

## Cytotoxic Activity and Molecular Docking of Indole Alkaloid Voacangine, Bis-indole Alkaloids Vobtusine, and Vobtusine Lactone from the Indonesian Plant: *Voacanga foetida* (Blume) Rolfe

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### Info Article

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### ABSTRACT

The purpose of this study was to evaluate the *in vitro* cytotoxic activity of two isolated compounds from *Voacanga foetida* (Blume) Rolfe against the A549 cell line. The *in silico* study was carried out to predict the cytotoxic mechanism of two isolated compounds (voacangine and vobtusine lactone) and previously reported compound (vobtusine) against pro-survival receptors (Bcl-2, Bcl-xL, Mcl-1), pro-activation receptor (Bax), and apoptotic execution receptor (caspase-3). This research is an experimental quantitative using column chromatography and HPLC methods in the isolation process, MTT assay in determining the cytotoxic activity, and molecular docking in determining the prediction of ligand-receptor interactions. The fractions (VFB-DA, VFB-DB, VFB-BuOH, VFB-DB4) and voacangine exhibited very strong cytotoxic activity, while vobtusine lactone showed moderate cytotoxic activity. The docking scores for voacangine, vobtusine, and vobtusine lactone against Bcl-2 were -9.93; -10.07; -9.03 kcal/mol, against Bcl-xL were -9.77; -11.69; -9.76 kcal/mol, against Mcl-1 were -10.70; -10.77; -9.53 kcal/mol, and for Bax were -8.99; -6.87; -6.99 kcal/mol, as well as against caspase-3 were -12.05; -12.21; -12.02 kcal/mol. The cytotoxic activity of voacangine, vobtusine, and vobtusine lactone was thought to cause cell death by suppressing Bcl-xL and Mcl-1 activities and also increasing Bax and caspase-3 activities.

**Key words:** *Voacanga foetida*, vobtusine lactone, *in vitro*, *in silico*

### INTRODUCTION

Cancer has an increasing incidence of morbidity and mortality. Current treatments include surgery, chemotherapy, radiation. All of which are aimed at killing cancer cells. However, the results are not satisfactory as side effects are alarming to sufferers. In the light of this, this study is to explore medicinal plants that are thought to kill cancer cells either by necrosis or apoptosis (Susanty *et al.*, 2020).

At the cellular level, cancer can occur if there is a failure to respond to normal apoptosis, failure

of proliferation inhibition, and loss of heterozygosity (LOH) (Kyzas *et al.*, 2004). The part of the chromosome that contains the tumor suppressor gene is LOH. The mechanism of cancer cell death can occur through two processes, necrosis and apoptosis. The one that plays an important role in the carcinogenesis process is the apoptosis process.

Alkaloids show strong biological activity. It is estimated that more than 2000 compounds are classified as alkaloids, some of them from the Apocynaceae family which are generally used as

anticancer agents (Macabeo *et al.*, 2009). As an anticancer, alkaloid acts on the cytotoxic pathway, anti-proliferation, induces apoptosis, mitotic inhibitors in the G2/M and subG1 phases of the cell cycle, microtubules inhibitor, inhibitor topoisomerase enzyme, anti-metastasis, induces caspase-cascade and anti-angiogenesis.

Cytotoxicity is the level of a substance's toxic activity against a cell. It is one of the most important indicators for biological evaluation in vitro studies. Cancer drugs have different cytotoxic mechanisms: the destruction of cell membranes, prevention of protein synthesis, irreversible binding to receptors, inhibition of polydeoxynucleotide elongation, and enzymatic reactions. The cytotoxic and cell viability tests are required to determine cell death caused by this damage. While the cytotoxic compound can damage normal cells and cancer cells, it is used to inhibit the growth of malignant tumor cells.

*Voacanga foetida* (Blume) Rolfe, (Apocynaceae) which are found in West Sumatra Indonesia known as "Tampa badak". They are a small tree of 20 meters long, with white sap, stem diameter reaches 40 centimeters. This plant is widely spread in Indonesia, Malaysia, and the Philippines. *V. foetida* leaves were generally used to treat various skin conditions such as wounds, itching, swelling, and have been used externally for skin disorders (Hadi *et al.*, 2019).

*V. foetida* is high in alkaloids (Susanty *et al.*, 2018). It is a source of the indole alkaloids lombine, voacangine, voacristine, and coronadine (Hadi *et al.*, 2019; Zocoler *et al.*, 2005). It is also known as a source of bis-indole alkaloids or alkaloid dimers vobtusine (Susanty *et al.*, 2020). The bis-indole alkaloid consists of two identical or distinct indole alkaloid monomer units held together by C-C, C-O-C, or C-N bonds. Previous researchers have also shown that the leaves and bark of these plants exhibited potential cytotoxicity activity.

This study is a continuation of our research on other Sumatran plants (Dachriyanus *et al.*, 2000; Hirasawa *et al.*, 2019; Amelia *et al.*, 2019; Susanty *et al.*, 2014). In this work, we reported the structural characterization of vobtusine lactone and the cytotoxic activity of two isolated alkaloids from *V. foetida* against the A549 cell line. The molecular docking study was also performed to predict their cytotoxic mechanism against several pro-survival receptors (Bcl-2, Bcl-xL, Mcl-1), pro-activation receptor (Bax), and apoptotic execution receptor (caspase-3).

