### Isolation and Antibacterial Activities of Actinomycetes from Rhizosphere Plant Cane (Saccharum officinarum) on Escherichia coli and Staphylococcus aureus

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#### Abstract

The phenomenon of increasing human bacterial infections and increased resistance to antibiotics encourages the exploration of more antibiotic producers. One source of antibiotics comes from the Actinomycetes bacteria, the source of isolation comes from the rhizosphere of sugarcane plants because it is known in the rhizosphere to produce exudates that are beneficial to bacteria. The purpose of this study was to obtain Actinomycetes isolates and determine the antibacterial abilities of selected isolates. Initial isolation was carried out using Starch Casein Agar (ScA) media, then color grouping, Gram test and cell morphology observation. The next stage is testing the antibacterial activity carried out using the well method. Data obtained in the form of basic characters Actinomycetes, inhibition zone diameter and antibacterial activity count. The results showed that 4 isolates had antibacterial ability against test bacteria, among others, ACB34a isolates, ACB44c isolates (9.62 mm2 / 10µl) and ACB54c (0.38 mm2 / 10µl), then antibacterial activity against S. aureus. among others, ACB44c (27.33 mm2 / 10µl), ACB55c (8.04 mm2 / 10µl), and ACB34a (8.04 mm2 / 10µl).

Keywords: Actinomycetes; Antibacterial; Characterization; Rizosphere of sugar cane

#### **1. INTRODUCTION**

The increased of disease infections and antibiotics resistance have encouraged the exploration of more antibiotic-producing sources. Actinomycetes are known to have antibacterial abilities and contribute a lot to produce antibiotics as well as bioactive molecules that influence the industrial world (Sateesh et al., 2011).

In the study of Krismawati et al. (2015) Streptomycetes members of Actinomycetes can be found in the rhizosphere and nonrhizosphere of magrove vegetation. Several isolates were found to be able to produce antibiotics against several test bacteria. The types of bacterial compounds found were thought to be Erytromycin, Tetracyclin, Rimfampicyn, Polymyxin and Chloramphenicol. According to Raningsih (2015), antimicrobial effects on plasma membranes affect their integrity. Increased permeability and leakage of cells due to damaged cell membranes causes the release of intracellular material.

According to Khanna et al. (2011), microbial diversity in the rhizosphere or around cultivated plants is a direct result of interactions and adaptations between plants and microbes. The results of interactions and adaptations are possible adaptations or resistance to plant toxins, have specific secondary metabolites producing certain Soils around the roots or antibiotics. rhizosphere have organic material that active encourage the development of microbes rather than ordinary soils or bulk soils (Geetanjali and Jain, 2016).

Rhizosphere of sugarcane (Saccharum officinarum) has loose soil structure with smooth air aeration, tolerant of pH 6-7.5 even up to pH 8 and perfect root development (Indrawanto al., 2010). This et soil condition affect the presence of (2008)Actinomycetes. Niswati et al. explained that exudates found in the rhizosphere affect the bacterial population. Sugarcane exudates are related to the presence of nitrogen, phosphate and amino acid composition. Exudates released depend

Bioedukasi Vol. XVII. No. 1 April 2019

Received 14 February 2019 | Received in revised form 15 March 2019 | Accepted 29 March 2019 | Published online 1 April 2019

on the type of plant, monocot plants can release more exudate than dicotyledonous plants (Kato et al., 1997). The presence of Actinomycetes in plant rhizosphere has been explored on rice plants (Ambarwati, 2012), mangrove (Krismawati et al., 2015), banana (Sudarma, 2010); eggplant, potatoes, corn, cabbage, coffee, teak (Susilowati et al., 2007). The purpose of this study was to isolate Actinomycetes from sugarcane rhizosphere and determine their antibacterial ability against *Escherichia coli* and *Staphylococcus aureus*.

# 2. RESEARCH METHOD

was an observational This study conducted from January to March 2019. Soil samples were obtained from the rhizosphere of the sugar cane plant (S. officinarum) in the Bangkal Kamal area. The implementation of isolation, initial screening, continued color grouping screening, antibacterial ability test and basic characterization were carried out in Microbiology Laboratory the of the Department of Biology.

