

Original Article

CYTOTOXIC ACTIVITY OF BRAZILEIN ISOLATED FROM SECANG (*CAESALPINIA SAPPAN* L.) AGAINST MCF7/DOX CELLS BY INHIBITION OF P-GLYCOPROTEIN

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ABSTRACT

Objective: This study was focused on isolation of brazilein from the dried heartwood of Secang (*Caesalpinia sappan* L.) followed by its characterization using Infrared (IR) spectroscopy, liquid chromatography-mass spectrometry (LC-MS) with electrospray ionization (ESI), proton (¹H), carbon-13 (¹³C) nuclear magnetic resonance (NMR) and two dimensional (2D)-NMR, evaluation the cytotoxic activity of brazilein in MCF-7 resistant doxorubicin (MCF-7/DOX) cells and evaluate the interaction between brazilein and ATP with P-glycoprotein (Pgp) *in silico* using molecular docking.

Methods: Brazilein was isolated and purified from ethyl acetate fraction by flash silica gel column chromatography, eluting with chloroform, ethyl acetate and methanol in gradient concentration. In the cytotoxicity assay, MCF-7/DOX cells were cultured in the presence of brazilein for 24 hour (h) and cell viability was evaluated by using 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide (MTT) assay. Interactions between brazilein and the target proteins were evaluated and calculated *in silico* by molecular docking using PLANTS.

Results: The infrared, mass spectra with a molecular weight of 284 and NMR signal confirmed that was brazilein. MTT assay showed a dose-dependent inhibition of MCF-7/DOX cell proliferation with brazilein IC₅₀ value of 43 μM. The docking score of brazilein was -71,45 kcal/mol and ATP value -96,23 kcal/mol.

Conclusion: Brazilein has a potent cytotoxic value on MCF-7/DOX and high affinity in Pgp protein target. Brazilein can be developed as anticancer especially in cancer resistance incidence. Further study must be established to evaluate the molecular mechanism of brazilein inhibit MCF-7/DOX cell proliferation *in vitro*.

Keywords: Isolation, Brazilein, Cytotoxic, MCF-7/DOX, Pgp

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INTRODUCTION

Brazilein has been already isolated and purified from methanolic extract heartwood of secang by preparative HPLC using ODS-3 column and acetonitrile: isopropanol (65:35 v/v) as mobile phase [1]. In this study, brazilein is isolated, purified by using flash column chromatography and characterized using UV-Vis, infrared, mass spectra and NMR signal. Flash column chromatography has several advantages than preparative HPLC such as less expensive, this method do not require the help of expensive machinery, low maintenance, cost effective and available in many laboratories. This finding method for isolation and purification of brazilein by flash column chromatography can be utilized for production of brazilein standard for the laboratory. Therefore flash column chromatography could be an alternative method to isolate this active compound with high purity.

Secang (*Caesalpinia sappan* L.) has been traditionally used as coloring agent in foods and beverages. Besides, secang has a lot of pharmacological effects such as anti-inflammation [2], antioxidant [3], antimicrobial activities [4], antiacne [5] and anticancer [6]. Several phenolic compounds are isolated from *C. sappan*, such as homo-isoflavonoid protosappanin A, protosappanin B, 4-O-methylsappanol, caesalpin J, brazilin, and brazilein [7]. Brazilin and brazilein are the major compounds of this plant proven to be responsible for its pharmacological effects. Brazilein was reported to have a cytotoxic effect on lung cancer cells, nasopharyngeal cancer cells and prostate cancer cells with IC₅₀ value 5-18 μM [8] and caused cell cycle arrest at the G1 phase in MCF-7 cells [9]. Brazilein inhibit HER-2 activation on MCF-7/DOX cells. MCF-7/DOX cells were cells model of breast cancer cell resistance to chemotherapeutic agents [10]. One of the chemotherapeutic agents that is common to

