

Biofungicide Producing Bacteria: an In Vitro Inhibitor of *Ganoderma boninense*

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ABSTRACT

Oil palm is widely known as one of vegetable oil sources and the main commodity in Indonesian agriculture because of the benefits in non-food and food industries. *Ganoderma boninense* attack results in considerable losses to agriculture. Chemical control creates a harmful effect on health and the environment. Biocontrol is required to take over the function of chemical control. This study aimed to select bacteria that produce bioactive compounds as biofungicide against *G. boninense* pathogenic fungi and identify bacteria producing biofungicide using molecular method. The stages of bacterial isolate selection were performed through the selected hemolysis and isolate tests in the antagonistic test. Bacteria were extracted using ethyl acetate and their extract activity were tested. Analysis of bioactive compounds was conducted using thin layer chromatography (TLC) and the identification was based on 16S rRNA gene. The result of bacterial pathogenic test was obtained from two selected bacterial isolates namely 11B LB and 11B MD. Both bacterial isolates showed antagonistic effects by forming an inhibitory zone against *G. boninense* growth with percentage of inhibitor of 81 and 75%. Activity test of bacterial extract showed that crude extract of bacterial isolate 11B MD had the highest inhibitor activity that is 88.34%. TLC analysis proved that the active extract of bacteria containing metabolite compounds had Rf value of 0.1, 0.28, and 0.38. Isolate bacteria 11B MD was identified as *Pseudomonas aeruginosa*.

1. Introduction

Oil palm (*Elaeis guineensis* Jacq.) is one of the vegetable oil sources that become the main commodity in agriculture in Indonesia. Production of palm oil is used in the food industries and chemicals. It also substitutes fuel oil which is currently mostly preferred for petroleum. Based on data from Food Agriculture Organization (FAO), Indonesia has the highest average area of oil palm plantation in 2008-2012 which amounted to 35.69% of the total area of oil palm plantation in the world. Malaysia ranks second for oil palm plantations and followed by Nigeria while Thailand ranks fourth (Indarti *et al.* 2014).

Productivity data for oil palm crops in 2008-2012 provided information on significant differences between oil palm plantation area and productivity

values, particularly in Indonesia and Thailand. Indonesia occupies the last position in terms of productivity compare to Thailand which has a productivity value of about 17.2 tons/ha (Indarti *et al.* 2014). The low value of palm oil productivity in Indonesia is caused by several factors and one of them is the attack of pathogenic microorganisms. One of the pathogens concerned in attacking oil palm crops is *Ganoderma boninense* which can cause basal stem rot (BSR) (Naher *et al.* 2012). Susanto *et al.* (2005) stated that basal stem rot disease known as BSR disease can affect oil palm production because it causes death up to 50%. In addition, BSR disease results in producing fewer fruits of palm oil plants or even do not produce any fruit at all, thereby this disease results in decreasing palm oil production.

Synthetic fungicides are used by farmers as attempts to suppress pathogen attacks. The use of synthetic fungicides can accelerate the occurrence of pathogenic races so that they are resistant. In addition,

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it can cause poison to humans and the residue can cause environmental pollution (Hadizadeh *et al.* 2009). Biocontrol is an effective alternative in controlling losses due to pathogenic fungi so as to reduce the impact of plant disease attack. The microorganisms that have the potential to be used as biocontrol agents against *G. boninense* attack are bacteria. Bacteria have a faster and more adaptable growth in less favorable environments. *Pseudomonas alcaligenes*, *Bacillus pumilus* and *Rhizobium sp.* can be used as biocontrol because it produces antifungal compounds that inhibit pathogenic fungi (Akhtar *et al.* 2010). Asaka and Shoda (1996) confirmed that several *Bacillus* strains has a variety of antifungal compounds such as cyclic lipopeptides which is effectively used as a biocontrol agent to overcome pathogens on plants.

The use of biocontrol agents that play a role in inhibiting the attack of *G. boninense* pathogenic fungal has not showed a significant effectiveness. Information on organisms that are potentially used as biocontrol agents is needed. Previous research results have obtained bacterial isolates from healthy oil palm plants in Lebak and Medan that are believed to have a role in resisting and inhibiting the growth of *G. boninense*. Identification of bacteria and analysis of metabolite compounds that have the ability as a biofungisida against pathogenic fungi have not been analyzed in this study using the bacteria isolates.

