Anti-cancer potential of sea grape (*Caulerpa racemosa*) extract by altering epithelial-mesenchymal transition and pro-apoptosis proteins expression in MCF-7 breast cancer cells

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**ABSTRACT**

Breast cancer is a neoplasm that is increasing globally based on its prevalence. *Caulerpa racemosa* is a type of sea grape that has anti-cancer activity based on previous studies. We aim to determine the anti-cancer activity of *C. racemosa* extract on Michigan cancer foundation-7 (MCF-7) breast cancer cells through the mechanism of apoptosis by observing cell viability and caspase-3 expression, also as an anti-metastatic agent on expression of Snail and Vimentin. A true experimental study was carried out using a randomized post-test-only control group design. MCF-7 cells were used as subjects with the intervention of *C. racemosa* extract at varying doses. Cell apoptotic was assessed by flow cytometry and cell viability test, while anti-metastatic activity was assessed by wound healing assay. Immunofluorescence staining was used to assess the expression level of caspase-3, Snail, and Vimentin. The results showed an increased activity of cell apoptosis, according to flow cytometry and cell viability tests. Furthermore, expression of caspase-3 was found to be significantly higher in the 400 g/ml dose group and MCF-7 cell migration inhibition by decreasing Snail and Vimentin protein expression. These results indicate a “novel mechanism” of *C. racemosa* in the MCF-7 model as an anti-cancer and anti-metastatic agent to inhibit breast cancer progression.

**INTRODUCTION**

Breast cancer is the most common cancer and the leading cause of death from cancer in females worldwide [1]. Despite the unfortunate outcome that gained global attention, breast cancer is still late diagnosed until the advanced stages due to women’s negligence in performing breast self-inspection and clinical examination [2]. Poor prognosis was associated with locally advanced and metastatic breast cancer, with an average survival rate of less than 5 years. Furthermore, 10% of newly diagnosed breast cancer progressed to a metastatic stage [3]. Hence, it set a limitation on the early stage first choice of treatment by surgical tumor resection [4]. Chemotherapy served as the second option after the surgery, though it came along with adverse side effects. Not only targeting malignant cells, but these drugs are also toxic to healthy growing cells. It faces a major drawback as some tumors have developed chemotherapy resistance [3]. New approaches and anti-tumor compounds are constantly sought, and natural sources such as plants provide an abundance of promising anti-cancer drugs [5].

Apoptosis, defined as programmed cell death, is an important body mechanism for removing excess, damaged, or harmful cells [6]. Apoptosis dysregulation, which is a hallmark of cancer, disrupts the established balance of cell proliferation and cell death [7]. Numerous molecules, particularly the
Caspase signaling cascade, play roles in regulating cell apoptosis [1]. Caspase-3 is a key pro-apoptotic caspase [6]. Aberrant Caspase-3 expression is linked with several cancer types [1]. Former research displayed that high caspase-3 expression is associated with a more favorable prognosis in hepatocellular carcinoma and nonsmall cell lung cancer [8].

The process of programmed cell death is crucial in the metastatic process. Cancer cells must be able to avoid various forms of cell death to metastasis [9]. Metastasis occurs when uncontrollable growth cells infiltrate adjacent body parts or spread to other distant body organs [10]. Diagnosed with metastatic cancer is associated with the worst prognosis [11]. Multiple cell processes, including epithelial-mesenchymal transition (EMT), extracellular matrix degradation, and tumor angiogenesis have been demonstrated to be significant parts of cancer metastasis [9].

EMT is a biological process that enables mesenchymal phenotype changes in polarized epithelial cells that normally interact with the basement membrane through the basal surface [12–14]. Epithelial cells lose cell-to-cell junctions, apical-basal polarity, and epithelial markers; however, the cells acquire cell motility, a spindle-cell shape, and mesenchymal markers [14,15]. Mesenchymal markers such as Vimentin, Snail, and N-cadherin were discovered, while epithelial markers such as E-cadherin, occludins, and desmoplakin were lost during EMT [15,16]. Overexpression of Vimentin and Snail has been linked to increased tumor growth, invasion, and poor prognosis, as well as being a predictor of an aggressive tumor phenotype [17,18].

