

BAB V

PENUTUP

A. Kesimpulan

Berdasarkan hasil penelitian yang telah dilakukan dapat disimpulkan bahwa:

Pertama, isolat bakteri endofit dari umbi talas (*Colocasia esculenta* L.) *Pseudomonas knackmussii* dan *Bacillus siamensis* memiliki aktivitas antibakteri terhadap bakteri *Escherichia coli* ATCC 25922.

Kedua, isolat bakteri endofit umbi talas (*Colocasia esculenta* L.) *Pseudomonas knackmussii* memiliki waktu optimum fermentasi pada hari ke-2 dengan diameter daya hambat 11,83 mm dan *Bacillus siamensis* yang memiliki waktu optimum fermentasi pada hari ke-5 dengan diameter daya hambat 10,89 mm.

Ketiga, isolat bakteri endofit umbi talas (*Colocasia esculenta* L.) *Pseudomonas knackmussii* memiliki potensi aktivitas antibakteri lebih besar dari *Bacillus siamensis*.

B. Saran

Pertama, perlu dilakukan penelitian lebih lanjut untuk mengetahui bagaimana kurva pertumbuhan bakteri endofit umbi talas (*Colocasia esculenta* L.) dan mengetahui senyawa yang terdapat dalam isolat bakteri endofit *Pseudomonas knackmussii* dan *Bacillus siamensis*.

Kedua, perlu dilakukan penelitian lebih lanjut terkait mekanisme kerja dari senyawa antibakteri dari bakteri endofit umbi talas (*Colocasia esculenta* L.) dalam menghambat bakteri patogen.

Ketiga, perlu dilakukan pengembangan lebih lanjut terkait pengembangan hasil senyawa bioaktif dari isolat bakteri endofit *Pseudomonas knackmussii* dan *Bacillus siamensis* untuk antibakteri.

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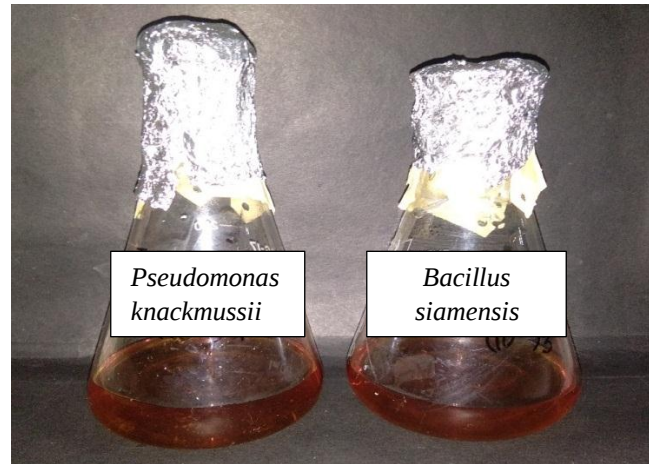
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Lampiran 1. Suspensi bakteri uji