2. Materials and Methods

2.1. Selection of Bacterial Isolates through Hemolysis Test

Five bacterial isolates (P10 LB, 11BLB, 1TSA LB, 11B MD, and 1TSA MD) from the collection of Agromicrobiology Laboratory in Agency for Assessment and Application of Technology, South Tangerang were selected by hemolysis test. The five bacterial isolates were scratched on the blood agar medium and incubated at 37°C for 24 hours. Selection was performed on the pathogenicity of the five bacterial isolates characterized by the formation of the hemolytic zone.

2.2. Bacterial Isolate Antagonistic Test Against *Ganoderma boninense*

The bacterial antagonistic test against *G. boninense* was performed on the basis of Bivi *et al.* (2010) that is double culture method (dual culture). *G. boninense* cultures were taken about 1 cm and placed on Potato Dextrose Agar (PDA) medium. Bacterial isolates tested were scratched in contrast to the fungus. Positive control was performed using nystatin (commercial

antifungal). Subsequently incubation was performed for 7 to 14 days at room temperature, and observed the antagonistic effect of bacterial isolate on growth of mycelium fungus *G. boninense*. Inhibitory activity was known by measuring the growth of mycelium fungus and see whether or not the inhibition zone was formed. The inhibitory level was calculated using the formula $IP = R1 - R2 / R1 \times 100\%$, where IP = inhibition percentage, R1 = growth of pathogen in control, and R2 = growth of pathogens on dual culture treatment.

2.3. Measurement of Growing Bacterial Curve

Stages of measurement of bacterial growth began with bacterial preculture. The result of bacterial preculture was reoccupied on Nutrient Broth (NB) medium and incubated on a shake machine with 150 rpm agitation. Measurement of bacterial growth was done every 4 hours for 24 hours. Total plate count (TPC) method is used in the measurement of bacterial growth. A total of 1 ml of bacterial culture was inserted into a test tube containing 9 ml of 0.85% NaCl physiologic solution. Serial dilution was made from 10^{-1} up to 10^{-8} . The 10^{-5} to 10^{-8} dilution series were taken as much as 0.1 ml to be inserted into a petri dish containing Nutrient Agar (NA). It was trimmed and incubated at room temperature were done.

2.4. Extraction of Bacteria Producing Antifungus Compounds

Bacterial isolates were grown on NB media and harvested as bacterial cells begun in the logarithmic growth phase until death phase. The bacterial cells obtained were centrifuged at 10.000 rpm for 10 minutes. The supernatant portion was added solvent ethyl acetate with a ratio of 1:1 (v/v). The mixture was inserted into a separating funnel and shaken for 15 minutes. The formed organic solvent was concentrated using a rotary vacuum evaporator at a temperature of $\pm 40^\circ\text{C}$ (Abdel-Raouf and Ibraheem 2008).

2.5. Activity Test of Extracted Bacterial Culture on *Ganoderma boninense*

This test used modified cylinder-plate method (Cui *et al.* 2012) that is by making wells on PDA media using cork borer. Supernatant extracts from bacterial cultures were fed into a well with a concentration of 10.000 ppm. Next, about 1 cm of fungus *G. boninense* was placed on PDA media beside the wells containing extracts. As a positive control, nystatin was used in the same concentration with those on the extract. Negative control was performed with no addition of extract. Measurement of inhibition rate of bacterial extract was done according to method Bivi *et al.* (2010).

2.6. Thin Layer Chromatography (TLC)

The TLC plate used was silica gel GF₂₅₄ while the eluent used was the ratio of ethyl acetate and n-hexane concentration. Bottling of bacterial extract on TLC plate was done using capillary pipe. The elution process was performed by placing the TLC plate vertically on the chamber wall containing the eluent. The TLC plates were removed from the chamber to be observed staining spots formed under UV light λ 254 and λ 366 nm. The calculation of Rf values corresponds to Wagner *et al.* (1984).