*Caulerpa racemosa*, widely addressed as sea grapes, is marine widely consumed seaweed. The high bioavailability of seaweeds displayed a potent bio-resource not only for functional food but also as medicine due to a broad range of biological activities [19,20]. These compounds, such as alkaloids, caulerpin, racemosins, and other interesting discoveries, gain interest as they possess various bioactivities: anti-inflammatory, cytotoxic, anti-proliferative, and anti-metastatic [19,21–23]. A previous study demonstrated *C. racemosa*’s anti-cancer and anti-metastatic potential in HeLa cells by inhibiting cancer cell migration and lowering Snail and Vimentin expression [24]. However, there has been limited research into *C. racemosa*’s potent anti-cancer activity against breast cancer. The activity of *C. racemosa* extract from Indonesia as an anti-cancer and anti-metastatic agent in apoptosis induction, upregulation of caspase-3 expression, and influences on Snail and Vimentin protein expression levels in breast cancer cells was investigated in this study.

**MATERIALS AND METHODS**

This study is in vitro true experimental study using breast cancer cell line culture, Michigan cancer foundation-7 (MCF-7) cell, conducted at the Biomedical and Biomolecular Laboratory, Faculty of Medicine, Brawijaya University, Malang, Indonesia. The research has been carried out since May 2022 after being ethically approved.

**MCF-7 breast cancer cell culture**

MCF-7 human breast cancer cells were obtained from the Biomedical Laboratory, Medical Faculty, Brawijaya University. The MCF-7 cells were cultured in 24-well plates containing Dulbecco’s Modified Eagle Medium (DMEM) media (Invitrogen, Massachusetts) supplemented with 10% fetal bovine serum (FBS), and antibiotic (100 U/ml Penicillin and 100 µl/ml Streptomycin), then maintained at a pH of 7.2–7.4 (Fig. 1). After the cultured cells reached 80% density, they were incubated at 37°C in an incubator with 5% CO₂. Cells were periodically harvested using trypsin-ethylenediaminetetraacetic acid (trypsin-EDTA) solution [25].

**Caulerpa racemosa extract preparation**

Sea grapes (*C. racemosa*) were obtained from the Mantehage Sea, North Sulawesi, Indonesia. Species identification and authentication were confirmed at the Pharmacology Department, Faculty of Mathematics and Sciences, Sam Ratulangi University, Manado, Indonesia. Finalization of the extract was conducted in the Biochemistry Laboratory, Faculty of Medicine, Brawijaya University. Raw *C. racemosa* was rinsed in water, dried at room temperature, and ground with an electric grinder. The coarse powder (1,000 g) was macerated in 96% ethanol for 72 hours and evaporated in a 40°C oven to produce a thick extract (with a 34% yield) and then dissolved in dimethyl sulfoxide. Each extraction process was carried out in triplicate. The crude extract was filtered with the Whatman 41 filter paper. The filtered crude extract was concentrated at 40°C with a rotary evaporator (RV 8, IKA Rotation Evaporator) under low pressure (100 mllibars) for 90 minutes. The crude extracts were stored in a refrigerator at 10°C [26]. Based on the previous study, the metabolite profile content of *C. racemosa* extract was examined using liquid chromatography-mass spectrometry and yielded a mass of 39,813,278 (39,712,257 m/z), indicating the presence of the antioxidant compound caulerpin in the extract [27]. Caulerpin is a secondary metabolite of *C. racemosa* that is a potent anti-cancer compound [27]. The 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl-2H-tetrazolium bromide (MTT) assay to investigate the safety profile of *C. racemosa*, and the LC50 value for a macerated extract of *C. racemosa* after 24 hours of incubation was 914.78 g/ml. The safety profile result indicates that *C. racemose* extract was safe to develop into various products.

**Trypan blue exclusion test**

The cultured cells were first transferred to a 24-well plate (Matsunami, Japan) and incubated for 24 hours. MCF-7 cells were administered with *C. racemosa* extract containing caulerpin at the concentration of 0 µg/ml as a control, 100 µg/ml, 200 µg/ml, and 400 µg/ml. Viable cells were examined using the trypan blue exclusion method after being treated with *C. racemosa* extract containing caulerpin. The cells on the well plate were placed into a tube and grouped according to the extract dose. They were centrifuged, while the supernatant was discarded. A mixture of 10 µl of cells, 10 µl of complete medium, and 20 µl of trypan blue was evaluated using a hemacytometer under a binocular microscope (Fig. 1) [28]. To minimize the risk of bias, the number of cell viability was calculated three times for each treatment group.

