Marine Cyclic Dipeptide Cyclo (L-Leu-L-Pro) Protects Normal Breast Epithelial Cells from tBHP-induced Oxidative Damage by Targeting CD151

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ABSTRACT
Background: Oxidative stress plays a key role in breast carcinogenesis. Cyclo (L-Leu-L-Pro) (CLP) is a homodetic cyclic dipeptide with 2,5-diketopiperazine scaffold isolated from marine actinobacteria. This study aimed to evaluate the protective activity of CLP and linear - (L-Leu-L-Pro) (LP) from tert-butyl hydroperoxide (tBHP)-induced damage using normal breast epithelial cell line model (MCF-12A).

Methods: The cytoprotective activity was evaluated by detecting the changes in intracellular ROS, mitochondrial superoxide, hydroxyl radical, hydrogen peroxide, and lipid peroxidation detection assays as well as cytotoxic assays of MTT, LDH assays and phase contrast microscopy. Genoprotective activity was evaluated by (Apurinic/Apyrimidinic) AP site, alkaline Comet, and 8-hydroxy-2-deoxyguanosine assays.

Results: The marine cyclic peptide, CLP, significantly protected MCF-12A cells by scavenging tBHP induced intracellular ROS such as super oxide, hydroxyl radicals and hydrogen peroxide, and by reducing the cytotoxicity and genotoxicity effect compared to LP. Moreover, the results showed that CD151 gene silencing by shRNA significantly reduced the overexpression of CD151, tBHP-induced ROS generation, cytotoxicity and genotoxicity in MCF-12A cells. The overexpression of CD151 caused increased levels of cytochrome P450, but was reduced following the application of CD151shRNA and CLP which led to elevated levels of intracellular ROS.

Conclusion: In the present study we noticed that CD151 gene silencing by shRNA and treatment with CLP have similar effects on reducing the intracellular ROS. This study uncovers the protective activity of CLP against a CD151-mediated oxidative stress-induced cellular damage. Our observations suggest that the anti-stress and anti-inflammation properties of CLP might have implications in cancer and are worth testing in cancer cell lines and tumor cells.

Introduction
Oxidative stress plays a key role in breast carcinogenesis. The prooxidant, tert-butyl hydroperoxide (tBHP), is extensively used as an exogenous stress inducer. tBHP mimics oxidative stress by resembling products of oxidative stress-mediated lipoperoxidation and has extensively been used as an inducer of ROS-mediated oxidative stress. Typically, two pathways, cytochrome P450 (CYP), and glutathione oxidase are involved in tBHP metabolism, and tBHP metabolites are known to reduce mitochondrial function by inducing oxidative stress. tBHP also induces oxidative stress through NF-kB pathway. Moreover, tBHP is a well-studied oxidative stress-mediated tumor promoter. The metabolites of tBHP are known to promote free radical
mediated carcinogenesis in skin cells, epidermal cells, fibroblasts and keratinocytes. Also, it promotes carcinogenesis by inducing the expression of oncoproteins in epidermal cells, breast cells, and promote lung tumors. TBP promotes cytotoxicity and genotoxicity in various cell lines. TBP-mediated oxidative stress triggers EGFR activation by promoting phosphorylation at its tyrosine residues which causes cellular transformation of epithelial cells.

Our previous study demonstrated that CD151, a member of a well-known multicyclicity-associated gene family, is associated with EGFR signaling. TSP-15, a human CD151, is linked with ROS generating dual oxidase. A study has reported that ROS generation system requires tetraspanin and they co-occur during evolution. Another study reported that tetraspanins are essential in the induction of H2O2 by dual oxidase via forming a complex at the cell surface. CD151 is one of the oncogenic members of the tetraspanin family that participates in cancer progression and metastasis by associating with ROS generating proteases, signaling enzymes, GPCRs, cadherins, and proteoglycans. Recently, we demonstrated that CD151 interacts with EGFR in breast cancer cells. EGFR is involved in the regulation of oxidative stress. Using the Chip assay, Chen et al. demonstrated that EGF/EGFR complex directly binds to the promoter of cytochrome P450 (CYP). The role of members of cytochrome P450 (CYP) has been well reported in the conversion of metabolites into potentially toxic compounds which can damage cells by forming DNA/protein adducts or by generating ROS. CYP mediates cytotoxicity and cell death induced by toxic compounds in epithelial cells and drug induced cell death by increasing the LDH leakage in hepatocytes, lipid peroxidation dependent cytotoxicity and apoptosis as well as arachidonic acid induced cytotoxicity and apoptosis. CYP genes are controlled by various regulatory networks and microRNAs, receptors and transcriptional factors at basal levels, heat shock proteins by stabilization, protein-protein interaction and organization in the lipid membranes. We speculate that CD151 may be associated with TEM establishment to facilitate cytochrome P450 mediated ROS generation in TBP-induced cytotoxicity in MCF-12A cells.

Natural products prevent disease progression by scavenging ROS and enhancing the body’s natural defense by inhibiting lipid peroxidation or direct interaction with key stress-related signaling molecules. Moreover, they can also control the initiation of carcinogenesis by protecting against DNA damage. Marine secondary metabolites find clinical applications owing to their antioxidant or protective activity. Actinobacteria contribute to 70% of the known drugs. Novel peptides with marine sources are under clinical trials due to their antibacterial, anti-inflammatory, and anti-cancer properties.