$$R_f = \frac{\text{Range of component}}{\text{Range of eluent}}$$

2.7. Identification using Molecular Method

Bacterial DNA was isolated using the Cetyl Trimethylammonium Bromide (CTAB) method (Ausubel *et al.* 1997). DNA amplification was done using universal primer namely primer forward 27F and 1492R (Frank *et al.* 2008). The amplification process occurred on a Polymerase Chain Reaction (PCR) machine with composition and PCR reaction conditions were based on a procedure performed by Zarei *et al.* (2012). The PCR reaction composition was 15 μ l PCR Master Mix (i TaqTM) 2X, 3 μ l DNA template, 3 μ l for primary forward, 3 μ l reverse primer, and 6 μ l nuclease free water (ddH₂O). The PCR conditions used were pre-denaturation (94°C, 5 min), 35 cycles of denaturation (94°C, 0.5 min), annealing (55°C, 0.5 min), extension (72°C, 1.5 min), and post extension (72°C, 5 min). The result of amplification (amplicon) was assessed by electrophoresis on 1% agarose gel and the result of 16S rRNA amplicon is purified and sequenced. The results of sequencing were then aligned with the Gene Bank data. The process was carried out on database of National Center for Biotechnology Information website (<http://www.ncbi.nlm.nih.gov>) using BLASTN (Basic Local Alignment Search Tool-Nucleotida) to identify bacterial isolates.

3. Results

3.1. Selection of Bacterial Isolates Through Hemolysis Test

The bacterial isolates used in this study were selected based on the presence or absence of pathogens possessed by the five unidentified bacterial isolates: P10 LB, 11B LB, 1TSA LB, 11B MD, and 1TSA MD. The results show that there were 3 bacterial isolates that had the ability to lyse blood cells (pathogenic). The pathogenicity of bacteria was measured from the formation of hemolytic zones around the bacterial colonies that grew on the gel agar medium after incubation for 1 x 24 hours. Two

other bacterial isolates, 11B LB and 11B MD showed no activity in lysing blood cells (Figure 1). Both bacterial isolates were used in subsequent tests.

3.2. Antagonistic Test of Bacterial Isolate Against *Ganoderma boninense*

The results of antagonistic test showed that bacteria isolates 11B LB and 11B MD had effective capability in inhibiting growth of *G. boninense* i.e. 81% and 75% respectively. The positive control used was nystatin dissolved in dimethyl sulfoxide (DMSO) or methanol solvent which showed the percentage of inhibition rate of 83 and 78% (Table 1). In addition, there was an inhibition zone formed by bacterial isolates (Figure 2) which proved the two selected bacterial isolates had the ability to inhibit the growth of *G. boninense*.

3.3. Measurement of Growing Bacterial Curve

The result indicated that bacterial isolates had a short adaptation phase so that the growth can directly enter the logarithmic phase (exponential phase). Both isolates were long enough in logarithmic phase, which presented from hours 0 to 16. In this phase, the bacterial cells in isolate 11B LB continued to increase in number from 1.5 x 10⁷ CFU/ml to 2.8 x 10⁹ CFU/ml. Bacterial cells in isolate 11B MD cells at the 0th hour was 2.1 x 10⁷ CFU/ml and it increased at 16th hour to 3.4 x 10⁹ CFU/ml. Stationary phase started after logarithmic phase. Stationary phase in both bacterial isolates was quite short and led to the phase of death. The death phase of both bacterial isolates

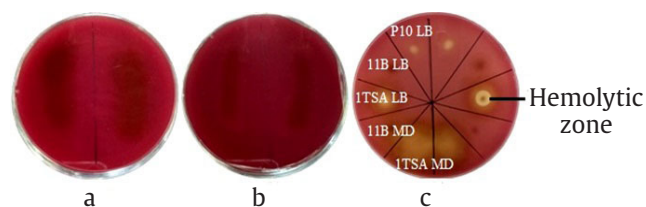


Figure 1. The result of hemolysis test of bacterial isolates on the blood agar media. (a) isolate 11B LB, (b) isolate 11B MD, and (c) the five bacterial isolates

Table 1. The percentage of inhibition of bacterial isolates against the growth *Ganoderma boninense* (%)

| Isolate | Percentage of inhibition (%) |
|---|------------------------------|
| Negative control (<i>G. boninense</i>) | 0 |
| 11B LB (Lebak) | 81 |
| 11 B MD (Medan) | 75 |
| Positive control (nystatin was dissolved into DMSO) | 83 |
| Positive control (nystatin was dissolved into methanol) | 78 |

11B LB and 11B MD occurred until the 24th hour. It was characterized by a decrease in the number of bacterial cells. Bacterial isolate 11B LB decreased in number of cells up to 24 hours with a value of 2.5×10^9 CFU/ml. Bacterial isolate 11B MD was 2.5×10^9 CFU/ml (Figure 3). This bacterial growth phase became a reference in determining suitable time in harvesting bacterial cells for the extraction process that is at 8th to 24th hour.