## MATERIAL AND METHODS

Lung adenocarcinoma (A549) cell line was obtained from Tohoku University, Japan. Chemical reagents used in this work include trypsin-EDTA 0.25% (Gibco), trypan blue (Invitrogen), penicillin (10,000 units/mL), Streptomycin (10,000 µg/mL), and RPMI 1640 (Wako). The equipment and instrument were used in this work include flask culture (diameter 25 cm<sup>2</sup>), corning 2 µm, 96 well-plate, 15 mL and 50 mL falcon tubes, laminar airflow, CO<sub>2</sub> incubator (HERA cell 150, Thermo electron corporation), Biological Safety Cabinet Class II Type A/B3 (Sanyo), Microplate Spectrometer (BIO-RAD), UV Ultra spec 2100 pro, NMR spectrometer (Bruker AV-400), mass spectrometer (ESI-MS Waters ZQ-2000), and CD J-820 spectropolarimeter.

### Extraction of *V. foetida* bark and isolation of alkaloids

The *V. foetida* barks were collected from the forest of biological education and research of Andalas University, West Sumatera, in January 2019. The specimen was identified at herbarium Andalas University. The dried bark of *V. foetida* (5.2kg) was ground and extracted for five days with methanol (10L). This maceration was repeated twice and the combined extracts were concentrated under reduced pressure. The concentrated extract was diluted with tartaric acid (3%, 500mL) and then partitioned with dichloromethane (5x500mL). The acidic layer was basified with sodium bicarbonate and then extracted with dichloromethane (5x500mL). The combined dichloromethane base extract was concentrated under reduced temperature to give a crude alkaloid fraction as a dark brown gum (11.3g).

A portion of crude alkaloid fraction (10g) was subjected to column chromatography on amino silica with the increasing amount of ethyl acetate in hexane as eluent. Fractions with contain similar patterns on TLC were combined and as much 100 µL of fraction solution (20 mg/mL) was subjected to HPLC ODS MS-II column (20x30mm) and a mixture of 3% ACN in H<sub>2</sub>O (0.1% formic acid) was used as mobile phase. Resolution factor (R) = 0.16. Flow rate = 8 mL/min. The similar HPLC fractions were combined, dried, and recrystallized to give a pure isolated compound. Then, the structure of the pure isolated compound was confirmed by spectroscopic analysis, including UV-Vis, IR, MS, XRD, 1D, and 2D NMR analysis. In addition, the optical rotation was determined.

Table I. SMILES data (Pubchem ID and CAS Number) were created using the MOE builder application.

Compunds	Identity	Structure "SMILES"
Vocangine	Pubchem ID 73255 CAS Number 510-22-5	<chem>CCC1CC2CC3(C1N(C2)CCC4=C3NC5=C4C=C(C=C5)OC)C(=O)OC</chem>
Vobtusine	Pubchem ID 301819 CAS Number 19772-79-3	<chem>COC1=CC=CC2=C1N3CC4(CC5C3(C26CCN7C6 C8(C5) CCO C8CC7)O)CN9CCC12C9C3(C40CC3)CC(=C1NC1=CC=CC=C21)C(=O)OC</chem>
Vobtusine lactone	CAS Number 19772-81-7	<chem>COC1=CC=CC2=C1N3CC4(CC5C3(C26CCN7C6C8(C5)CC(=O)OC8CC7)O)CN9CCC12C9C3(C40CC3)CC(=C1NC1=CC=CC=C21)C(=O)OC</chem>
Abt-199	Pubchem ID 49846579	<chem>CC1(CCC(=C(C1)C2=CC=C(C=C2)Cl)CN3CCN(C C3)C4=CC(=C(C=C4)C(=O)NS(=O)(=O)C5=CC(=C(C=C5)NCC6CCOCC6)[N+](=O)[O-])OC7=CN=C8C(=C7)C=CN8)C</chem>
Abt-702	Pubchem ID 10341154	<chem>C1COCCN1C2=NC=C(C=C2)C3=NC4=NC=NC(=C4C(=C3)C5=CC(=CC=C5)Br)N</chem>
Kaempheritrin	Pubchem ID 5486199	<chem>CC1C(C(C(C(O1)OC2=CC(=C3C(=C2)OC(=C(C3=O) OC4C(C(C(C(O4)C)O)O)O)C5=CC=C(C=C5)</chem>

### Cytotoxic assay

The Cytotoxic assay was determined using MTT assay by following the previous procedure (Susanty *et al.*, 2020). Approximately 5000 cells/well of A549 cell line were plated in a 96-well plate (90µL/well) and incubated at 37°C in CO<sub>2</sub> incubator. The cells were cultured in 5% CO<sub>2</sub> for 24h. Various concentrations of the samples in DMSO (10µL) were added to the cell culture (20µg/mL for screening and four various concentrations of 5, 10, 20, and 40µg/mL for IC<sub>50</sub> calculation) and were incubated for 48h. Vinblastine solution was used as a positive control with a concentration of 100ng. As much as 15µL of MTT (3-(4,5-Dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide, SIGMA) solution (5 mg/mL) was added and the mixtures were incubated in a CO<sub>2</sub> incubator (37°C, 5% CO<sub>2</sub>) for two hours. After confirming the formation of formazan, 100µL of Stop Solution (10% SDS-HCl) was added to dissolve formazan. Optical density was measured using an ELISA microtiter plate reader at 550nm and 700nm (background). The cell viability (%) was determined by the following formula.

$$\text{Cell viability (\%)} = \frac{(A-B)}{(C-B)} \times 100\%$$

In this case, A = absorbance of media + cell + sample, B = absorbance of media, C = absorbance of media + cell.

