The biological prospective of red-pigmented bacteria cultured from contaminated agar media

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Abstract. Arifiyanto A, Afriani H, Putri MH, Damayanti B, Riyanto CL. 2021. The biological prospective of red-pigmented bacteria cultured from contaminated agar media. Biodiversitas 22: 1152-1159. Contaminated agar media was often depleted due to the growth of microbes, which is undesirable for culture. However, the contaminating microbes usually have a distinctive morphology. This research aimed to identify the potential of red-pigmented bacteria originated from the contamination of Drosophila melanogaster larvae. Fruit fly larvae that grow on contaminated tryptic soy agar were accompanied by the appearance of the red-pigmented bacteria colony. The bacterial colonies were purified by the re-streaking method on tryptic soybean agar. This strain was characterized morphologically, biochemically, and molecularly. Results showed that the MBC1 strain was identified as Serratia marcescens. The various metal susceptibility tests at 25 ppm did not affect the growth of the MBC1 strain. Meanwhile, it was also able to inhibit the growth of certain pathogens such as Aspergillus niger, Candida sp., Fusarium sp., and Rigidoporus sp. Strain MBC1 was able to produce antioxidant compounds. The lipolytic and amyloytic activity can be developed for bioremediate agriculture waste and biosurfactant production.

Keywords: Contaminated culture, Drosophila melanogaster, red-pigmented bacteria, prospect, Serratia

INTRODUCTION

Microbial contamination was an unexpected problem in cell culture. Re-production of fermentation must be pursued if the culture media was contaminated. Longer time and additional media became inefficiency constraints due to contamination. The presence of non-target microbe growth was one of the causes of contamination. Repeated use of glassware was may also increase the likelihood of contamination. The source of contamination frequently also arose from media culture. The airborne particulates from the environment also increased the chances of contamination (Mirjalili et al. 2005).

The nutritional content in the contaminated media often attracts insects. Insects have been known as a microbial vector. Genera Staphylococcus and Weisella were dominantly found at microbial communities of the house fly Musca domestica (Park et al. 2019). Acetobacter thailandicus identified as bacteria that were mutually associated with Drosophila melanogaster (Pais et al. 2018). D. melanogaster was also reported as a host of certain genera of yeast like Candida, Cryptococcus, Hanseniaspora, Hanusa, Kloeckera, Kluyveromyces, Pichia, Rhodotorula, Saccharomyces, Saccharomycopsis, and Torulopsis (Broderick and Lemaitre 2012). Lactic acid bacteria Enterococci played a role in modulating Drosophila to lay eggs on decaying food. Females were lay eggs at rotting material, trash, fruits, and human food. These microorganisms were developed and attract other females to perch and lay eggs (Liu et al. 2017).

In the present investigation, we study the contaminated culture on which D. melanogaster larvae were grown. Increasing the number of growing larvae was able to inhibit the growth of fungi, which grow in the contaminated media. From several piles of Petri overgrown with larvae, the pigmented bacterial colony showed a zone of inhibition against nearby molds. Thus, we proposed to investigate the biological prospect of red-pigmented bacteria cultured from contaminated agar media.

MATERIALS AND METHODS

Bacteria culture

All contaminated culture media in Petri dishes were closed without sealing so that wild D. melanogaster larvae developed on them. The colony of red-pigmented microbes that inhibited fungal growth isolated and purified into ISP1, ISP2, Nutrient agar, Potato Dextrose Agar (PDA), Trypticase soy agar, or tryptone soya agar (TSA) media (HiMedia®). Microbial growth was observed and characterized according to Suryawanshi et al. (2014). The isolate was identified by morphological, physiological, biochemical (Abdullah et al. 2017), and molecular approaches. Molecular identification of isolate was tested using primers 27F-5’AGAGTATTTGATCCTGCTAG-3’ and 1492R-5’GGTTACCTTGTACGACTT3’ (Macrogen). PCR (25 μL final volume) was carried out by mixing 2.5 μL 10X PCR buffer, 0.5 μL mixtures of dNTP, 1.0 μL of each PCR
primer, 0.5 μL of Taq DNA polymerase, and 1.0 μL of each template DNA. Initial denaturation was carried out for 25 seconds at a temperature of 94 °C. Then it was reconstructed at 94 °C for 10 seconds for 35 cycles. Annealing was taken at 46 °C for 30 seconds. The elongation was continued at 72 °C for 1.5 minutes. Meanwhile, the last extension occurred at 72 °C for 10 minutes. The PCR results were further sequenced and the results compared with other bacterial sequences in GenBank.