3.4. Activity Test of Extracted Bacterial Culture on *Ganoderma boninense*

The crude extract of viscous fluid was obtained from the supernatant extraction of bacterial culture in logarithmic growth phase to death phase. Extracted bacterial culture at the 8th, 12th, 16th, 20th, and 24th hour were tested to see their activity on *G. boninense*. The results of activity test explained that only crude extracts from bacterial culture 11B MD at 8th and 24th hour had inhibitory activity. The inhibition values were 86.67 and 88.34% (Table 2) respectively. Treatment of crude extract of bacterial isolate 11B LB did not show

any inhibition on growth of mycelium *G. boninense*. A positive control using nystatin showed an inhibition with a value of 60% while in the negative control there was no inhibitory activity (Figure 4).

3.5. Thin Layer Chromatography (TLC)

Extract of supernatant bacterial culture at the 8th and 24th hour indicated the presence of inhibition. The compound content was analyzed by TLC method.

Table 2. The inhibition percentage of bacterial supernatant extract in inhibiting the growth of *G. boninense* (%)

| Bacterial crude extract | Percentage of inhibition (%) |
|---|------------------------------|
| Negative control (without treatment) | 0 |
| Positive control (nystatin was dissolved in methanol) | 60 |
| 11 B MD (Medan) from the culture 8 th hour | 86.67 |
| 11B MD (Medan) from the culture 24 th hour | 88.34 |

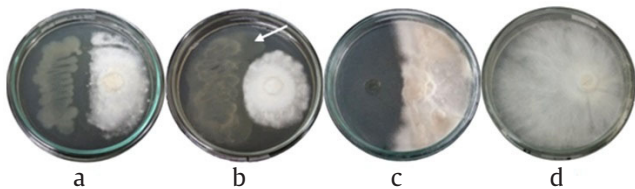


Figure 2. The effect of inhibition of bacterial isolates against of *G. boninense* with incubation time for 14 days at room temperature. (a) isolate bacteria 11B LB, (b) isolate 11B MD, (c) nystatin (positive control), and (d) growth of *G. boninense* (negative control)

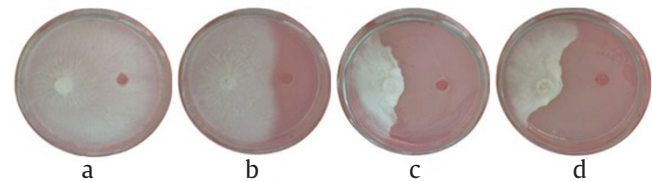


Figure 4. Effect of bacterial supernatant extract of isolate 11B MD on the growth of *G. boninense* with incubation time for 14 days at room temperature. (a) without crude extract (negative control), (b) nystatin (positive control), (c) supernatant extract of bacterial culture 11B MD at 8th hour, and (d) at 24th hour