be used in breast cancer therapy is doxorubicin. However, several problems come up after the use of doxorubicin such as its toxicity to normal tissues, severe side effects, and developed resistance. Doxorubicin induces the expression of Pgp [11]. Pgp is a transporter belong to ATP-binding cassette (ABC) family [12]. These transporters function as efflux pumps of chemotherapy agent to extracellular with ATP binding [13]. In MCF-7 breast cancer cell cases, overexpression of Pgp could be reduced the chemotherapy agent concentration intracellular that influence decreasing its cytotoxicity activity through efflux mechanism [14]. Pgp is an important transporter that plays role in resistance breast cancer mechanism. For that reason, these transporter is one of protein target therapy to overcome the resistance breast cancer cases. To our knowledge and literature survey, there is no discussion and evaluation about the cytotoxic activity of brazilein on MCF-7/DOX cells through Pgp inhibition. This study was conducted to isolate brazilein from the dried heartwood of secang and to evaluate its cytotoxic effect on resistant breast cancer cells using MCF-7/DOX cells by inhibition of pgp.

MATERIALS AND METHODS

Sample

Brazilwood or 'kayu secang' (*Caesalpinia sappan* L.) was obtained in the form of dried heartwood powder from Center for Research and Development of Medicinal Plants and Traditional Medicines in Tawangmangu Indonesia.

Extraction and fractionation of kayu secang

The dried heartwood powder of secang was extracted with methanol (Merck) at a volume ratio of 1:10 (for 48 h, at room temperature). The methanolic extract was then concentrated under

reduced pressure by rotary evaporator. The extract was partitioned with n-hexan and ethyl acetate (Sigma Aldrich co. USA.) by liquid-liquid extraction (LLE). The ethyl acetate fraction was evaporated using a rotary evaporator to yield crystal.

Isolation and purification of brazilein in ethyl acetate fraction of kayu secang by silica gel column chromatography

The ethyl acetate fraction of kayu secang was separated by flash silica gel column chromatography using chloroform-ethyl acetate-methanol eluent with gradient concentration (varying from non-polar until polar eluent). The fraction was collected every 30 ml and each fraction was evaluated and monitored by thin layer chromatography (TLC) profile using silica gel as the stationary phase and chloroform: ethyl acetate: methanol (1:7:2) as the mobile phase. The crude fraction was then purified using flash silica gel column chromatography again with the same mobile system. The pure fraction was obtained if gave one red spot in TLC profile and evaluated by TLC densitometry and Liquid Chromatography gave one peak area.

Characterization of brazilein

The purified brazilein was characterize using UV-Vis spectrophotometry, Infrared spectrophotometry, LC-MS, ¹³C-NMR, ¹H-NMR and 2D NMR (¹H-¹H COZY, ¹H-¹³C HMQC and ¹H-¹³C HMBC).

Cell culture

The MCF-7/DOX cells used are the collection of Cancer Chemoprevention Research Center (CCRC), Universitas Gadjah Mada, transferred from Nara Institute of Science and Technology (NAIST), Japan. The cells were cultured in DMEM (Gibco) Culture Media containing 10% Fetal Bovine Serum (FBS) (Gibco), 1% penicillin-streptomycin, and 0.5 % fungizone. Trypsin-EDTA 0.25% (Gibco) was used in cell harvest [15].

Cytotoxicity and cell viability assay: 3-(4, 5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay

MTT cytotoxicity assay was used to examine the effect of an isolated compound as a sample on MCF-7/Dox cells. Cells were distributed to 96-well plate with density of 5x10³ cells/well and incubated in 37°C with 5% CO₂ for 24 h. Certain concentrations of the sample were then applied. After 24 h incubation, 3-[4, 5-dimethylthiazole-2-yl (-2, 5-diphenyltetrazolium bromide)] (MTT) dissolved in PBS as MTT reagent was applied, followed by 4 h incubation. Stopper reagent, a solution of 10% v/v Sodium Dodecyl Sulphate (SDS) (Merck) dissolved in 0.1 N HCl (Merck), was then applied. The plate was then kept with protection from light overnight, continued with absorbance determination (λ 595 nm) using ELISA reader (Bio-Rad). During the process, Phosphate Buffer Saline (PBS) pH 7.4 containing KCl (HPLC grade, Sigma), NaCl (HPLC grade, Sigma), Na₂HPO₄ (HPLC grade, Sigma) and KH₂PO₄ (HPLC grade, Sigma) dissolved in aquadest was used washing reagent.

