BUKTI KORESPONDENSI

Kaempferol And Quercetin Isolated From The Leaves Of Atingia Excelsa To Arrest Cell Cycle In G0/G1 Phase Human Tongue Cancer Sp-C1 Cell Lines

1.	Bukti pengiriman artikel ke jurnal	22 januari 2020
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3.	Permintaan revisi sesuai arahan reviewer	24 April 2020
4.	Permintaan perbaikan hasil cek plagiasi	22 juli 2020
5.	Pengiriman naskah hasil perbaikan	5 Agustus 2020

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KAEMPFEROL AND QUERCETIN ISOLATED FROM THE LEAVES OF Atingia excelsa TO ARREST CELL CYCLE IN G0/G1 PHASE HUMAN TONGUE

• CANCER SP-C1 cell lines

ABSTRACT

The leaves of Altingia excelsa were found to strongly inhibit SP-C1 human tongue cancer cell lines. This study was focused on identifying the antiproliferative compound found in A. excelsa leaves and assessing its mechanism of action. The active compound was isolated using column chromatography and identified by the spectroscopic method and was also tested for its anti-proliferative properties and the cell cycle analisis analysis in SP-C1 cells by flowcytometry analysis. This work resulted in the isolation of a flavonoid, which was identified to be kaempferol and quercetine quercetin. The compounds inhibited SP-C1 cell proliferation in a time- and dose-dependent manner with IC₅₀ values of 0.72 μ g/mL and 0. 70 μ g/mL for the 24 hours treatments, respectively. Furthermore, the flowcytometry analysis suggested that the compounds exerted its anticancer activities by inhibited inhibiting cell cycle. These results suggested that compounds found in A. excelsa providies a basis for its potential use in cancer disease management.

Keywords: Altingia excelsa, cancer, cell cycle, kaempferol, quercetine-quercetin

INTRODUCTION

More than 95% of the carcinomas of the oral cavity are of squamous cell type, in nature. They constitute a major health problem in developing countries, representing a leading cause of death. The survival index continues to be small (50%), as compared to the progress in diagnosis and treatment of other malignant tumors. According to World Health Organization, carcinoma of the oral cavity in males in developing countries, is the sixth commonest cancer after lung, prostrateprostate, colorectal, stomach and bladder cancer, while in females, it is the tenth commonest site of cancer after breast, colorectal, lung, stomach, uterus, cervix, ovary, bladder and liver.¹ Oral squamous cell carcinoma (OSCC) is the most common malignant tumor of this anatomic site, and in approximately 80% of cases, it is associated with extrinsic factors such as the use of tobacco, alcohol or both.² OSCC is a very difficult disease to treat because of multidisciplinary and diverse treatment strategies and the varied natural behavior of the cancer. Local invasion and frequent regional lymph node metastases together with relative resistance to chemotherapeutic. The conventional strategies of OSCC management still depend on surgery, radiotherapy, chemotherapy and targeted therapy.³ The poor outcome of chemotherapy to OSCC contributes to the poor prognosis for OSCC.⁴ Therefore, novel, effective therapy for OSCC treatment is still needed. Due to this high incidence, the identification of novel compounds that inhibit cancer development has become a crucial objective for scientists. Of the hundreds of chemicals that have been and are being evaluated for their anti-cancer activities, natural products derived from medicinal plants rank among the most promising.⁵ In an effort To identify novel agents that may inhibit cancer development, we have focused our investigations on discovering bioactive compounds from plants.⁶

In our previous study, we found that the leaves of the Hammamelidaceae family, demonstrated anti-tumor properties.^{7,8} These preliminary studies suggest that *A. excelsa* may be

further developed as a source of anti-cancer agents. Thus, in this study, we investigated and characterized the inhibited cell cycle activities of *A. excelsa* leaf extracts.

MATERIALS AND METHODS

Plant materials. A. Excelsa leaves were collected from Wayang Windu Mountain, Pangalengan, West Java, Indonesia. The plant species was were identified by the Department of Biology, Faculty of Mathematics and Natural Sciences, University of Padjadjaran, Indonesia.