### Molecular docking study

All ligands used in this study have been reported as anticancer, even Susanty *et al.* (2020) recently reported that the anticancer mechanism of vobtusine is to suppress the expression and function of the Bcl-2 protein, in turn triggering an increase in the expression of Bax and caspase-3. Likewise, the Abt-199, Abt-702, and kaempheritrin which were used as controls have also been reported to have their respective functions in suppressing Bcl-2 expression, increasing Bax and caspase-3 expression.

The structures of ligands were built from SMILES data (Pubchem ID and CAS Number) using the MOE builder application (Table I), while receptors were downloaded from [www.pdb.org](http://www.pdb.org). These receptors include pro-survival protein Bcl-2 (PDB ID 2W3L), Bcl-xL (PDB ID 2YXJ), Mcl-1 (PDB ID 3KZ0); pro-apoptotic protein Bax (PDB ID 1F16); and apoptotic execution protein caspase-3 (PDB ID 1GFW).

The ligands and receptors were prepared and their 3-dimensional structures were optimized by adding hydrogen, removing water molecules, adding partial charges, and minimizing energy. Then, the binding site was selected using the default MOE system and the ligands were docked onto the receptors. The docking complex was stored in pdb format, while the docking data were saved in mdb format and the ligand-receptor interaction was visualized.

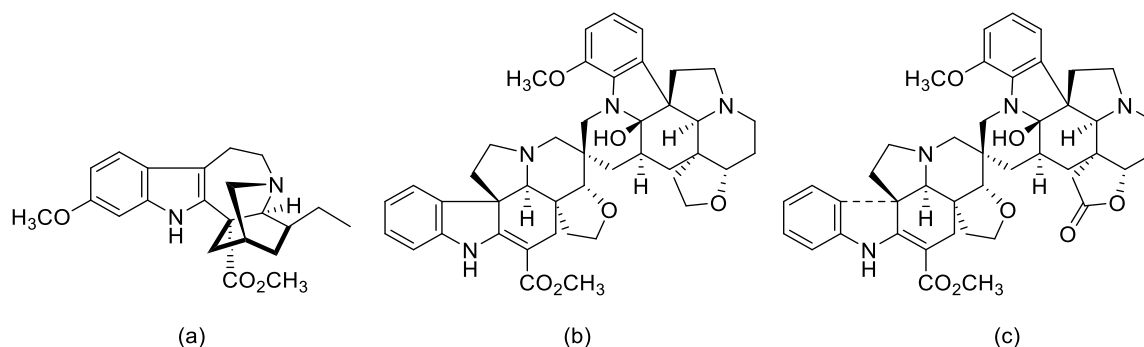


Figure 1. The structure of isolated compounds, (a) voacangine (Zocoler *et al.*, 2005), (b) vobtusine (Susanty *et al.*, 2020), and (c) vobtusine lactone ( $C_{43}H_{48}N_4O_7$ ) from DCM base fraction of *V. foetida* (VFB-DB2&3).

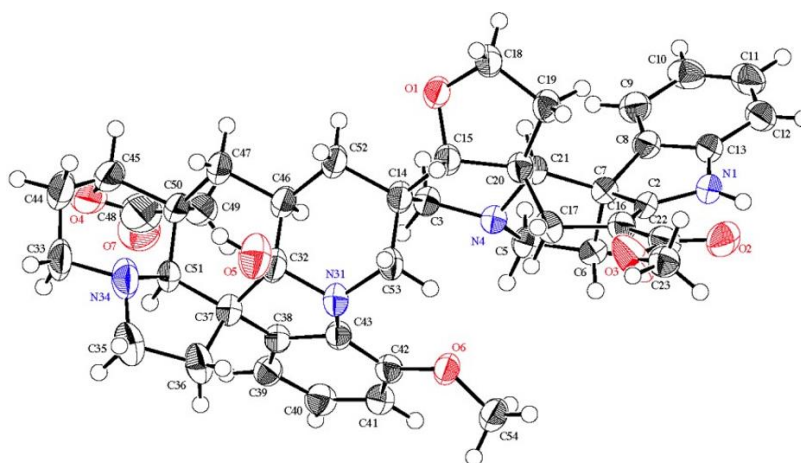


Figure 2. Crystallography structure of vobtusine lactone obtained by XRD analysis [Flack parameter 2776 Friedel pairs were 0.3].

## RESULT AND DISCUSSION

The structure and the crystallography structure of vobtusine lactone in this work were depicted in Figure 1 and Figure 2. The compound was isolated from the DCM base fraction of *V. foetida* (VFB-DB2&3).

### Characteristic of an isolated compound

The similar fractions were combined, dried, and recrystallized using a mixture of *n*-hexane/ethyl acetate to give pure vobtusine lactone (10.9mg). The compound was obtained as a colorless platelet crystal which decomposed at 300°C and  $[\alpha]_{26D} = -2.41$  (c. 0.1, MeOH).