Molecular docking

In this study, the docking of brazilein to pgp was observed, using ATP as a comparison. Ligands preparation were done by using Marvin Sketch. The structure of pgp (1MV5) proteins were taken from Protein Data Bank (PDB). Protein preparation was done by using YASARA. Molecular docking was conducted by using PLANTS (Protein-Ligand Ant System) Software, giving docking score as a result.

RESULTS

Extraction and fractionation of kayu secang

The methanolic extract of kayu secang was obtained 10.24% b/b and then this extract was applied to fractionation using liquid-liquid extraction (LLE) with n-hexan and ethyl acetat. Hexan was used to reduce and eliminate the lipid, resin and non-polar component in the methanolic extract of kayu secang. The ethyl acetat fraction was obtained after concentrated using a rotary evaporator to reddish yield crystal as much 2.3 % b/b.

Isolation and purification of brazilein in kayu secang by silica gel column chromatography

Vacuum column chromatography by using silica gel and chloroform-ethyl acetate-methanol (varying concentration) as a mobile phase was performed to separate the components in the ethyl acetate fraction. The mobile phase was chosen based on the different polarities of each eluent, hopefully, the fraction that obtained from this method more pure. Each fraction was collected at volume 30 ml and obtained 115 fractions. This fraction was evaluated by TLC profile for each fraction and fraction that have same chromatographic profile was collected into one bottle. From this process, 115 fractions could be separated further into 10 fractions. Fraction II-VI was collected into one bottle because fractions have same TLC profile. One of the several spots in the same track on TLC profile gave a red spot dominantly and it was predicted as brazilein.

This crude fraction was chromatographed again by using the same system. The fraction was collected every 30 ml and evaluated the TLC profile. This purification process obtains 16 fractions. Fraction no II gave one red spot. It was evaporated by using a rotary evaporator to reddish crystal as much as 0.007 % b/b.

To ensure that fraction no II from purification process was really pure, the purification assay was performed. The purified component was applied in TLC method using silica gel 60 F₂₅₄ with three different mobile phase system (n-hexan: chloroform: methanol: acetat acid = 2:7:1:1 v/v (1); chloroform: methanol: acetate acid = 5:1:1 v/v (2); and chloroform: methanol: aquadest: acetate acid = 6:3:1:1 v/v (3)). This compound gives one spot to all system that used. Chromatogram result showed that hRf of this compound in the mobile system no 1 was 40 and hRf in the mobile system no 2 was 60 and the last system gave hRf value as much as 87.5. TLC-densitometry profile of this compound showed one dominant peak in λ 329 nm. It was subjected to further characterization.

Characterization of brazilein

Isolate compound that obtained from purification process, was characterized by UV-Vis spectrophotometer, FTIR, ESI mass spectrometer and NMR spectrometer.

Its UV-Vis spectra (fig. 1) showed isolate in methanol has maximum absorption in wavelength 287.5 nm, 368 nm, 435.5 nm and 588.5 nm. Very strong absorption at 287.5 nm revealed that isolate has an aromatic group with $\pi \rightarrow \pi^*$ transition [16]. Brazilein have one benzene ring that been fusion with dihydropirane ring. Peak at 368 nm indicates as an unsaturated keton that conjugated [16]. Brazilein also has its functional group. Isolate also has an absorption at 435.5 and 588.5 nm that means isolate has a long chromophore and auxochrome in its structure [16,17]. Auxochrome bind to chromophore system that makes the compound transition into bathochromic transition.

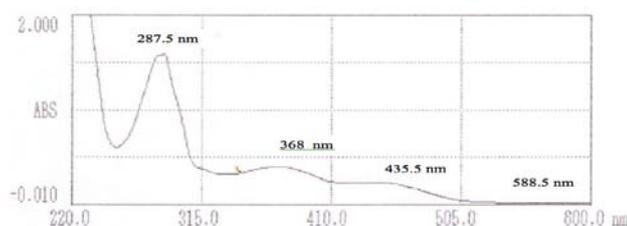


Fig. 1: UV-Vis spectra of isolate in methanol solvent by using spectrophotometer milton roy spectronic 3000 array

Spectra (fig. 2) showed a wide peak at 3307.9 cm^{-1} . It indicated isolate has a-OH (hydroxyl) functional group, very strong absorption at 1662.6 cm^{-1} showed the component has a carbonyl functional group (C=O). C=O group is an important infrared spectra characterization of brazilein that could be distinguished by brazilin, the other major compound of kayu secang [18]. The other absorption with strong and sharp absorption at 1604.8 cm^{-1} showed isolate has a C=C functional group from the aromatic ring. Some functional group that explained which present in the structure of brazilein as a predicted component. However, this estimation needs further confirmation by ESI-MS instrument to confirm its molecular weight.