Extraction and isolation. The dried leaves of *A. excelsa* (2.5 kg) were extracted with methanol (3x24 h) at room temperature. The solvent was subsequently evaporated under reduced pressure at 50°C to yield a concentrated extract. The methanol extract (280 g) was fractionated *n*-hexane and water to obtain an *n*-hexane extract (86 g) and a water layer. The water layer was then extracted with ethyl acetate to obtain an ethyl acetate fraction (120 g) and a water fraction (90 g). The cytotoxicity of the fractions was assessed on SP-C1 tongue cancer cells using the methyl thiazolyl tetrazolium (MTT) assay. The ethyl acetate fraction, which was the most active fraction, was chromatographed on Wakogel C-200 (Wako Pure Chemical, Japan) with a mixture of *n*-hexane, ethyl acetate and methanol with increased polarity. The major compounds were then isolated and purified using silica G-60 with sulfuric acid-ethanol (1:9) and were identified by spectroscopic methods including ultra violet ultraviolet and infrared spectrometry (UV-IR), and nuclear magnetic resonance (NMR).

Cell culture and treatment. The SP-C1 human tongue cancer cell line used in this study were cultured in RPMI-1640 medium (Sigma, St. Louis, MO, USA) supplemented with 10% fetal bovine serum and antibiotics (100 U/mL penicillin and 100 μ g/mL streptomycin). For cell treatments, various concentrations of the sample were added to the cell culture medium. After 24 h, the cells were released from treatment, the medium was replaced, and cells were subsequently collected at the indicated times.

Cell cycle analysis using flow cytometry. The cells were grown in 24-well plates at 37 °C under 5% CO2 until 80% confluence were was reached. The medium was subsequently changed, and the flavonoids were added to the indicated concentrations. Next, the cells were incubated at 37°C for 48 h. After incubation, the cells were harvested and washed three times with ice-cold PBS (pH 7.4). The supernatant was removed, and the cells were washed with 1 mL of PBS and centrifuged at 4 °C. Finally, the supernatant was removed, and 200 µL of 70% ice-cold ethanol and 200µL of PBS was added to the cells and stored at -20 °C until further use. For use in the flow cytometry experiments, the cell pellet was washed two more times with PBS. The cell pellet was suspended in 0.5 mL of staining reagent (50µg/mL PI, 50 U/mL RNase, 0.1 mM EDTA,0.1% Triton X-100, and PBS) and incubated for 30 min at 37 °C in the dark. The DNA fluorescence was measured using a Becton Dickinson (Franklin Lakes, USA) FACScanto II flow cytometer with an excitation wavelength of 488 nm and emission wavelength of 585 nm. Pulse width area signals were used to discriminate between G2 cells and cell doublets. The data were analyzed using FACSDiva Software (Beckton Dickinson). The relative distribution of 10⁴ events for each sample was analysed for background aggregates and debris, an indicatorof apoptosis and the G0/G1-, S-, and G2/M-phases of the cell cycle. The control treatments consisted of a culture medium supplemented with FBS. Serum-deprivation treatment was used as an inducer of G0/G1 cell cycle arrest.

RESULTS

Ethyl acetate fraction of *A. excelsa* leaves inhibits SP-C1 cell proliferation. Treatment MTT assay with the *A. excelsa* extract methanol was found to inhibit the proliferation of SP-C1 human tongue cancer cells (IC_{50} 75.41 µg/mL) (Fig. 1). The methanol extract was then fractionated based

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on polarity, using *n*-hexane, ethyl acetate and water. The fractions were then individually applied to SP-C1 cells and were found to inhibit cell proliferation with an IC₅₀ value of 44.85 μ g/mL for the *n*-hexane fraction, 12.85 μ g/mL for the ethyl acetate fraction and 18.02 μ g/mL for the water fraction. Due to its low IC₅₀ value, we then explored the ethyl acetate fraction for its anti-cancer potential.



Kaempferol and quercetin are major compounds of the ethyl acetate fraction. The principle active compounds of the ethyl acetate fraction of *A. excelsa* extract were isolated and purified. The compound exhibited a melting point of 152.7-153 °C and a molecular ion peak at m/z 432 in the LC-MS spectrum. Based on hydrogen and carbon content, the molecular ion peak and ¹H and ¹³C NMR profiles indicated that the compound had a molecular formula of C₁₅H₁₀O₇. The UV spectrum produced maximal absorbance peaks at λ_{max} 265 and 342 nm, which were characteristic of a flavonoid with a flavone skeleton. The addition of NaOH produced a bathochromic shift in the absorption bands, indicating the presence of hydroxyl groups in the skeleton, one of which was attached to C-5, as indicated by a further bathochromic shift following the addition of H₃BO₃. Absorption bands at 1,675 and 3,197 nm of the IR spectrum indicated where the molecule harbors conjugated carbonyl and hydroxyl groups, respectively.