Cyclic dipeptides are modest peptide derivatives, usually present in nature and are normally more stable and potent than other cyclic peptides in terms of drug efficacy. Cyclic dipeptides with different biological activities were isolated from marine organisms. They are known to exhibit antiinflammatory, antitumor, antiviral, and antibacterial activities. For instance, cyclo (D-Tyr-D-Phes) exhibits antioxidant and anti-cancer properties. Cyclo(His-Pro) was found to activate caspase-3 and enhance poly(ADP-ribose) polymerase (PARP) cleavage and DNA fragmentation. The cyclic dipeptides, azonazine, cyclo (His-Pro), 2,5-diketopiperazines, petrocidin A are well known for targeting cancer cells. Nilov et al. (2018) demonstrated that cyclo (L-Ala-L-Ala) and cyclo (L-Ala-D-Ala) had sensitise DNA damaging effects of chemotherapeutics in drug resistance cancer cells.

Cyclo(L-Leu-L-Pro) (CLP), a homodetic cyclic peptide belongs to the class of 2,5-diketopiperazines, isolated from marine actinobacteria strain marine Streptomyces, sponge and halobacterium. It is noted for vast biological actions including antimicrobial, antifouling, antioxidant, cytotoxic, anti-mutagenic activities, anti-carcinogenic and anti-migratory potential of TNBC cells. CLP's biological role was reported as a nutrient by the human metabolic database, although cytoprotective activity was not observed. Therefore, we evaluated CLP's cytoprotective and genoprotective activities against TBP-induced damages in MCF-12A cell line model targeting CD151-EGFR signaling pathway.

**Methods**

**Culture and Treatment**

MCF-12A cells were obtained from ATCC and maintained in DMEM medium containing 10% FBS at 37 °C in an incubator under 5% CO2 atmosphere. To evaluate radical scavenging activity, MCF-12A cells (1x10^4 cells/well) in a 96-well plate were pretreated overnight with TBP (0.25mM) and then with CLP, ascorbic acid (AA) and linear L-Leucyl-L-Proline (LP) (0, 20, 40, 60, 80 and 100 μM) for 24 hours.

TBP, CLP, LP and AA were prepared in 0.1% DMSO in DMEM. To evaluate cytoprotective and genoprotective activities, MCF-12A cells were pretreated with CLP (0, 20, 40, 60, 80 and 100 μM) overnight and then with TBP (0.25 mM) for 24 hours.

The concentration of CLP that we used was based on our previous studies on TNBC’s. AA and LP were used as positive controls to compare the radical scavenging, cytoprotective and genoprotective activities of CLP.
Intracellular ROS detection assay

The intracellular ROS was computed using DCF-DA by cellular ROS detection assay kit as per the manufacturer’s instructions (Abcam, Cambridge, USA, ab113851). Briefly, the cell samples were rinsed with PBS, and stained with DCF-DA (20 μM) for 30 min. After staining, the samples were washed again with PBS and the fluorescence emitted by the cells was measured at the Ex/Em wavelength 485/535 nm using a spectrofluorometer. The tBHP-induced intracellular ROS scavenging ability was reported as percentage of tBHP treated control.13

Mitochondrial superoxide (O2−) detection assay

The mitochondrial superoxide was measured with a mitochondrial superoxide detection assay kit according to the manufacturer’s instructions (Abcam, Cambridge, USA, ab219943). The cells were rinsed with PBS and incubated with 100 μL of MitoROS-580 staining solution for 1 hour. Next, the fluorescence intensity was measured as outlined above, at the Ex/Em wavelength 540/590 nm, and tBHP-induced mitochondrial superoxide scavenging activity was expressed as percentage of tBHP treated control.14

Mitochondrial hydroxyl radical detection assay

The mitochondrial hydroxyl radicals were measured using a mitochondrial hydroxyl radical detection assay kit (Abcam, Cambridge, USA, ab219931). The cells were incubated with assay buffer (100 μL) and OH580 staining solution (100 μL) for 1 hour at 37°C after rinsing with PBS. Then, the fluorescence intensity was measured as outlined above. The scavenging ability of mitochondrial hydroxyl radical was expressed as percentage of tBHP treated control.

Intracellular hydrogen peroxide assay

Intracellular hydrogen peroxides was determined using an H2O2 cell-based assay kit (Caymen, Michigan, USA, Cat# 600050). Next, the cells were rinsed with PBS and 10 μL of each assay buffer and enzyme reaction solution containing horseradish hydrogen peroxidase and hydrogen peroxide detector (ADHP) was added. Within 5 minutes, the fluorescence emitted by the cells was measured at Ex/Em of 530/590 nm. The tBHP-induced intracellular hydrogen peroxide scavenging activity was expressed as percentage of tBHP control.15

Lipid peroxidation (MDA) assay

The lipid peroxidation was evaluated using a lipid peroxidation (MDA) assay kit (Abcam, Cambridge, USA, ab118970). Two hundred μL of the treated cell lysis solution was added to each well, homogenized by placing on ice, and centrifuged at 13,000 rpm for 10 minutes. Then the cell lysate (200 μL) was incubated with 200 μL of TBA reagent for 1 h at 95°C. Next, the absorbance was measured at 532 nm using a microplate reader. The inhibition of tBHP-induced lipid peroxidation was expressed as percentage of tBHP treated control.16

MTT assay

The treated cells were incubated with MTT reagent (5 mg/mL of PBS) for 2 hours. Next, the formazan crystals formed were solubilized by adding 200μL of DMSO. The absorbance was computed using an ELISA reader at 595 nm. The cytotoxic effect was expressed by comparing cytotoxicity with tBHP treated control.