Susceptibility test
A susceptibility test toward antibiotic was conducted using the disk diffusion method. The antibiotics used consisted of nystatin, chloramphenicol, clindamycin, ampicillin, streptomycin, griseofulvin, fluconazole. The dose tested was 10 and 20 μg/L (Bengtsson-Palme and Larsson 2016).

Hemolytic test
The hemolysis ability of the isolates was tested using the blood agar method. It was incubated for 1–2 days at room temperature. The activity of the tested bacteria was observed according to hemolysis type (Arifiyanto et al. 2020).

Hydrolase enzyme test
The hydrolase enzymes consisting of lipase, chitinase, mannanase, cellulase, amylase, protease, were tested on the test isolate. Test conducted in qualitative method and measured for enzyme index to compare. Isolate Streptomyces AB8 was used as a control (Agustina et al. 2019; Arifiyanto et al. 2020; Garcia-Orozco et al. 2019; Rosa et al. 2020; Sumardi et al. 2020).

Fermentation media
The fermentation media was optimized by culturing isolate into certain fermentation media (TSA, TSA+yeast, Nutrient broth, Mueller Hilton Broth, ISP4, ISP9, Trypton water, Tauge extract media, Sucrose solution, rice washing water, yeast-starch broth, and gelatine). Culture with good color formation was selected (Chambers et al. 2019; Priyatno et al. 2011; Trudgeon et al. 2020). The starter was made by adding 1 mL of isolate (cell turbidity adjusted to 0.5 McFarland standard) into 9 mL of fermentation medium at room temperature (180 rpm) and incubating for 3 days. The 3-day old starter was put into 990 mL of fermentation medium and incubated for 6 days. Growth curves were made by taking 1 mL of culture and measured at a wavelength of 600 nm twice a day. The cultures were harvested and centrifuged at 3000 rpm for 30 minutes. The filtrate obtained was an extracellular pigment and pellets were extracted using methanol to obtain intracellular pigments. The filtrate was extracted with 70% methanol (1:1) added and left for 12 hours to obtain pure pigment (Asnani et al. 2016). In this study, the extracellular pigment was obtained.

Antioxidant test
Antioxidant activity was measured using the DPPH method (Castro et al. 2019). The materials used were DPPH (2.2 diphenyl-1-pircylylhidrazyl) (Sigma). 4 mg of DPPH was dissolved in 25 mL of methanol in a volumetric flask as a stock solution (equivalent to a concentration of 160 mg / L). The resulting solution was stored in a dark room and protected with aluminum foil. Stock solutions were diluted with concentrations of 20, 40, 60, 80, and 100 mg / L. The absorbance was measured using a spectrophotometer at 515 nm. A total of 20 μL of samples were mixed with 180 μL of DPPH (concentration of 150 μM) in dark conditions and allowed to stand for 30 minutes then measured the absorbance at λ 515 nm. The blank solution used consisted of 20 μL methanol in 180 μL DPPH measured at the same wavelength (Ab). Trolox was used as a positive control. The DPPH test treatment was carried out in three repetitions (triplicate). During the reduction process by antioxidants, the DPPH radical solution changed color from purple to pale yellow. Sample measurement can be seen in equation 1.