The tools used were soil tester. incubator, test tube, haemocytometer, centrifuge, vortex, oven, petri dish, ose, cotton, microscope, glass cover, object glass, measuring cup, beaker glass, spirtus lamp, aluminum foil, water bath, woll yarn, filter paper and tissue paper, pestle mortar, analytical scales. The materials used in this studv include. ScA media. Oatmeal. Escherichia coli **FNCC** 0091 and Staphylococcus aureus FNCC 0047, NaCl, crystal violet, methylene blue, iodint, safranin, sterile distilled water, 75% alcohol and 96% alcohol, Nyastin (Fungicide) ), Nutrient Agar, Nutrient Broth, starch, K2HPO4, KNO3, Casein, MgSO4.7H2O, Agar, 3% H2O3, sucrose.

The procedure carried out in the study included making Oatmeal media with composition (10 g soluble starch, 2 g K2HPO4, 2 g KNO3, 0.3 g Casein, 0.05 g MgSO4.7H2O, 0.02 g CaCO3, 0.01 g FeSO4.7H2O, 15 g Agar, 1000 mL aquadest water and pH 7 (Mohseni et al., 2013), Sca media with composition (50 g oatmeal, 5 g sucrose, 20 g agar and 1000 mL aquades), NA, NB and test bacterial preparation for *E. coli* and *S. Aureus*.

Bacterial preparation was done by growing E. coli dan S. aureus on NB then the bacteria was counted its cell density using haemocytometer. The next stage was taking soil samples with five sampling points forming zigzag patterns in sugar cane gardens, measuring soil temperature, pretreatment (heating 50 ° C soil samples) in the oven to kill gram negative bacteria and other non-target spores, isolating isolates which grows in ScA media and color grouping. Color grouping is a grouping of isolates grown on Oatmeal media so that it is based on the color of the mycelium substrate, the color of the aerial mycelium and mycelium pigmentation.

The stages were continued with purification of isolates which showed the key character of Actinomycetes. Kev characteristics of members the of Actinomycetes include the character of the colonies in the ScA media, appearance of the mycelium in the agar media Oatmeal, cell morphology and gram staining. The suspected isolates then undergo antibacterial ability tests using difusion method. The isolates used in this test were suspected Actinomycetes on NB medium for 48 hours which were centrifuged. As much as 10 µL of supernatant applied in the diffusion method. The zone of inhibition formed around the well then observed and measured (Usmati and Marwati, 2007).

### 3. RESULT AND DISCUSSION

The results of the initial isolation were successfully obtained 37 isolates and 23 of them were thought to be Actinomycetes. The results of measurements of soil temperature using soil tester showed a temperature of  $30^{\circ}$  C with a pH of 6.5.