Lactate dehydrogenase (LDH) assay

Treated cells were rapidly homogenized in LDH assay buffer on ice and cell debris was removed by sedimentation at 10,000 rpm for 15 minutes at 4°C. The soluble fraction was used for measuring LDH activity by LDH activity assay Kit (Sigma-Aldrich, USA). The protective effect was expressed by comparing the LDH levels with tBHP treated control.

Phase-contrast microscopy

Cells (3x103/well) in a 96-well plate were treated with tBHP (0.25mM) alone or in combination with CP (84.21 μM) or LP (176 μM) for 24 hours. After treatment, the cells were washed thrice with serum-free media, and images were captured at 40X resolution under the inverted phase-contrast microscope.

Detection of AP Sites

The cellular DNA was extracted from treated and untreated cells. The reaction mixture [1 μg of DNA, 200 μl of 10 mM Tris (pH 9), 15 μl of 5 M NaCl] was incubated with 30 μl of avidin-HRP for 60 min at room temperature (RT). The DNA HRP was extracted by treating with 65 μl of 1 mM DAPER for 5 minutes followed by centrifugation for 5min at 12,500 × g at 4 °C. The pellet was washed with 1.4 ml of wash buffer and suspended in 500 μl ice-cold 50 mM Na-citrate (pH 5.3) by sonication. The HRP activity was determined showing AP sites in DNA–HRP by ELISA method.

Detection of 8-hydroxy-2′-deoxyguanosine (8-OHdG)

Competitive ELISA was performed using an 8-OHdG ELISA kit, by the protocol of the Cayman Chemicals (USA). The purified DNA was subjected to enzymatic digestion using nuclease P1 at 50 °C for 1 h and with alkaline phosphatase at 37 °C for 30 minutes. The digested DNA was boiled for 10 min and placed on ice for 5 minutes. The hydrolyzed DNA was measured by reading the absorbance at 412 nm. The levels of 8-OHdG were measured and expressed as µg of 8-OHdG/ml.

Competitive ELISA was performed using an 8-OHdG ELISA kit, by the protocol of the Cayman Chemicals (USA). The purified DNA was subjected to enzymatic digestion using nuclease P1 at 50 °C for 1 h and with alkaline phosphatase at 37 °C for 30 minutes. The digested DNA was boiled for 10 min and placed on ice for 5 minutes. The hydrolyzed DNA was measured by reading the absorbance at 412 nm. The levels of 8-OHdG were measured and expressed as µg of 8-OHdG/ml.
Alkaline Comet assay
The treated and untreated cells were plated on a slide previously coated three times with low melting agarose (0.75%). The slide was then placed in lysing solution at 4 °C for 1 hour followed by electrophoresis by setting the voltage to 20 V for 20 min. The slide was then soaked in neutralizing buffer followed by ethanol for 5 minutes. Then the slide was stained with ethidium bromide (40 μl). The tail length was measured, and the olive tail movement (OTM) was computed as: (head mean) x tail % DNA/100.

Cell death by Annexin V ELISA method
After treatment, cell samples were washed with PBS, and apoptotic cells were determined using Annexin V ELISA method. Briefly, cells were incubated with 50 μl of annexin V antibody at 25 °C for 1 hour. The apoptosis rate was determined as per the manufacturer’s instructions (Abcam, USA).

Cell-based ELISA for quantification of Cd151
After treatment, the cells were washed with PBS and incubated with CD151 primary antibody overnight at 4 °C. After washing off the unbound primary antibody, cells were treated with HRP-conjugated secondary antibody specific to CD151 (100 μL) and incubated at 37 °C for 1h. Following washing the unbound secondary antibody, the cells in each well were incubated with TMB One-Step Substrate Reagent (100 μL) at 37 °C in the dark for 30 min. After stopping the reaction by adding stop solution (50 μL), the absorbance was computed using a microplate reader set to 450 nm.

CD151 gene silencing using shRNA
After overnight seeding of MCF-12A cells (5x104) in a 6-well plate, the cells were transfected with CD151 shRNA overnight as described by Gayatri et al. After transfection, the cells were treated with tBHP (0.25 mM) for 24 hours and intracellular ROS was measured using DCF-DA using cellular ROS detection assay kit, extracellular LDH using LDH assay kit and DNA damage using alkaline comet assay as described above.

CD151 overexpression in MCF-12A cells
After overnight seeding of MCF-12A cells (5x10⁴) in a 6-well plate, the cells were transfected with 10 μg untagged PrecisionShuttle mammalian plasmid encoding CD151(CAT#: SC319271, OriGene, USA) using TurboFectin Transfection Reagent (TF81001, OriGene) following the manufacturer’s protocol. After 24 hours of transfection, the cells were treated with CLP and LP for 24 hours. Then CD151 expression levels were determined by cell-based ELISA, and intracellular ROS, LDH and DNA damage were determined using their specific methods outlined above.