The ¹H NMR spectrum of the compound showed two aromatic hydrogen signals with 'meta coupling' at δ 6.35 (1H, d, J = 2.2 Hz) and 6.18 (1H, d, J=2.2 Hz), which was predicted by the hydrogens at C-6 and C-8 of the A ring of the flavone skeleton. Accordingly, this compound was suggested to have a hydroxyl group at C-5 and C-7. Furthermore, its 1H NMR spectrum revealed two signals with 'ortho coupling' at δ 6.92 (2H, d, J = 6.7 Hz) and 7.74 (2H, d, J = 6.7 Hz), the signals of which were approximated from the hydrogens at C-3', C-5' and C-6' of the B ring. The absence of a specific signal for an olefinic hydrogen at C-3 and the presence of an anomeric hydrogen signal at δ 5.37 (1H, d, J = 7.2 Hz) suggested that the compound was a flavonol glycoside. The appearance of an anomeric carbon signal at δ 94.9 in the ¹³C NMR spectrum indicated the presence of a sugar moiety. Due to a correlation between the anomeric hydrogen signal (δ 5.37) and the anomeric carbon signal (δ 94.9) that was revealed by analysis of the HMBC spectral data, the position of the sugar moiety was assigned to the C-3 hydroxyl group, The methyl signal.



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Chemical structure of Quercetin

Kaempferol and quercetin inhibits SP-C1cell proliferation in a dose-dependent manner. The effects of kaempferol and quercetin on the viability of SP-C1 cells were evaluated. The treatment of cancer SP-C1 cell lines with kaempferol and quercetin resulted in a dose-dependent inhibition of cell growth, as demonstrated by the MTT assay. Twenty-four hours of treatment with kaempferol and quercetine quercetin inhibited the proliferation of SP-C1 cells with IC_{50} values of 0.72 and 0.70 µg/mL, respectively. Subsequent cell cycle analysis, based investigation applying the IC_{50} dose of 0.39, 0.78, and 1.56 µg/mL kaempferol and quercetin was performed on SP-C1 cells.

Kaempferol and quercetin inhibited the SP-C1 cell cycle in the G0/G1 phase. To determinate the percentage of SP-C1 cells present in different phases of the cell cycles G0/G1, S, and G2/M, the cells were first synchronized with serum deprivation prior to before the treatment with or without quercetin and kaempferol at various concentrations (C, K 0.39, K 0.78, K 1.56, Q 0.39, Q 0.78, and Q 1.56 μ g/mL) for 24 h period, serum-deprived cells were progressively accumulated in the G0/G1-phase, where as whereas serum-supplied cells were present in the S- and G2/M-phases (Figure 2). Treatment with K (all concentrations) and Q (all concentrations) significantly increased the percentage of cells in the G0/G1-phase and G2/M-phase as compared with the control. Notably, the percentage of cells in the G0/G1-phase increase of cells in the G0/G1 – phase increase compared to the control after K treatment. Percentage of cells in the G0/G1 – phase increase compared to the control after Q treatment but the percentage of cells in the G0/G1 – phase increase in concentration.



Figure 2. Flowcytometry analysis SP-C1 cancer cell using treated some concentration of kaemperol and quercetin.

DISCUSSION

The ¹H- and ¹³C-NMR spectra of compounds from *A. Excelsa* exhibited resonances due to aromatic systems. The ¹H-NMR spectrum of them showed the presence of two doublet signals corresponds to four aromatic protons in ring B, characteristics for the 1',4'-disubstituted flavone. The ¹³C-NMR signals of them were assigned with the help of a DEPT experiment. A total of fifteen carbon signals were observed in the ¹³C-NMR spectrum. The degree of unsaturation was accounted for eight out of the total eleven double bond equivalents. The ¹³C-NMR spectrum of them showed the presence of 15 aromatic carbon signals. So, from ¹H and ¹³C NMR profiles indicated that the compound had a molecular formula of C₁₅H₁₀O₇.

