

In silico Study on the Promising Active Components of Terpenoid and Fucoïdon from *Sargassum* sp. in Inhibiting CGRP and TNF- α

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ABSTRACT

Introduction: The new discovery of the active substance in *Sargassum* sp marks the new era for drug industry as it is very effective as the new migraine medication compared to analgesics which have already been popular previously in treating migraine. By using the *in silico* methods, this study intended to identify the preventive effect of the active substance in *Sargassum* sp within the stage of pain and inflammation development in migraine. In migraine pathophysiology, the clinical findings would build and verify the role of CGRP and TNF- α . **Methods:** This research applied a one-shot experimental study and by employing the potential test through PubChem (<https://pubchem.ncbi.nlm.nih.gov/>), the result of this study proved that tannins, terpenoids and fucoidone were contained in the active substance of *Sargassum* sp leading to the possession of potential as the drug to treat migraine. **Results:** Terpenoids and tannin binding affinity value is higher than other substances. Terpenoids and fucoidon had similar amino acid residues with controls. Seaweed metabolites have great potential as inhibitors of CGRP and TNF- α because the binding affinity score is close to control. **Conclusion:** The active substance in *Sargassum* sp has an inhibitory effect on the occurrence of CGRP and TNF- α in migraine based on *in silico* studies.

Key words: CGRP, Fucoidone, *In silico*, *Sargassum* sp., Terpenoid, TNF- α .

INTRODUCTION

Generally, seaweed carries three phyla consisting of *Chlorophyceae* (green algae), *Phaeophyceae* (brown algae), and *Rhodophyceae* (red algae). In every phylum, it comprises with thousands of various species. In Asian countries, seaweed has been familiarly utilized as both foods and traditional remedy ages ago by many people. Known as the source of functional metabolites, the people's attention has grown greatly toward it^{1,2}. Some of its functional metabolites possessed by seaweed are its potentials as antibacterial, antiviral, anticancer, anti-hypersensitive, anti-diabetic and anti-inflammatory. Thus, the 2020 database from Food and Agriculture Organization (FAO) recorded the increase production of seaweed has been tripled from 10.6 million tons in 2000 reaching to 32.4 million tons in 2018.

Florotanin, fucoxanthin, fucoidan, alginate, fukosterol, merodit- erpenoids and phenolic acids are the dominant types of bioactive compounds in *Sargassum* sp. The biological activity of the bioactive compound *Sargassum* sp. namely antioxidant, anti-cancer, antitumor, antiallergic, anti-inflammatory, antihypertensive, antiobesity, antidiabetic, antimicrobial, anti-browning, neuroprotective and hypocholesterolemic.^{1,2} Terpenoids can influence various mechanisms of inflammation arising in response to various etiological factors because comprehensive studies of terpenoid mechanisms have shown therapeutic effects on inflammation.³

In many previous studies, the role of *calcitonin gene-related peptide* (CGRP) in migraine pathophysiology has been identified and confirmed by clinical findings. Capi's study revealed that trigeminal ganglia neurons secrete CGRP then multiply its transcription under the conditions

of mimicking the neurogenic inflammation and pharmacotherapy in migraine by impeding CGRP transcription and lessening the CGRP release as well as tumor necrosis factor α (TNF- α) which is an endogenous inflammatory mediator implicated in migraine.⁴

In silico method can be effectively employed in finding the interplay of active substances with *Sargassum* sp. up to the molecular level. *In silico* study is highly considered as the primary step in discovering the candidate of treatments or remedy prior to both *in vitro* or *in vivo* investigations.^{5,6} This investigation intended to examine and analyze the preventive effects in the active substances of *Sargassum* sp in contrast to the development of CGRP and TNF- α in migraine.

MATERIALS AND METHODS

This investigation employed a one-shot experimental study type applying an *in silico* method which was conducted at the INBIO Biomolecular & Bioinformatics Laboratory located in Lowokwaru, Malang, East Java, Indonesia.