The UV absorption bands of isolated compound (vobtusine lactone) at max 239.5nm ( $\epsilon=20042.2$ ), 267.1nm ( $\epsilon=14186.2$ ), and 321.9nm ( $\epsilon=20042.2$ ) indicated two indole chromophores. The IR spectrum showed an absorption band at 3370 $cm^{-1}$ . This absorption band indicates the

presence of N-H and O-H bonds. The absorption band at 1600 $cm^{-1}$  indicates the presence of a C=O bond and the absorption band at 2905 $cm^{-1}$  indicates the presence of an aliphatic C-H bond. The ESIMS+ spectra showed the molecular ion peak at  $m/z=731.5$  and it was matched with the molecular formula of vobtusine lactone,  $C_{43}H_{48}N_4O_7$  (Kunesch *et al.*, 1976).

The 1D and 2D NMR analyses were also performed to ensure the structure of the isolated compound. The  $^1H$ NMR spectra indicated the presence of an aromatic methoxy proton at 3.73ppm. Based on the HMBC analysis, the proton correlates with an oxy-aryl carbon (C12') at 145.3ppm. The  $^1H$  NMR spectra also indicate the presence of carbomethoxy proton at 3.79ppm. Based on the HMBC analysis, the proton correlates with the carbonyl of ester at 168.7ppm. In addition, the presence of N-H proton in indole moiety was indicated by the singlet signal at 8.93ppm.

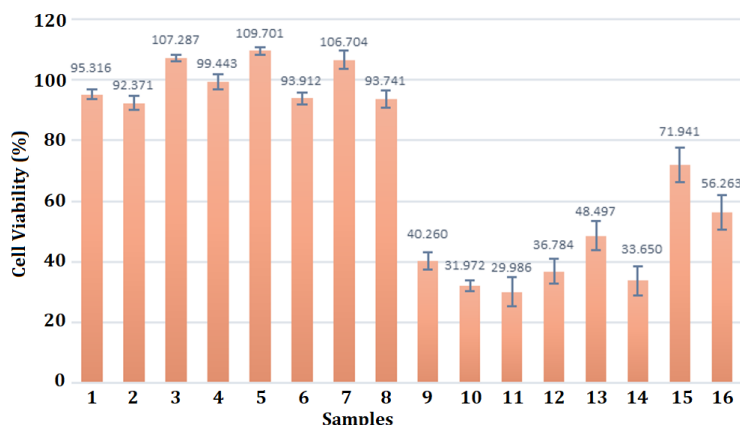


Figure 3. Percentage of cell viability of extracts, fractions, and isolated compounds from *V. foetida* (tested concentration = 20 $\mu$ g/mL). Sample 1 (VFL-Hex), 2 (VFL-EtOAc), 3 (VFL-EtOH), 4 (VFL-MeOH), 5 (VFL-DA), 6 (VFL-DB), 7 (VFL-BuOH), 8 (VFB-MeOH), 9 (VFB-DA), 10 (VFB-DB), 11 (VFB-BuOH), 12 (Voacangine), 13 (Vobtusine lactone), 14 (VFB-DB 4), 15 (VF-F19/20), and 16 (Vinblastine, 100ng). VFL = *Voacanga foetida* leaves, VFB = *Voacanga foetida* bark, Hex = *n*-hexane, EtOAc = ethyl acetate, EtOH = ethanol, MeOH = methanol, DA = DCM acid, DB = DCM base, BuOH = *n*-butanol

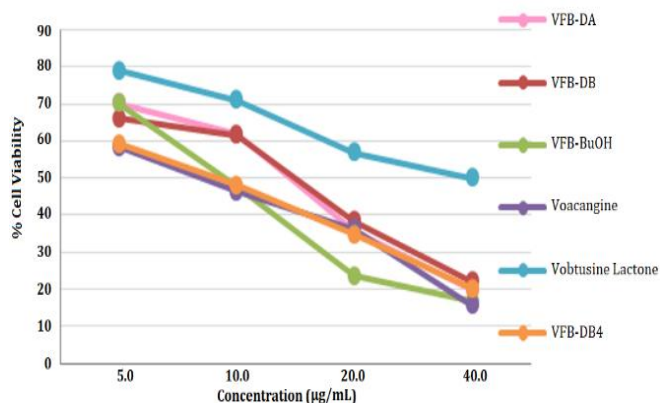


Figure 4. Percentage of cell viability of VFB-DA, VFB-DB, VFB-BuOH, Voacangine, Vobtusine lactone, and VFB-DB4 from *V. foetida* at various concentrations against A549 cell line at 48 hours incubation.

Then, the  $^{13}\text{C}$  NMR spectra also indicated the presence of carbonyl of lactone at 175.5ppm. This signal is important to determine whether the isolated compound is vobtusine or vobtusine lactone. Overall, the correlations in HSQC and HMBC spectra showed a suitable correlation with the structure of vobtusine lactone, and have high similarity with previously reported NMR spectra of vobtusine lactone isolated from *Tabernaemontana sphaerocarpa* Blume is a member of the Apocynaceae family (Zaima *et al.*, 2009).

Vobtusine lactone was crystallized from MeOH to give colorless platelet crystal. Crystal

data:  $a = 11.3328 (2) \text{ \AA}$ ,  $b = 12.1330 (2) \text{ \AA}$ ,  $c = 13.9244 (4) \text{ \AA}$ ,  $\beta = 110.3716 (4)^\circ$ ,  $V = 1794.87(10) \text{ \AA}^3$ ,  $Z = 2$ ,  $FW = 750.89$ ,  $D_{\text{calc}} = 1.389 \text{ g/cm}^3$ ,  $\mu$  for  $\text{Cu K}\alpha$  radiation is  $7.841 \text{ cm}^{-1}$ ,  $T = -180 \pm 1^\circ \text{C}$ . The structure was solved by direct methods and expanded using Fourier techniques. The non-hydrogen atoms were refined anisotropic, Hydrogen atoms were refined using the riding model. The final cycle of full-matrix least-squares refinement on F2 was based on 3441 observed reflections and converged with unweighted and weighted agreement factors of  $R1 = 0.0677$  [ $I > 2.00\sigma(I)$ ] and  $wR2 = 0.1967$ .