The result of mass spectra analysis of isolate was shown in fig. 3. The spectra were in positive ion mode. There was 2 peak with high intensity in m/z 287 and m/z 285. Peak in m/z 285 suitable with $[M+H]^+$ ion formula, by calculating the formula, we found the mass of isolate was 284. Brazilein with molecular structure $C_{16}H_{12}O_5$ have molecular weight 284. This observation confirmed that isolate was brazilein, the major dyestuff component in kayu secang extract with molecular weight 284. This result was suitable with the other literature that brazilein is known to have a molecular weight of 284 [19-20].

The other highly intensity peak, m/z 287 was suggested that an artefact with $[M+3H]^+$ ion formula. In mass spectra, there was the reaction of molecule sample because of availability of the energy. Brazilein could have reduction process by binding 2 hydrogen atoms with still be neutral molecules and then this reduction molecule could receive 1 proton H^+ , that could be seen in m/z 287 as $[M+3H]^+$.

Identification of isolate could be held by using one dimension of NMR Spectroscopy (1H -NMR, ^{13}C -NMR, DEPT) and two dimensions (HMQC, COSY and HMBC). DEPT spectra of isolate shown in table 1. Isolate had five olefinic methane groups signal in δ 132.3; 112.9; 112.5; 110.0 and 104.3 and two methylene groups' signal in δ 70.9 and 43.0. Based on HMQC spectra (table 1), two protons in methylene groups were not equivalent to each other. This phenomenon was caused by *crossing peak* between δ 70.9 carbon

resonance signal and proton resonance signal in δ 3.92 and δ 3.68. The other signal was carbon resonance signal δ 43.0 crossing peaks with proton resonance signal in δ 3.01 and δ 2.76.

The 1H NMR spectra of isolate showed that isolate had seven different proton environment, five was an olefinic methane proton and two proton was a methylene signal, δ 3.68 (1H) and 3.92 (1H) were assigned to H-6. Signal at δ 2.76 (1H) and 3.01 (1H) as a H-7. The olefinic proton signal at δ 6.46 was ortho coupled with the signal at δ 7.18 ($J = 8.0\text{ Hz}$) and was meta coupled with δ 6.28 ($J = 2.6\text{ Hz}$). Therefore, these three olefinic signals were assigned to H-2, H-1 and H-4, respectively. H-8 and H-11 were assigned by 2D NMR such as HMQC, HMBC and 1H - 1H COZY correlations.

H-8 and H-11 were assigned by HMBC correlations as shown in fig. 4. The proton at δ 6.59 (1H) as H-11. H-11 has a long-range correlation with C-12 at δ_c 137.5. C-12 was correlated with a proton at δ 3.92 (H-6 b) and 2.76 (H-7 c). Four olefinic protons were known as H-1, H-2, H-4 and H-11 that means the remained olefinic proton as H-8 at δ 6.70 (1H).

Brazilein performed cytotoxic activity on MCF-7/DOX cells

The cytotoxicity of brazilein on MCF-7/DOX cells was determined by using MTT assay. This assay is based on the reduction of the soluble yellow tetrazolium salt to a purple insoluble formazan product by mitochondrial succinic dehydrogenase which can be measured spectrophotometrically after dissolving in SDS solution [21]. The IC_{50} value was $43\text{ }\mu\text{M}$ (fig. 5). A potent cytotoxic activity was exhibited by brazilein against MCF-7/DOX cells.

Brazilein potentially inhibits the protein target pgp

Validation result from Pgp protein showed molecular docking protocol for that protein target could be accepted with RMSD value $< 2\text{ }\text{\AA}$ (table 2) and molecular docking brazilein to Pgp could be continued. The result of molecular docking showed the binding energy of brazilein to Pgp was -71.45 and ATP as a native ligand had binding energy with Pgp was -96.23 (table 2). Interaction between brazilein and ATP with Pgp (PDB ID 1MV5) showed in fig. 6.

