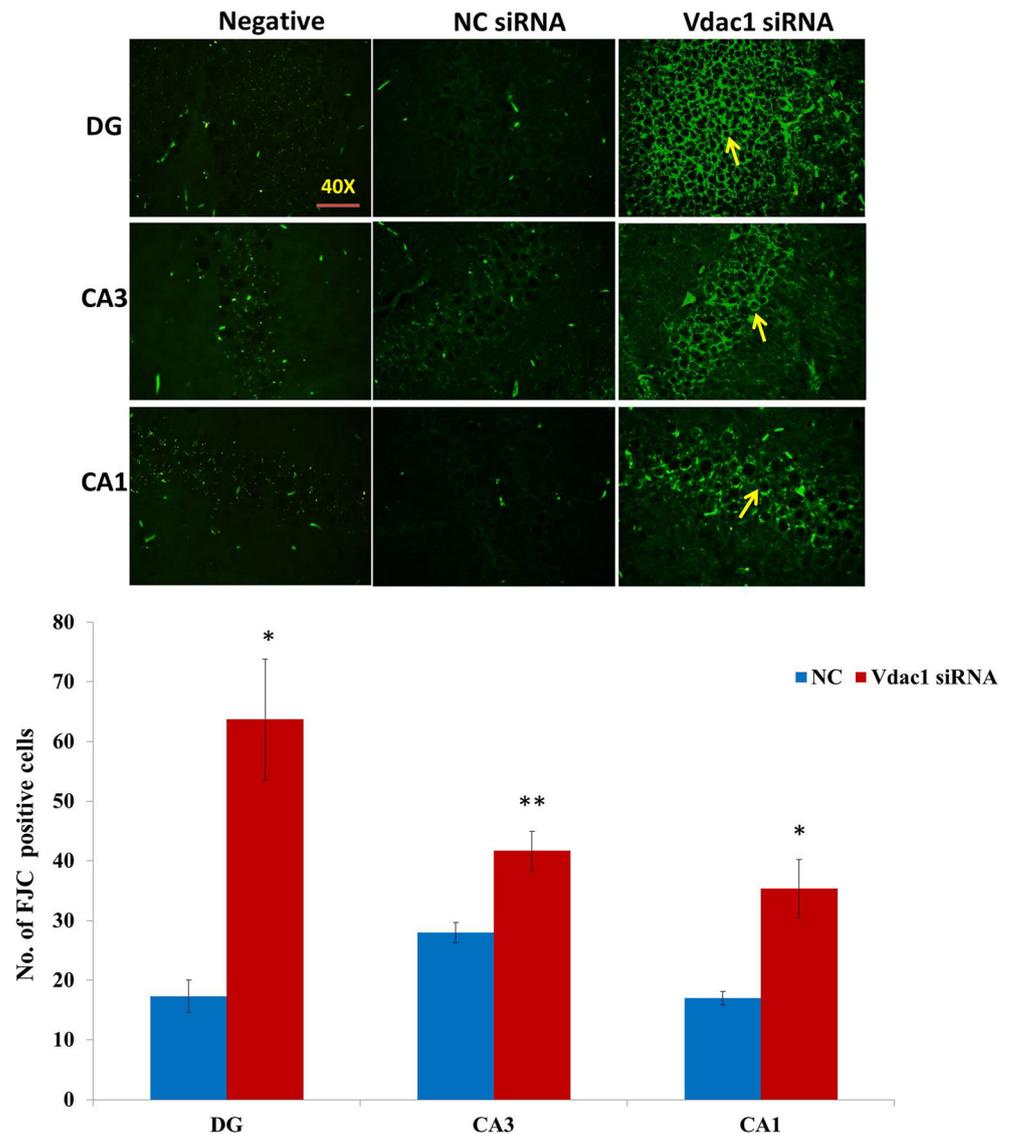
As the downstream function of Vdac1 is affected in scopolamine-induced amnesia, we checked whether downregulation of Vdac1 increased neurodegeneration. Therefore, we planned to silence Vdac1 protein in the hippocampus of normal young mice. The level of Vdac1 protein and total ATP decreased significantly in siRNA-infused hippocampus as compared to NC (Fig. 6a). Also, mitochondrial membrane integrity was altered and mitochondrial membrane potential was decreased in Vdac1-silenced mice as compared to control (Fig. 6c). Further, we found that Vdac1 silencing significantly increased the degenerated FJC-positive neuronal cells in DG, CA3, and CA1 subregions of the hippocampus of mice as

compared to NC (Fig. 7). These results suggested that downregulation of Vdac1 might cause depletion of ATP, mitochondrial disintegration, and increase in the number of degenerated neuronal cells in the hippocampus of scopolamine induced-amnesic and Vdac1-silenced mice.

