

Manufacture of Local Microorganism (MOL) from Vegetable Waste with Nutrition Source Supply Variation

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Abstract—Local Microorganisms (MOL) is a bioactivator consisting of a collection of various kinds of microorganisms that function as agents for the decomposition of organic matter. Vegetable waste is a collection of various kinds of vegetables after being sorted because they are not suitable for sale. Almost all vegetable waste will undergo lactic acid fermentation carried out by lactic acid bacteria (LAB). Lactic acid bacteria (LAB) are gram-positive bacteria, do not form spores, are catalase negative, are resistant to acidic conditions, and are facultative anaerobes. The purpose of this study was to determine the correct composition of rice washing water and granulated sugar for the highest number of microbes in MOL. MOL optimization was carried out by means of anaerobic fermentation for 14 days. The results of this study indicate that the microbes contained in MOL are *Bacillus subtilus*, *Bacillus cereus*, *Lactobacillus acidophilus*, *Spirillum*, *Streptovercillium*, and *Leuconostoc mesenterousdes*. The highest total population of microorganisms in variations in the composition of rice washing water and sugar is in sample C1 which is 91×10^5 Colonies/100 mL in MOL vegetable waste, which is 650 mL of rice washing water and 100 grams of sugar.

Index Terms—About: MOL, local microorganism, rice washing water, vegetable waste.

I. INTRODUCTION

The existence of waste as a material that is wasted or disposed of from sources resulting from human and natural activities that do not yet have economic value has become a crucial environmental problem (Sunarsih, 2018).

Organic waste is a type of waste composed mostly of organic compounds such as plant

residues, animals or feces. By nature, organic waste can be classified as degradable waste, which is a type of waste with natural properties that are easy to degrade by living things, especially microbes (Waluyo, 2018). In the process of degradation or decomposition of organic matter, microbes act as inoculums or starters (activators) that accelerate the decomposition process (Manullang 2017).

Various kinds of decomposing microorganisms in nature which are often referred to as local microorganisms (MOL) can be used as bioactivators. This type of microbe can be cultured using various sources of organic matter. Vegetable waste can be a source of organic matter and a good medium for the proliferation of decomposing microorganisms, and can be used as a bioactivator (Suwatanti, 2017). In addition to being a source of organic matter which after rotting can be a source of bacteria for moles, in its manufacture, moles also require carbohydrates in rice washing water and glucose in granulated sugar which act as sources of energy or nutrition for microorganisms in mole solution (Lindung, 2015).

In a previous study conducted by Suwatanti (2017), it was not known the type and number of microbes contained in the mole solution of vegetable waste that were capable of being a bioactivator. In this study, before being used as moles, the fermented solution was only physically tested to determine whether moles were successfully formed or not.

Because of the theoretical foundations above and the results from previous research, we want to conduct research on the manufacture of moles from vegetable waste with variations in the provision of nutritional sources to determine the types of microbes contained in moles and the composition of the right nutritional sources for the results of moles with the highest number of microbes. The purpose of this study was to determine the types of microbes contained in the MOL of vegetable waste and to determine the composition of rice and sugar washing

waterappropriate sand to yield the highest number of microbes in the MOL of vegetable waste.

II. RESEARCH METHODOLOGY

The research was carried out at the Chemical Engineering Laboratory of ITN Malang. The research phase is literature study, research preparation, research implementation, data collection, data analysis, evaluation and finally report generation.

A. Research variable

1. Controlled Variable
 - Salt Weight :45 grams
 - Vegetable Waste Weight : 500grams
 - TimeMol Fermentation :14 days
2. Independent Variable
 - Water VolumeRice Wash:600, 625, 650 ml
 - Sugar Weight :100, 125, 150, 175, 200 gram

B. Tools and Materials

The tools needed in this research include used bottles, fermentors, measuring cups and plastic hoses. Meanwhile, the materials needed include Aquadest, rice washing water, salt, sugar, and vegetable waste (cabbage, mustard greens and tomatoes).