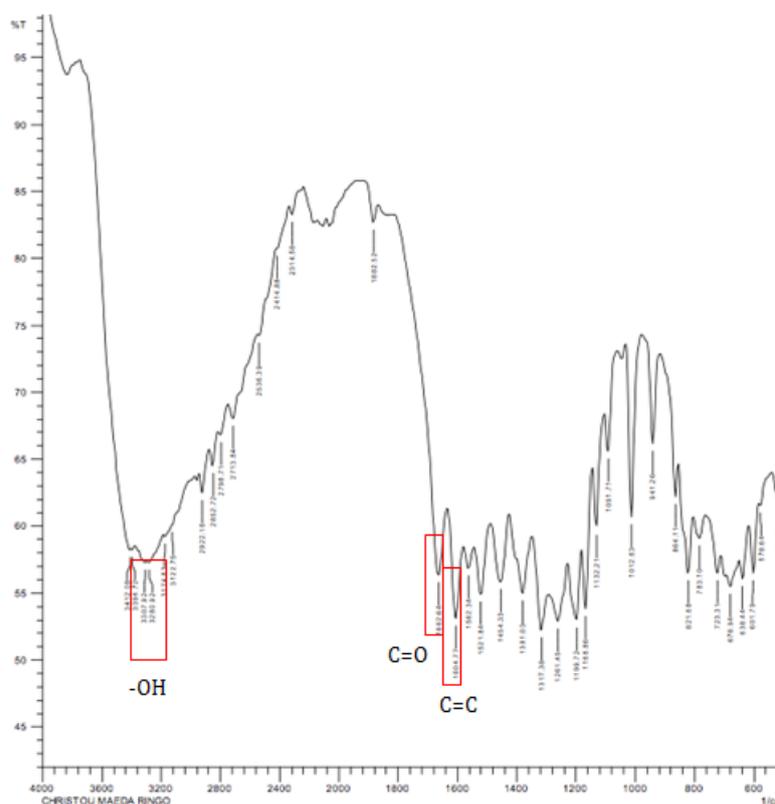


Fig. 2: Infrared spectra of isolate that isolated from ethyl acetate fraction by silica gel column chromatography

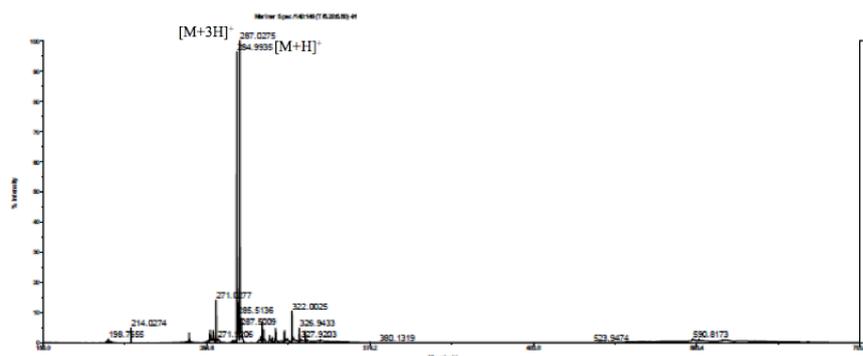


Fig. 3: Mass spectra of isolate

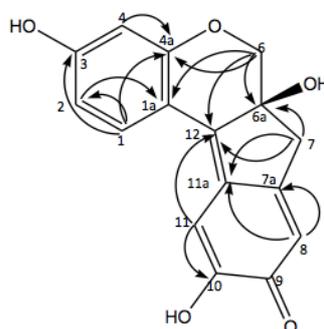


Fig. 4: HMBC correlations of isolate

Table 1: NMR spectra of brazilein by using CD₃-OD solvent and DELTA2 instrument with 500 MHz