Protein-protein interaction by ClusPro and PyDOCK webserver
The ClusPro is a widely used fully automated protein-protein docking server, which uses direct docking of two interacting proteins (https://cluspro.bu.edu). It uses PIPER16 for the rigid body docking program based on the Fast Fourier Transform (FFT) correlation approach. PyDOCK web server, which uses rigid-body docking orientations generated by FTDock and evaluation is based on electrostatics, de-solvation energy and limited van der Waals interactions. Using ClusPro and PyDOCK webserver, the interaction between CD151 and cytochrome p450 was predicted.

Measurement of cytochrome P450 levels by ELISA
After treatment, cell samples were collected by scraping into ice-cold PBS, sedimented by centrifugation at 3000 rpm for 5min. The cell pellet was suspended in a cell lysis buffer, and the cells were lysed by pipetting up and down 5-10 times. Then the lysate was cleared by removing cell debris using centrifugation at 10,000 rpm for 5min at 4 °C. Next, cytochrome P450 levels in the cell lysate were determined by human quantitative sandwich cytochrome P450 ELISA kit (CUSABIO Technology LLC, USA) using cytochrome p450 antibody precoated microplate following the manufacturer’s instructions. After removing the unbound substances by washing, the plate was incubated by adding cytochrome P450 antibody conjugated to biotin, followed by HRP conjugated avidin. Finally, the color was developed by incubating with TMB substrate solution. After stopping the color development using a stop solution, the optical density was measured at 450 nm using a microplate reader.

Statistical analysis
The data from the experiments were analyzed statistically and represented graphically using Microsoft Excel. To ensure the consistency of the results, each experiment was performed three times, and values were represented as mean ± SD (n=3). One-way variance analysis was performed to calculate the means of the dependent variable (response) and independent variable (concentration) using NumPy (v 1.1.2) and Google Colab. Student t-test was performed to compare mean values. The statistical significance was set with a confidence level of 95% and p value <0.05. IC₅₀ values were calculated using MS Excel ProPlus (Version 2016).

Results
Intracellular ROS scavenging ability of cyclo(L-Leu-L-Pro)
tBHP (0.25mM) stimulates augmented oxidative stress in MCF-12A cell line which helps to understand the protective role of CLP in regulating CD151-mediated oxidative stress. The scavenging
activity of CLP applied at increasing concentrations (0–100 µM) against tBHP-induced intracellular ROS production was compared with that of linear dipeptide (LP) and radical scavenger, AA (Figure 1a). The proxidant tBHP triggered considerable oxidation of DCFH to DCF and enhanced intracellular fluorescence intensity in controls. However, CLP treatment decreased the tBHP-induced fluorescence intensity in a dose-dependent manner. Cyclic dipeptide CLP diffuses into the cells through the cell membrane, where it may prevent the generation of ROS required to oxidize intracellular DCFH2 to the fluorescent DCF. tBHP stimulated intracellular ROS generation, scavenge by treatment with CLP. The percent of radical scavenging activity are 17.8±2.3, 32.6±2.2, 41.7±2.3, 54.2±3.4 and 70.2±3.3% with CLP, and the positive control AA exhibited 21.6±2.5, 39.4±2.3, 52.4±2.4, 59.3±3.4 and 75.3±3.2% of radical scavenging activity, while for another positive control LP, the radical scavenging activity was 8.2±1.2, 12.2±1.3, 21.2±1.3, 32.2±2.4 and 38.1±2.4% at 20, 40, 60, 80 and 100 µM, respectively. The IC₅₀ values of CLP, AA and LP for scavenging intracellular ROS were 70.6, 130 and 60.5 µM, respectively. (Figure 1a) This study demonstrates the concentration-dependent intracellular ROS scavenging activity of the CLP like ascorbic acid, which is higher than that of LP.

Superoxide ion (O₂⁻) is the most acceptable candidate for causing oxidative stress damage to the cell. The oxidative stress causes the escape of electrons from mitochondrial ETC and directly generates the O₂ by reacting with molecular oxygen. The present study observed that tBHP-induced intracellular O₂⁻ was scavenged by 20.8±2.3, 42.6±2.3, 52.5 ±3.4, 63.4±3.3 and 85.4±4.3% with CLP and 24.6±2.2, 46.3±2.4, 56.2±3.3, 68.5±4.3 and 89.4±4.4% with AA and 10.8±1.8, 26.3±2.4, 33.2±3.4, 43.9±4.1 and 59.6±4.8% with AA at 20, 40, 60, 80 and 100 µM as shown in Figure 1b. The IC₅₀ of the CLP was 56.0 µM, LP was 86.4 µM and AA was 50.7 µM, indicating the potential O₂⁻ scavenging ability of CLP close to AA. Among the reactive oxygen centered species, hydroxyl radicals (OH) cause serious damage to proteins, polyunsaturated fatty acids and DNA and are implicated in radical-mediated pathology. The intracellular OH scavenging action of CLP was related to intracellular antioxidant ability. The results depicted in Figure 1c show that tBHP- induced intracellular OH scavenging activity of the CLP was 19.2±2.3, 28.3±2.3, 41.4±2.3, 55.4±3.4 and 81.5±4.4% whereas AA showed 21.4±2.3, 32.2±2.3, 48.4±3.3, 62.4±3.4 and 82.8±4.4% and LP exhibited 9.8±1.8, 22.6±2.1, 30.5±2.9, 39.4±3.1 and 56.4±3.8% at 20, 40, 60, 80 and 100 µM, respectively. The IC₅₀ of CLP was 66.3 µM, LP was 93.1 µM and AA was 60.68 µM, indicating potential intracellular OH scavenging activity of CLP, similar to AA. One of the interesting developments in the free radical-mediated pathology is the formation of hydroxyl radicals by the interaction between O₂ and H₂O₂. Hence, we evaluated the intracellular H₂O₂ scavenging activity of the CLP. The study observed that tBHP-induced intracellular H₂O₂ scavenging activity with the CLP was 16.9±2.3, 34.6±2.3, 45.7±3.3, 56.2±3.4 and 72.1±4.3%, whereas 21.6±2.5, 38.4±2.3, 55.4±3.5, 62.3±3.4 and 76.4±4.3% with AA, and 8.6±1.2, 16.6±1.8, 28.5±2.1, 36.3±2.9 and 52.1±3.2% with LP at 20, 40, 60, 80 and 100 µM, respectively, indicating that intracellular H₂O₂ scavenging activity of CLP is similar to that of AA as shown in Figure 1d. The significant radical scavenging ability may recommend CLP for the treatment of stress associated cancers.