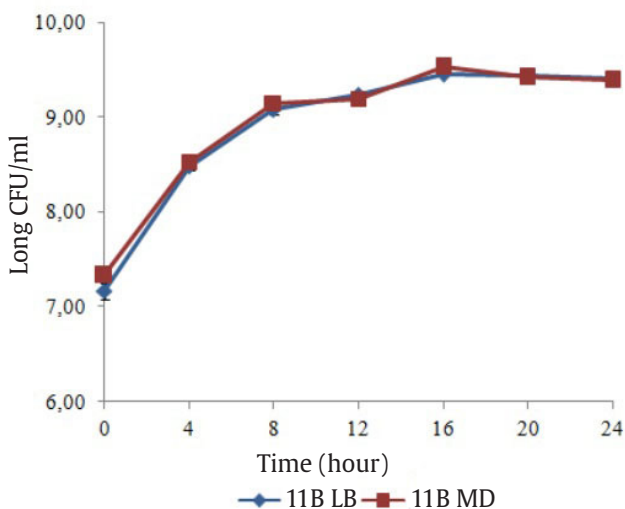


Figure 3. The growth curve of bacterial isolates of 11B LB and 11B MD

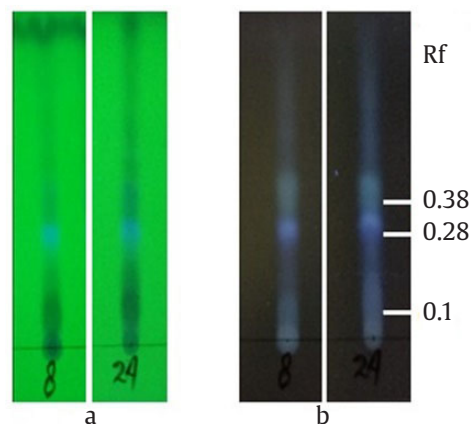


Figure 5. Result of TLC of bacterial crude extract of isolate 11B MD from culture at 8th and 24th hour. (a) observation under 254 nm UV light and (b) 366 nm UV lamp

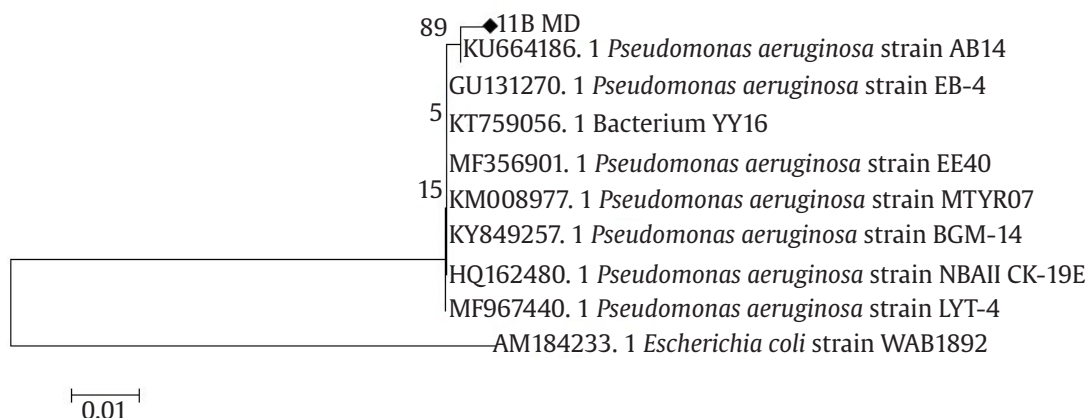


Figure 6. Phylogenetic tree analysis of bacterial isolate 11B MD based 16S rRNA gene