Traditional medicinal plants traditionally have long been regarded as a source of potential therapeutic agents, and the search for new drugs or leads are usually based on that approach.^{9,10} In drug discovery, we have recently applied a new approach of to selecting plants based on the Hammamelidaceae family.^{7,8} In our previous study, we found that the extracts of the *A. excelsa* leaves were strongly cytotoxic to the SP-C1 Human tongue cancer cell lines. Thus, these extracts had the potential for further investigation. The present study was focused on identifying an antiproliferative compound from the *A. excelsa* leaves. This work resulted in the isolation of two flavonoids, kaempferol and quercetin, which strongly inhibited the SP-C1 cell lines proliferation in a time- and dose-dependent manner. This compound has not been reported before in connection with its cytotoxicity in these cancer cell lines

In this study, kaempferol and quercetin with some concentration to arrest G0/G1 phase cell cycle. Cell growth is an addition additional amount of cells from the cell cycle process. The cell cycle consists of 4 phases, that is G_1 , S, G_2 , and M. G_1 (Gap 1) is a phase when the cell will synthesis the DNA or go out from cell cycle reversibly or irreversibly to differentiation. The cell that in the G_1 phase will easily controlling cell eyele cycle at a point, that is in the restriction point (R) that will determine the cell to come in back to the cell cycle, go out from cell cycle entering G_0 phase or differentiating.¹¹

When the cell cycle phase flow in restriction point dan enter S phase controlled by cyclindependent kinase (Cdks) and D, E, and A cyclin. D cyclin function as a growth factor that the expression more depend on the extracellular signal than the cell position on the cycle.¹² When the cell entering the G₀ cycle, on or more of the D cyclin (D1, D2 and D3) were inducted as a part apart from the first response of a growth factor stimulation, protein synthesis and fomer former complex with catalytic subunit (Cdk4, Cdk6) depend on the mitogenic stimulation.¹³ In reverse, if the mitogenic substance were removed, so the D Cyclin synthesis will stop. D cyclin is an unstable protein and the enzymatic activity quickly disappear, so that the cell quickly go out from the cell cycle. A specific inhibitor of Cdk4 and Cdk6 known as Ink4 can directly stop the activity of D cyclin/cdk 4/6 and causing G₁ phase arrest (rest of G₁ phase).¹³

Cell cycle inhibition of tongue cancer (SP-C1 after kaempferol and kuersetin quercetin intervention as the results of Rasmala leaves isolation. There is a system that controlled the cell cycle. The controlling system known as a checkpoint. There are 4 checkpoints that found on cycle cell, that is G₁, S, G₂, and M. Checkpoint in a cell cycle onvolving involving protein cyclin groups and cyclin-dependent kinase (cdk).¹⁴ Protein groups that involved on the cell cycle checkpoint determine cell cycle to stop. DNA repair, or apoptosis if the DNA repair doesn't occur.¹⁵ Kaempferol (4) can inhibit cell cycle on G_0 - G_1 phase at 0.39 µg/mL, 0,78 µg/mL and 1,56 µg/mL concentration. This cell cycle inhibition suspected because of kaempferol (4)/kuersetin quercetin (5) that can inhibit the work of c-myc. We know that the function from c-myc transcription factor is increasing D cyclin and E cyclin, so that there are increasing activity of G1-CDK (D1-CDK cyclin) and G1/S-CDK (E-CDK6 cyclin). Thus, the kuersetin-quercetin in this research can inhibit tongue cancer SP-C1 cell cycle on G₀-G₁ phase at 0,39 µg/mL, 0,78 µg/mL dan 1,56 µg/mL concentration. This results analogueanalog with the research who is conducted by Mocanu et al, in 2013 years that shows kuersetin quercetin can inhibit Epidermoid Cancer Cell line A-431 cell cycle at 5 and 10 µM concentration in 24 hours on G₀/G₁ phase and inhibit Mamary SK-BR-3 cell cycle at 50 and 75 µM concentration in 24 hours on G₀-G₁ phase. Research belonging to Chen et al, in 2011 years shows

that kuersetin quercetin can inhibit oral cancer OSCC SCC-25 cell cycle at 50 and 75 μ M concentration in 12 hours incubation on the G₁ phase. Cell cycle inhibition which is conducted by kuersetin quercetin on G₀-G₁ phase possibly because kuersetin quercetin will synergize with cyclin-dependent kinase inhibitor (CDK1) p21 that that function as inhibitor D-CDK4/6 cyclin activity.¹⁶ With inhibition of D-CDK4/6 cyclin activity so the cell will be obstructed to enter S phase, so that the cell will stop on the G₀G₁ phase or p21 beside binding cdk and PCNA (proliferating cell nuclear antigen), that is a DNA polymerase α subunit, which needed for replication or DNA reparation. P21 will inhibit cell cycle on the G₁ phase as a result of cell-contact inhibition and because of TGF β .¹⁷

CONCLUSION

In conclusion, our results suggest that kaempferol and quercetine inhibited the growth of SP-C1 cells through inhibited cell cycle in the G0/G1 phase.