Cytoprotective activity of CLP against tBHP-mediated cytotoxicity

The current study evaluated the protective ability of the CLP on tBHP stimulated lipid peroxidation in MCF-12A cells. Malondialdehyde (MDA) is the most widely cited lipoperoxidation product instigating from unsaturated lipids of membranes during oxidative stress. The protective effect of CLP was confirmed by lipid peroxidation assay. The percentage of inhibition of tBHP-induced lipid peroxidation by serial concentrations of CLP increased in a dose dependent manner to a maximum of 89.8±5.3%. This figure stood at 91.8±7.3 % with AA and 57.8±4.1% with LP (Figure 2a). This study also evaluated the impact of the CLP on tBHP-mediated cytotoxicity in MCF-12A cells. The results show that percent of viable cells increased with increasing concentration. Treatment with CLP showed 68.6±3.2% viability, which indicates that CLP efficiently enhanced tBHP treated MCF-12A cells viability. Treatment with positive controls like AA and LP showed maximum viability of 74.9±4.3 and 33.4±2.8%, respectively (Figure 2b). Further, the CLP’s cytoprotective activity was evaluated by measuring LDH leakage from cell, an indicator of the cell membrane damage. The CLP efficiently blocked tBHP-induced LDH release from MCF-12A cells. The percentage of inhibition of LDH release increased with increasing concentration from 20 to 100 µM. The maximum inhibition of LDH release was measured with CLP (62.8±3.4%), AA (71.5±4.3%) and LP (28.4±2.1%) with IC₅₀ of 82.9 µM, 72.7 µM and 174 µM, respectively (Figure 2c). Further, the effect of CLP on tBHP-induced morphology of MCF-12A cells was monitored under phase-contrast microscopy (Figure 2d). The results show that tBHP (0.25 mM) induced morphological changes in MCF-12A cells. However, pretreatment with CLP (84.2 µM) significantly protected the MCF-12A cells from tBHP-induced morphological changes better than LP (174 µM) and AA (76.6 µM). These results suggest the potential cytoprotective activity of the CLP. Table 1 shows the data related to
Figure 1. Protective effect of CLP from tBHP-induced intracellular ROS levels in MCF-12A. Cells (1x10^4/well) in a 96-well plate were pretreated overnight with tBHP (0.25 mM) and subsequently with serial concentrations of CLP (0, 20, 40, 60, 80 and 100 µM) for 24 h. Only tBHP treated cells were served as untreated control. The fluorescence generated upon conversion of DCFH2 to DCF due to the CLP scavenging activity was measured at the Ex/Em wavelength 485/535 nm using a spectrofluorometer and expressed as % control. The percent control is defined as: Fluorescence change in CLP treated cells / tBHP treated cells x 100. The same concentrations of ascorbic acid (AA) and linear dipeptide, L-Leucyl-L-Proline (LP) were used as positive controls. The graphs show the protective effect of CLP on tBHP-induced a) intracellular ROS production, b) mitochondrial superoxides, c) mitochondrial hydroxyl radicals and d) intracellular hydrogen peroxide. The cumulative data of each assay were collected from three independent experiments and shown as means ±SEM (n=3).

Figure 2. Protective activity of CLP against tBHP-induced cytotoxicity in MCF-12A cells. Cells were pretreated overnight with CLP (0, 20, 40, 60, 80 and 100 µM), subsequently with tBHP (0.25mM) and incubated for 24 h. Only tBHP treated cells were served as untreated control. AA and LP were used as positive controls under similar concentrations. a) Protective effect of CLP on tBHP-induced lipid peroxidation in MCF-12A cells. b) Protective effect of CLP on the tBHP-induced cytotoxicity of MCF-12A cells. The results were expressed as % viability of tBHP treated control. c) Protective effect of CLP on tBHP induced cell membrane damage. The results were expressed as % inhibition of LDH leakage. d) Protective effect of CLP on tBHP induced morphological changes by phase-contrast microscopy. The cumulative data of each assay were obtained from 3 independent experiments and shown as means ±SEM (n=3).
Table 1. Cytoprotective effect of CLP on MCF-12A cells

