

The Effect of Different Starter Dosage to Organoleptic Value of Kefir Cow Milk Products

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Abstract. The study of "The Effect of Different Starter Dosage to Organoleptic Value of Kefir Cow Milk Products" was carried out from May to July 2016 at the Microbiology Research Laboratory, Department of Biology, Faculty of Mathematics and Natural Sciences, Andalas University, Padang. The purpose of this study was to determine the organoleptic value (aroma, taste and organoleptic) of cow's milk kefir products from three different starter doses, and the results were with Wilcoxon Level Test. The results showed that organoleptic assessment of the aroma of cow milk kefir products with a treatment dose of 15% starter (3.00) was the most preferred dose for the panelist, while kefir with a treatment dose of 5% (2.27) is a dose that is less preferred by panelists. The taste assessment of the organoleptic of cow's milk kefir products, the starter dose of 10% (2.87) was the most preferred by the panelists, but the dose of 15% (2.33) was the least. Similar to the taste, the consistency of cow milk kefir with the dose of 10% (3.07) was most preferred, while the dose of 15% (2.33) was least preferred by the panelists.

Keywords: dosage, cow milk kefir, organoleptic assessment

I. INTRODUCTION

Kefir is one of the fermented milk products which contain of bacteria and yeast that work symbiotically and produce organic acid compounds, CO₂ and alcohol. Kefir contains 0.65-1.33 g / l CO₂, 3.16-3.18% protein, 3.07-3.17% fat, 1.8 to 3.8% lactose, 0.5 - 1.5 % ethanol and 0.7-1.0% lactic acid [1].

The kefir was produced from fermented milk by lactic acid producing bacteria, acetic acid producing bacteria and yeast. The taste of kefir is dominated by sour taste caused by the activity of acid-producing bacteria [2], which originally from the Caucasus mountain region between the Black Sea and the Caspian Sea, Southwest Russia. Kefir has also known as kepi, kippe, kapov, kephir and kiaphir, and has been widely consumed in several Asian and Scandinavian countries [3]. In Indonesia this type of fermented milk has not been publically known.

Kefir was produced from milk which is fermented by lactic acid bacteria such as *Lactobacillus lactis*, *Lactobacillus delbrueckii* subsp. *bulgaricus* and yeast that produces acid and alcohol. At the end of the ripening process, it should be covered to form carbonates [4]. The bacteria contained in kefir are homo-fermentative lactic acid bacteria which produce 90% lactic acid [5], and dominated by *Lactobacillus* groups [6]. *Lactobacillus bulgaricus* and *Streptococcus thermophilus* are bacteria that can be found in warm to hot, tropical and subtropical regions [7].

Kefir fermentation can be lasts for 18-24 hours at the temperature of 22°C [8]. Two groups of microorganisms, lactic acid bacteria and *Candida* yeast, were work in symbiosis in kefir fermentation. Lactic acid was produced by lactic acid bacteria from glucose breakdown which stimulates yeast growth. While yeast is important in kefir fermentation process, because it produces ethanol compounds and flavor-forming components, so produce a distinctive taste [3]. Organoleptic properties are food

compounds which were assessed using the five senses of subjective judgments [9]. Organoleptic tests are commonly used for commodity quality checks, process control during processing and as a method of measurement and measurement of quality properties in research. The value was assessed by the sense of smell, taste and stimulation of the mouth [10]. So far, there is no study has been done on organoleptic values (aroma, taste and organoleptic) of cow's milk kefir products that assessed on three different starter doses (5%, 10% and 15%). Therefore this study was conducted to investigate the effect of three different doses of starters to the organoleptic of kefir produced.

II. RESEARCH METHOD

The research was conducted from May to July 2016 at the Microbiology Research Laboratory, Department of Biology, Faculty of Mathematics and Natural Sciences, Andalas University, Padang. The data obtained was analysed in Wilcoxon tests. Before organoleptic assessment, cow milk kefir was made using a 10% starter per volume of ultra-high temperature (UHT) cow milk that was fermented for 36 hours in the following sequences: 24 hours was fermented at 25 ° C, then 12 hours was fermented at 5 ° C [11]. For this study, kefir was made of three doses (5%, 10%, and 15%) per volume of the UHT cow milk. These tree different kefir contents were tested on 15 panelists. Control was also performed using commercial kefir product.

III. RESULTS AND DISCUSSION

Results

Algasida EhEp Activity Level

The level of algae activity can be known by calculating the percentage of treatment algae density compared to controls. Evaluation is done on the last day of observation. Calculation of the percentage of bacterial

consortium algal activity using the equation in [19]. Episymbiont bacteria consortium showed low algal activity in all target plankton, except *Nitzschia* sp. and *Porphyridium* sp (Table 1).

Plankton Morphology

Plankton growth was tested to be low, except for *Nitzschia* sp. During the five days of observation of plankton microstructure conditions, there were various symptoms of cell damage that could be recognized from cell deformation and damage to cell components. The cytoplasm is damaged with shrinkage symptoms > 50% of its original size (Figure 1) *Plankton density*

Dinoflagellate density both control and treatment were evaluated daily for five days. *Porphyridium* sp. Plankton cell density. and the BG treated with EhEp showed a thin increase in density compared to the control (Figure 2). This also occurs with the growth of diatomic plankton groups, the pattern of growth follows control (Figure 3).