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Vol. 14 No.1 Februari 2020

ISSN : 1907-5987

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RESEARCH ARTICLE

Kaempferol and Quercetin Isolated from The Leaves of Atingia Excelsa to Arrest Cell Cycle in G0/G1 Phase Human Tongue Cancer SP-C1 Cell Lines

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Online submission : 23 January 2020 Accept Submission : 10 March 2020

ABSTRACT

The leaves of Altingia excelsa were found to strongly inhibit SP-C1 human tongue cancer cell lines. This study was focused on identifying the antiproliferative compound found in A. excelsa leaves and assessing its mechanism of action. The active compound was isolated using column chromatography and identified by the spectroscopic method and was also tested for its antiproliferative properties and the cell cycle analysis in SP-C1 cells by flowcytometry analysis. This work resulted in the isolation of a flavonoid, which was identified to be kaempferol and quercetin. The compounds inhibited SP-C1 cell proliferation in a time- and dose-dependent manner with IC_{50} values of 0.72 µg/mL and 0. 70 µg/mL for the 24 hours treatments, respectively. Furthermore, the flowcytometry analysis suggested that the compounds found in A. excelsa providies a basis for its potential use in cancer disease management.

Keywords: Altingia excelsa, cancer, cell cycle, kaempferol, quercetin.

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INTRODUCTION

Approximately 300.373 new cases of oral squamous cell carcinoma (OSCC) are annually reported around the world, what makes oral cancer the sixth most common cancer worldwide. The term oral cancer is referred to as a subgroup of head and neck malignant neoplasms affecting the lips, the anterior twothirds of tongue, the salivary glands, the gingiva, the floor of the mouth, the oral mucosal surface and the palate, with the tongue being the most common location.¹ Oral squamous cell carcinoma (OSCC) is the most common malignant tumor of this anatomic site, and in approximately 80% of cases, it is associated with extrinsic factors such as the use of tobacco, alcohol or both.² OSCC is a very difficult disease to treat because of multidisciplinary and diverse treatment strategies and the varied natural behavior of the cancer. Local invasion and frequent regional lymph node metastases together with relative resistance to chemotherapeutic. Management of OSCC varies consider-ably; small cancers of the oral cavity are usually managed by surgery alone, whereas advanced oral cancers are usu-ally treated with primary radical surgery followed by ra-diation chemoradiation or and targeted therapy.³ The poor outcome of chemotherapy to OSCC contributes to the poor prognosis for OSCC.⁴ Therefore, novel, effective therapy for OSCC treatment is still needed. Due to this high incidence, the identification of novel compounds that inhibit cancer development has become a crucial objective for scientists. Of the hundreds of chemicals that have been and are being evaluated for their anti-cancer activities, natural products derived from medicinal plants rank among the most promising.⁵ To identify novel agents that inhibit may cancer development, have focused we our investigations discovering on bioactive compounds from high plants level.⁶

In our previous study, we found that the leaves of the Hammamelidaceae family, demonstrated anti-tumor properties.^{7,8} These preliminary studies suggest that *A. excelsa* may be further developed as a source of anti-cancer agents. Thus, in this study, we investigated and characterized the inhibited cell cycle activities of *A. excelsa* leaf extracts.

MATERIALS AND METHODS

Plant materials. A. Excelsa leaves were gatted from Wayang Windu Mountain, Pangalengan, West Java, Indonesia. The plant species were identified by plant taxonomy laboratorium of Biology Departement, FMIPA, Padjadjaran University, Indonesia.