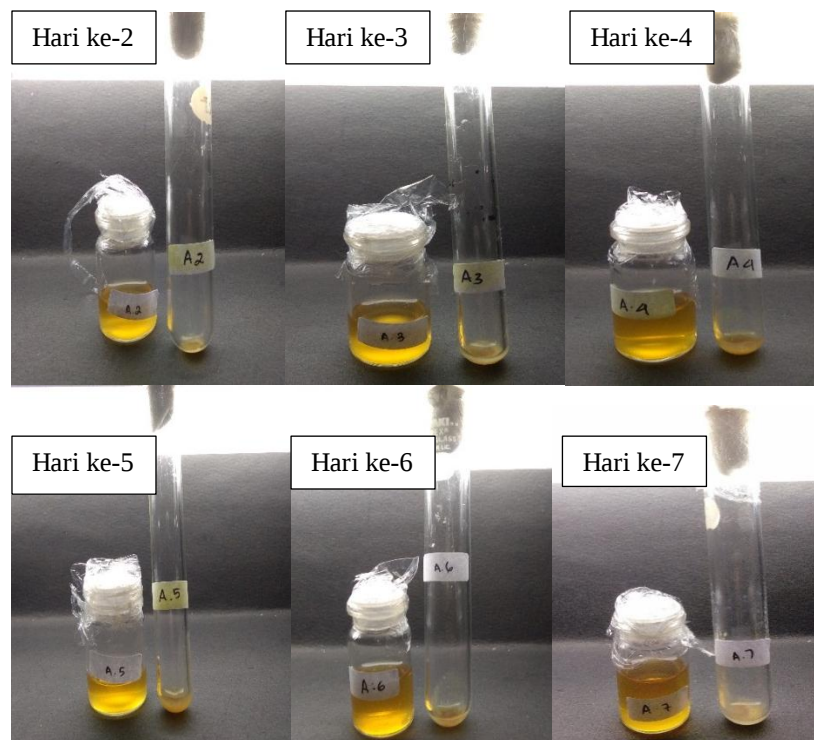


Suspensi bakteri *Escherichia coli*

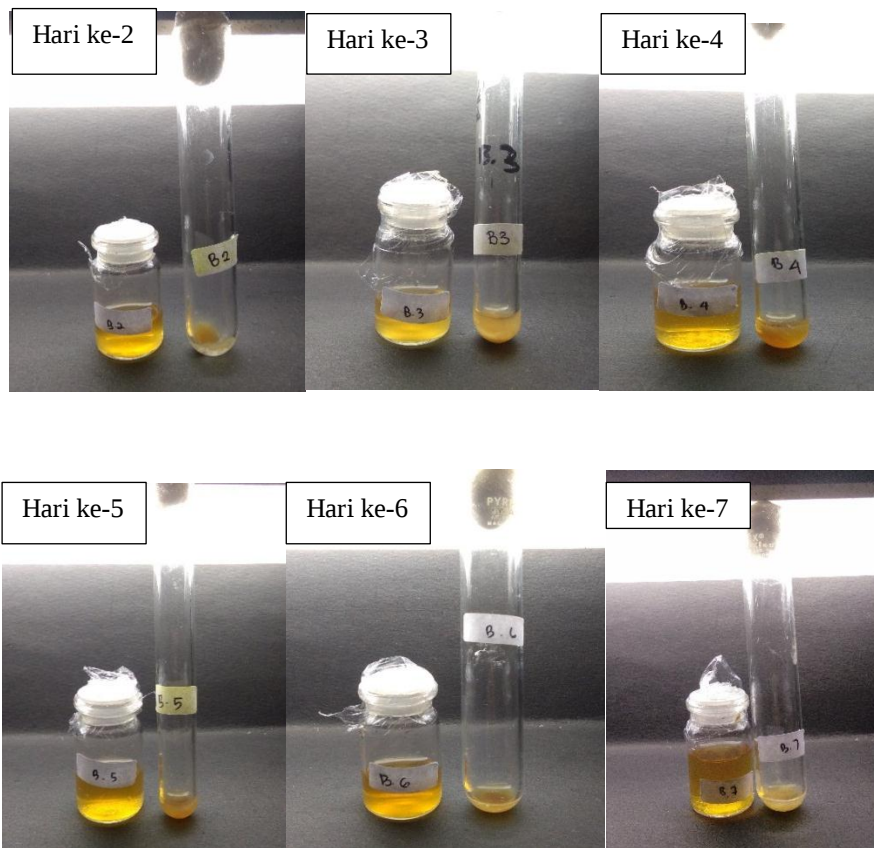
Lampiran 2. Fermentasi isolat bakteri endofit *Pseudomonas knackmussii* dan *Bacillus siamensis*