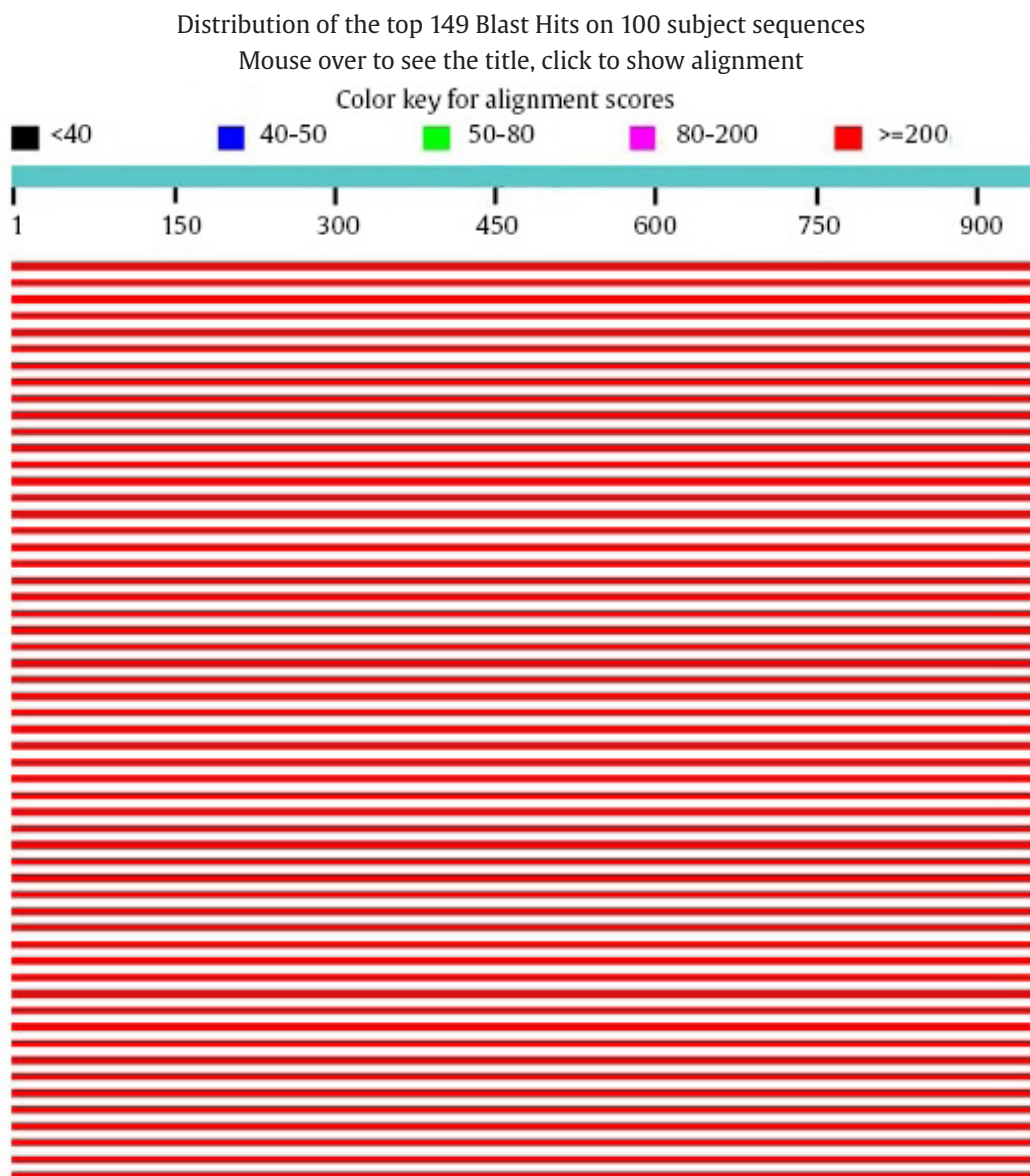


Figure 7. Result of BLAST-N sequence of 16S rRNA gene with the Gene Bank data isolate 11B MD

Chromatographic analysis proved the presence of metabolite compounds that contained bacterial supernatant extract but had not been identified. The compounds contained in the bacterial extracts were separated in the 3:1 eluent ratio (ethyl acetate: n-hexane). The results observed from the stains found on the chromatogram. In addition, stains and Rf values of supernatant extracts from bacterial cultures at 8th and 24th hours showed similarity. The retention factor (Rf) value of the stain was 0.1, 0.28, and 0.38 (Figure 5).

3.6. Identification using Molecular Method

The analysis of metabolite compounds proved that bacteria isolate 11B MD has several bioactive compounds and subsequently performed molecular identification of the isolates. Molecular identification using 16S rRNA gene was aimed to identify the species name of bacteria isolate 11B MD. PCR results from sequences of 16S rRNA after assembling at BLAST showed that the bacterial isolates were believed to be *Pseudomonas aeruginosa* with 99% of similarity (Figure 6).

4. Discussion

Knowing the stages of bacterial isolate selection was aimed to select potential bacterial isolates which serve better biocontrol agents. Selected bacterial isolates did not form clear zones or hemolytic zones on blood agar media. This means that it is not pathogenic. Pathogenic bacteria are capable of producing hemolysin which causes disease for humans (Figueroa-Lopez *et al.* 2016). Pathogenic bacteria are able to produce hemolysin that lyses blood cells as components on the agar media. Blood agar media is a differential medium used to distinguish the type of hemolytic in bacteria. Pathogenic bacterial isolates show different hemolytic types. Bacterial isolate P10 LB and ITSA LB showed this by forming a clear zone around the bacterial colony. However, bacterial isolates ITSA MD formed greenish color on blood agar media. This fact is in line with what McKane and Kandel (1998) said that bacteria with β -hemolysis capabilities will be able to lyse blood cells perfectly characterized by clear zones around the growing bacterial colonies. Bacteria that have a type of α -hemolysis can not lyse the blood cells perfectly marked by the change of media color to greenish.