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Koloni

Elevation

Convex

Convex

Convex

Ranset

Convex

Convex

Surface

Rough

Rough

Rough

wrinkled

Rough

Rough

Rough

Optic

Opaque

Opaque

Opaque

Opaque

Opaque

Opaque

Characteristics

Edge

Entire

Entire

Entire

Filament

Entire

Entire

ACB25b ACB25c ACB26a ACB26b ACB26c ACB27 ACB34a ACB34b ACB35a ACB35b ACB35b ACB35c ACB35d ACB36 ACB44a	+ + + + + + + + + + + + + + + +	Kream Kream White White White White White	Circular Circular Circular Non ba	Entire Entire Entire	Convex Convex Convex Convex Actinomycete Convex Convex Convex	Rough Rough	Opaque Opaque Opaque Opaque Opaque Opaque
ACB26a ACB26b ACB26c ACB27 ACB34a ACB34b ACB35a ACB35b ACB35c ACB35d ACB35d	+ + + + + + + + + + + + +	Kream White White White White	Circular Circular Non Circular Circular Circular Non ba	Entire Entire bacterial A Entire Entire Entire	Convex Convex Actinomycete Convex Convex	Rough Rough es (slimy) Rough Rough	Opaque Opaque Opaque Opaque
ACB26b ACB26c ACB27 ACB34a ACB34b ACB35a ACB35b ACB35c ACB35d ACB35d	+ + + + + + + + + + + +	White White White White	Circular Non I Circular Circular Circular Non ba	Entire bacterial A Entire Entire Entire	Convex actinomycete Convex Convex	Rough es (slimy) Rough Rough	Opaque Opaque Opaque
ACB26c ACB27 ACB34a ACB34b ACB35a ACB35b ACB35c ACB35d ACB35d ACB36	+ + + + + + + + + + +	White White White	Non Circular Circular Circular Non ba	bacterial A Entire Entire Entire	ctinomycete Convex Convex	es (slimy) Rough Rough	Opaque Opaque
ACB27 ACB34a ACB34b ACB35a ACB35b ACB35c ACB35d ACB35d ACB36	+ + + + + + + + +	White White	Circular Circular Circular Non ba	Entire Entire Entire	Convex Convex	Rough Rough	Opaque Opaque
ACB34a ACB34b ACB35a ACB35b ACB35c ACB35d ACB36	+ + + + + + +	White White	Circular Circular Non ba	Entire Entire	Convex	Rough	Opaque
ACB34b ACB35a ACB35b ACB35c ACB35d ACB36	+ + + + +	White	Circular Non ba	Entire			
ACB35a ACB35b ACB35c ACB35d ACB36	+ + + +		Non ba		Convey		
ACB35b ACB35c ACB35d ACB36	+ + + +	White			CONVEX	Rough	Opaque
ACB35c ACB35d ACB36	+ +	White	NT 1	cterial Ac	tinomycetes	(slimy)	
ACB35d ACB36	+	White		cterial Ac	tinomycetes	(slimy)	
ACB36			Circular	Entire	Pulvinate	Rough	Opaque
	+			No growth	during re-cu	ıltur	
ACB44a		White	Circular	Entire	Convex	Rough	Opaque
	+			0	uring re-cult		
ACB44b	+			o growth d	uring re-cult		
ACB44c	+	White	Circular	Entire	Convex	Rough	Opaque
ACB45a	+	Kream	Circular	Entire	Convex	Rough	Opaque
ACB45b	+			bacterial A	ctinomycete	es (slimy)	
ACB46a	+	Kream		Entire	Convex	Rough	Opaque
ACB46b	+			No growth	during re-cu	ıltur	
ACB54a	+	White	Circular	Entire	Convex	Rough	Opaque
ACB54b	+	White	Circular	Entire	Convex	Rough	Opaque
ACB54c	+	Krim	Circular	Entire	Convex	Rough	Opaque
ACB55a	+		1	No growth	during re-cu	ıltur	
ACB55b	+			bacterial A			
ACB55c	+	White	Circular	Entire	Convex	Rough	Opaque
ACB55d	+						
ACB56a	+		No	growth d	uring re-cult	tur	
					uring re-cult		
	+	White			Convex	0	Opaque
ACB57b	+		Non	bacterial A	ctinomycete	es (slimy)	
d the appearant ate, aeral and particular showed a	nce or color of n pigmentation. As a color appearan	nycelium much as ce, those	ACB	45a, AC			ACB44c, CB55c dan
	ACB45b ACB46a ACB46b ACB54a ACB54a ACB54b ACB54c ACB55a ACB55b ACB55c ACB55d ACB56a ACB56a ACB56b ACB57a ACB57b e stage of cc d the appearant ate, aeral and p lates showed a	ACB45b+ $ACB46a$ + $ACB46b$ + $ACB54a$ + $ACB54b$ + $ACB54c$ + $ACB55a$ + $ACB55b$ + $ACB55c$ + $ACB55d$ + $ACB56a$ + $ACB56b$ + $ACB57a$ + $ACB57b$ $ACB57b$ $ACB57b$ + $ACB57b$ $ACB57b$ $ACB57b$ + $ACB57b$ <	ACB45b+ACB46a+ACB46a+ACB46b+ACB54a+ACB54b+ACB54b+ACB55a+ACB55b+ACB55c+ACB55d+ACB56a+ACB56b+ACB56b+ACB57a+	ACB45b+NonACB46a+KreamCircularACB46b+NACB54a+WhiteCircularACB54b+WhiteCircularACB54c+KrimCircularACB55a+NonACB55b+NonACB55c+WhiteACB55d+NonACB55d+NonACB55d+NonACB56a+NoACB56b+NoACB57a+NoACB57b+NoACB57b+NoACB57b+ACBACB57b+ACBACB57b+ACBACB57b+ACBACB57b+ACBACB57b+ACB	ACB45b+Non bacterial AACB46a+KreamCircularEntireACB46b+No growthACB54a+WhiteCircularEntireACB54b+WhiteCircularEntireACB54c+KrimCircularEntireACB55a+No growthACB55b+Non bacterial AACB55c+WhiteCircularACB55d+No growth dACB55d+No growth dACB56a+No growth dACB56b+No growth dACB57a+WhiteACB57b+Non bacterial AACB57b+Non growth dACB57b+ACB26a, AACB57b+ACB26a, AACB57a.+ACB45a, ACACB57a.+ACB57a.AcB57a.+ACB57a.	ACB45b+Non bacterial ActinomyceteACB46a+KreamCircularEntireConvexACB46b+No growth during re-cuACB54a+WhiteCircularEntireConvexACB54a+WhiteCircularEntireConvexACB54b+WhiteCircularEntireConvexACB54c+KrimCircularEntireConvexACB55a+No growth during re-cuACB55b+No growth during re-cuACB55b+Non bacterial ActinomyceteACB55c+No growth during re-culACB55d+No growth during re-culACB56a+No growth during re-culACB56a+No growth during re-culACB56b+No growth during re-culACB57a+WhiteCircularEntireConvexACB57b+Non bacterial ActinomyceteACB57a+Non bacterial Actinomycetee stage of color grouping, the dataACB26a, ACB34a, ACB45a, ACB46a, ACACB45a, ACB46a, ACa the appearance or color of mycelium ate, aeral and pigmentation. As much as lates showed a color appearance, thoseACB57a.	ACB45b+Non bacterial Actinomycetes (slimy)ACB46a+KreamCircularEntireConvexRoughACB46b+No growth during re-culturACB54a+WhiteCircularEntireConvexRoughACB54b+WhiteCircularEntireConvexRoughACB54c+KrimCircularEntireConvexRoughACB55a+No growth during re-culturACB55b+Non bacterial Actinomycetes (slimy)ACB55c+WhiteCircularEntireConvexRoughACB55d+Non bacterial Actinomycetes (slimy)ACB55d+No growth during re-culturACB56a+No growth during re-culturACB56a+No growth during re-culturACB57a+WhiteCircularACB57b+No growth during re-culturACB57b+No growth during re-culturACB57b+No growth during re-culturACB57b+No growth during re-culturACB57b+No bacterial Actinomycetes (slimy)e stage of color grouping, the data te, aeral and pigmentation. As much as lates showed a color appearance, thoseACB57a.