Protein structure and ligand

The database of PubChem (<https://pubchem.ncbi.nlm.nih.gov/>) was utilized as the source in which the active substances from *Sargassum* sp was obtained from as shown in Table 1. The The three-dimension structure of CGRP and TNF- α a was generated from the database of INBIO Laboratory or from Research Collaboratory for Structural Bioinformatics Protein Data Bank (RCSB PDB) (<https://www.rcsb.org/>).⁷

Molecular docking

Autodock Vina on the PyRx 9.5 program was utilized for molecular docking process. Table 2 presents the targets of molecular docking process in the form of proteins.

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Table 1: Ligand used from *Sargassum* sp.

PUBCHEM 16133892	Tannin
PUBCHEM 92023653	Fucoidan
PUBCHEM 451674	Triterpenoid

Table 2: Protein Target.

ID	Control Inhibitor
PDB 3N7I	CGRP Receptor Olcegepant
PDB 2AZ5	TNF- α 6,7-DIMETHYL-3-[(METHYL{2-[METHYL({1-[3-(TRIFLUOROMETHYL)PHENYL]-1H-INDOL-3-YL}METHYL)AMINO]ETHYL}AMINO)METHYL]-4H-CHROMEN-4-ONE

From the molecular docking process, it was derived the forecast of the interaction potency between receptor and ligand in regard to their binding affinity. More negative scores indicated stronger relation. The indication that the antagonist activity happened from the examined substance toward the target protein when the tested substance displayed the score which was close to the control one. Olcegepant and 6,7-DIMETHYL-3-[(METHYL{2-[METHYL({1-[3-(TRIFLUOROMETHYL)PHENYL]-1H-INDOL-3-YL}METHYL)AMINO]ETHYL}AMINO)METHYL]-4H-CHROMEN-4-ONE became the protein target of this docking process. The visualization of molecular docking result was derived by using PyMol 2.3.1. LigPlot 2.1. was applied to generate the interaction of amino acid. The bond between the receptor and the control inhibitor was imitated using a grid box as a special docking process performed in this investigation.

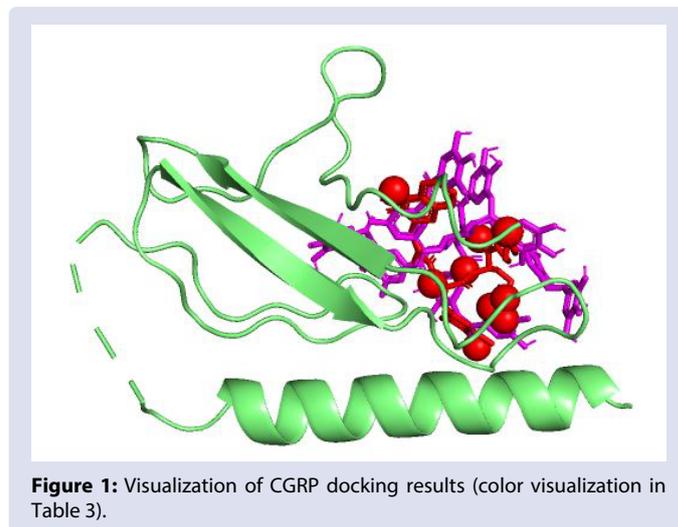
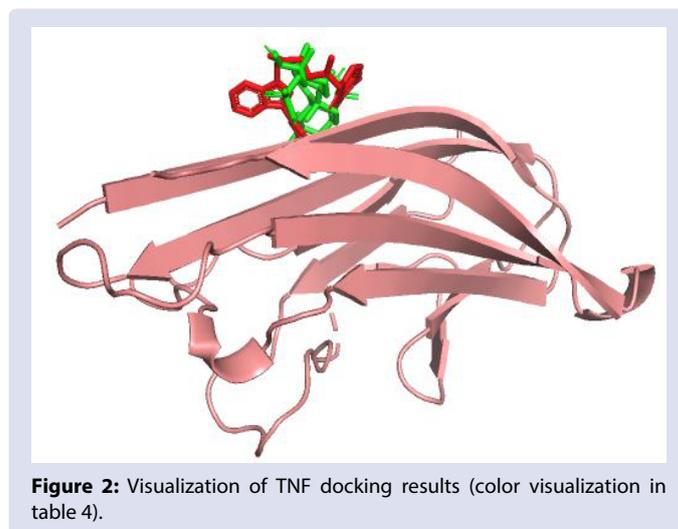
RESULTS