Fermentasi isolat bakteri *Pseudomonas knackmussii* dan *Bacillus siamensis*

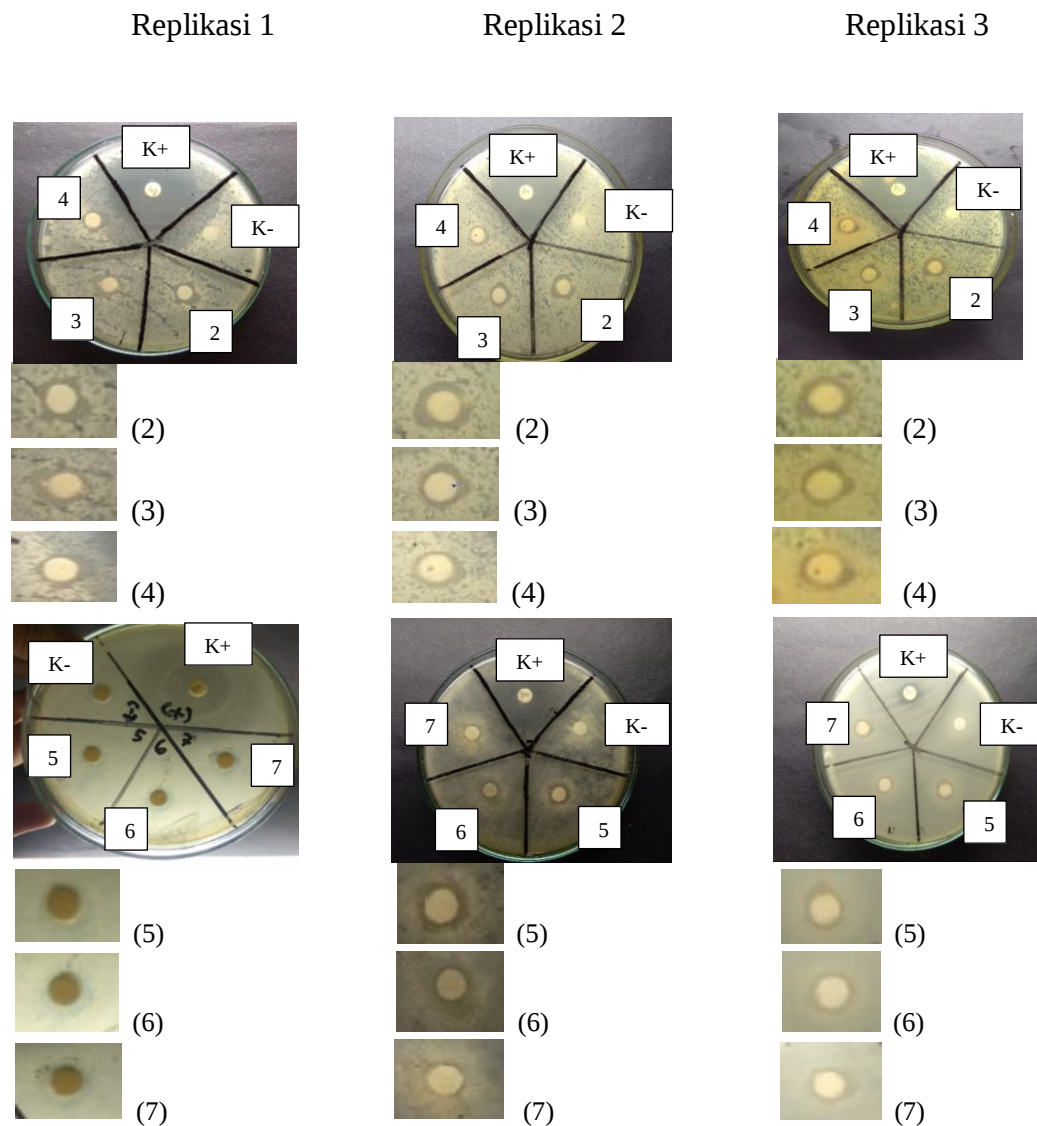


Supernatan isolat bakteri endofit *Pseudomonas knackmussii*



Supernatan isolat bakteri endofit *Bacillus siamensis*

Lampiran 3. Uji Aktivitas antibakteri hasil isolat bakteri endofit *Pseudomonas knackmussii* dan *Bacillus siamensis* dari umbi talas (*Colocasia esculenta* L.) terhadap *Escherichia coli* ATCC 25922



Keterangan :

+ : Kontrol positif (Antibiotik ciprofloxacin)

- : Kontrol negatif

2 : Fermentasi hari ke 2

3 : Fermentasi hari ke 3

4 : Fermentasi hari ke 4

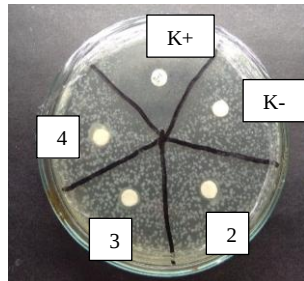
5 : Fermentasi hari ke 5

6 : Fermentasi hari ke 6

7 : Fermentasi hari ke 7

Uji aktivitas antibakteri isolat bakteri endofit *Pseudomonas knackmussii* terhadap *Escherichia coli*

Replikasi 1



(2)

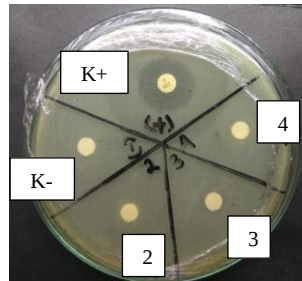


(3)



(4)

Replikasi 2



(2)

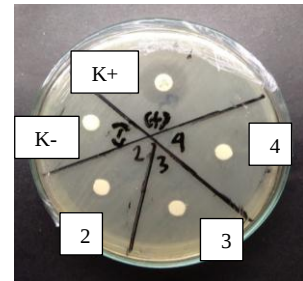


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Replikasi 3



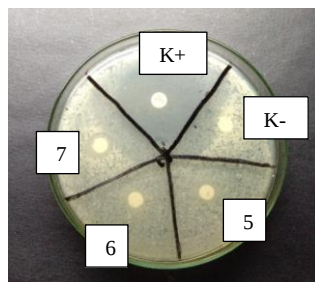
(2)



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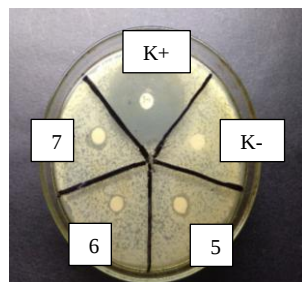
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(6)



(7)



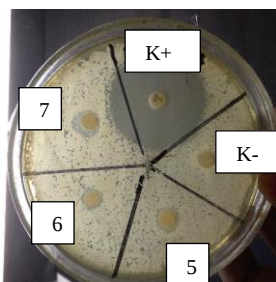
(5)



(6)



(7)



(5)



(6)



(7)

Keterangan :

+ : Kontrol positif (Antibiotik ciprofloxacin)

- : Kontrol negatif

2 : Fermentasi hari ke 2

3 : Fermentasi hari ke 3

4 : Fermentasi hari ke 4

5 : Fermentasi hari ke 5

6 : Fermentasi hari ke 6

7 : Fermentasi hari ke 7

Uji aktivitas antibakteri isolat bakteri endofit ECE-4 terhadap *Escherichia coli*

Lampiran 4. Hasil pencarian dugaan protein antibakteri yang disintesis oleh bakteri endofit *Pseudomonas knackmussii* pada laman *Universal Protein Resource (UniProt)*

UniProtKB results

UniProtKB consists of two sections:

- Reviewed (Swiss-Prot) - Manually annotated**
Records with information extracted from literature and curator-evaluated computational analysis.
- Unreviewed (TrEMBL) - Computationally analyzed**
Records that await full manual annotation.