 Table 1. Results of Isolation and Initial Screening of Actinomycetes Bacteria (Starch Casein Agar) and Colony Morphology)

Shape

Circular

Circular

Circular

Circular

Circular

Circular

Color

White

White

White

White

White

White

Growth in ScA Medium

+

+

+

+

+

+

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No

1.

2.

3.

4.

5. 6. Isolates

ACB14a

ACB14b

ACB15

ACB16

ACB24

ACB25a

Table 2. Advanced Screening Color Grouping Actinomycetes (Oatmeal Agar), Gram Tests	,
and Cell Morphology)	_

		Characteristics							
			Oat	meal Agar			o Gram		
No	Isolates	Growth	Substrat Mycellium	Aeral Mycellium	Pigmentation of mycelium	Cell Morpho			
1.	ACB14a	+	Pink	Pink	-	Filamen	+		
2.	ACB14b	+	White	Green	purple	Filamen	+		
3.	ACB15	+	White	Grey	-	Filamen	-		
4.	ACB16	+	Slimy	Green	-	Filamen	-		
5.	ACB24	+	White	White	-	Filamen	-		
6.	ACB25a	+	White	White	-	Filamen	-		
7.	ACB25b	+	Greyish pink	Greyish pink	-	Filamen	+		
8.	ACB25c	+	White	Grey	Golden yellow	Filamen	+		
9.	ACB26a	+	White	-	-	Filamen	+		
10.	ACB26b	+	White	Grey	-	Filamen	-		
11.	ACB27	+	Pink	Pink	-	Filamen	-		
12.	ACB34a	+	Pink	Pink	-	Filamen	+		
13.	ACB34b	+	White	White	-	Filamen	-		
14.	ACB35c	+	White	Pink	-	Filamen	-		
15.	ACB36	+	Pink	Pink	-	Filamen	+		
16.	ACB44c	+	Pink	White	-	Filamen	+		
17	ACB45a	+	White	White	Pink	Filamen	+		
18.	ACB46a	+	White	Grey	-	Filamen	+		
19.	ACB54a	+	White	Grey	-	Filamen	-		
20.	ACB54b	+	White	Grey	-	Filamen	-		
21.	ACB54c	+	Grey	Grey	Golden yellow	Filamen	+		
22.	ACB55c	+	White	Greyish pink	-	Filamen	+		
23.	Acb57a	+	Greyish green	White	-	Filamen	+		

The suspected Actinomycetes were then tested for antibacterial abilities. The test was carried out by calculating the inhibition zone (Usmati and Marwati, 2007). Four isolates that showed positive result were ACB34a, ACB44c, ACB54c and ACB55c. Table 3 and 4 showing antibacterial ability and activity of the selective isolates.