Effects of Algal Activity on Growth

Student t-test results showed the algal activity of EhEp episymbiont bacteria had almost no effect on plankton growth. This is indicated by no significant difference between the control and treatment at 5% significance level, except in *Nitzschia* sp. The algal activity shows a significant effect. This means that the episymbiont bacteria consortium has no significant effect on the growth of dinoflagellates and diatoms (Table 2).

Discussion

Many bacterial isolates have been known to have the ability as algicides. The source of these isolates are planktonic bacteria. Some even develop along with algae growth explosions. The mechanism is a natural counterweight for the plankton group. Some bacteria that are known to be antagonistic in plankton are *Flavobacterium* sp. 5N-3 which can inhibit the growth of *Gymnodinium* sp. *Bacillus* sp. SY-1 isolated from Masan waters in South Korea was able to suppress the growth of *Cochlodinium polykrikoides* [20]. Bacterial isolates obtained from Japan's Kagosima Bay *Saprospira* sp. SS98-5 is even able to reduce the growth of the diatom group *Chaetoceros ceratosporum* [21].

Researcher [22] succeeded in isolating *P. haloplanktis* AFMB-08041 from the surface water in Masan Bay, Korea. These bacteria are able to inhibit the growth of *Prorocentrum minimum* up to 90% within 5 days. Observations show that plankton is directly attacked by this bacteria. The algal compound produced is active *beta-glucosidase*. Other capabilities of these bacteria can suppress the growth of various species of HABs including *Alexandrium tamarense*, *Akashiwo sanguinea*, *Cochlodinium polykrikoides*, *Gymnodinium catenatum*, and *Heterosigma akashiwo*. These bacteria tend to attack plankton with morphology similar to *P. minimum* such as *P. dentatum*. Therefore, the bacteria is prepared to control HABs in the natural environment.

This study uses bacterial isolates from EhEp. In the algal activity test against *Porphyridium* sp. and BG (dinoflagellate) found that episymbiont bacteria consortium was less able to inhibit algal growth. The EhEp bacterial

consortium showed <40%, with a five-day inoculation time. The algal activity was confirmed also by the student t-test results which stated that there was no significant difference in plankton growth density between control and treatment.

EhEp bacterial consortium on *Nitzschia* sp. showed algal activity tended to be significant. Its growth density showed an increase from 15,967 cells.mL⁻¹ on the first day to 43,245 cells.mL⁻¹. This number is smaller than the control on the fifth day which reached 54,400 cells.mL⁻¹. The effect of algal activity on *Nitzschia* sp. can be observed in cell damage, characterized by cell deformation and cytoplasmic shrinkage. It is also owned by *Saprospira* sp. SS98-5 was able to reduce the growth of the diatom group *Chaetoceros ceratosporum* [21].

The *Nitzschia* genera consists of many species and can live in fresh, brackish and sea waters. The genera has very fast reproductive abilities and can adapt to various environmental conditions. Environmental factors that influence its development are the presence of abundant nutrients, lack of predators, high DO, high temperatures and high light intensity. Some of the species have the ability to produce neurotoxins (domoic acid). In tropical waters its can be very high abundance, if the predator is a little. High abundance can trigger the appearance of species that produce toxins. Researcher [23] succeeded in isolating *Nitzschia navis-varingica* from Vietnamese waters which had the ability to produce domoic acid.

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Observations show that plankton trapped by cytoplasmic biofilms shrink. EhEp consortium bacteria can produce secondary metabolites which can seep into plankton cells. This also occurs in diatoms, where cells undergo deformation and the cytoplasm of the cell undergoes shrinkage. Researcher [24] used biosurfactant glycolipid biosurfactant sophorolipid isolated from yeast *Turolopsis gabarata* as algal activity for *Alexandrium tamarense*, *Heterosigma akashiwo* and *Cochlodinium polykrikoides*. These metabolites damage the cytoplasm and lyse cell membranes. Researcher [25] found *Streptomyces* sp. L74 produces secondary metabolites which can disrupt the mechanism of the microcystis aeruginosa cyanobacter antioxidant. These mechanisms include indirect inhibition by producing secondary metabolites. Researcher [26] found that bacteria capable of suppressing plankton growth have the ability to produce chitinase enzymes. The enzyme can damage the diatom cell wall, so that the diatom undergoes lysis. The destruction of the cell wall is a direct attack mechanism.

IV. CONCLUSION

The EhEp bacterial consortium showed less significant algal activity against diatoms and dinoflagellates, except *Nitzschia* sp. This fact makes the episymbiont of *E. acoroides* not yet potentially used as a source of algal activity. The ability of seagrass bacteria consortium requires further study. The aspects that can be examined are episymbiont interaction patterns with other planktonic microorganisms and with the seagrass. Ecological aspects that play a role in this interaction pattern can be investigated to find out the influential environmental factors, which require further study. Further things that can be developed are episymbiont interaction patterns with other planktonic microorganisms and with the seagrass.

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Abbreviations

EhEd	: <i>Enhalus acoroides</i>	endosymbion consortium bacteria
ThEd	: <i>Thalassia hemprichii</i>	endosymbion consortium bacteria
BG	: the dinoflagellate which was isolated from Jakarta Bay when it was blooming in 2015 and had not been identified	
IMK Daigos's	: medium for plankton	
MPC-200	: Plankton counter	
%AA	: percent of algacidal activity	
T	: Treatment growth	
C	: Control Growth	
t	: the observation period	
K	: control	
rP	: treatment	