| No | Concentration (µM) | Inhibition of lipid peroxidation (%) | Cell viability (%) | LDH leakage (%) |
|----|-------------------|-------------------------------------|-------------------|----------------|
|    | CLP+ tBHP         | AA+ tBHP                            | LP+ tBHP          | CLP+ tBHP      | AA+ tBHP      | LP+ tBHP      |
| 1  | 20                | 48.8±3.3                            | 55.8±4.3          | 20.1±1.9       | 4.5±0.9       | 5.2±0.9       |
| 2  | 40                | 52.2±4.2                            | 62.2±5.2          | 26.2±2.1       | 10.8±1.1      | 14.4±1.2      |
| 3  | 60                | 68.6±4.1                            | 72.6±6.1          | 39.6±2.9       | 26.4±1.9      | 32.8±2.3      |
| 4  | 80                | 85.2±5.2                            | 88.2±6.2          | 51.2±3.1       | 42.8±2.1      | 49.9±3.3      |
| 5  | 100               | 89.8±5.3                            | 91.8±7.3          | 57.8±4.1       | 68.6±3.2      | 74.9±4.3      |

Cytoprotective activity of CLP (Cyclo(L-Leu-L-Pro)) was measured in terms of inhibition of lipid peroxidation, cell viability and inhibition of LDH leakage against tBHP-induced cell damage in MCF-12A cells. AA (Ascorbic acid) and LP (L-Leucyl-L-Proline) were used as control. The results were expressed in percent of inhibition of lipid peroxidation, percent cell viability and percent of inhibition of LDH release. Each experiment was performed three times, and values were represented as mean ± SD (n=3).

In vitro comet assay is a sensitive and frequently used method to confirm the genoprotective activity. Therefore, DNA damage protecting activity of CLP, AA and LP in tBHP treated MCF-12A cells was assessed using in vitro Comet assay, and the results...
Table 2: Genoprotective effect of CLP on MCF-12A cells

| No | Concentration (µM) | AP sites/10^5bp | 8-OHdG (µg/ml) | Decrease in OTM (%) | Dead cells (%) |
|----|-------------------|----------------|----------------|---------------------|---------------|
|    | CLP+ tBHP | AA+ tBHP | LP+ tBHP | CLP+ tBHP | AA+ tBHP | LP+ tBHP | CLP+ tBHP | AA+ tBHP | LP+ tBHP | CLP+ tBHP | AA+ tBHP | LP+ tBHP |
| 1  | 20        | 38±4         | 34±4         | 32±3        | 3.5±0.4       | 2.5±0.1   | 2.1±0.1    | 22±1.8     | 24±1.1   | 12±0.9     | 65±4.3    | 76±4.1   | 71±3.2   |
| 2  | 40        | 32±3         | 30±3         | 25±2        | 2.9±0.1       | 1.9±0.09  | 1.5±0.09   | 34±1.9     | 39±2.4   | 21±1.1     | 53±3.2    | 60±3.6   | 52±3.1   |
| 3  | 60        | 26±2         | 24±2         | 18±2        | 2.1±0.1       | 1.1±0.06  | 0.9±0.02   | 53±2.7     | 62±2.8   | 33±1.4     | 41±2.1    | 59±2.3   | 35±2.9   |
| 4  | 80        | 18±2         | 16±2         | 9±2         | 1.7±0.03      | 0.7±0.03  | 0.5±0.01   | 64±2.8     | 74±2.9   | 41±2.1     | 33±2.1    | 47±3.1   | 43±1.8   |
| 5  | 100       | 12±2         | 10±2         | 5±2         | 0.8±0.02      | 0.3±0.01  | 0.2±0.01   | 76±3.3     | 83±3.1   | 53±2.9     | 21±1.9    | 36±2.2   | 31±1.6   |

The results showed that the number of dead cells decreased with CLP was 31.6±1.6% and with positive controls AA (36.3±2.2%) and LP (31.6±1.6%), respectively with an increase in concentration to 100 µM compared to tBHP control, which is considered as 100%. Table 2 lists the data of genoprotective effect of CLP, LP or AA in MCF-12A cells.

Figure 4. CLP protects tBHP treated MCF-12A cells by targeting CD151.

CLP-induced CD151 levels were reduced with CLP and CD151shRNA (a). MCF-12A cells were pretreated overnight with CLP (70.6 µM) and ascorbic acid (60.5µM) or transfected with CD151shRNA, subsequently with tBHP (0.25 mM) and incubated for 24 h. Untreated cells were served as control. The expression of CD151 was determined by cell-based ELISA assay. The results were expressed in OD at 450nm. After silencing the CD151 gene using shRNA, the intracellular ROS (b), extracellular LDH levels (c) and OTM (d) were evaluated in tBHP treated MCF-12A cells. Means ±SEM (n=3), p<0.05. MCF-12A cells (5x10⁴) in a 6-well plate were transfected with 10 µg untagged Precision Shuttle mammalian plasmid encoding CD151 using Turbo Fectin Transfection Reagent. After 24h of transfection, cells were treated with CLP or LP for 24h. The intracellular ROS was measured using DCF-DA by cellular ROS detection assay kit (e), extracellular LDH using LDH assay kit (f) and DNA damage by alkaline comet assay (g) as described above. The cumulative data of each assay were obtained from 3 independent experiments, and results were shown as means ±SEM (n=3),* p<0.05