C/H	δ_c (ppm)(DEPT)	δ H(ppm) (Nh, m, j)	COSY	$^1\text{H}-^{13}\text{C}$ HMBC
1	132,3(CH)	7,18 (1H, d, 8,0)	H1-H2	H1-C3/C4a
1a	115,6(Cq)	-	-	-
2	110,0 (CH)	6,46 (1H, dd, 8,0,2,6)	H2-H1	H2-Cl a/C4
3	158,0 (Cq)	-	-	-
4	104,3 (CH)	6,28 (1H, d, 2,6)	-	H4-Ca
4a	155,8 (Cq)	-	-	-
6	70,9 (CH ₂)	3,68 (1H,d, 11,65) 3,92 (1H, d, 11,65)	H6'-H6"	H6"-Cl a/C4a/C6a/C12
6a	78,2 (Cq)	-	-	-
7	43,0 (CH ₂)	2,76 (1H,d, 16,2) 3,01 (1H,d, 16,2)	H7'-H7"	H7"-C6/C6a/11a H7"-C6/C6a/11a/C12
7a	145,8 (Cq)	-	-	-
8	112,5 (CH)	6,70 (1H,s)	-	H8-C7a/C11a
9	209,3 (Cq)	-	-	-
10	145,5 (Cq)	-	-	-
11	112,9 (CH)	6,59 (1H,s)	-	H11-C10/C12
11a	131,4 (Cq)	-	-	-
12	137,5 (Cq)	-	-	-

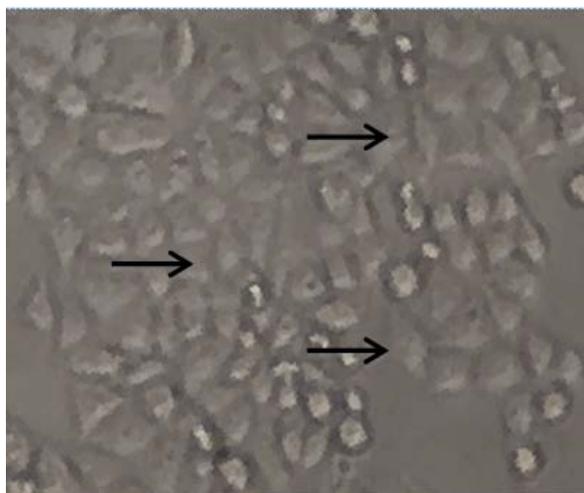
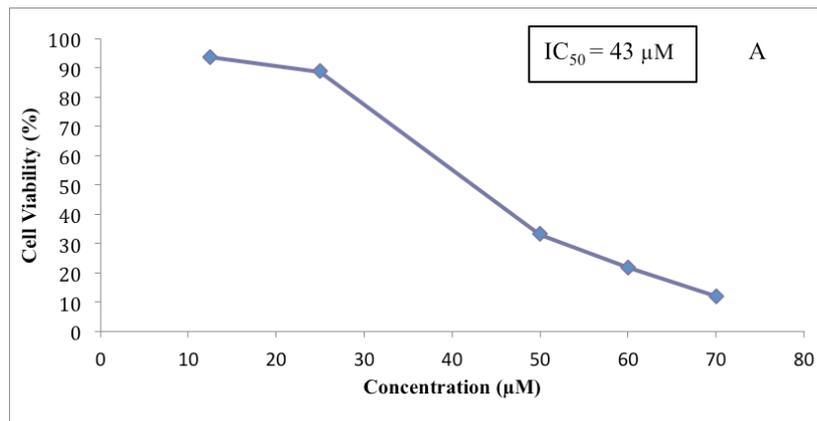
Cq = Quarternary carbon, J = Coupling constant, H = Amount of proton

Table 2: Docking score obtained after docking of brazilein to HER-2

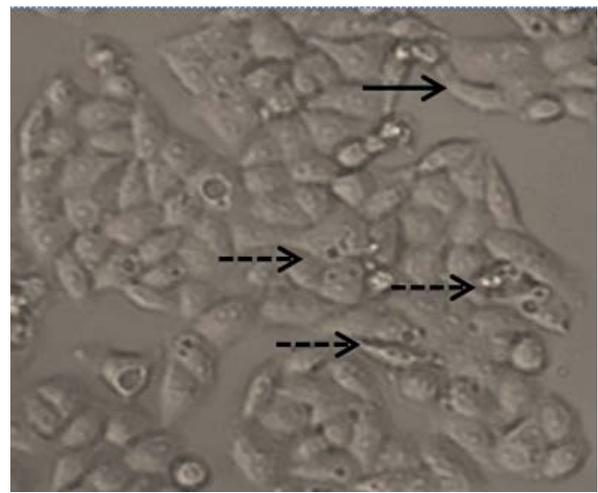
	Score docking Pgp (1MV5)
RMSD (Root Mean Square Deviation)	1.25° A
ATP	-89.72
Brazilein	-77.73