**Annexin-V/PI staining**

Utilizing the Annexin-V-FITC-PI kit (Sigma-Aldrich, St. Louis, MO), flow cytometry was applied to assess the extent
of cell apoptosis. After being washed twice with cold cell
staining buffer, MCF-7 cells were resuspended in Annexin-V
binding buffer at a concentration of 0.5–1.0 × 10^5 cells/ml. The
cell suspension was subsequently transferred to a 5 ml test tube
along with 5 l of FITC Annexin-V and 10 l of a propidium iodide
solution at a concentration of 20 g/ml. Following that, the cells
were slowly vortexed and incubated in the dark for 15 minutes at
room temperature (25°C). After that, 400 l of Annexin-V staining
buffer was added to each tube, which was subsequently analyzed
through a flow cytometer set to the suitable machine settings
(Fig. 1). Cell Pro Quest software was used to analyze the data.

Wound healing assay

MCF-7 cells were cultured in DMEM media until they
formed monolayers in 24-well plates. A micropipette tip was
used to scratch a single line to form a space between cells in each
well. DMEM medium was used to wash away the dislodged
cells. After the scratch generated a space between cells, the cells
were grown in starvation media (DMEM with FBS 0.1%) to
ensure the cells could not proliferate and could only migrate
to close the space. Then, varying doses of extract (0, 100,
200, and 400 µg/ml) were added. Every 3 hours, the wound-
closing process was monitored until the scratch in the control
group was completely closed. ImmunoRatio ImageJ application was
used to calculate the scratch width between MCF-7 cells.

Observation of caspase-3, Vimentin, and Snail expression by
immunofluorescent assay

Immunofluorescence staining was used to look at the
expression of caspase and other apoptosis-related proteins in
cell samples at the same time. This method was also used to
assess the expression level of Snail and Vimentin proteins.
Cell cultures were grown in 24-well plates with a circular glass
cover with an 18-mm diameter fitted to the bottom of each well
(Matsunami, Japan). Cells were washed twice with Phosphate-
buffered saline (PBS) before being fixed in 4% formaldehyde
in PBS for 15 minutes at room temperature. Following that,
the cells were washed twice with PBS and permeabilized for
5 minutes in PBS with 2 ml 0.1%–0.5% Triton X-100 at 4°C.
Then, triton X-100 was aspirated, and cells were washed three
times with PBS. Afterward, a 200 µl blocking buffer (10% goat
serum, 2% bovine serum albumin, 0.2% Triton-X) was added
to the slide. Then, the slides were incubated for 2 hours at room
temperature. After incubation, the slides were rinsed once again
with PBS.

A primary antibody with a ratio 1:100 was applied
to each slide. The primary antibodies used were anti-rabbit
cleaved caspase-3 (Cell Signaling Technology, Danvers, USA)
to stain caspase-3 protein, anti-rabbit Vimentin Cell Signaling
Technology, Danvers, MA) to stain Vimentin protein, and anti-
mouse FITC Snail-1 (Santa Cruz, CA) to stain Snail protein.
Slides without primary antibodies were prepared as negative
controls. After that, the slides were incubated in a humid room
overnight at 4°C. The next day, the slides were irrigated with PBS
three times for 10 minutes and then dried at room temperature.
The slides were then treated with 100 µl of secondary antibody
diluted 1:1,000 in PBS. The secondary antibodies used were
Alexa Fluor-488 goat anti-rabbit IgG (Invitrogen, Carlsbad,
CA) and Alexa Fluor-488 goat anti-mouse IgG (Invitrogen,
Carlsbad, CA). The nucleus was stained with 4,6-diamidino-2-
phenylindole (DAPI; Sigma-Aldrich, St. Louis, MO) 1 g/ml for
15 minutes and then washed 6 times with PBS. The evaluation
was performed using an Olympus IX71 inverted fluorescence
microscope with a magnification of 200× [25,29].

Data collection and statistical analysis

ImageJ software was used for the result quantification
of cleaved caspase-3, Snail, and Vimentin protein expression.
The data outcome was the intensity of these proteins expressed
as pixels. Statistical analysis of scratch width, cleaved
caspase-3, Snail, and Vimentin proteins expression data was
analyzed using GraphPad Prism 8 software and IBM SPSS
version 26 for MacBook. Shapiro-Wilk test was performed to
evaluate data distribution. If the data were normally distributed
(p > 0.05), a One-Way ANOVA test was conducted to examine
the difference in average between treatment groups. Otherwise,
the Kruskal-Wallis test would be performed if the data were not
normally distributed (p < 0.05). If the test results are significant
(p < 0.05), the data will be analyzed further using Dunn’s
Post hoc test to determine the significance between groups.
Subsequently, post-tests and correlation tests were carried out.
The data were presented in the form of a diagram.