Table II. IC<sub>50</sub> value and cytotoxic classifications of fractions and isolated compounds from *V. foetida* against A549 cell line

Samples	IC <sub>50</sub> (µg/mL)	Cytotoxic classifications
VFB-DA	12.3	Highly Cytotoxic Activity
VFB-DB	12.4	Highly Cytotoxic Activity
VFB-BuOH	9.5	Highly Cytotoxic Activity
Voacangine	8.2	Highly Cytotoxic Activity
Vobtusine Lactone	37.4	Moderate Activity
VFB-DB4	8.5	Highly Cytotoxic Activity

### Cytotoxicity Activity

The results of the cytotoxic assay of extracts, fractions, and isolated compounds were performed in triplicate and the results were depicted in Figure 3. The figure explained that four fractions (VFB-DA, VFB-DB, VFB-BuOH, and VFB-DB4), dan two isolated compounds (voacangine and vobtusine lactone) exhibited a lower percentage of cell viability than vinblastine as a positive control at tested concentration. It means that some mentioned fractions and both isolated compounds have a higher ability to reduce the number of A549 cells.

The results of the cytotoxic assay as depicted in Figure 4 were performed in four variations of concentrations (5; 10; 20 and 40 µg/mL). The figure showed that voacangine (highlighted in purple line) exhibited the lowest percentage of cell viability at the lowest and highest tested concentrations, while the vobtusine lactone (highlighted in blue line) exhibited the highest percentage of cell viability at all various tested concentrations. Based on the result of IC<sub>50</sub> calculation as shown in Table II, vobtusine lactone has moderate cytotoxic activity with IC<sub>50</sub> of 37.4 µg/mL, while voacangine and four fractions exhibited very highly cytotoxic activity against A549 cell line.

The results of this study are in line with the study of Chen *et al.* (2016). Based on the study, voacangine also exhibited very high cytotoxic activity against HEPG-2, A375, MDA-MB-231, SH-SY5Y, and CT26 cancer cells with IC<sub>50</sub> values of 10; 14; 8; 7.7 and 11µg/mL, respectively. Then, the cytotoxic activity of vobtusine lactone has also been reported by Zaima *et al.* (2018). Vobtusine lactone exhibited IC<sub>50</sub> of 8.0; 18.0; 7.6; 17.6 and 22.5µM against HL-60, RPMI8226, NCI-H226, HCT116, and MCF7 cancer cells, respectively.

The difference in cytotoxic activity between fractions and isolates is thought as a result of

voacangine and vobtusine lactone have anticancer mechanisms, not in the cytotoxic pathway but the apoptotic pathway, this estimate is supported by previous research conducted by Macabeo *et al.* (2005) which states that the fractions of plants *V. globasa*, (same family with *V. foetida*) exhibited strong cytotoxic activity by apoptotic induction through the mitochondrial pathway. This study is also in line with our study. Seeing this reality, this *V. foetida* plant has the potential to be developed as a natural source of both bioactive fractions and compounds to fight cancers by apoptotic induction.

### Molecular docking study

The interaction model of ligand-receptor is further explained in silico via molecular docking techniques. The results of this study show that all ligands (tested compounds and control) have an affinity for all receptor targets. This is characterized by the release of energy when forming complexes with these receptors, but the strength of the affinity is different. The more negative the docking score obtained shows the stronger affinity with the receptor.

The pro-survival protein receptors showed that all three tested ligands (voacangine, vobtusine, and vobtusine lactone) had stronger affinity than controls (Abt-199). The strongest affinity was shown by vobtusine with a docking score of -10.04 kcal/mol, -11.69 kcal/mol, and -10.77 kcal/mol against Bcl-2, Bcl-xL, and Mcl-1 receptors, respectively (Table III).

In contrast to the pro-apoptotic receptor "Bax", all three tested ligands have weaker affinity than controls. The weakest affinity was shown by vobtusine with a docking score of -6.86 kcal/mol, then vobtusine lactone with -6.99 kcal/mol, and voacangine with -8.99 kcal/mol. Abt-702 as the control ligand shows the strongest affinity with a docking score of -9.95 kcal/mol (Table IV).

Table III. Docking score of ligands with pro-survival receptors

Receptors	$\Delta G$ (kcal/mol)				Binding site
	Voa.	Vob.	Vob. Lac.	Cont.	
Bcl-2	-9.93	-10.07	-8.31	-9.03	Arg 86, Phe 97, Ala 90, Val 93, Glu 94, Trp 135, Glu 138, Tyr 139, Arg 142, His 143
Bcl-xL	-9.77	-11.69	-10.29	-9.76	(Thr 115, Pro 116, Gly 117, Thr 118, Ala 119, Tyr 120, Leu 162, Trp 169)1; (Tr 120, Phe 123, Glu 124, Val 127, Asn 128, Trp 169, Thr 172, Tyr 173, Asp 176, His 177)2
Mcl-1	-10.70	-10.77	-10.02	-9.53	(Glu 16, Ans 17, Tyr 20, Tyr 21)1: (Val 216, Gly 219, Val 220, Asn 223, His 224)2