C. Research procedure

The stages of the research procedure "Manufacture of Local Microorganisms (MOL) from Vegetable Waste with Nutrition Source

Supply Variation " are as follows:

1. prepare vegetable waste (a mixture of cabbage, mustard greens, tomatoes) as much as 500 grams and then chop it into small pieces
2. put vegetable waste into a fermentor with a modified 1.5 liter size with a plastic intermittent cover
3. connecting a plastic hose to a 600 ml plastic bottle filled with lime water which serves to observe the presence or absence of CO2 gas
4. adding granulated sugar to vegetable waste in the fermentor as much as 100, 125, 150, 175, 200 grams respectively for fermentors 1, 2, 3, 4, 5
5. added leri water (rice washing water) from the first wash to the vegetable waste in the fermentor as much as 600 ml each for fermentors A1-A5, 625 ml for fermentors B1-B5 and 650 ml for fermentors C1-C5
6. add 45 grams of salt for each fermentor
7. anaerobically ferment vegetable waste(fermentor is tightly closed) for 14 days
8. observing the presence or absence of changes in water color, physical pieces of rotting vegetable waste, and the presence or absence of CO2 gas as indicated by the presence or absence of lime deposits formed in plastic bottles filled with lime water. on
9. open the bottle cap fermentor after 14 days of fermentation and then perform an analysis of the type of microbe and the total microbe on the sample of the mole solution formed.

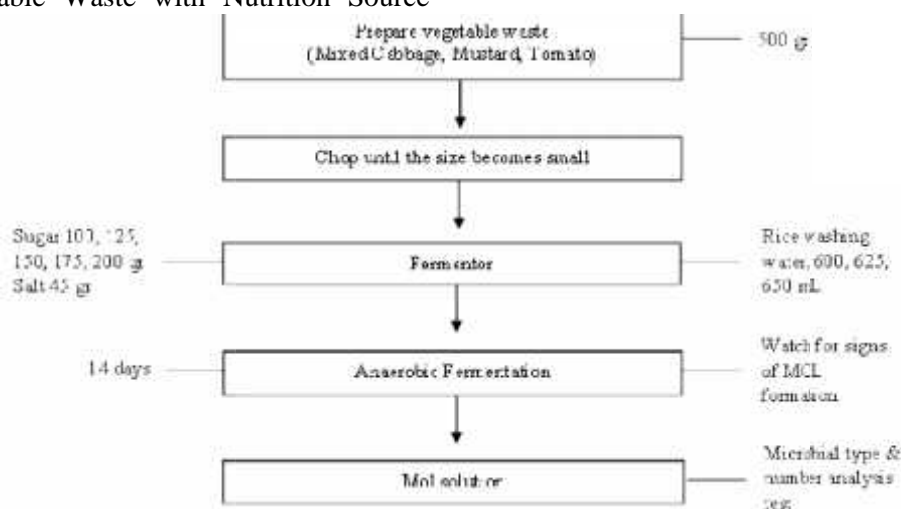


Figure 1: Research Flowchart

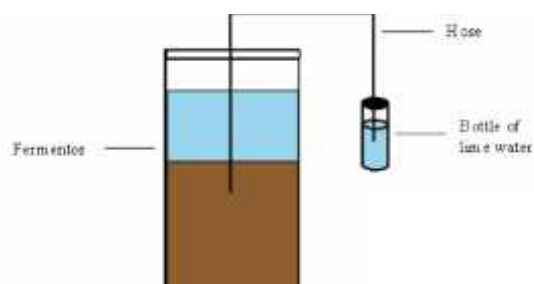


Figure 2: Fermentor Circuit

III. RESULT AND DISCUSSIONS

A. Microbial Type Analysis Results

The results of the analysis of the type of microbes carried out at the Chemical Engineering Microbiology Laboratory, ITN Malang.

Table 1
Microbial Type Analysis Results

Sample	Rice Wash Water (mL)	Sugar (grams)	Types of Microbes
A1		100	<i>Bacillus Subtillus</i> , <i>Lactobacillus Acidophilus</i>
A2		125	<i>Bacillus Cereus</i>
A3	600	150	<i>Spirillum</i>
A4		175	<i>Bacillus Subtillus</i> , <i>Streptovercillium</i>
A5		200	<i>Bacillus Subtillus</i> , <i>Bacillus Cereus</i>
B1		100	<i>Bacillus Subtillus</i> , <i>Spirillum</i>
B2		125	<i>Bacillus Subtillus</i>
B3	625	150	<i>Bacillus Subtillus</i>
B4		175	<i>Bacillus Subtillus</i> , <i>Spirillum</i>
B5		200	<i>Spirillum</i> , <i>Bacillus Subtillus</i>
C1		100	<i>Spirillum</i>
C2		125	<i>Bacillus Subtillus</i> , <i>Leuconostoc Mesenterous des</i>
C3	650	150	<i>Leuconostoc Mesenterous des</i> , <i>Lactobacillus Acidophilus</i>
C4		175	<i>Bacillus Subtillus</i> , <i>Streptovercillium</i>
C5		200	<i>Bacillus Cereus</i> , <i>Bacillus Subtillus</i>