RESULTS

Results of cell viability test using Trypan blue exclusion
method

The Trypan blue viability test revealed that the 400
µg/ml dose group had the least amount of viable MCF-7
cancer cells compared to the other dose groups. Furthermore,
there was a significant difference in post-treatment viable cells
compared to the control group (p < 0.05). Although there was
no significant difference in cell viability between the 100, 200,
and 400 µg/ml dosage groups, a consistent decrease in cell
viability was observed. In general, the mean and SD of viable cell amounts in each group were as follows: 0 µg/ml (990 × 10^3 ± 26.46), 100 µg/ml (836 × 10^3 ± 55.08), 200 µg/ml (796 × 10^3 ± 153.7), and 400 µg/ml (546 × 10^3 ± 58.59). The viability test results demonstrated that a dose of 400 g/ml was the best dose with the lowest number of cancer cell viability, indicating high apoptotic activity (Fig. 2). There was a significant negative correlation between the dose and the viability of MCF-7 cells (r = -0.877). Thus, the higher concentration of C. racemosa displayed stronger activity to reduce the viability of MCF-7 cells.

**Cleaved caspase-3 expression level after administration of C. racemosa extract**

Figure 3 shows that increasing the dose of C. racemosa extract was associated with increased expression of cleaved caspase-3 in MCF-7 cells. Caspase-3 expression was significantly higher in cells treated at 400 g/ml compared to the control group (p < 0.05) (Table 1). In line with previous findings, the dose-dependent relationship between C. racemosa concentration and caspase-3 expression revealed a dose-dependent effect, indicating that the higher the dose of C. racemosa, the higher the caspase-3 expression in MCF-7 cancer cells. The lowest mean and SD were in the control group (0 ± 0) and the highest was in the 400 µg/ml group (44.83 ± 19.18). In addition, Figure 3 can also be observed qualitatively, the high intensity of caspase-3 can be seen from the red scattering of immunofluorescent staining at doses of 200 and 400 µg/ml.

Data showing an increase in the expression level of caspase-3 protein through immunofluorescence staining using the anti-rabbit cleaved caspase-3 primary antibody after administration of C. racemosa extract was strengthened by data from flow cytometry of MCF-7 cells (Fig. 4). The results of MCF-7 cell flow cytometry showed an increase in the percentage of total late apoptosis in accordance with increasing of extract concentration (Table 2). This shows a correlation that increasing the level of cleaved caspase-3 protein expression after administration of the extract will induce apoptosis of MCF-7 cells.

**Cell migration assessment using wound healing assay**

After 24 hours of incubation, the scratch test (wound healing assay) was performed to determine the rate of cell migration activity. ImageJ software was used to measure the width of the scratches. After 24- and 48-hour interventions, the width (in pixel units) between scratch cells was measured (Table 3). There were significant differences in cell gap width between the 0 g/ml group and the 100, 200, and 400 g/ml groups after 24 hours of intervention (p < 0.001). This scratch test revealed that an increase in C. racemosa concentration was associated with a wider gap between two MCF-7 cell populations that had previously been separated by the scratch. There was an inhibition of MCF-7 cancer cell migration rate as a consequence of the treatment (Fig. 5). A similar result was obtained after 48 hours of intervention, in which statistically significant differences were seen between all groups of 0, 100, 200, and 400 µg/ml (p < 0.0001). A strong positive correlation (r = 0.808) between C. racemosa extract dose and a wide range was displayed from the scratch test 24- and 48-hours of MCF-7 cancer cells.