The UniProt Knowledgebase (UniProtKB) is the central hub for the collection of functional information on proteins, with accurate, consistent and rich annotation. In addition to capturing the core data mandatory for each UniProtKB entry (mainly, the amino acid sequence, protein name or description, taxonomic data and citation information), as much annotation information as possible is added.

Filter by:

Unreviewed (2)
PSEKB (2)

View by
Results table

Taxonomy

Quote terms: "pseudomonas knackmussii"

Entry	Entry name	Protein names	Gene names	Organism	Length
A0A024HE19	A0A024HE19_PSEKB	Tol-Pal system protein TolQ	tolQ PKB_1389	Pseudomonas knackmussii (strain DSM 6978 / LMG 23759 / B13)	231
A0A024HDR2	A0A024HDR2_PSEKB	TonB, C-terminal	PKB_1391	Pseudomonas knackmussii (strain DSM 6978 / LMG 23759 / B13)	349

1 to 2 of 2 Show 25

Display

Entry

Protein | **Tol-Pal system protein TolQ**

Gene | **tolQ**

Organism | *Pseudomonas knackmussii* (strain DSM 6978 / LMG 23759 / B13)

Status | Unreviewed - Annotation score: ●●●○○○ - Protein inferred from homologyⁱ

Functionⁱ

Part of the Tol-Pal system, which plays a role in outer membrane invagination during cell division and is important for maintaining outer membrane integrity.

GO - Biological processⁱ

- bacteriocin transport Source: InterPro
- cell cycle Source: UniProtKB-KW
- cell division Source: UniProtKB-UniRule

View the complete GO annotation on QuickGO ...

Keywordsⁱ

Biological process Cell cycle, Cell division UniRule annotation

Names & Taxonomyⁱ

Protein namesⁱ Recommended name: **Tol-Pal system protein TolQ** UniRule annotation

Gene namesⁱ Name: **tolQ** UniRule annotation Imported

ORF Names: PKB_1389 Imported

Organismⁱ *Pseudomonas knackmussii* (strain DSM 6978 / LMG 23759 / B13) Imported

Display

Entry

Protein | Submitted name: **TonB, C-terminal**

Gene | **PKB_1391**

Organism | *Pseudomonas knackmussii* (strain DSM 6978 / LMG 23759 / B13)

Status | Unreviewed - Annotation score: ●○○○○○ - Protein predictedⁱ

Functionⁱ

GO - Molecular functionⁱ

- transporter activity Source: InterPro

View the complete GO annotation on QuickGO ...

GO - Biological processⁱ

- bacteriocin transport Source: InterPro

View the complete GO annotation on QuickGO ...

Names & Taxonomyⁱ

Protein namesⁱ Submitted name: **TonB, C-terminal** Imported

Gene namesⁱ ORF Names: PKB_1391 Imported

Organismⁱ *Pseudomonas knackmussii* (strain DSM 6978 / LMG 23759 / B13) Imported

Taxonomic identifierⁱ 1301098 [NCBI]

Taxonomic lineageⁱ Bacteria > Proteobacteria > Gammaproteobacteria > Pseudomonadales > Pseudomonadaceae > Pseudomonas > **Tol-Pal system protein TolQ**

Proteomesⁱ UP000025241 Component: Chromosome

Protein Tol-Pal system protein TolQ dan TonB, C-terminal

Lampiran 5. Hasil pencarian dugaan protein antibakteri yang disintesis oleh bakteri endofit *Bacillus siamensis* pada laman *Universal Protein Resource (UniProt)*

The screenshot shows the UniProt website interface. The search bar at the top contains the query 'bacillus siamensis bacteriocin'. The results section shows a single entry: A0A268DTK5, A0A268DTK5_9BACI, Bacteriocin, CHH79_09480, CWD84_04735, Bacillus siamensis, 111. The detailed view for this entry is shown below, including the protein name, gene name, organism, and taxonomic lineage.

UniProtKB results

UniProtKB consists of two sections:

- Reviewed (Swiss-Prot) - Manually annotated**
Records with information extracted from literature and curator-evaluated computational analysis.
- Unreviewed (TrEMBL) - Computationally analyzed**
Records that await full manual annotation.

Filter by: Unreviewed (1) TrEMBL

Popular organisms
Bacillus siamensis (1)

View by: Results table

Display: Entry, Publications, Feature viewer, Feature table

Names & Taxonomy:

- Protein names: Submitted name: **Bacteriocin**
- Gene names: ORF Names: CHH79_09480, CWD84_04735
- Organism: **Bacillus siamensis**
- Taxonomic identifier: 659243 [NCBI]
- Taxonomic lineage: Bacteria > Firmicutes > Bacilli > Bacillales > Bacillaceae > Bacillus > Bacillus amyloliquefaciens group
- Proteomes: UP000234366 Component: Chromosome, UP000215585 Component: Unassembled WGS sequence

Subcellular location:

Topology:

Feature key	Position(s)	Description	Actions	Graphical view	Length
Transmembrane	65 - 93	Helical	Sequence analysis		29

GO - Cellular component:

Protein bacteriocin

UniProtKB results

UniProtKB consists of two sections:

- Reviewed (Swiss-Prot) - Manually annotated**
Records with information extracted from literature and curator-evaluated computational analysis.
- Unreviewed (TrEMBL) - Computationally analyzed**
Records that await full manual annotation.