\[
\text{% inhibition} = \left[1 - \frac{A_s - A_b}{A_b}\right] \times 100 \quad (1)
\]

Where:
As: absorbance sample
Ab: absorbance control

Antimicrobial activity test
The pathogen Aspergillus niger, Candida sp., Escherichia coli, Fusarium sp., Rigidoporous sp. Dickeya sp. var 5., Dickeya sp. var 10., Trichoderma sp., were prepared as much as 100 μL with a concentration of 10⁸ CFU/mL to be grown on Mueller Hinton Agar (MHA) media. The inhibition test was carried out using the paper disc method. 10 μL extract was dropped on 6mm filter paper. The same treatment was carried out on methanol which was used as a control. The cultures were incubated for 24 hours at room temperature and the diameter of the inhibition zone was measured (Arifiyanto et al. 2020).

RESULTS AND DISCUSSION

Identification
Based on the identification result (Tabel 1) strain MBC1 had certain characters. The cells were rod-shaped, motile, and Gram-negative bacteria. It showed an oxidase negative and catalase-positive test.

Culturing in various media was conducted to figure out appropriate media for fermentation. MBC1 were produced red pigment in TSA, MHA, tryptone water, gelatine, tauge extract media (TEM). The result of Gram staining was suggested that the MBC1 isolate belongs to the genus Serratia, molecular identification using 16S rRNA obtained a similar result (Figure 1).
pattern of the MBC1 strain of bacteria tends to be stable at no more than 0.3 OD at the measured cell density using a spectrophotometer. On the other hand, the AB8 strain bacteria showed a peak increase on day 7th reached 0.8 OD. These data indicate that the period required for the two strains of bacteria for optimum growth was different. The MBC1 strain bacteria grew optimally on the 3rd day while AB8 required a week to grow. This data used in the fermentation process to obtain pigment metabolite.

### The potency of cells activity

**Hydrolase enzyme**

Based on the screening of the enzyme activity index (Figure 3), the MBC 1 strain was found superior to AB8 (Streptomyces control) in the enzyme activity of lipase and amylase. On the other hand, AB8 strains showed only slightly superior results in chitinase, cellulase, and protease enzymes.

The results were compared with the antioxidant assay by the influence of heavy metals. Trolox as control was detected to reach a lower peak than metal treatment. Heavy metals have been shown to increase the absorbance and antioxidant ability of the pigment *Serratia marcescens* strain MBC1 (figure 4). On the other hand, a greater heavy metal dose was given as it reduced the antioxidants activity.

### Antimicrobial activity

In addition to the antioxidant activity, efforts were also made to identify the antimicrobial potential of the MBC1 and AB8 strains. The results of antimicrobial activity are presented in figure 6. Bacteria strain AB8 was able to inhibit the growth of Dickeya sp. var 5 and var10 and *Aspergillus niger* only. On the other hand, the MBC1 strain was able to inhibit several pathogens, namely *Aspergillus niger*, *Candida* sp., *Fusarium* sp. and *Rigidoporus* sp.

### Discussion

Pigments or dyes are commonly encountered in higher organisms such as animals, plants, and even molds. However, it was not uncommon also to be found in microorganisms such as bacteria. The pigmented bacteria groups include *Agrobacterium aurantiacum* (Pink-red), *Staphylococcus aureus* (golden yellow), *Chromobacterium violaceum* (purple), *Serratia marcescens* (red), *Bacillus* spp. (creamy white), *Flavobacterium* sp (yellow). Even pigments were keys to identifying species (Abdulkadir 2017). Pigments are used in industry to make a wide variety of colors, some of which are water-soluble. Pigments produced by many microorganisms are non-toxic, so that they are environmentally friendly and can be used as dyes, foodstuffs, pharmaceuticals, cosmetics, and other industrial needs. Bacterial pigments show antioxidant, and antimicrobial activity (Artifiano et al. 2020). They can also be used as additives, color enhancers, and anticancer (Azman et al. 2018). *Micrococcus roseus* and *Rhodotorula glutinis* were produced red pigments beside the member of genera *Serratia* sp (Rostami et al. 2016).