**Snail protein expression level after administration of C. racemosa extract**

The result of the Kruskal-Wallis test showed that the independent variable (dose of C. racemosa extract) and the dependent variable (Snail protein expression level) were statistically significant with a p-value of 0.003 (p < 0.05) (Table 1). Dunn’s Post hoc test showed a significant p-value of 0.003 (p < 0.05) between cells treated at a dose of 400 µg/ml compared...
Vimentin protein expression level after administration of *C. racemosa* extract

The ANOVA test revealed a significant relationship between Vimentin expression and *C. racemosa* extract, with a *p*-value of 0.000 (*p* < 0.05) (Table 1). A significant difference was found when the overall treatment dose groups to control. The Spearman correlation test revealed a negative association between *C. racemosa* extract dose with Snail expression with a very strong degree of negative correlation (*r* = −0.958). According to this correlation test, the higher the dose of *C. racemosa* extracts administered, the lower the amount of Snail expression (Fig. 6).

### Table 1. The difference in caspase-3, Snail, and Vimentin expression area between varying doses of *C. racemosa* extract.

<table>
<thead>
<tr>
<th><em>Caulerpa racemosa</em> extract dose</th>
<th>0 μg/ml (Control group)</th>
<th>100 μg/ml</th>
<th>200 μg/ml</th>
<th>400 μg/ml</th>
<th><em>p</em>-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Caspase-3 expression area (pixel)</td>
<td>0 ± 0</td>
<td>24.13 ± 3.98</td>
<td>30.67 ± 0.37</td>
<td>44.83 ± 19.15</td>
<td>0.003</td>
</tr>
<tr>
<td>Snail expression area (pixel)</td>
<td>61.27 ± 15.42</td>
<td>35.56 ± 4.23</td>
<td>12.66 ± 2.71</td>
<td>9.41 ± 1.34</td>
<td>0.003</td>
</tr>
<tr>
<td>Vimentin expression area (pixel)</td>
<td>68.48 ± 17.10</td>
<td>39.41 ± 9.46</td>
<td>14.48 ± 3.75</td>
<td>15.53 ± 5.36</td>
<td>0.000</td>
</tr>
</tbody>
</table>

### Table 2. Apoptosis activity rate in MCF-7 cells between varying doses of *C. racemosa* extract.

<table>
<thead>
<tr>
<th><em>Caulerpa racemosa</em> extract dose</th>
<th>0 μg/ml (Control group)</th>
<th>100 μg/ml</th>
<th>200 μg/ml</th>
<th>400 μg/ml</th>
<th><em>p</em>-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Apoptosis activity (%)</td>
<td>4.86 ± 1.07</td>
<td>5.14 ± 1.58</td>
<td>5.46 ± 0.54</td>
<td>7.1 ± 0.16</td>
<td></td>
</tr>
</tbody>
</table>

### Table 3. Post 24- and 48-hours scratch test shows cell scratch width (pixels).

<table>
<thead>
<tr>
<th><em>Caulerpa racemosa</em> extract dose</th>
<th>0 μg/ml (Control group)</th>
<th>100 μg/ml</th>
<th>200 μg/ml</th>
<th>400 μg/ml</th>
<th><em>p</em>-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>24 hours</td>
<td>535 ± 45</td>
<td>702 ± 58</td>
<td>759 ± 66</td>
<td>760 ± 45</td>
<td>0.001</td>
</tr>
<tr>
<td>48 hours</td>
<td>95 ± 22</td>
<td>484 ± 50</td>
<td>575 ± 48</td>
<td>625 ± 67</td>
<td>0.0001</td>
</tr>
</tbody>
</table>

### Figure 4. Increasing the apoptotic activity of MCF-7 cells after 24 hours of incubation with *C. racemosa* extract. Total apoptosis in MCF-7 cells was measured using flow cytometry with annexin-V/PI staining. Annexin-V/PI−: MCF-7 viable cells; Annexin-V/PI+: necrotic cells; Annexin-V+/PI−: early apoptosis; Annexin-V+/PI+: late apoptosis.

### Figure 5. Diagram of comparison of scratch test between treatment groups and MCF-7 cells migration activity by wound healing assay or scratch test. **** *p* < 0.001 (24 hours); ***** *p* < 0.0001 (48 hours); ** *p* < 0.05. Vimentin protein expression level after administration of *C. racemosa* extract

The ANOVA test revealed a significant relationship between Vimentin expression and *C. racemosa* extract, with a *p*-value of 0.000 (*p* < 0.05) (Table 1). A significant difference was found when the overall treatment dose groups...
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were compared using Tukey Multiple Comparison ($p < 0.05$). Pearson correlation test revealed a negative association between *C. racemosa* extract dose with Vimentin expression with a very strong degree of negative correlation ($r = -0.914$). According to this correlation test, the higher the dose of *C. racemosa* extracts administered, the lower the amount of Vimentin protein expression (Fig. 7).