Table 3: Interaction brazilein and ATP to pgp protein with amino acid residue in pgp

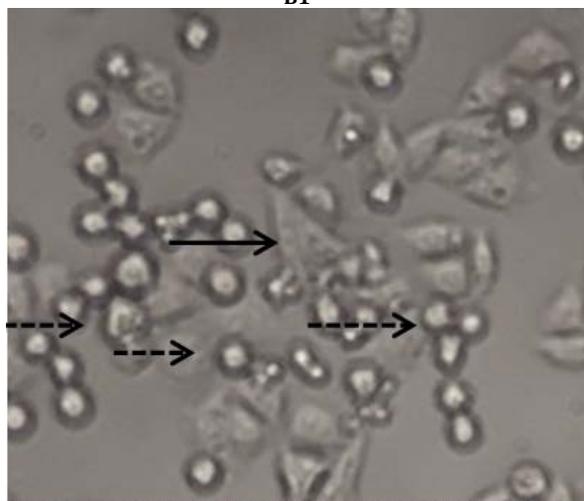
Target proteins	Compound	Amino acid residue	Bonding formation	distance
Pgp (1MV5)	ATP	Phe 421	Hydrogen Bond	2.65
		Leu 533	Hydrogen Bond	3.82
	Brazilein	Ser 423	Hydrogen Bond	1.65
		Asp 505	Hydrogen Bond	2.64
		Leu 504	Hydrogen Bond	2.20



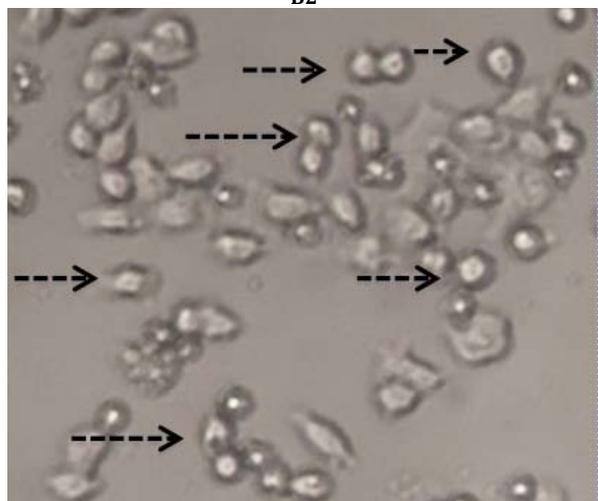
B1



B2



B3



B4

Fig. 5: Effect of Brazilein on MCF-7/DOX Cell Viability (A). Cell viability diagram was obtained from average \pm standard of error (SE) with $n=3$. The cytotoxic effect images of untreated control MCF-7/DOX cells (B1); after 24 h Brazilein 10 μM (B2); Brazilein 25 μM (B3); Brazilein 50 μM treated MCF-7/DOX cells (B4), \rightarrow normal cell morphology, \dashrightarrow change in cell morphology

DISCUSSION

According to Teng *et al.* (2005) [22], compounds with IC_{50} values below 50 μM had a potent cytotoxicity against cancer cells, meaning brazilein had cytotoxic effect in MCF-7/DOX cells and brazilein performed as a potent chemopreventive agent. This effect was supported by the morphological changing such as cell nuclei appear shrunken, irregular morphology and some cells undergo membrane blebbing.

The cytotoxic effect of brazilein on MCF-7/DOX cells might be caused by cell cycle arrest or apoptosis induction. Therefore, further study must be conducted to evaluate the molecular mechanism that contributes to brazilein's potency as an anticancer agent against MCF-7/DOX cells by using *in silico*. This study has been held to describe the molecular mechanism of brazilein and Pgp protein as protein target by evaluating the score docking. Score docking shows interaction strength of active compound and

protein target. Lower score docking means higher binding affinities between them. Molecular docking analysis of brazilein

was a bond in ATP binding site of Pgp. Pgp is an energy (ATP)-dependent protein transporter [23].