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**Table 1. Morphology and biochemical characters of isolate MBC1**

| Test                  | Result     |
|-----------------------|------------|
| Gram/ KOH             | +/-        |
| Indole                | +          |
| Aerobic fermentation  | +          |
| Lipase                | +          |
| Oxidase               | -          |
| Cellulases            | +          |
| Protease              | +          |
| Arginine              | +          |
| Casein                | +          |
| Amylase               | +          |
| Chitinase             | +          |
| Mannanase             | +          |
| Catalase              | +          |
| Motility              | +          |

**Antibiotics**

- Nystatin -
- Chloramphenicol -
- Clindamycin -
- Ampicillin -
- Streptomycin -
- Griseofulvin -
- Fluconazole -

**Metals**

- MgSO₄ +
- AgNO₃ +
- CuSO₄ +
- BaCl₂ +
- HgCl₂ +
- FeSO₄ +
- PbNO₃ +

**C sources**

- Fructose +
- Galactose +
- Glucose +
- Glycerol +
- Lactose +
- Maltose +
- Mannitol +
- Starch +
- Sucrose +
- Cellulose +

**Media**

- TSA Red
- TSA + yeast No color
- NA No color
- MHA Red
- ISP 4 No color
- ISP 9 No color
- Tryptone Water Red
- Gelatine Red
- TEM Red
- Sucrose solution No color
- Rice washing water No color
- YSB No color
- Blood agar γ
- NA+ urea Red

Bacterial cell growth was observed for 10 days density measurements carried out in the morning and evening (Figure 2). The data in Figure 2 performed that the growth...
Phylogenetic tree result indicated that the MBC1 strain was closely related to *Serratia ureilytica*. *Serratia ureilytica* was a sub-species of *Serratia marcescens* that used urea as a source of nitrogen for its growth (Bhadra et al. 2005). *Serratia* was Gram-negative bacteria, a member of the γ-Proteobacteria subclass and grouped by the *Enterobacteriaceae* family (Bhadra et al. 2005). Several genera of the genus, namely species belonging to this genus. *Serratia marcescens*, *Serratia liquefaciens*, *Serratia plymuthica*, *Serratia rubidea*, *Serratia fonticola*, *Serratia marmorubra*, *Serratia proteamaculans*, and *Serratia odorifera* were reported as pathogens by (Horinouchi et al. 2010). *Serratia* has a wide range of habitats. They had frequently been isolated from soil, water, and animals (García-Fraile et al. 2015). They are also often found in association with plants and insects. *Serratia marcescens AL2-16* that associated with *Achyranthes aspera* L was reported to enhance plant growth-promoting properties (Devi et al. 2016). *Serratia ureilytica* species were found from extreme environmental habitats (Filippidou et al. 2019). *Drosophila melanogaster* had known to be a source of microbial association host, *Serratia* was included (Chambers et al. 2019; Gilbert et al. 2020). Prodigiosin was a red pigment which was the characteristic of this genus (Azman et al. 2018).

![Phylogenetic tree](image1)

**Figure 1.** Phylogenetic tree of the *Serratia marcescens* MBC1 strain with the access code MT797950

![Growth curves](image2)

**Figure 2.** The growth curves of AB8 and MBC1
Figure 3. The enzyme index of the bacteria strains AB8 and MBC1

Figure 4. The effect of lead and mercury on the cell density of the pigment of *Serratia marcescens* strain MBC1. Linear equation for Pb (y = -0.219x + 3.98) and Hg (y = -0.3626x + 3.0435) to determine lethal concentration (LC50).