**DISCUSSION**

The most common cancer in women is breast cancer [30]. Varying previous research identified secondary metabolites and crude extracts of the *Caulerpa* species as promising anti-cancer candidates [19,24,27]. Caulerpin, the major constituent of *C. racemosa* demonstrated *in-vitro* anti-tumor activity [7]. Caulerpenyne (Cyn), possesses anti-neoplastic, anti-mitotic, and anti-proliferative activity. It was found to be cytotoxic to several human tumor cells [19]. Breast cancer action is mediated by estrogen hormone and estrogen receptors (ERs) [31]. The presence of ER alpha is critical for a favorable prognosis and response to hormonal therapy. Molecular *in silico* studies of the ER protein with caulerpin and the standard drug, mitoxantrone, revealed lower binding (−11.18 kcal/mol) compared to standard drugs (−8.45 kcal/mol) [32]. The findings of this study support the possibility of developing this extract as a breast cancer treatment.

A study of *C. lentillifera* extract in human breast cancer MCF-7 cells revealed the role of caulerpin in the activation of caspase-3 and caspase-9, which subsequently induced apoptosis [7,19]. Polyphenols and flavonoids derived from *Caulerpa* sp. influence the expression of proteins involved in the cell cycle, resulting in cell cycle arrest [5]. The antioxidant activity of phenols and flavonoids as chemoprevention is capable of neutralizing free radicals (ROS) to prevent cell damage and further progression into malignancy [5]. However, “there has been no study” of *C. racemosa* extract as an apoptosis inducer by altering the expression level of caspase-3 and anti-metastatic by altering EMT’s protein regulators for breast cancer, particularly in the MCF-7 cell model.

This study revealed the potent anti-cancer effect of crude *C. racemosa* extracts in MCF-7 human breast cancer cells by inducing apoptosis. Administration of various doses of the crude extract increased the percentage of apoptotic activity in the MCF-7 cells shown in the cell viability test (Fig. 2). Enhanced apoptotic activity along with increasing crude extract dose was evidenced by correlation value $r = 0.877$ and $p < 0.05$. This experiment’s results are consistent with previous studies that demonstrated the anti-apoptotic activity of *C. racemosa* crude extract in neuroblastoma, cervical, and colon cancer cell lines [19,21,33]. Based on the biomolecular view of apoptosis activity, the mechanism of apoptosis involves various proteins to regulate and operate the apoptosis process. The most important protein for apoptosis, thus, as a marker for apoptotic activity is a caspase protein.

Caspase proteins are currently classified into three types: inflammatory caspase, initiator caspase, and effector caspase [18]. Caspase-3 is an effector protein for an apoptotic process that directly leads a cell death [1,8,18]. As an inactive dimer, caspase is activated by proteolytic cleavage which turns caspase-3 zymogen into cleaved caspase-3 [6]. The pro-apoptotic
activity of C. racemosa extracts was linked to an increase in the expression of the pro-apoptotic protein caspase-3 (Fig. 8). Increased cleaved caspase-3 indicated apoptosis induction [8]. Cleaved caspase-3 was evaluated with immunofluorescence staining (Fig. 3). There was an increase in caspase-3 expression, respectively, to C. racemosa extracts dose increment in MCF-7 cells ($r = 0.783$, $p < 0.05$). Caspase-3 expression was also dose-dependent, with a higher dose of C. racemosa extracts inducing higher caspase-3 expression in MCF-7 cancer cells (Table 1). Based on the experiment work, it can be concluded that administration of a crude extract from C. racemosa is able to enhance the expression of a caspase-3 protein that has a role as an effector protein to leading cell death in MCF-7 cells. It is in accordance with another experiment result that showed an administration of a crude extract could induce an apoptosis activity proven by decreasing MCF-7 cancer cell viability (low cell amount). Based on the findings, we can conclude that crude extract of C. racemosa is capable of inducing apoptosis in MCF-7 cells by increasing caspase-3 expression.