Analysis molecular docking

A specific docking process performed in this investigation with the imitation of the bond between the receptor and the inhibitor control drug using a grid box. The prediction of the interaction's potency between the receptor and the ligand in regard to the binding affinity score was estimated using the docking process. A stronger interplay existed between the receptor and the ligand was displayed by more negative value. It also could indicate that an activity as the inhibitor of target protein by metabolite which was close to the capabilities of drug inhibitor when the examined metabolites possessed the score near the control one.

Sargassum sp comprised with active substances which included tannin, triterpenoid and fucoidan. The binding affinity for each substance with CGRP and TNF- α was indicated using the molecular docking. The indication of stronger relation was revealed by the more negative score of binding affinity. The crystal structure complex of CGRP with olcegepant as the control inhibitor, also with TNF- α with 6,7-DIMETHYL-3-[(METHYL{2-[METHYL({1-[3-(TRIFLUOROMETHYL)PHENYL]-1H-INDOL-3-YL}METHYL)AMINO]ETHYL}AMINO)METHYL]-4H-CHROMEN-4-ONE as the control inhibitor was the target of this molecular docking. Figure 1 and 2 display the visualization of molecular docking's result by PyMol 2.3.1. Table 3 presents that the most negative score was revealed in tannin with the strongest affinity compared to CGRP, followed by triterpenoid, control and fucoidan. Meanwhile, Triterpenoid displayed the strongest affinity with the most negative score compared to TNF- α , followed by control, and fucoidan as shown in Table 4. Based on the docking results, seaweed metabolites have great potential as inhibitors of CGRP and TNF- α because the binding affinity score is close to control. CGRP and TNF- α increased in migraine conditions, it is hoped that the administration of *Sargassum* sp can help normalize the protein levels.

Four hydrophobic bonds and one hydrogen bonds in amino acid residue were revealed in triterpenoid which was similar to the control as presented in Table 5. Meanwhile, fucoidan displayed three hydrophobic bonds in amino acid residue that was also similar with the control. The interaction between the metabolite of *Sargassum* sp could be seen from the visualization results of LigPlot with the identical position of target protein with the control as both control and metabolite had the same amino acid residues.

**Figure 1:** Visualization of CGRP docking results (color visualization in Table 3).**Figure 2:** Visualization of TNF docking results (color visualization in table 4).**Table 3: The result of molecular docking between CGRP and ligand.**

Substance	Binding Affinity (kcal/mol)	Color
Tannin	-8.1	Violet
Fucoidan	-6.2	Yellow
Triterpenoid	-7.1	Green
Control	-7.4	Red

Table 4: The result of molecular docking between TNF α and ligand.

Substance	Binding Affinity (kcal/mol)	Color
Tannin	-0.8	Violet
Fucoidan	-4.3	Yellow
Triterpenoid	-5.7	Green
Control	-6.7	Red

Table 5: LigPlot visualization results.

Receptors	Compound	Hydrophobic Bond	Hydrogen Bond
TNF- α	Control	GLN61, TYR119 , LEU120 , LEU57 , GLY121, SER60, TYR59	TYR151
	Triterpenoid	TYR59, LEU57, ILE118, LYS98, LEU120, TYR151	TYR119
CGRP	Control	MET42, ASP70 , GLY71, ARG119 , TRP124, TRP121 , PHE92, PHE95, ILE41	THR122, TRP72
	Fucoïdon	ASP70, TYR124, TRP72	ARG119, THR122, TRP121

Note: The same amino acid residue between control and seaweed metabolite ligands is displayed by the bold sign.