### Discussion

Earlier, we reported that the level of 18 proteins was altered in the hippocampus of scopolamine-induced amnesic mice. Among these proteins, Vdac1 level was maximally downregulated and showed common interaction with other altered proteins as assessed by *in silico* analysis. Moreover, silencing of Vdac1 in the hippocampus of normal young mice showed

**Fig. 7** Effect of Vdac1 silencing on neurodegeneration: The panels show representative picture of Fluro-Jade C staining in DG, CA3, and CA1 subregions of hippocampus. Histogram represents increased number of FJC-positive neuronal cells in the hippocampal subregions of Vdac1-silenced mice. Arrows indicate FJC-positive cells. Data are presented as mean  $\pm$  SEM of three animals in each group. Statistically significant difference is represented as one ( $p < 0.05$ ) and two ( $p < 0.01$ ) asterisks in Vdac1 siRNA group as compared to NC



impairment in recognition memory. These findings suggested that downregulation of Vdac1 in the hippocampus might be involved in memory impairment during scopolamine-induced amnesia and Vdac1 silencing in mice. Vdac1 is an energy transporter protein present on outer mitochondrial membrane. So, its downregulation might be affecting its associated function in the hippocampus leading to memory impairment. Therefore, in the present study, we explored Vdac1 downstream function during scopolamine-induced amnesia.

As we observed downregulation of Vdac1, this might impair ATP/ADP transportation leading to depletion of ATP level in the hippocampus of scopolamine-induced amnesic mice. Earlier, it has been reported that cannabinoid induces memory loss by impairing the mitochondrial function and energy metabolism [12]. Reduced ATP level is detrimental to the cellular homeostasis and might trigger oxidative damage, disruption of mitochondrial membrane potential, and ultrastructure leading to

apoptosis during amnesia [28]. It has been elucidated that oxidation of metabolites and production of ATP level are altered in Vdac1-deficient animals [29] and cause defects in energy production pathway and structural and functional integrity of mitochondria [30, 31]. A recent study showed that metformin, which is used as a drug to reduce glucose level in type 2 diabetic patients, acts as energy disruptor and causes loss of mitochondrial integrity through disorganization of cristae [32]. We noted decreased mitochondrial membrane potential, disorganized cristae, and increased small mitochondria in the hippocampus of scopolamine-induced amnesic mice. Such damage to cristae, disruption of membrane potential, and more defective small size mitochondria might cause apoptosis and neurodegeneration [33]. It has been suggested that deregulation of fission, a process of mitochondrial dynamics, takes place by which small-sized mitochondria are accumulated during the apoptotic process [34]. In addition, we also observed very high calcium levels in

mitochondria of scopolamine-induced amnesic mice. This accumulation of calcium in the mitochondria might cause blackish appearance as observed in transmission electron micrograph of scopolamine-induced amnesic mice. An earlier report has shown that calcium precipitates with phosphate and sequesters in the form of granules which give blackish appearance to mitochondria [35, 36]. Accumulation of calcium in the mitochondria induces cytochrome c release and triggers apoptosis [37].

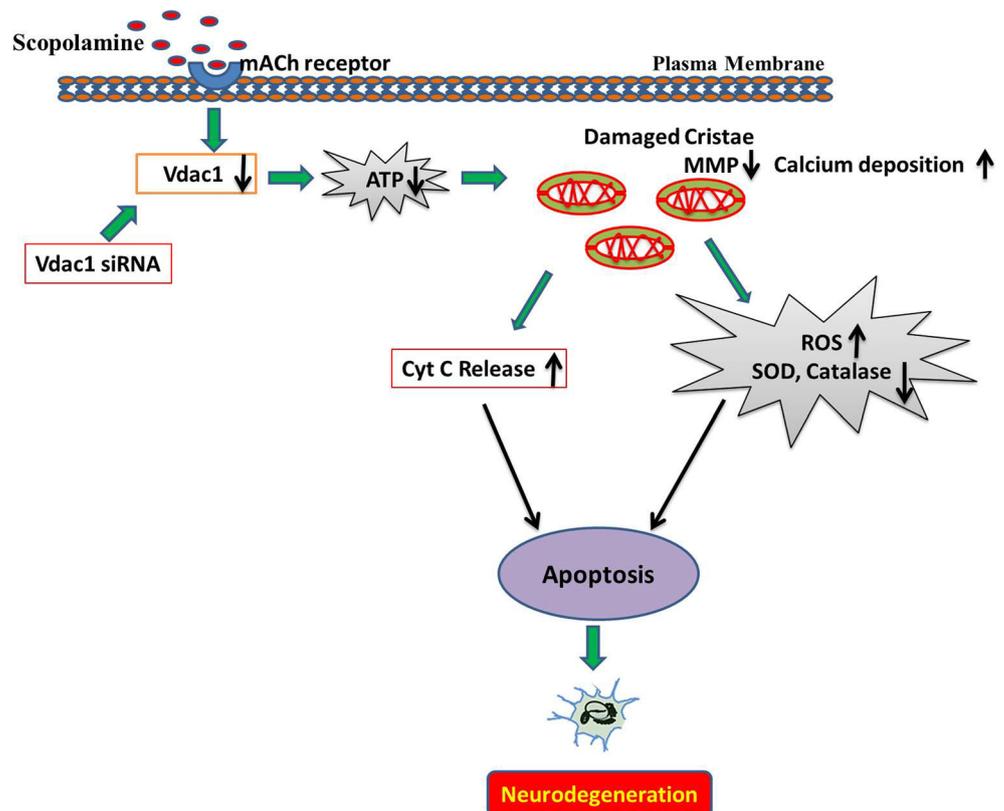
As mitochondria are the major source of ROS, defects in mitochondria raise ROS level, release of cytochrome c to cytosol, and reduction of antioxidant enzyme activity, leading to mitochondria-mediated apoptosis during aging and neurodegenerative diseases [38–40]. Similarly, we also found increased ROS generation and reduced antioxidant enzyme activity in the hippocampus of amnesic mice. These results suggest that depletion of ATP, elevation of ROS level, and reduction of antioxidant enzyme activity might trigger mitochondria-mediated apoptosis leading to neurodegeneration in the hippocampus of amnesic mice.