Table 3. Antibacterial ability

	Antibacterial ability							
No	Isolates	E. coli	Diameter Clear zone	Category*	S. aureus	Diameter Clear zone	Category*	
1.	ACB14a	-	-	-	-	-	-	
2.	ACB14b	-	-	-	-	-	-	
3.	ACB25b	-	-	-	-	-	-	
4.	ACB25c	-	-	-	-	-	-	
5.	ACB26a	-	-	-	-	-	-	
6	ACB34a	-	-	-	+	3,2 mm	weak	
7.	ACB36	-	-	-	-	-	-	
8.	ACB44c	+	3,5 mm	weak	+	5,9 mm	mediocre	
9.	ACB45a	-	-	-	-	-	-	
10.	ACB46a	-	-	-	-	-	-	
11.	ACB54c	+	0,7 mm	weak	+	5,8 mm	mediocre	
12.	ACB55c	-	-	-	+	3,2 mm	weak	
13.	ACB57a	-	-	-	-	-	-	
*Categorization according David and Stout (1971)								

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Received 14 February 2019 | Received in revised form 15 March 2019 | Accepted 29 March 2019 | Published online 1 April 2019

Table 4. Antibacterial activity							
Antibacterial activity							
Isolate	E. coli	Inhibition Activity (mm²/10µl)	S. aureus	Inhibition Activity (mm <sup>2</sup> /10µl)			
ACB34a	-	-	+	8,04 mm²/10μl			
ACB44c	+	9,62 mm²/10µl	+	27,33 mm²/10µl			
ACB54c	+	0,38 mm²/10µl	+	26,41 mm²/10µl			
ACB55c	-	-	+	8,04 mm²/10μl			

Isolation of Actinomycetes from the rhizosphere sugarcane (Saccharum of officinarum) carried out with five was sampling points (zig-substance pattern). Based on the data, 13 out of 37 isolated bacteria were suspected Actinomycetes. These 13 isolates were ACB14a, ACB14b, ACB25a, ACB26a, ACB45a. ACB34a. ACB36, ACB44c, ACB46a, ACB54c, ACB55c and ACB57a.

Several screening stages were carried out to obtain all of 13 isolates. Those stages consist of 50° C pre-treatment to kill nonbacteria or non Actinomycetes target (Sembiring, 2000), screening on ScA media and mycelium screening on the Oatmeal (Krismawati medium et al., 20015); (Retnowati et al., 2017). The appearance of colonies, mycelium, tests on initial screening has a distinctive appearance that can differentiate from other bacteria. These characteristics are related to colonies that appear rough, dry, gram-positive, color mycelium, distinctive pigmentation and cell morphology in the form of filaments.

Environmental factors showed soil temperature of 30°C, pH 6.5, and clay texture. Environmental factors such as soil conditions support the presence of Actinomycetes to live (Hasyim et al., 2013; Goodfellow et al., 1988). In addition, according to Niswati et al. (2008); Lagos et al. (2015); Sinma et al. (2015) Actinomycetes and plants have reciprocity or mutual benefit.

Sugar cane produce exudates containing amino acids, sugar, vitamins, tannins and others substances that affect the Actinomycetes or rhizobacteria. On the other hand, Actinomycetes provide benefits in the form of active degradation of organic matter, suppress plant pathogen infections and provide phosphate nutrition.

Adaptations and interactions to certain toxic substances lead to secondary metabolites production to specific antibiotics (Khanna et al., 2011). This open up the opportunity to utilize a metabolite product in other fields, in this study against E. coli and Staphylococcus aureus.

Four isolates of Actinomycetes were able to inhibit the growth of test bacteria. Actinomycetes are a group of bacteria that produce extracellular secondary metabolites so that they are utilized in the source of antibacterial compounds without going through cell breaks (Sharma et al., 2014). Actinomycetes bacteria inhibit cell wall synthesis from test bacteria which causes a decrease in cell osmotic pressure so that test bacteria become inhibited (Jawetz et al., 2007).

A centrifugation at 3000x rpm for 15 minutes was carried out before application.

The supernatant is used as a stock of antibacterial sources that are implanted in wells. Isolates that show antibacterial ability by showing clear zones around the well.

Isolates AcB34a showed а weak inhibition toward S. aureus with 3.2 mm clear zone, while isolate ACB44c showed weak inhibition toward E. coli with 3.5 mm clear zone. However ACB44c showed moderate ability in inhibit S. aureus with 5.9 mm inhibition zone. Similar result showed in ACB54c isolate that showed poor activity towards E.coli (0,7 mm) but moderate to S. aureus (5.8 mm). The last isolate, ACB55c, considered as weak againts S. aureus with a 3.2 mm inhibition zone.