DISCUSSION

Based on the docking results, seaweed metabolites have great potential as inhibitors of CGRP and TNF because the binding affinity score is close to control. CGRP and TNF α increased in cephalgia conditions, it is hoped that the administration of *Sargassum sp.* can help normalize the protein levels. Similar results were found in a study by Handajani and Prabowo where brown seaweed extract (*Sargassum sp.*) at a dose of 400 mg/kg was able to significantly reduce TNF- α expression and repair joint damage. The results showed that the expression of TNF- α in the adjuvant arthritis group with the treatment of brown seaweed extract and cold stressor decreased significantly up to 48.4% and the repair of joint damage in the metatarsophalangeal tissue was marked by the repair of chondrocyte cells.⁸

Tannins in the ethanol extract of *Sargassum sp.* act as free radical scavengers that can inhibit the formation of superoxide O₂⁻ anions and convert them into more stable products, thereby increasing the enzyme superoxide dismutase (SOD) and reducing oxidative stress. Another study by Aulanni'am reported that *Sargassum sp.* was able to reduce free radicals in animal models of inflammatory bowel disease.^{9,10}

Activity of anti-inflammatory and antinociceptive both *in vitro* and *in vivo* was found in Terpenoid from *Cannabis*. The impacts operate independently of TNF- α . Terpenoids in *Cannabis* exhibit only momentary immunosuppression, thus, by using it, the acute inflammation can be overcome. Terpenoids can be abundantly found in plants as the rich source. The development of chronic joint swelling is able to be inhibited by terpenoids. Furthermore, terpenoids contribute to the various mechanisms correlated to any inflammation appearing as the reaction various etiologic factors.^{3,11}

In addition, previous result of Hu C's investigation exhibited that the fall of mechanical allodynia and thermal hyperalgesia can be the result of fucoïdon. The inhibition of neuroimmune activation can be impeded by fucoïdon fundamentally indicated with the glial activation, cytokine production of TNF- α , IL-1 β , IL-6 and the activation of extracellular signal-regulated protein kinase. In previous study, any analgesic effect from intrathecal fucoïdon in rats treated with spinal nerve ligation is related to the inhibition of neuroimmune activation which is also in line with the maintenance of neuropathic pain. Moreover, fucoïdon's inhibition on the expression of inflammatory cytokines reciprocates to its declined hypersensitivity.¹²

CONCLUSION

There are three substances tannin, terpenoids and fucoïdon in *Sargassum sp.*, which have the potential as anti-migraine. Terpenoid and Tannin binding affinity value is higher than other substances. Terpenoids and fucoïdon had similar amino acid residues. Therefore, the active substance in *Sargassum sp.* has an inhibitory effect on the occurrence of CGRP and TNF- α in migraine based on *in silico* studies.

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DISCLOSURE STATEMENT

The authors have no conflicts of interest to declare.

ABBREVIATIONS

CGRP: *Calcitonin Gene-related Peptide*; TNF α : Tumor Necrosis Factor α ; FAO: Food and Agriculture Organization; O₂⁻: Superoxide; SOD: Superoxide dismutase; GLN: Glutamine; TYR: Tyrosine; LEU: Leucine; GLY: Glycine; SER: Serine; ILE: Isoleucine; LYS: Lysine; MET: Methionine; ASP: Aspartic Acid; ARG: Arginine; TRP: Tryptophan; PHE: Phenylalanine; THR: Threonin.

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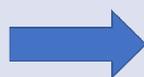
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GRAPHICAL ABSTRACT



Sargassum sp.



PUBCHEM 16133892 - Tannin
PUBCHEM 92023653 - Fucoïdan
PUBCHEM 451674 - Triterpenoid



Calcitonin gene-related peptide (CGRP)
Tumor necrosis factor α (TNF- α)

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