Antagonistic test showed that selected bacterial isolates had inhibitory activity against *Ganoderma boninense* growth. Inhibition value of bacterial isolates proves that both selected bacterial isolates have the ability to inhibit *G. boninense* growth in vitro. This is

in accordance with Bivi *et al.* (2010) that endophytic bacteria isolated from oil palm plants has inhibition value greater than 50% against *G. boninense*. Ramli *et al.* (2016) confirmed that *Pseudomonas aeruginosa* isolated from oil palm plants has inhibitory activity against *G. boninense* valued at 70%. Inhibitory capability is due to the metabolism produced by both bacterial isolates such as bioactive compounds that can inhibit the growth of *G. boninense*. Macroscopic observation showed that bacterial isolate 11B MD was able to produce inhibition zone against *G. boninense*. Inhibition zone is a form of resistance performed by bacteria so that there is no direct contact between bacterial cells with fungal cells. Formed zone of inhibition on bacterial isolates is influenced by competition in nutrition as well as environmental factors (Nawangsih *et al.* 2010).

The presence of phases that occur during bacterial growth was presented by bacterial growth curve. The growth curve showed that both isolates 11B LB and 11B MD had a short adaptation phase. The condition is caused due prior to observing the growth curve performed before preculture that accelerate the growth of bacteria for leading logarithmic phase. Schultz and Kishony (2013) explained that the phase lag/adaptation of bacterial growth is influenced by the new environmental conditions and the amount of inoculum. The duration of logarithmic phase that occurs in both bacterial isolates showed an activity increase in number of bacterial cells up to 16th hour. *Bacillus amyloliquefaciens* SAHA 12.07 and *Serratia marcescens* KAHN 15.12 has logarithmic phase up to 6th hour in which this phase primary metabolite was produced in the form of chitinase that capable of lysing the cell walls of fungi that interfere with the growth of *G. boninense* (Azizah *et al.* 2015).

The stationary phase experienced by both bacterial isolates started from the 16th hour and it has entered the phase of death at 20th hour. In the stationary phase, secondary metabolite products are produced such as antibiotics or other bioactive compounds that are toxic to microorganisms. Secondary metabolite products can be exploited by bacteria as a form of defense against unfavorable conditions. This is confirmed by Vater *et al.* (2002) that bacterial growth produce metabolite products in the stationary phase of antifungal compounds that can inhibit the growth of pathogenic fungi. In the phase of death, there was a decrease in the number of cells due to excessive accumulation of toxins and depletion of nutrients so that more bacterial cells were dead.

The extraction was conducted using ethyl acetate as a semipolar solvent had a wide polarity range. The polarity range of the solvent can attract the antifungal

compounds contained in the supernatant bacterial culture. This is confirmed by Prapagdee *et al.* (2012) that the use of ethyl acetate solvent can extract the antifungal compounds found in bacterial cultures. The extract activity test showed that the crude extract of supernatant bacterial culture 11B LB was inactively inhibited the growth of mycelium *G. boninense*. The condition is caused by the difference of solubility level in the bacterial metabolite so that not all metabolite compounds can be drawn by semipolar solvent. As a result, at least there is a metabolite product that can be extracted in a supernatant bacterial culture. Therefore, the extract test showed a low inhibitory activity against *G. boninense* growth.

Extracts that have inhibitory activity against *G. boninense* are crude extracts from supernatant bacterial culture 11 B MD at 8 and 24 hours. Other crude extracts (extracts from 12th, 16th, and 20th cycle cultures) have a low inhibitory activity. This is caused by the concentration of metabolite products produced greater at the 8th and 24th hour. Metabolite product that were produced is antifungal because the extract test indicates the presence of inhibitory activity against the growth of the fungus. Factors that affect the growth of bacteria are nutrients contained in the media and environmental conditions of bacterial growth such as temperature, agitation, pH and so forth. The process of bacterial metabolism is very sensitive to change in bacterial growth factors that can affect the synthesis of metabolites at the time of bacterial growth phase.