Table IV. Docking score of ligands with pro-apoptotic receptor

Receptor	$\Delta G$ (kcal/mol)				Binding site
	Voa.	Vob.	Vob. Lac	Cont.	
Bax	-8.99	-6.87	-6.99	-9.95	Tyr 164, Phe 165, Gly 166, Thr 167, Thr 169, Trp 170, Thr 172, Val 173

Table V. Docking score of ligands with apoptotic execution receptor

Receptor	$\Delta G$ (kcal/mol)				Site Binding
	Voa.	Vob.	Vob. Lac	Cont.	
Caspase-3	-12.05	-12.21	-12.02	-14.25	(Thr 62, Ser 65, Gly 66, His 121, Cys 163)1; (Tyr 204, Ser 205, Trp 206, Arg 207, Asn 208, Ser 209, Phe 256)2

Likewise, with the caspase-3 protein receptor, all three tested ligands showed weaker affinity than the docked ligand (kaempheritrin). Each releases energy of -12.05 kcal/mol (voacangine), -12.21 kcal/mol (vobtusine), -12.02 kcal/mol (vobtusine lactone). Kaempheritrin as a control ligand releases greater energy (-14.25 kcal/mol) (Table V).

Hydrogen bonding sufficiently affects the level of ligand affinity when interacting with the pro-survival protein receptor (Bcl-2) (Figure 5). The interaction of the vobtusine with the Bcl-2 receptor showed that the presence of 2 hydrogen bonds on the Tyr-139 and Arg-142 residues obtained an affinity of -10.07 kcal/mol. The presence of 1 hydrogen bond between the vobtusine lactone and the Bcl-2 receptor on the Arg-142 residue obtained an affinity of -8.31 kcal/mol. The presence of 1 hydrogen bond of the Abt-199 control ligand with the Bcl-2 receptor on the Arg-142 residue obtained an affinity of -9.03 kcal/mol (Table III). Thus, the more hydrogen bonds formed, the greater the affinity.

However, the amount of affinity is not only determined by the presence of hydrogen bonds. The voacangine does not show the formation of a hydrogen bond with the Bcl-2 receptor, but the strength of the affinity is quite large, reaching -9.93 kcal/mol. The voacangine has a simpler structure, this is what makes it easier for voacangine to enter the binding site of the Bcl-2 receptor. This statement is supported by (Saffari-Chaleshtori *et al.*, 2019) who reported the inhibitory activity of thymoquinone against Bcl-2. Even though when interacting with the Bcl-2 receptor, this compound does not form hydrogen bonds at all, this compound has a simple spatial structure that allows easy entry to the Bcl-2 receptor binding site.

Likewise, the Bcl-xL receptor shows that the number of hydrogen bonds is not a dominant factor in increasing of the ligand affinity for the Bcl-xL receptor (Figure 6). The voacangine ligand formed one hydrogen bond when interacting with Bcl-xL at residue Arg-165 with an affinity of -9.77 kcal/mol.