Extraction and isolation. Simplicia leaves of A. excelsa (2.5 kg) were extracted with methanol (3x24 h) at 20-25°C (Room temperature). The solvent was subsequently evaporated under reduced pressure at 50°C to produce a concentrated extract. 280 g of methanol extract was fractionated by n-hexane and water to gained an 86 g n-hexane extract and the layer of water. The water layer then extracted with ethyl acetate to gained an 120 g ethyl acetate fraction and 90 g water fraction. Fractions cytotoxicity was assessed on SP-C1 tongue cancer cells using methyl thiazolyl tetrazolium (MTT) assay. Ethyl acetate fraction, which was the most active fraction, was chromatographed by Wakogel C₂₀₀ (Wako Pure Chemical, Japan) with a mixture of n-hexane, ethyl acetate and methanol with raised the polarity. Main compounds were then isolated and purified by silica G₆₀ with sulfuric acidethanol (1:9) were identified and by of mass spectroscopic methods consist spectroscopy (MS), ultraviolet (UV), infrared spectrometry (IR), and nuclear magnetic resonance (NMR).9

Cell culture and treatment. SP-C1 human tongue cancer cell line was used in this research were used RPMI-1640 as cultured medium (Sigma, St. Louis, MO, USA) added with 10% FBS and antibiotics among them 100 U/mL penicillin and 100 μ g/mL streptomycin. Cell's treatments, several concentrations of the sample were added to the cell culture medium. After 24 h, the cells were released from treatment, the medium was replaced, and cells were subsequently gathered at the indicated times.¹⁰

Cell cycle analysis using flow cytometry. The cells were grown in 24-well plates at 37° C with CO₂ from 5% until 80% encounter was reached. Afterwards, medium was changed, and flavonoids were added to the indicated concentrations. Furthermore, cells were incubated for 48 h at 37°C. After incubation, the cells were harvested and rinsed three times with ice-cold PBS (pH 7.4). The supernatant was eliminated, and the cells were rinsed with 1 mL of PBS and centrifuged at 4 °C. Finally, the supernatant was eliminated, and 200 µL of 70% ice-cold ethanol and 200µL of PBS was added to the cells and stored at -20 °C for further use. Utilization in the flow cytometry experiments, the cell pellet was rinsed twice with PBS. Cell pellet was suspended in 0.5 mL of staining reagent (50µg/mL PI, 50 U/mL RNase, 0.1 mM EDTA,0.1% Triton X-100, and PBS) and incubated for 30 min at 37 °C in the dark. Measuring DNA fluorescence was used Becton Dickinson (Franklin Lakes, USA) FACScanto II flow cytometer with 488 nm excitation wavelength and 585 nm emission wavelength. Pulse width area signals were used to discriminate between G2 cells and cell doublets. Data analyzed was used FACSDiva Software (Beckton Dickinson). Relative distribution of 10⁴

events for each sample was analyzed for background aggregates and debris, apoptosis indicator and the G0/G1-, S-, and G2/M-phases of the cell cycle. Control treatments consisted of a culture medium supplemented with FBS. Serum-deprivation treatment was used as an inducer of G0/G1 cell cycle arrest.¹¹

RESULTS

Fraction of Ethyl acetate A. excelsa leaves inhibits SP-C1 cell proliferation. Treatment MTT assay shown that A. excelsa methanol extract was inhibit the proliferation of SP-C1 human tongue cancer cells (IC₅₀ 75.41 µg/mL) (Fig. 1). Methanol extract then fractionated based on polarity, using *n*-hexane, ethyl acetate and water. Afterward the fractions individually applied to SP-C1 cells and inhibit cell proliferation with an IC₅₀ value of 44.85 μ g/mL for the *n*-hexane fraction, 12.85 μ g/mL for the ethyl acetate fraction and 18.02 µg/mL for the water fraction. Due to its low IC₅₀ value, we then explored the ethyl acetate fraction for its anticancer potential.



Figure 1. IC₅₀ value some extracts *A. excelsa* Nornha

Main compounds of the ethyl acetate fraction are Kaempferol and guercetin. Isolation and purification of the ethyl acetate fraction A. excelsa extract to gathered bioactive compounds. The compound exhibited a melting point at 152.7-153°C and a molecular ion peak at m/z 432 in the LC-MS spectrum. According to hydrogen and carbon content, the molecular ion peak and ¹H and ¹³C NMR profiles indicated that the compound has a molecular formula of C₁₅H₁₀O₇. Ultraviolet spectrum provides maximal absorbance peaks at λ_{max} 265 and 342 nm, which were characteristic of a flavonoid with flavone skeleton. The addition of NaOH produced bathochromic shift in the absorption bands, indicating the presence of hydroxyl groups in the skeleton, one of which was attached to C-5, as indicated by further bathochromic shift following the addition of H₃BO₃. Absorption bands at 1,675 and 3,197 nm of the IR spectrum indicated where the molecule harbors conjugated carbonyl and hydroxyl groups, respectively.