Supernatant culture 11B MD extract at 8th hour was from bacterial cultures harvested in the logarithmic growth phase. The 24th hour extract was derived from the culture of death phases. In general, primary metabolites such as enzymes can be produced in the logarithmic phase. On the other hand, it is still possible to produce secondary metabolite products which are commonly formed in stationary phase. Petatan-Sagahon *et al.* (2011) explained that the culture of *P. fluorescens* 16 in the logarithmic phase can inhibit the growth of pathogenic fungi in plants. This is confirmed by Zvanych *et al.* (2014) that said secondary metabolite compounds can be produced in the mid-logarithmic and stationary-phase. Crude extracts from the 24th hour supernatant culture also demonstrated the ability to inhibit *G. boninense* fungi. This is due to the presence of metabolite compounds contained in the extract. Ramachandran *et al.* (2014) stated that in the mid and final conditions of the stationary phase, bacterial growth is produced secondary metabolites. It explains that the compounds produced by bacteria isolate 11B MD are able to inhibit the growth of *G. boninense* as it is similar to the case with positive control treatment. This result is proved by giving

the same concentration between crude extract and nystatin by showing similarity as antifungal. Therefore, compounds of bacterial isolate 11B MD can be used as antifungal because they have inhibiting properties such as nystatin which is a commercial antifungal.

The crude extracts from supernatant bacterial culture showing the presence of inhibitory activity were then analyzed using thin layer chromatography (TLC). This method is performed to determine the metabolite compounds contained in an extract. In the chromatogram seen a stain formed when observed under UV light. The presence of a fluorescent stain indicates the presence of a compound having a chromophore group. The chromophore group may interact with UV light to form a stain on the chromatogram and the chromophore has a conjugated double bond. Laniado-Laborin and Cabrales-Vargas (2009) stated that antifungal compounds with conjugated double bonds may disrupt the permeability of the fungal cell membranes resulting in damage. This damage inhibit the growth of the fungus as a consequence of the incapability of transport process, selective, and protection of the cell membranes.

Fluorescent stains under UV light prove the presence of certain compounds that can absorb UV light at wavelengths of 254 and 366 nm. Gupta *et al.* (2012) explained that TLC results from bacterial extracts containing metabolite compounds may glow after being observed in UV light 254 and 366 nm. Rf values showed that the compounds contained an extract. The Rf values obtained (Figure 5) are consistent with the results of a study conducted by Caldeira *et al.* (2007) that bioactive antifungal compounds has Rf value which is equal to 0.22, 0.28, and 0.38. The result of Rf value obtained shows that the group of compounds in bacterial extract are suspected to be phenol compounds. Group of phenol compounds proposed by Gomathi *et al.* (2012) shows Rf value of 0.06, 0.13, 0.21, 0.28, 0.36, 0.38, 0.44, 0.73, 0.84, and 0.94. The statement was also reinforced by Chong *et al.* (2012) that syringic acid which is a kind of phenol compounds has the ability to inhibit the growth of *G. boninense* causes basal stem rot of oil palm plantation.

Molecular identification of bacterial isolate 11B MD was performed based on the encoding gene of 16S rRNA. The 16S sequence of rRNAs has a stable (conservative) base sequence and undergoes very slow evolutionary changes. In addition, it can be used to track the relationship of kinship on each bacteria so it is appropriate to be used in identification (Figure 6). Identification using a gene encoding 16S rRNA showed that isolate 11B MD was assumed to be *P. aeruginosa* with a similarity value of 99%. This is reinforced by Drancourt *et al.* (2000) statement that the data are

considered as the same species if they have similarity values greater than 99% based on sequences of 16S rRNA gene. Tests conducted has proved that the bacteria *P. aeruginosa* has the ability to inhibit the growth of mycelium *G. boninense*. Suryadi et al. (2014) explains that *P. veronii* can produce chitinase enzymes and in *P. aeruginosa* bacteria can produce glucanase. Therefore, they potentially are biocontrol agents to reduce pathogenic fungi in plants. Both enzymes are enzymes that can degrade the cell wall of the fungus. The inhibitory capability of the bacteria is also reinforced by Parvin et al. (2016) statement that *P. aeruginosa* isolated from the palm oil plant rhizosphere produces metabolite products such as phenazine in order to inhibit the growth of *G. boninense*.

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