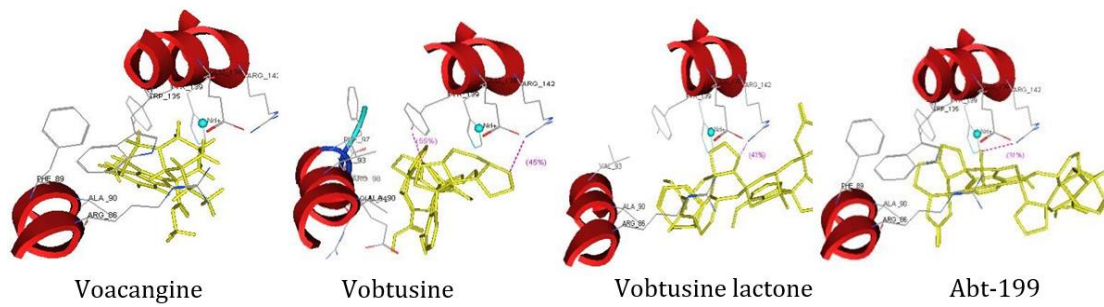


Figure 5. Models of ligand interactions with Bcl-2 receptor

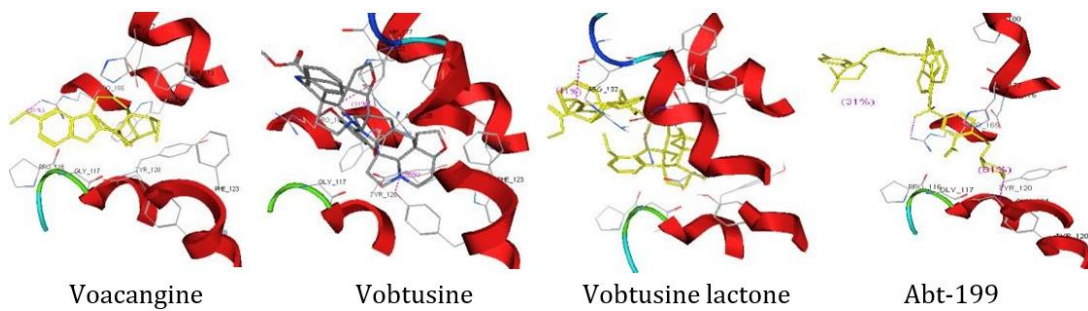


Figure 6. Models of ligand interactions with Bcl-xL receptor

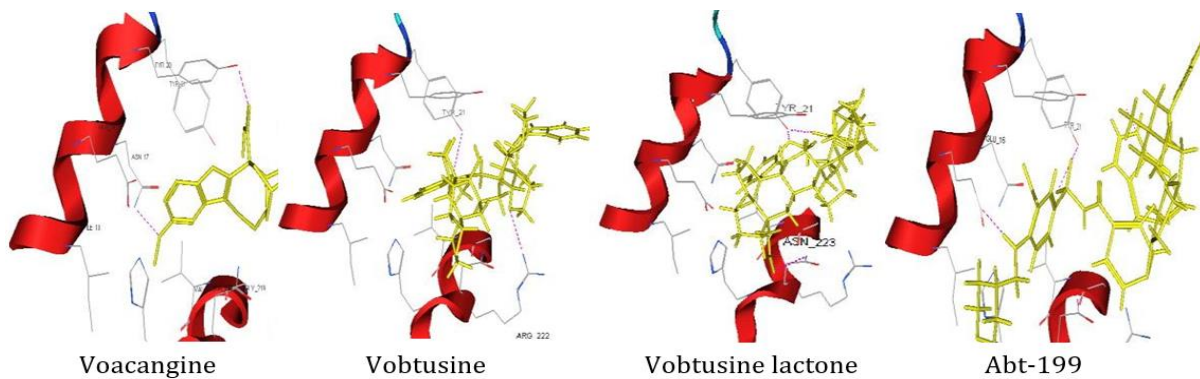


Figure 7. Models of ligand interactions with Mcl-1 receptor

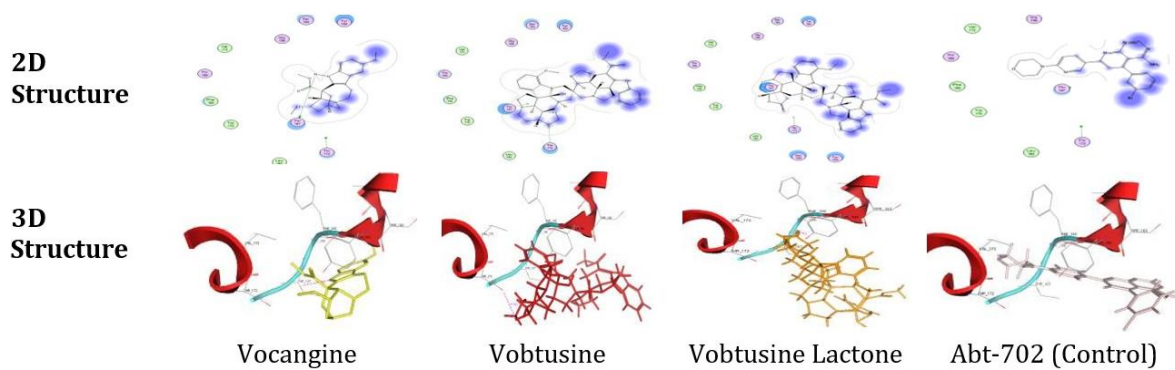


Figure 8. Models of ligand interactions with Bax receptor



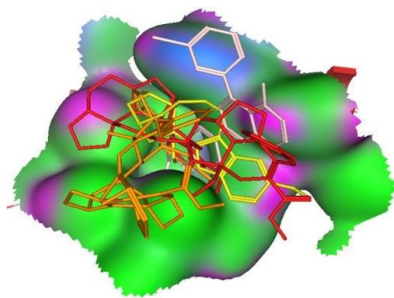


Figure 9. Interaction model of ligands [voacangine (yellow), vobtusine (red), vobtusine lactone (orange), and control (white)] vs Bax receptor with a narrow gap of non-polar “hydrophobic groove” (Blue = semi-polar residue, Purple = polar residue, and Green = non-polar residue).

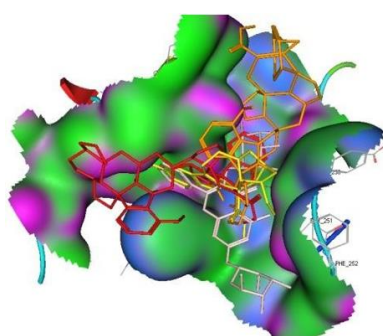


Figure 10. Interaction model of ligands [voacangine (yellow), vobtusine (red), vobtusine lactone (orange), and control (white)] with caspase-3 receptor. The binding site area having polar and semi-polar residues (Blue = semi-polar residue, Purple = polar residue, and Green = non-polar residue).