¹H NMR spectrum of the compound shown two hydrogen of aromatic signals with 'meta coupling' at δ 6.35 (1H, d, J = 2.2 Hertz) and 6.18 (1H, d, J=2.2 Hertz), which was predicted by hydrogens at C-6 and C-8 of the A ring of the flavone skeleton. Hence, this compound was suggested to have a group of hydroxyl in C-5 and C-7. Furthermore, ¹H NMR spectrum revealed two signals with 'ortho coupling' at δ 6.92 (2H, d, J = 6.7 Hz) and 7.74 (2H, d, J = 6.7 Hz), the approximation signals from the hydrogens at C-2', C-3', C-5' and C-6' of the B ring. Lack of a specific signal for olefinic hydrogen at C-3 and existence of an anomeric hydrogen signal at δ 5.37 (1H, d, J = 7.2 Hz) suggested that the compound was a flavonol glycoside. Appearance of anomeric carbon signal at δ 94.9 in the ¹³C NMR spectrum indicated the availability of a sugar moiety. In consequence to a correlation between the anomeric hydrogen signal (δ 5.37) and the anomeric carbon signal (δ 94.9) that revealed by analysis of the HMBC spectral data, sugar

moiety position was established to the C-3 hydroxyl group, signal of methyl.



Chemical structure of Kaempferol



Chemical structure of Quercetin

Quercetin and Kaempferol inhibits SP-C1 cell proliferation in a concentrationdependent manner. Kaempferol and guercetin effects on the viability of SP-C1 cells were evaluated. Treatment of cancer SP-C1 cell lines with kaempferol and quercetin shown that concentration-dependent inhibit of cell growth, as demonstrated by the MTT assay. 24 hours of treatment with kaempferol and quercetin inhibited the proliferation of SP-C1 cells with IC₅₀ values 0.72 and 0.70 µg/mL, respectively. Subsequent cell cycle analysis, based investigation applying the IC₅₀ concentration of 0.39, 0.78, and 1.56 µg/mL kaempferol and quercetin was applied on SP-C1 cells.

Kaempferol and quercetin inhibited the SP-C1 cell cycle in the G0/G1 phase. To determinate the percentage of SP-C1 cells present in different phases of the cell cycles G0/G1, S, and G2/M, first cells were synchronized with serum deprivation before treatment with or without quercetin and kaempferol at several concentrations (C, K 0.39, K 0.78, K 1.56, Q 0.39, Q 0.78, and Q 1.56 µg/mL) for 24 h period, serum-deprived cells were progressively accumulated in the G0/G1-phase, whereas serum-supplied cells were present in the S- and G2/M-phases (Figure 2). Treatment with K (all concentrations) and Q (all concentrations) significantly increased the

percentage of cells in the G0/G1-phase and decrease in S and G2/M-phase as compared with the control. Particularly, Cell percentage in G0/G1-phase increased an appropriate increase concentration compared to the control after K treatment. Percentage of cells in the G0/G1 – phase increase compared to the control after Q treatment but the percentage of cell decrease a long increase in concentration.



Figure 2. Flowcytometry analysis SP-C1 cancer cell using treated some concentration of kaemperol and quercetin.

DISCUSSION

¹H-¹³C-NMR spectra The and of compounds from Α. Excelsa exhibited resonances due to aromatic systems. The ¹H-NMR spectrum of them showed the presence of two doublet signals corresponds to four aromatic protons in ring B, characteristics for the 1',4'disubstituted flavone. The ¹³C-NMR signals of them were assigned with the help of a DEPT experiment. A total of fifteen carbon signals were observed in the ¹³C-NMR spectrum. The degree of unsaturation was accounted for eight out of the total eleven double bond equivalents. The ¹³C-NMR spectrum of them showed the presence of 15 aromatic carbon signals. So,

from ¹H and ¹³C NMR profiles indicated that the compound had a molecular formula of $C_{15}H_{10}O_7$.

Rural medicinal plants since a long time as sources of potential therapeutic medication, and find out of novel medication or leads compound are usually based on that approach.^{12,13} In drug invention, researcher have recently applied a new approach to selecting plants based on the Hammamelidaceae family.^{7,8} In previous research, we found that the *A. excelsa* leaves extracts were strongly cytotoxic to the SP-C1 Human tongue cancer cell lines. Therefore, these extracts potential for further investigation. Present study focused on identifying antiproliferative compound from the *A. excelsa* leaves. This study was isolated of two flavonoids, kaempferol and quercetin, which strongly inhibited proliferation of SP-C1 cell lines in a time- and concentration-dependent manner. There is no compound that reported before in relation with its cytotoxicity in these cancer cell lines.