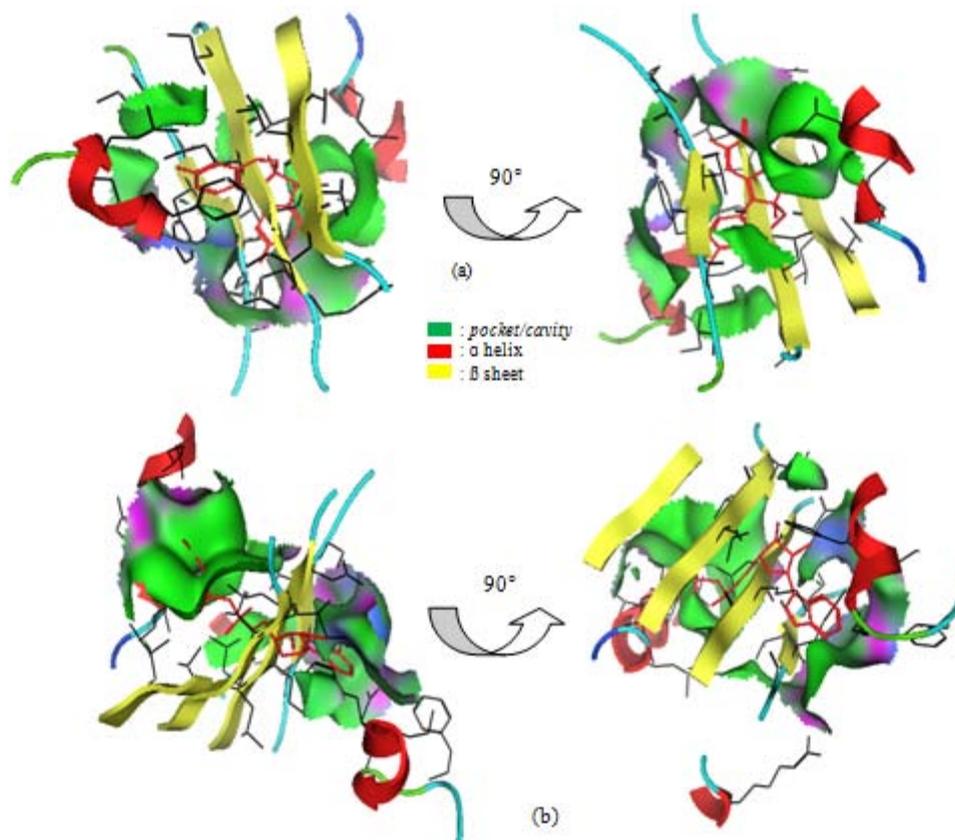


Fig. 6: Docking representations of brazilein and ATP interaction in ATP binding site to Pgp target proteins using PLANTS. The 3D interaction between brazilein and ATP binding site of Pgp (1MV5) (a); Interaction between ATP and ATP binding site of Pgp (1MV5) (b)

Here, this study demonstrated that brazilein exhibited promising anticancer activity on resistance cancer cases through ATP binding inhibition in ATP binding site of Pgp (fig. 6). This inhibition disrupts Pgp activity because there was not enough energy to be used in active transport process by Pgp [24]. That interaction happens through hydrogen bonding between O atom in carbonyl group of C₉ atom of brazilein with H atom in Ser 423, Leu 504 and Asp 505 amino acid residu of Pgp protein target (table 3). ATP interacted in ATP binding site of Pgp (fig. 6b) associated between hydrogen bonding of H atom in ATP with O atom in Phe 421 and Leu 533 amino acid residu of Pgp (table 3). The further study must be held to explore the molecular mechanism of brazilein *in vitro* associated with inhibition of brazilein in Pgp activation.

CONCLUSION

Fraction II which has the intense reddish color crystal, identified as brazilein based on the characterization of the purified component obtained from vacuum column chromatography result and brazilein had a potent cytotoxicity against MCF-7/DOX cells with IC₅₀ value was 43 μM by inhibition of P-glycoprotein (pgp).

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AUTHORS CONTRIBUTION

Linda Laksmiani isolated brazilein using a flash column with guidance by Ratna Susidarti. Evaluation of brazilein cytotoxic

activity as well as its molecular mechanism with guidance and advice by Edy Meiyanto. All authors contributed ideas and thoughts to the writing of this paper.

CONFLICT OF INTERESTS

Declared none

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