Figure 5. The antioxidant activity of the pigment *Serratia* sp on lead and Mercury.
Antioxidant activity was induced by the presence of heavy metals (Gjorgieva et al. 2013). The result showed that antioxidants were higher in the stress treatment of Hg than Pb. The reason is that Mercury was more toxic rather than lead (Arifiyanto et al. 2017). According to Tristantini et al. (2018), a compound that has IC50 below 50, might be categorized as a strong antioxidant. The linear equation in Figure 5 shows that the LC 50 is less than 50. Using at 10 mg/L concentration DPPH and ABTS were completely hunted by prodigiosin (Arivizhivendhan et al. 2018). The pigment was a higher number on Hg compared to Pb. This means that bacteria need more antioxidants that come from pigments. Both Serratia marcescens strain MBC1 and Streptomyces AB8 showed a decrease antioxidant activity when metal concentration was higher than 60 ppm. Gjorgieva et al. (2013) explained that high metal levels treatment also caused DNA damage.

Serratia marcescens strain MBC1 has the potential to act as a biocontrol agent. In addition to being antioxidant, Arivizhivendhan et al. (2018) reported that the prodigiosin pigment was also able to inhibit the growth of pathogens including S. aureus, E. coli, B. cereus, C. botulinum, V. vulnificus, and S. enteritidis.

The result of the susceptibility test exhibited that, Serratia sp strain MBC1 was not resistant to nystatin, chloramphenicol, clindamycin, ampicillin, streptomycin, griseofulvin, and even flucanazole. Hemolysis results also showed the type of gamma (no hemolysis). The γ-Hemolysin group was related to certain subunit combinations of the protein. One of the proteins regarding γ-Hemolysin or γ-toxin was beneficial for microbes to form colonization, while there was no information about pathogenicity. Thus γ-Hemolysin was not related to pathogens (Oliveira et al. 2018).

Lipolytic activity was interesting because it showed a significant index. Extracellular-lipolytic activities arose as a strategy for bacteria to survive in extreme environments (Cherif et al. 2011). Previously it has been proved that lipase activity is one of the screening steps in biosurfactant activity (Arifiyanto et al. 2020). Biosurfactants were defined as compounds that have much more efficient and stable, better in toxicity, biocompatibility, and biodegradability than chemical surfactants. The compound also able to obtained from waste fermentation using cassava flour wastewater according to (Aratijo et al. 2019). It was supporting our data. Besides lipolytic activity, Serratia marcescens strain MBC1 has shown good amylase enzyme performance. It was also found by (Purkayastha et al. 2018) that the genus Serratia capable of producing chitinase, mannanase, and cellulase. It should be suitable if utilized in agricultural and marine bioremediation applications.

Consumption of various types of foodstuffs and higher emission rates also contribute to high free radicals. Free radicals are also known as reactive oxygen species (ROS). Antioxidants act to provide a protective effect on human health from oxidative damage caused by ROS (Tan et al. 2018). ROS can be derived from hydroxyl groups, superoxides, and peroxy radicals that attack macromolecules including membrane lipids, proteins, and DNA (Bray 2000) which then cause serious health complications such as cancer (Farinati et al. 2010; Notarte et al. 2019) diabetes mellitus (Maritim et al. 2003) and neurodegenerative (Sun and Chen 1998) and chronic inflammatory diseases (Federico et al. 2007).

The characterization results in the red color pigment of the MBC1 strain was Serratia sp. The lipolytic and amylolytic activity could be developed to recycle agriculture waste and biosurfactant production. Their pigment was to be developed as protection of oxidants. This strain also showed the ability to inhibit some pathogens, so it has been suggested as biocontrol for the pathogen of crops.

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Figure 6. Antimicrobial activity of the MBC1 and AB8 1) Escherichia coli, 2) Candida sp., 3) Fusarium sp., 4) Aspergillus niger, 5) Rigidoporous sp. 6) Dickeya sp. var 5, 7) Dickeya sp. var 10, and 8) Trichoderma sp.
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