In this study, kaempferol and quercetin with some concentration to arrest G0/G1 phase cell cycle. Cell growth is an additional amount of cells from the cell cycle process. The cell cycle consists of 4 phases, that is G₁, S, G₂, and M. G₁ (Gap 1) is a phase when the cell will synthesis the DNA or go out from cell cycle reversibly or irreversibly to differentiation. The cell that in the G₁ phase will easily controlling cell cycle at a point, that is in the restriction point (R) that will determine the cell to come in back to the cell cycle, go out from cell cycle entering G₀ phase or differentiating.¹⁴

When the cell cycle phase flow in restriction point dan enter S phase controlled by cyclindependent kinase (Cdks) and D, E, and A cyclin. D cyclin function as a growth factor that the expression more depend on the extracellular signal than the cell position on the cycle.¹⁵ When the cell entering the G₀ cycle, on or more of the D cyclin (D1, D2 and D3) were inducted as apart from the first response of a growth factor stimulation, protein synthesis and former complex with catalytic subunit (Cdk4, Cdk6) depend on the mitogenic stimulation.¹⁶ In reverse, if the mitogenic substance were removed, so the D Cyclin synthesis will stop. D cyclin is an unstable protein and the enzymatic activity quickly disappear, so that the cell quickly go out from the cell cycle. A specific inhibitor of Cdk4 and Cdk6 known as Ink4 can directly stop the activity of D cyclin/cdk 4/6 and causing G1 phase arrest (rest of G₁ phase).¹⁶

Cell cycle inhibition of tongue cancer (SP-C1 after kaempferol and quercetin intervention as the results of Rasmala leaves isolation. There is a system that controlled the cell cycle. The controlling system known as a checkpoint. There are 4 checkpoints that found on cycle cell, that is

G₁, S, G₂ and M. Checkpoint in a cell cycle involving protein cyclin groups and cyclindependent kinase (cdk).¹⁷ Protein groups that involved on the cell cycle checkpoint determine cell cycle to stop, DNA repair, or apoptosis if the DNA repair doesn't occur.¹⁸ Kaempferol (4) can inhibit cell cycle on G_0 - G_1 phase at 0.39 µg/mL, 0,78 µg/mL and 1,56 µg/mL concentration. This cell cycle inhibition suspected because of kaempferol (4)/ quercetin (5) that can inhibit the work of c-myc. We know that the function from cmyc transcription factor is increasing D cyclin and E cyclin, so that there are increasing activity of G1-CDK (D1-CDK cyclin) and G1/S-CDK (E-CDK6 cyclin). Thus, the quercetin in this research can inhibit tongue cancer SP-C1 cell cycle on G₀-G₁ phase at 0,39 µg/mL, 0,78 µg/mL dan 1,56 µg/mL concentration. This results analog with the research who is conducted by Mocanu et al, in 2013 years that shows quercetin can inhibit Epidermoid Cancer Cell line A-431 cell cycle at 5 and 10 µM concentration in 24 hours on G₀/G₁ phase and inhibit Mamary SK-BR-3 cell cycle at 50 and 75 µM concentration in 24 hours on G₀-G₁ phase. Research belonging to Chen et al, in 2011 years shows that quercetin can inhibit oral cancer OSCC SCC-25 cell cycle at 50 and 75 µM concentration in 12 hours incubation on the G₁ phase. Cell cycle inhibition which is conducted by quercetin on G₀-G₁ phase possibly because quercetin will synergize with cyclin-dependent kinase inhibitor (CDK1) p21 that that function as inhibitor D-CDK4/6 cyclin activity.¹⁹ With inhibition of D-CDK4/6 cyclin activity so the cell will be obstructed to enter S phase, so that the cell will stop on the G_0/G_1 phase or p21 beside binding cdk and PCNA (proliferating cell nuclear antigen), that is a DNA polymerase a subunit, which needed for replication or DNA reparation. P21 will inhibit cell cycle on the G1 phase as a result of cell-contact inhibition and because of TGF β .²⁰

CONCLUSION

In conclusion, our results suggest that kaempferol and quercetin inhibited the growth of

SP-C1 cells through inhibited cell cycle in the G0/G1 phase.

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