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Filter by: BLAST Align Download Add to basket Columns

Quote terms: "bacillus siamensis"

Entry	Entry name	Protein names	Gene names	Organism	Length
A0A268DMZ8	A0A268DMZ8_9BACI	Surfactin synthetase	CHH79_18810	Bacillus siamensis	434

View by: Results table Taxonomy

Display: BLAST Align Format Add to basket History Other tutorials and videos Help video Feedback

Entry: Submitted name: **Surfactin synthetase**

Gene: **CHH79_18810**

Organism: **Bacillus siamensis**

Status: Unreviewed - Annotation score: - Protein predicted

Function:

Caution: The sequence shown here is derived from an EMBL/GenBank/DBJ whole genome shotgun (WGS) entry which is preliminary data. Imported

Names & Taxonomy:

Protein names	Submitted name: Surfactin synthetase Imported
Gene names	ORF Names: CHH79_18810 Imported
Organism	Bacillus siamensis Imported
Taxonomic identifier	659243 [NCBI]
Taxonomic lineage	Bacteria > Firmicutes > Bacilli > Bacillales > Bacillaceae > Bacillus > Bacillus amyloquelaceus group
Proteomes	UP000215585 Component: Unassembled WGS sequence

Family & Domains:

Domains and Repeats:

Feature key	Position(s)	Description	Actions	Graphical view	Length

Prtein Surfactin

UniProtKB results

UniProtKB consists of two sections:

- Reviewed (Swiss-Prot) - Manually annotated**
Records with information extracted from literature and curator-evaluated computational analysis.
- Unreviewed (TrEMBL) - Computationally analyzed**
Records that await full manual annotation.

The UniProt Knowledgebase (UniProtKB) is the central hub for the collection of functional information on proteins, with accurate, consistent and rich annotation. In addition to capturing the core data mandatory for each UniProtKB entry (mainly, the amino acid sequence, protein name or description, taxonomic data and citation information), as much annotation information as possible is added.

Filter by: Unreviewed (4) TREMBL

Popular organisms: Bacillus siamensis (4)

View by: Results table, Taxonomy, Keywords

Quote terms: "bacillus siamensis"

Entry	Entry name	Protein names	Gene names	Organism	Length
A0A385CIC8	A0A385CIC8_9BACI	Iturin A synthetase A	ituA	Bacillus siamensis	278
A0A385CIG7	A0A385CIG7_9BACI	Iturin A synthetase B	ituB	Bacillus siamensis	170
A0A385CIS0	A0A385CIS0_9BACI	Iturin A synthetase C	ituC	Bacillus siamensis	197
A0A385CIT6	A0A385CIT6_9BACI	Iturin A synthetase D	ituD	Bacillus siamensis	216

Display: Entry, Publications, Feature viewer, Feature table

Protein | Submitted name: **Iturin A synthetase A**
Gene | **ituA**
Organism | **Bacillus siamensis**
Status | Unreviewed - Annotation score: ●○○○○ - Protein predicted²

Function¹

GO - Molecular function²

- catalytic activity ¹ Source: InterPro
- phosphopantetheine binding ¹ Source: InterPro

View the complete GO annotation on QuickGO ...

Names & Taxonomy¹

Protein names ¹	Submitted name:
Iturin A synthetase A	Imported
Gene names ¹	Name: ituA
Bacillus siamensis	Imported
Taxonomic identifier ¹	659243 [NCBI]
Taxonomic lineage ¹	Bacteria > Firmicutes > Bacilli > Bacillales > Bacillaceae > Bacillus > Bacillus amyloquelaciens group ²

PTM / Processing¹

Amino acid modifications

Protein Iturin A

Lampiran 6. Formulasi dan pembuatan media

1. Media *Endo Agar*

Peptone 10 g
Di-photasium hydrogen phosphate 2,5 g
Lactose 10 g
Sodium sulfit anhidrate 3,3 g
 Fuchsin 0,3 g
 Agar-agar 12,5 g
 pH $7.4 \pm 0,2$ g

Dilarutkan 39 g bahan diatas dalam 1 liter aqua destilata kemudian disterilisasi dengan autoclave 121°C selama 15 menit (Merck 2005).

2. *Pseudomonas Selektive Agar (PSA)*

Pepton from casein 10,0 g
 Pepton from meat 3,5 g
 Laktosa 10,0 g
 Sodium sulfit 2,5 g
 Agar-agar 12,5 g
 pH 7,4

Dilarutkan 45,3 g bahan diatas dalam 1 liter aqua destilata ditambah dengan 10 mL gliserin, kemudian disterilkan dengan autoclave 121°C selama 15 menit.

3. *Nutrien Agar (NA)*

Pepton 10 g
 Ekstrak daging sapi / ekstrak ragi 10 g
 Agar 15 g
 NaCl 5 g
 pH 7,4

Dilarutkan 23 g bahan diatas dalam 1 liter aqua destilata kemudian disterilisasi menggunakan autoclave pada suhu 121°C selama 15 menit.