The experiment also demonstrated that crude extract has anti-metastatic properties by inhibiting cell migration. The inhibitions of migration along with the increase of the extract doses given were evidenced by the correlation value $r = 0.808$ and $p < 0.001$ (Fig. 5). MCF-7 was treated with 400 μg/ml doses is the most effective in inhibiting cell migration as evidenced by the difference in the width of the scratch between pre- and post-treatment, which means that cell migration is successfully inhibited when compared to cells without treatments (Table 3). Metastasis of breast cell cancer involves a process called EMT [19,21]. EMT will alter the characteristics of epithelial cells and give rise to mesenchymal cells. Besides that, polarity and attachment of epithelial cells to the basal membrane are lost, and then, it makes the cell apart from basal to circulation and metastasis occurs [24]. Based on several pieces of literature, there are three proteins that are involved during the EMT process such as Vimentin and Snail [25,34–36].

Snail protein is a zinc finger protein family that has a main role in regulating cancer cell metastasis [25,37]. Snail protein is able to bind with a gene promoter which encodes a cadherin protein and acts as an inhibitor protein, thus the transcription rate of cadherin can be decreased through disruption of RNA polymerase binding [25,38,39]. cCadherin protein is a class of transmembrane glycoproteins that serve as anchors for cells in tissues or cells with basement membranes [40]. According to the statement, the cadherin protein is required for cell attachment in tissue. In a cancer cell, the level expression of Snail protein is higher than in the normal cell, thus it influences the decreasing expression level of cadherin and makes the cells detach each other easily [25,37,39]. Snail protein expression level was evaluated with immunofluorescence staining (Fig. 6). Treating MCF-7 cells using crude extract of C. racemosa is able to decrease the expression level of Snail protein (Table 1). This result is strengthened with a significant $p$-value ($p = 0.003$) and $r$-value ($r = -0.958$) in statistical analysis. Decreasing of Snail expression level leads to an increase in the cadherin expression level [25]. Eventually, the adhesion between cells is strengthened, thus, migration activity becomes lower (Fig. 8). It is in accordance with the wound healing assay or scratch test result that shows the decrease of cell migration along increasing in extract dose (Fig. 5).

According to recent research, carcinoma cells that have completed their EMT process will develop a mesenchymal phenotype and express mesenchymal markers such as Vimentin [25,34,35]. Vimentin levels will eventually rise as a result of increased EMT activity in carcinoma cells [25,35]. Treatment of C. racemosa extract is able to decrease Vimentin expression in MCF-7 cells (Table 1). This expression was observed using immunofluorescence and Vimentin expression was seen to be significantly decreased when treated with C. racemosa extract, especially in the dose of 400 μg/ml (Fig. 7). This result is strengthened with a significant $p$-value ($p = 0.000$) and $r$-value ($r = -0.914$) in statistical analysis. According to these findings, C. racemosa extract can inhibit EMT in breast cancer cells indicating with decrease in the presence of Vimentin protein in the cells (Fig. 8). This is a preliminary study that reveals caulerpin of C. racemosa extract has potent pro-apoptotic activity against MCF-7 human breast cancer cells. However, further steps of the experiment include continuing to in-vivo study to strengthen the findings of this experiment before entering the clinical trial. Furthermore, the toxicity and adverse effects of C. Racemosa could be investigated further.

CONCLUSION
The C. racemosa extract exhibited a potential anti-cancer agent through pro-apoptotic activity in MCF-7 breast cancer cells. The extract reduced MCF-7 cell viability by increasing the expression of pro-apoptotic protein, cleaved caspase 3. Furthermore, an anti-metastatic activity of C. racemosa extract was demonstrated in the experiment through wound healing assay by inhibition of Snail protein expression level and also decrease of Vimentin expression level indicating that the extract is capable of inhibiting the progression of MCF-7 cells during the EMT process for their metastasis.

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AUTHORS’ CONTRIBUTIONS
HKP, NA, FRQ, and NIT conducted an experiment, analyzed data, conceptualized, and designed the study, and wrote the manuscripts. DIH, ADS, VMY, SAN, WR, and WB contributed to processing the pictures, graphical abstracts, data interpretation, writing manuscripts, and editing. All authors have read and approved the final manuscript.

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CONFLICT OF INTEREST
All authors and contributors (HKP, NA, FRQ, DIH, ADS, NIT, VMY, SAN, WR, and WB) to the study declare no conflict of interest.
ETHICAL APPROVALS
This experiment has passed the ethical test of the Health Research Ethic Committee Faculty of Medicine Brawijaya University with ethical exemption letter number 18/EC/KEPK/01/2022 on January 31, 2022.

DATA AVAILABILITY
All data generated and analyzed are included in this research article.

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REFERENCES

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