Therefore, in the later part of the study, we checked the molecular markers of apoptosis. We found that the cytosolic cytochrome c level was increased, pro-apoptotic proteins (Bax, Bad, caspase 3) were upregulated, and anti-apoptotic protein Bcl-2 was downregulated in the hippocampus of scopolamine-induced amnesic mice. Interaction of Bcl-2 with Vdac1 regulates mitochondrial pore complex and membrane potential, thus preventing the release of cytochrome c and

apoptosis [41–43]. Our study suggests that downregulation of Vdac1 and Bcl-2 might affect their interaction to prevent apoptosis and might help in apoptosis. Moreover, loss of mitochondrial membrane potential causes pore formation and increases release of cytochrome c leading to apoptosis. Cytochrome c further forms apoptosome and activates the caspases which lead to DNA fragmentation and apoptosis, resulting in neurodegeneration. In addition, scopolamine-induced apoptosis also increased the FJC-positive degenerated neuronal cells in DG, CA3, and CA1 subregions of the hippocampus of amnesic mice and thus increased neurodegeneration. An earlier report has also demonstrated that dopamine administration causes disruption in mitochondrial membrane potential through downregulation of Vdac1 and subsequently neurodegeneration in human neuroblastoma cells. However, overexpression of Vdac1 is rescued from the toxic effect of dopamine [15]. These experimental evidences suggest that downregulation of Vdac1-mediated mitochondrial disintegration causes apoptosis and neurodegeneration in the hippocampus of scopolamine-induced amnesic mice.

Further, to check whether Vdac1 downregulation is involved in neurodegeneration during scopolamine-induced amnesia, we silenced Vdac1 expression in the hippocampus of normal young mice and observed reduction in total ATP level, mitochondrial membrane potential, and increased FJC-positive degenerated neuronal cells in the hippocampal subregions. Loss of membrane potential of mitochondria leads to

**Fig. 8** Proposed mechanism for altered energy metabolic pathway in the hippocampus of scopolamine-induced amnesic mice: Vdac1 downregulation causes depletion of ATP level, mitochondrial disintegration, ROS elevation, and reduction in antioxidant enzyme activity leading to apoptosis and neurodegeneration in the hippocampus of scopolamine-induced amnesic and Vdac1-silenced mice



disintegration which subsequently triggers molecular events of apoptosis. Earlier, Arif et al. [44] have demonstrated that Vdac1 silencing in different cancer cell lines inhibits cell growth through decrease in the mitochondrial membrane potential and ATP level. Deregulation of Vdac-mediated mitochondrial dysfunction is also associated with neurodegenerative diseases and cancer [45]. However, our findings suggest that downregulation of Vdac1 leads to depletion of ATP, mitochondrial disintegration, apoptosis, and neurodegeneration in the hippocampus of scopolamine-induced amnesic mice. In addition, hippocampal Vdac1-silencing also showed ATP depletion, mitochondrial disintegration, and neurodegeneration leading to impairment in recognition memory of normal young mice similar to scopolamine treatment.

Taken together, our present and previous findings show that downregulation of the energy transporter protein Vdac1 and impairment of its downstream function resulted in the depletion of ATP level, mitochondrial disintegration, and increased number of degenerated neuronal cells in the hippocampus of scopolamine-induced amnesic mice. Further, hippocampal Vdac1-silencing in normal young mice showed reduction in ATP level and mitochondrial membrane potential, which in turn increased the number of degenerated neuronal cells. Thus, our findings suggest that downregulation of Vdac1 is associated with ATP depletion and mitochondrial disintegration leading to neurodegeneration in the hippocampus of scopolamine-induced amnesic and Vdac1-silenced mice as presented in Fig. 8. Further studies are warranted to elucidate energy-dependent calcium transport in mitochondria during scopolamine-induced amnesia. This will establish the role of mitochondria in hippocampal neurodegeneration and help in understanding the mechanism of energy impairment in neurodegenerative diseases.

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## Compliance with Ethical Standards

**Conflict of Interest** The authors declare that they have no conflict of interest.

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