4. *Mueller Hinton Agar (MHA)*

Ekstrak daging 2.0 g
Casein hydrolisate 17,5 g

Tepung kanji 1,5 g

Agar-agar 17,0 g

pH $7.3 \pm 0,2$ g

Dilarutkan 38 g bahan diatas dalam 1 liter aqua destilata dan disterilisasi menggunakan autoclave pada suhu 121°C selama 15 menit (Merck 2005).

5. **Brain Heart Infusion (BHI)**

Ekstrak otak, ekstrak hati, dan pepton 27,5 g

Dekstrosa 2 g

Sodium chloride 5 g

Dy-sodium hydrogen phosphate 2,5 g

Agar-agar 15 g

pH $7.4 \pm 0,2$

Dilarutkan 52 g bahan diatas dalam 1 liter aqua destilata dan disterilisasi dengan autoclave pada suhu 121°C selama 15 menit.

6. **Sulfide Indol Motility (SIM)**

Peptone from casein 20.0 g

Peptone from meat 6. g

Ammonium iron(II)sitrate 0.2 g

Sodium thiosulfate 0.2 g

Agar-agar 3.0 g

pH: 7.3 ± 0.2

Bahan tersebut dilarutkan dalam 1 liter aqua destilata dan dan disterilisasi menggunakan autoclave pada suhu 121°C selama 15 menit (Merck 2005).

7. **Kligler's Iron Agar (KIA)**

Peptone from casein 15.0 g

Peptone from meat 5.0 g

Meat extract 3.0 g

Yeast extract 3.0 g

Sodium chloride 5. g 0

Lactose 10.0 g

*D(+)*glucose 1.0 g

Ammonium iron(III) sitrate 0.5 g

Sodium thiosulfate 0.5 g

Phenol red 0.024 g

Agar-agar 12.0. g

pH 7.4 ± 0.2

Bahan tersebut dilarutkan dalam 1 liter aqua destilata dan dan disterilisasi menggunakan autoclave pada suhu 121°C selama 15 menit (Merck 2005).

8. Lisyne Iron Agar (LIA)

Peptone from meat 5.0 g

Yeast extract 3.0 g

*D(+)*glucose 1.0 g

L-lysine monohydrochloride 10.0 g

Sodium thiosulfate 0.04 g

Ammonium iron(III) sitrate 0.5 g

Bromocresol purple 0.02 g

Agar-agar 12.5 g

pH: 6.7 ± 0.2 g

Bahan tersebut dilarutkan dalam 1 liter aqua destilata dan dan disterilisasi menggunakan autoclave pada suhu 121°C selama 15 menit (Merck 2005).

9. Simmon Sitrat Agar

Ammonium dihydrogen phosphate 1.0 g

Di-potassium hydrogen phosphate 1.0 g

Sodium chloride 5.0 g

Sodium sitrate 2.0 g

Magnesium sulfate 0.2 g

Bromothymol blue 0.08 g

Agar-agar 13.0. G

pH: 6.6 ± 0.2

Bahan tersebut dilarutkan dalam 1 liter aqua destilata dan dan disterilisasi menggunakan autoclave pada suhu 121°C selama 15 menit (Merck 2005).

Lampiran 7. Hasil ANOVA One Way isolat *Pseudomonas knackmussii*

Tests of Normality ^a							
Isolat	Waktu Fermentasi Isolat Bakteri Endofit	Kolmogorov-Smirnov ^b			Shapiro-Wilk		
		Statistic	df	Sig.	Statistic	df	Sig.
Diameter daya hambat	Kontrol positif	0,338	3	.	0,853	3	0,247
	Fermentasi hari ke 2	0,351	3	.	0,827	3	0,181
	Fermentasi hari ke 3	0,338	3	.	0,853	3	0,247
	Fermentasi hari ke 4	0,240	3	.	0,975	3	0,694
	Fermentasi hari ke 5	0,276	3	.	0,942	3	0,537
	Fermentasi hari ke 6	0,278	3	.	0,940	3	0,529
	Fermentasi hari ke 7	0,191	3	.	0,997	3	0,900

Test of Homogeneity of Variances

Diameter daya hambat				
Levene Statistic	df1	df2	Sig.	
8,792	7	16	0,000	

ANOVA

Diameter daya hambat					
	Sum of Squares	Df	Mean Square	F	Sig.
Between Groups	2008,444	7	286,921	544,652	0,000
Within Groups	8,429	16	0,527		
Total	2016,873	23			

Diameter Daya Hambat

Waktu Fermentasi Isolat		Subset for alpha = 0.05		
Bakteri Endofit	N	1	2	3
Kontrol Negatif	3	,0000		
Fermentasi Hari ke 2	3		11,8333	
Fermentasi Hari ke 3	3		12,2133	
Fermentasi Hari ke 5	3		12,2600	
Fermentasi Hari ke 7	3		12,2600	
Fermentasi Hari ke 6	3		12,2733	
Fermentasi Hari ke 4	3		12,3133	
Kontrol Positif	3			35,3367
Sig.		1,000	,479	1,000

Means for groups in homogeneous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3,000.