It can be seen from the result that isolates which have the highest values to the lowest in sequence include ACB44c isolates (9.62 mm2 / 10µl) and ACB 54c (0.38 mm2 / 10µl), against E. coli, then ACB44c isolates (27.33 mm2 / 10µl), ACB54c (26.41 mm2 / 10µl), ACB55c (8.04 mm2 / 10µl), and ACB34a (8.04 mm2 / 10µ1) against S. aureus. The value of the calculation showed the compounds had

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Received 14 February 2019 / Received in revised form 15 March 2019 / Accepted 29 March 2019 / Published online 1 April 2019

capabilities that were classified as moderate to low.

According to Kumala et al. (2015), clear zones which appear to be bactericidal regions, namely areas where extracts or supernatants containing a compound that can kill test bacteria (E. coli and S. aureus). Whereas zones formed outside the clear zone or bactericidal area are regions of the clear zone which only inhibit the test bacteria (Kumala et al., 2015). In this study the only visible zone is the clear zone which is the extract area of the supernatant which kills the test bacteria.

This finding is supported by previous research conducted by Ambarwati (2012) showing the antibacterial ability of Actinomycetes from rhizosphere isolation of rice plants to E. coli. There were 3 isolates from 11 isolates who had the ability to inhibit *E. coli* with the details of one isolate in the strong category, one isolate in the moderate category and one isolate in the weak category.

According to Meklat et al. (2011), Actinomycetes antibiotics were coded by the polyketide synthetase (PKS) gene precisely PKS-I, PKS-II and enedyne-PKS. The gene code that produces Actinomycetes antibiotics can be assumed to produce various antibiotic Goodfellow et al. compounds. (1988), explained that each type of strain of Actinomycetes can produce a typical antibiotic compound. Examples are actinohordin, streptomycin, erithromycin, bialaphos, tetracenomycin, rifampicin. Pure actinomycetes can show various chemical others. structures, among anthraevclines. aminoglyuises, glycopeptides, βlactams, polynes, macrolides. tetracyclins and polyether. These compounds have important function the health field.

The mechanism of inhibition for bacteria depends on the type of compound. It is known that Actinomycetes produce a lot of metabolites harmful to bacteria. There are many compounds that have role to inhibit E. coli and S. aureus. The compounds produced by plant rhizosphere Actinomycetes erytromycin, tetracycline, rifampicin with peptide compounds, flavonoids and alkaloids. The mechanism involving the attacks on RNA (protein synthesis) and cell wall structure (cell lysis). This is based on research conducted by Krismawati et al. (2015) and Masda (2018) with similarity of isolation sources, namely

plant rhizosphere and the same test bacteria (E. coli and S. aureus).

In the study of Krismawati et al., (2015) the isolation of Actinomycetes in the mangrove rhizosphere tested on E. coli and S. aureus showed that several isolates produced metabolites such as ervtromycin, tetracycline, through Layer rifampicin Thin Chromatography (TLC) methods. Then in the Masda study (2018)the isolation of Actinomycetes in the rhizosphere of bitter plants produced isolates of several isolates which had the ability to inhibit the bacteria E. coli and S. aureus. At the stage of the research, TLC was carried out and the groups of peptides, flavonoids and alkaloids were found. In general, these compounds interfere with the instability of hydrogen bonds with the complexes formed so that permeability fusion of bacterial cells is disrupted and lysis occurs. The diameter of the inhibitory zone in each selected isolate on test bacteria were different. Several factors affect the diameter of the clear zone are the media composition, temperature, agar thickness (Brady and Katz, 1990); Suriani et al, 2013); (Mujahid et al., 2016). The incubation temperature also affects the enzyme activity related to denaturation and optimal bacterial growth temperature. Bacteria do not grow when they are above the maximum or below the minimum (Suriani et al., 2013). Then the composition of growth media such as carbon sources, nitrogen and special additives to the media influence bacterial or zone of

thickness affect the visibility of the clear zone (Brady and Katz, 1990).

growth (Mujahid et al., 2016). The medium

# 4. CONCLUSION

As much as 13 Actinomycetes isolates were obtained from, 4 among them showed the inhibition activity againts E.coli and S.aereus which range in low to moderate category.

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Received 14 February 2019 | Received in revised form 15 March 2019 | Accepted 29 March 2019 | Published online 1 April 2019

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