B. Results of Analysis of the Number of Microbes

The results of the analysis of the number of microbes carried out at the Chemical Engineering Microbiology Laboratory, ITN Malang.

Table 2
Results of Analysis of the Number of Microbes

Sample	Rice Wash Water (mL)	Sugar (grams)	Number of Microbes (Colonies / 100 mL)
A1		100	61 x 10 ⁴
A2		125	36 x 10 ⁵
A3	600	150	6 x 10 ⁵
A4		175	93 x 10 ⁴
A5		200	29 x 10 ⁴
B1		100	27 x 10 ⁴
B2		125	173 x 10 ⁴
B3	625	150	61 x 10 ⁴
B4		175	12 x 10 ⁴
B5		200	12 x 10 ⁴
C1		100	91 x 10 ⁵
C2		125	15 x 10 ⁴
C3	650	150	36 x 10 ⁴
C4		175	15 x 10 ³
C5		200	3 x 10 ⁴

C. Discussion of the Results of the Analysis of Microbial Types

From the results of the analysis of the types of microbes in the sample solution, it is known that the microbes contained in local microorganisms from vegetable waste are types of *Bacillus Subtillus*, *Bacillus Cereus*, *Lactobacillus Acidophilus*, *Spirillum*, *Streptovercillium*, and *Leuconostoc Mesenterous des*.



Figure 3: *Bacillus Subtillus*

Bacillus Subtillus is a type of microbe that can be used as a decomposer because it is proteolytic so it is able to hydrolyze protein into amino

acids. In its utilization, *B. subtilis* is used to increase the nitrogen and phosphorus content (Alamsjah, 2011).

In addition, according to Ruhnayat, et al. (2007), *B. Subtilis* is also widely found in bio fertilizers because it is a type of phosphate solubilizing microbe. These microbes can grow on insoluble phosphate (insoluble phosphate) and can also convert natural phosphate into a soluble form.



Figure 4: *Bacillus Cereus*

Bacillus Cereus is one type of microbe *Bacillus Sp* which has the ability to degrade organic matter (Khoiriyah, 2017). According to Sari, et al. (2017), besides being able to degrade organic matter, *B. cereus* also has the potential to degrade organic acids and decrease COD and BOD levels. With its ability to hydrolyze starch, this microbe is able to increase the average quality of nitrogen.



Figure 5: *Lactobacillus Acidophilus*

Lactobacillus Acidophilus is a type of microbe that in fermenting carbohydrates and sugars can produce lactic acid. In addition, *L. Acidophilus* is also able to inhibit the growth rate of pathogenic bacteria because it is a type of probiotic microbe (Adriani, 2008).



Figure 6: *Spirillum*

Spirillum is one type of denitrifying microbe that is able to reduce nitrate and nitrite to nitrogen. As a denitrifying microbe, *Spirillum* is also a facultative microbe that can live in both aerobic and anaerobic conditions (Supono, 2015). According to Kunarso (1988), *Spirillum*

belongs to the class of reducing microbes or decomposers that are able to decompose nutrient and mineral materials into nutrient sources of energy.



Figure 7: *Streptovercillium*

Streptovercillium is one type of microbial *Actinomycetes* that plays a role in forming humus and expanding the growth area with its long chain spores. These microbes can be found in soils with high organic matter content as well as in plant rhizoforous parts (Sektiono, 2016). According to Nugroho, et al. (2019), *Streptovercillium* is also called *Streptomyces* and belongs to the microbial transglutaminase group, which is a type of microorganism capable of producing transglutaminase. In addition, *Streptomyces* is also a type of root fertilizing microbe that produces nutrients and growth hormones and as a biocontrol agent in biological organic fertilizers that can improve soil respiration (Antonius, 2011).