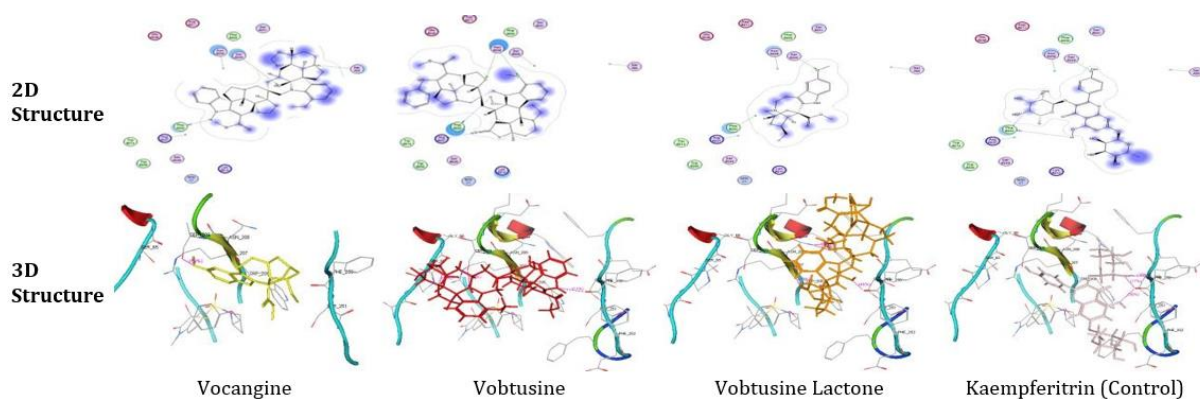


Figure 11. Models of ligand interactions with caspase-3 receptor

The vobtusine lactone formed one hydrogen bond when interacting with Bcl-xL at residue Arg-132 with an affinity of -10.29 kcal/mol. Meanwhile, the vobtusine formed two hydrogen bonds when interacting with Bcl-xL at Tyr-120 and His-177 residues with the strongest affinity of -11.69 kcal/mol. Likewise, the Abt-199 as control ligand

formed two hydrogen bonds when interacting with Bcl-xL at Tyr-120 and Arg-165 residues but with the weakest affinity of -9.76 kcal/mol (Table III).

The Mcl-1 receptor showed a ligand interaction pattern that was not different from the Bcl-2 or Bcl-xL receptors (Figure 7). Although vobtusine has two hydrogen bonds (Asn-12 and

Tyr-21) which gives the strongest affinity of -10.77 kcal/mol, the vobtusine lactone has three hydrogen bonds (2 Tyr-21 and Arg-222) but gives a lower affinity with the value of -10.02 kcal/mol. Likewise, voacangine formed three hydrogen bonds (2 Tyr-21 and As-223) with affinity -10.70 kcal/mol, and the Abt-199 as control ligand also with three hydrogen bonds (Glu-16 and 2 Tyr-21) but the affinity is even the weakest of -9.53 kcal/mol (Table III).

The same interaction pattern is shown by the pro-apoptotic protein receptor "Bax" (Figure 8). The strongest affinity value was indicated by Abt-702 as a control ligand (-9.95 kcal/mol) with a simpler structure, the same was seen in the voacangine has one hydrogen bond with the second strongest affinity value of -8.99 kcal/mol. Furthermore, the vobtusine and vobtusine lactone have lower affinity values of -6.87 and -6.99 kcal/mol, respectively (Table IV).

Interestingly, the Bax receptors have a "hydrophobic groove" gap for the ligand to enter the binding site (Figure 9). Interaction model of ligands [voacangine (yellow), vobtusine (red), vobtusine lactone (orange), and control (white)] vs Bax receptor with a narrow gap of non-polar "hydrophobic groove" (blue = semi-polar residue, purple = polar residue, and green = non-polar residue).

This gap greatly determines the affinity of the ligands that interact with it (Liu *et al.*, 2016). The smaller molecular weight ligands can infiltrate the narrow gap of the "hydrophobic groove" to interact with these Bax receptors. The structure of the Abt-702 control ligand with two benzoyl ring branches can reach a wider area of the binding site so that it has a stronger affinity than the vobtusine and vobtusine lactone with larger structures.

The caspase-3 protein receptor shows that the binding site is dominated by polar (purple) and semi-polar (blue) residues (Figure 10). The interaction model of ligands [voacangine (yellow), vobtusine (red), vobtusine lactone (orange), and control (white)] with the caspase-3 receptor. The binding site area having polar and semi-polar residues (Blue = semi-polar residue, Purple = polar residue, and Green = non-polar residue).

This binding site pattern allows the control ligand of kaempheritrin, a flavonoid derivative that has three benzoyl side chains to reach three sides of the caspase-3 binding site with bonds to the semi-polar residue. Besides, there are three hydrogen bonds formed between the kaempheritrin ligand and the Phe-250 residue. The

ability of the kaempheritrin ligand to reach the three sides of the caspase-3 binding site also contributes to the strength of the affinity to be the strongest, which is -14.24 kcal/mol.

The vobtusine and vobtusine lactone with four hydrogen bonds also showed strong affinities with -12.21 kcal/mol and -12.02 kcal/mol, respectively, but they were unable to reach all sides of the caspase-3 binding site. Hydrogen bonding in the residue (Table V) was reported as a typical description of the potential of the ligand as a caspase-3 protein activator (Jhansi Lakshmi *et al.*, 2009). The interaction models were further illustrated in Figure 11.

## CONCLUSION

Indole and bis-indole alkaloids were isolated from *V. foetida*. Two compounds (voacangine and vobtusine) have been reported by previous workers and one compound (vobtusine lactone) was first isolated from this plant species. The result of the molecular docking study and *in vitro* cytotoxic evaluation explains that all fractions (VFB-DB4, VFB-BuOH, VFB-DA, VFB-DB) and isolated compounds (voacangine, vobtusine, and vobtusine lactone) exhibited potential cytotoxic activity. The result suggests that this plant has the potential to be developed as a natural source of both bioactive fractions and compounds to fight lung cancer by apoptotic induction via suppressing Bcl-xL and Mcl-1 and activating Bax and caspase-3. However, toxicity evaluation is required to ensure their safety as natural anticancer agents.

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