Genoprotective activity of CLP (Cyclo(L-Leu-L-Pro)) was measured in terms of AP sites/105bp, inhibition of 8-OHdG, decrease in OTM (%) and percent of dead cells against tBHP-induced cell damage in MCF-12A cells. AA (Ascorbic acid) and LP (L-Leucyl-L-Proline) were used as control. Each experiment was performed three times, and values were represented as mean ± SD (n=3).
Pathway analysis of CD151 mediated t-BHP induced cytotoxicity

Kučera et al. reported that t-BHP is metabolized by cytochrome P450 and glutathione peroxidase mediated pathways. CD151-dependent tetraspanin-enriched microdomains (TEM) correlated with hepatocarcinoma. Previously, we showed that CLP reduced the CD151 expression and its interaction with EGFR. In the current study, we found that tBHP stimulated a 10-fold increase in the expression of CD151 compared to untreated control in MCF-12A cells. However, CLP and LP treatment reduced the 4.0 (p>0.05) and 2.1-folds of tBHP-induced expression of CD151, respectively, and CD151 shRNA treatment 7.1-folds (p>0.05) in MCF-12A cells (Figure 4a), but AA did not affect the CD151 expression (Data not shown). CD151 gene silencing using shRNA reduced the intracellular ROS (4.3-folds) (Figure 4b), extracellular LDH levels (6.2-folds) (Figure 4c), and OTM (8.3-folds) (Figure 4d) in tBHP treated MCF-12A cells. Further, CLP and LP treatment reduced the intracellular ROS levels 2.9 and 1.4-folds, respectively (Figure 4e), LDH levels 4.7 and 1.3-folds, respectively (Figure 4f) and OTM levels 5.2 and 1.2-folds, respectively (Figure 4g) in CD151 overexpressed MCF-12A cells, indicating that CLP protects the MCF-12A cells by targeting Cd151.

Protein-protein interactions are important for understanding cellular function and organization. To find the downstream mediator of CD151 in tBHP-induced cytotoxicity, we analyzed the interaction between CD151 and cytochrome P450, which is an important mediator of tBHP metabolism and membrane damage. By using Cluspro docking method, we interpreted that CD151 interacts with cytochrome P450 strongly (Figure 5a) with docking score of -993 kcal/mol. We have verified the results using another docking study, PyDOCK webserver (Figure 5b). The docking score of CD151 with P450 was -38.3 kcal/mol. Thus, P450 shows a high binding tendency for CD151 protein.

To further focus our investigation on the protection exerted by CLP in CD151 mediated cytotoxicity in tBHP treated MCF-12A cells, we determined the cytochrome p450 levels in MCF-12A cells treated with tBHP, CD151 shRNA and CD151 clone using quantitative sandwich ELISA (Figure 5c). The results showed that CD151 clone enhanced the cytochrome P450 levels by 8-folds, and CD151 shRNA reduced the cytochrome P450 levels by 9-folds. However, tBHP-induced cytochrome P450 levels by only 3-folds. Further, treatment with CLP and LP significantly reduced the tBHP-induced cytochrome P450 levels.

Figure 5. In Silico interaction of CD151 with P450.
a) In silico docking of CD151 (receptor) with P450 (ligand) was performed using ClusPro web server. The receptor and ligand were represented as ribbon form. The binding energy of CD151 with P450 was calculated as -990 kcal/mol. b) In silico docking of CD151 (receptor) with P450 (ligand) was performed using PyDOCK webserver. The receptor and ligand were represented as a solid model. The binding energy of CD151 with P450 was calculated as -38.3 kcal/mol. c) The impact of tBHP, CD151 shRNA and CD151 clone on cytochrome p450 levels was determined using quantitative sandwich ELISA. d) Effect of CLP and LP on tBHP-induced cytochrome p450 levels. The cumulative data of each assay were obtained from 3 independent experiments, and results were shown as means ±SEM (n=3), *p<0.05
cytochrome P450 levels (Figure 5d), but not considerably by AA (Data Not shown). These results indicate that CD151 is an upstream mediator of tBHP-induced cytotoxicity in MCF-12A cells.

Discussion

In normal physiological conditions, the homeostasis of ROS is maintained by cellular antioxidant defense system but during oxidative stress this homeostasis is lost which causes disturbance in the metabolism of free radicals and their detoxification. Excessive generation of radicals causes oxidative damage in proteins, fatty acids, and DNA which leads to inflammation and cancer. The stimulation of the antiradical mechanism is one of the most significant determinants of cytoprotective ability against oxidative stress-induced diseases.

CLP is a biologically active homodetic cyclic peptide, which we reported to exhibit significant inhibition of tBHP-induced oxidative stress in MCF-12A cell line. CLP's scavenging potential against intracellular ROS and mitochondrial superoxide and hydroxyl radicals was comparable to that of Ascorbic acid, a well-known antioxidant. This study emphasizes the significant cytoprotective as well as genoprotective activity of the CLP against tBHP-induced cellular stress in MCF-12A cells and results were compared to linear dipeptide (LP). Previously, a large amount of data pertaining to dipeptides, including aspartame (L-aspartyl-L-phenylalanine methyl ester) as well as L-alanyl-L-glutamine as cytoprotective agents was documented. Kaul et al. provided evidence suggesting that proline (Pro) has the ability to scavenge free radicals in vitro. Besides the role of proline as antioxidant, ameliorating metal toxicity has also been reported recently. Research by Phang et al. emphasized proline as a "stress substrate" and suggested it as a potential anti-cancer agent. Also, Pro has wound healing potential. In addition, Pro was reported as an activator of mTOR signal pathway in concert with leucine.