Lampiran 8. Hasil ANOVA One Way isolat *Bacillus siamensis*

Isolat	Waktu Fermentasi Isolat Bakteri Endofit	Tests of Normality ^a			Shapiro-Wilk		
		Kolmogorov-Smirnov ^b Statistic	df	Sig.	Statistic	df	Sig.
Diameter daya hambat	Kontrol positif	0,311	3	.	0,898	3	0,378
	Fermentasi hari ke 2	0,264	3	.	0,955	3	0,590
	Fermentasi hari ke 3	0,307	3	.	0,903	3	0,394
	Fermentasi hari ke 4	0,381	3	.	0,759	3	0,020
	Fermentasi hari ke 5	0,177	3	.	1,000	3	0,968
	Fermentasi hari ke 6	0,344	3	.	0,841	3	0,217
	Fermentasi hari ke 7	0,312	3	.	0,896	3	0,372

Test of Homogeneity of Variances

Diameter daya hambat			
Levene Statistic	df1	df2	Sig.
3,731	7	16	0,014

ANOVA

Diameterdayahambat					
	Sum of Squares	Df	Mean Square	F	Sig.
Between Groups	2063,487	7	294,784	1556,717	0,000
Within Groups	3,030	16	,189		
Total	2066,517	23			

Diameter Daya Hambat

Duncan^a

Waktu Fermentasi Isolat		Subset for alpha = 0.05				
Bakteri Endofit	N	1	2	3	4	5
Kontrol Negatif	3	,0000				
Fermentasi Hari ke 2	3		9,4033			
Fermentasi Hari ke 3	3			10,1700		
Fermentasi Hari ke 6	3			10,5933		
Fermentasi Hari ke 4	3			10,7767		
Fermentasi Hari ke 5	3			10,8867	10,8867	
Fermentasi Hari ke 7	3				11,6033	
Kontrol Positif	3					35,0133
Sig.		1,000	1,000	,081	,061	1,000

Means for groups in homogeneous subsets are displayed.

a. Uses Harmonic Mean Sample Size = 3,000.

Lampiran 9. Hasil ANOVA Two Way isolat bakteri endofit *Pseudomonas knackmussii* dan *Bacillus siamensis*

Tests of Normality						
	Kolmogorov-Smirnov ^a			Shapiro-Wilk		
	Statistic	df	Sig.	Statistic	df	Sig.
Standardized Residual for hasil	,180	48	,041	,914	48	,052

a. Lilliefors Significance Correction

Between-Subjects Factors

		Value Label	N
isolat bakteri endofit	1,00	pseudomonas knackmussii	18
	2,00	bacillus siamensis	18
	3,00	kontrol positif	6
	4,00	kontrol negatif	6
waktu fermentasi	1,00	Fermentasi hari ke-2	6
	2,00	Fermentasi hari ke-3	6
	3,00	Fermentasi hari ke-4	6
	4,00	Fermentasi hari ke-5	6
	5,00	Fermentasi hari ke-6	6
	6,00	Fermentasi hari ke-7	6
	7,00	kontrol positif	6
	8,00	kontrol negatif	6

Descriptive Statistics

Dependent Variable: diameter daya hambat

isolat bakteri endofit	waktu fermentasi	Mean	Std. Deviation	N
pseudomonas knackmussii	Fermentasi hari ke-2	11,8333	1,00620	3
	Fermentasi hari ke-3	12,2133	1,58658	3
	Fermentasi hari ke-4	12,3133	,12662	3
	Fermentasi hari ke-5	12,2600	,03606	3
	Fermentasi hari ke-6	12,2733	,20108	3
	Fermentasi hari ke-7	12,2600	,16523	3
	Total	12,1922	,67338	18
bacillus siamensis	Fermentasi hari ke-2	9,4033	,23029	3
	Fermentasi hari ke-3	10,1700	,80517	3
	Fermentasi hari ke-4	10,7767	,48789	3
	Fermentasi hari ke-5	10,8867	,34005	3
	Fermentasi hari ke-6	10,5933	,57239	3
	Fermentasi hari ke-7	11,6033	,23245	3
	Total	10,5722	,80697	18
kontrol positif	kontrol positif	35,1750	,55001	6
	Total	35,1750	,55001	6
kontrol negatif	kontrol negatif	,0000	,00000	6
	Total	,0000	,00000	6
Total	Fermentasi hari ke-2	10,6183	1,48245	6
	Fermentasi hari ke-3	11,1917	1,58707	6
	Fermentasi hari ke-4	11,5450	,90002	6
	Fermentasi hari ke-5	11,5733	,78268	6
	Fermentasi hari ke-6	11,4333	,99697	6
	Fermentasi hari ke-7	11,9317	,40236	6
	kontrol positif	35,1750	,55001	6
	kontrol negatif	,0000	,00000	6
	Total	12,9335	9,34254	48

Levene's Test of Equality of Error Variances^a

Dependent Variable: diameter daya hambat

F	df1	df2	Sig.
7,747	13	34	,058

Tests the null hypothesis that the error variance of the dependent variable is equal across groups.