Figure 8: *Leuconostoc Mesenterousdes*

Leuconostoc Mesenterousdes is one type of lactic microbe that has the ability to remodel complex compounds into simpler compounds that produce lactic acid (Aliya, 2015). In addition, according to Kusmiati, et al. (2002), *L. Mesenterousdes* can change the quality of fermented products.

D. Discussion of the results of the analysis of the number of microbes

The results of the analysis of the number of microbes from the mole solution of vegetable waste showed the highest results for the number of microbes in sample C1 which was 91×10^5 Colonies/100 mL and the smallest results for the number of microbes in sample C4 was 15×10^3 Colonies/100 mL.

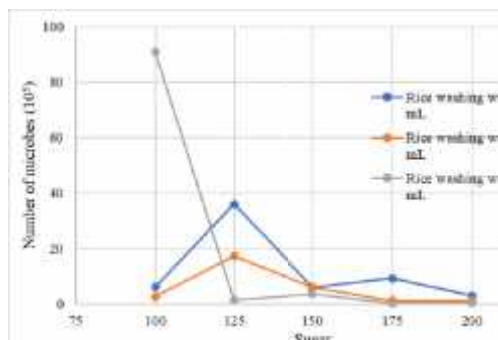


Figure 9: Graph of Relationship Between Amount of Sugar and Number of Microbes in Various Amounts of Rice Washing Water

To increase the amount of rice washing water, it was shown that there was an increase in the number of microbes in treatment 1 while in treatments 2, 3, 4, and 5 there was a decrease in the number of microbes.

This increase is because the carbohydrate content in rice washing water is a source of nutrients needed by microbes to grow (Budiyani, 2016). Carbohydrates are a type of staple food for anaerobic microbes as a source of energy (Indriani, 2013). According to Marsiningsih, et al. (2015), the amount of nutrients available in the mole solution is a factor that affects the availability of the number of microbes in the mole. So with the increasing number of nutrients, the number of microbes present in the mole solution also increases.

While the decrease in the number of microbes occurs because in the early stages of fermentation, certain types of microbes break down carbohydrates or starch to produce glucose, so that glucose increases but carbohydrate or starch levels decrease. The decrease in starch content as an energy source for certain types of microbes results in a decrease in the number of surviving microbes.

To increase the amount of granulated sugar, it was shown that there was an increase and decrease in the number of microbes in treatment A, B, and a decrease in the number of microbes in treatment C.

This increase is because sugar as a food source provides energy which causes optimal microbial growth. The increase was also because the microbes were in the growth log phase which caused the microbial cell defense to multiply.

While the decrease in the number of microbes

because there are microbes that die in the competition for nutritional sources. Competition as an interaction that occurs affects the ability of microbes to grow and live.

In addition, the decrease in the number of microbes also occurs because some microbes are in a static phase then the death phase (Irpan, 2018).

According to Budiyani, et al. (2016), in addition to sources of nutrition or energy for microbial growth, there are several other factors regarding operating conditions that affect microbial growth in the fermentation process, namely temperature, pH and oxygen. The decrease in the number of microbes that occurred was suspected because the anaerobic fermentation process was not maintained so that there was a leak in the fermentor cover which resulted in oxygen entering and making the microbes die.

In the fermentation process, carbohydrates will break down into simpler sugars, namely dextrose, mannose and sucrose which are used by lactic acid bacteria as an energy source and produce acidic compounds such as lactic acid and other volatile compounds that cause an acidic atmosphere. low product pH (Azizah, 2014). This causes bacteria to work more extra in the process of breaking down carbohydrates into simpler sugars so that bacteria have an excess of energy sources or carbohydrates, where excess carbohydrates can cause death in fermentor microorganisms, this is due to the results of too much alcohol so that it can damage microorganisms (Hidayah, 2019).

IV. CONCLUSION

From the research results of Making Local Microorganisms (MOL) from Vegetable Waste with Variations in Provision of Nutrient Sources, the following conclusions are obtained:

The types of microbes found in local microorganisms in vegetable waste are the types of *Bacillus subtilus*, *Bacillus cereus*, *Lactobacillus acidophilus*, *Spirillum*, *Streptovercillium*, and *Leuconostocmesenterousdes*.

The correct composition of rice washing water and sugar for the highest number of microbes in sample C1 is 91 x 10⁵ Colonies/100 mL in MOL of vegetable waste, which is 650 mL of rice washing water and 100 grams of sugar.

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