Cyclic dipeptides can bind to diverse targets due to conformationally constrained scaffolds and a vast number of tailoring enzymes. Yan et al. reported that hydrophobic amino acids and proline are critical for cyclic dipeptides' biological activity. The peptide bonds in proteins normally have a planar trans configuration. However, due to neighboring substituents' steric requirements, the cis form is more stable than the trans form when the Pro (imino acid) occurs at the bond's carboxyl-terminal side. Further, the Pro ring's restricted mobility might be related to cyclic dipeptides. The cis-trans isomerism of the N-alkyl amide bond in the Pro was found to be involved in receptor-mediated biological activity. The connection between the configuration of Pro and its physical activity might be important in explaining the mechanism of the inhibitory activity of cyclodipeptides.

Because of these properties, cyclic dipeptides consist of hydrophobic amino acids like leucine, and terminal amino acid proline are products of rational drug design. CLP significantly improved the viability of tBHP treated MCF-12A cells in a concentration-dependent manner. Further, CLP's cytoprotective effect in tBHP-induced cellular leakage of LDH was significantly high compared to other dipeptides like β-alanine-histidine and carnosine. Besides, pretreatment with CLP protected MCF-12A cells from tBHP-induced morphological changes better than LP. These results indicate the efficacy of marine natural product, the CLP as a cytoprotective agent.

Later, we studied CLP's effect on genoprotective activity against tBHP-induced stress in MCF-12A cell line. The results showed that CLP exhibited remarkable protection against DNA damage in terms of levels of 8-OHdG and tailing in comet assay. Limited reports are available on genoprotective activity of dipeptide. Thus, we have made an attempt to study the effect of CLP as a cytoprotective and genoprotective agent against stress-induced model. The present study showed that CLP is an efficient dipeptide, which can regulate cytoprotective cellular mechanism in addition to providing protection to DNA damage caused by oxidative stress.

Apart from metastatic activity, CD151-dependent TEM mediates hepatocarcinoma and liver fibrosis, and has also been proposed as CD151 potential therapeutic target of liver fibrosis. CD151 is a membrane protein that transduces intracellular signaling and regulates cellular functions. Previously, we showed that CLP reduced the CD151 expression and its interaction with EGFR. Bae et al. demonstrated that EGF induces H₂O₂ generation via EGFR in cancer cell lines. The present study observed increased levels of CD151 in tBHP treated MCF-12A cells. However, tBHP-induced CD151 levels were reduced with CD151 shRNA and CLP treatment which are comparable. However, LP did not affect CD151 expression. In addition, CD151 gene silencing decreased the tBHP- induced ROS and LDH levels in tBHP treated MCF-12A cells, indicating the involvement of CD151 in tBHP- induced cytotoxicity. In addition, CD151 gene silencing reduced the tBHP induced OTM in MCF-12A cells. Further, CLP reduced the intracellular ROS, extracellular LDH levels, and genotoxicity (OTM) in CD151 overexpressed MCF-12A cells.

High docking score in the interaction study of CD151 with cytochrome P450 by Cluspro docking method and PyDOCK web server indicated strong interaction CD151 with cytochrome P450. Further, overexpression of CD151 increased the CYP levels but reduced with CD151 shRNA with a moderate increase with tBHP. Also, treatment with CLP and
LP significantly reduced the tBHP induced cytochrome P450 levels, but not AA. Additionally, CD151 gene silencing using CD151 shRNA reduced the cytochrome P450 levels in tBHP-induced MCF-12A cells. These results indicate that CD151 is an upstream mediator of tBHP-induced cytotoxicity in MCF-12A cells and CLP protecting the MCF-12A cells by targeting Cd151.

In the present study, we have reported the cytoprotective role of CLP against various ROS types using MCF-12A cell line model. CLP increases the cellular antioxidant status in MCF-12A cells and scavenges the tBHP-induced intracellular ROS levels. This study also observed that the CLP protected MCF-12A cells from tBHP-induced lipid peroxidation, LDH leakage, and morphological changes. Moreover, a significant genoprotective activity of the CLP was observed against tBHP-induced DNA damage in MCF-12A cell line compared to positive control LP. Furthermore, CLP also reduced the tBHP-induced DNA damage by decreasing AP sites, 8-OHdG levels, and OTM in a concentration-dependent manner. There was a significant effect of CLP on expression of CD151 which can be compared to CD151 shRNA mediated gene silencing. Protein-protein interaction study by Cluspro docking method and PyDOCK web server indicated strong interaction CD151 with cytochrome P450. An increase in cytochrome P450 levels with CD151 overexpression and reduction with CD151 gene silencing by shRNA, as well as reduction with CLP and LP, indicates the involvement of CD151 gene via cytochrome P450 in tBHP-induced cytotoxicity in MCF-12A cells. These results also suggest that CLP protects the MCF-12A cells more significantly by targeting CD151 compared to its linear counterpart, LP. In conclusion, based on the intracellular ROS scavenging ability, cytoprotective, and genoprotective activities, CLP can be used as an efficient agent against oxidative damage mediated pathological diseases like cancer and inflammation.

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Ethical statements
None.

Conflict of Interest
Authors declare no potential conflicts of interest.

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