a. Design: Intercept + isolat + fermentasi + isolat * fermentasi

Tests of Between-Subjects Effects

Dependent Variable: diameter daya hambat

Source	Type III Sum of Squares	df	Mean Square	F	Sig.
Corrected Model	4090,687 ^a	13	314,668	921,085	,000
Intercept	8374,008	1	8374,008	24512,071	,000
Isolat	23,620	1	23,620	69,138	,000
Fermentasi	5,924	5	1,185	3,468	,012
isolat * fermentasi	2,752	5	,550	1,611	,184
Error	11,615	34	,342		
Total	12131,574	48			
Corrected Total	4102,302	47			

a. R Squared = ,997 (Adjusted R Squared = ,996)

Multiple Comparisons

Dependent Variable: diameter daya hambat

						95% Confidence Interval		
			Mean Difference (I-J)	Std. Error	Sig.	Lower Bound	Upper Bound	
LSD	(I) isolat bakteri endofit	(J) isolat bakteri endofit						
		pseudomonas knackmussii	bacillus siamensis	1,6200*	,19483	,139	1,2241	2,0159
			kontrol positif	-22,9828*	,27553	,087	-23,5427	-22,4228
			kontrol negatif	12,1922*	,27553	,040	11,6323	12,7522
	bacillus siamensis	pseudomonas knackmussii	-1,6200*	,19483	,355	-2,0159	-1,2241	
			kontrol positif	-24,6028*	,27553	,076	-25,1627	-24,0428
			kontrol negatif	10,5722*	,27553	,037	10,0123	11,1322
	kontrol positif	pseudomonas knackmussii	22,9828*	,27553	,525	22,4228	23,5427	
			bacillus siamensis	24,6028*	,27553	,312	24,0428	25,1627
			kontrol negatif	35,1750*	,33746	,021	34,4892	35,8608
	kontrol negatif	pseudomonas knackmussii	-12,1922*	,27553	,940	-12,7522	-11,6323	
			bacillus siamensis	-10,5722*	,27553	,348	-11,1322	-10,0123
			kontrol positif	-35,1750*	,33746	,021	-35,8608	-34,4892

Based on observed means.

The error term is Mean Square(Error) = ,342.

*. The mean difference is significant at the ,05 level.

diameter daya hambat

	isolat bakteri endofit	N	Subset			
			1	2	3	4
Duncan ^{a,b,c}	kontrol negatif	6	,0000			
	bacillus siamensis	18		10,5722		
	pseudomonas knackmussii	18			12,1922	
	kontrol positif	6				35,1750
	Sig.		1,000	1,000	1,000	1,000

Means for groups in homogeneous subsets are displayed.

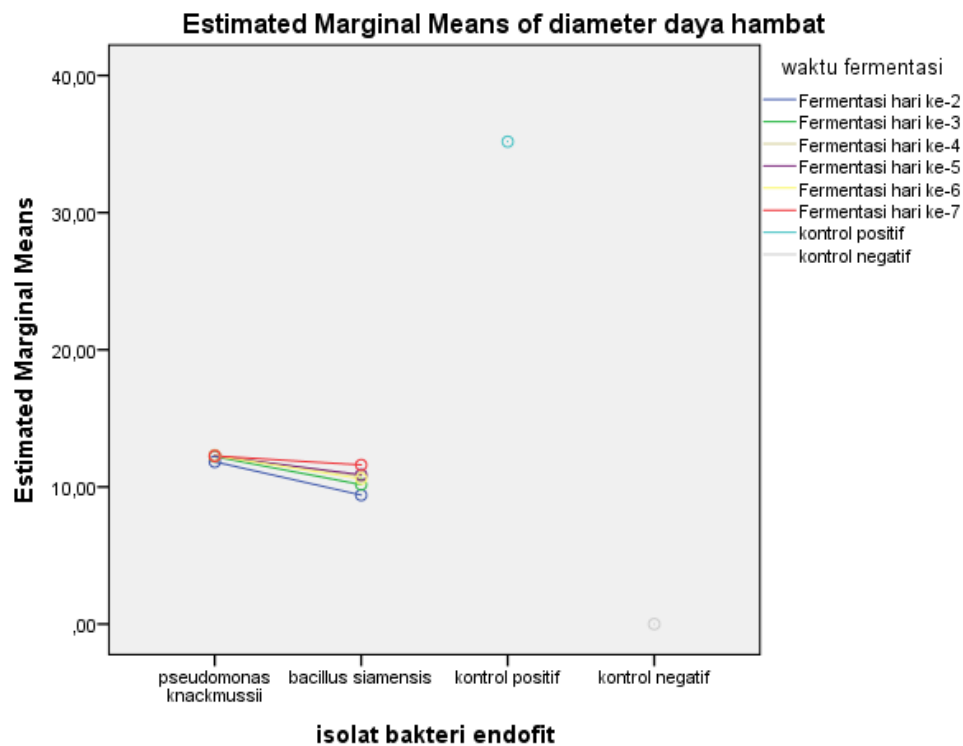
Based on observed means.

The error term is Mean Square(Error) = ,342.

a. Uses Harmonic Mean Sample Size = 9,000.

b. The group sizes are unequal. The harmonic mean of the group sizes is used. Type I error levels are not guaranteed.

c. Alpha = ,05.



Non-estimable means are not plotted