Research articles

Exploration of indigenous soil bacteria producing-exopolysaccharides for stabilizing of aggregates land potential as biofertilizer

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Abstract: Steady soil aggregation is important for agricultural land which is formed by the micro-aggregate to become a macro-aggregate. This formation is mediated by organic material and various kinds of macro-organisms such as fungi, worms, ants and insects. An organic agency involved in soil aggregation stability is exopolysaccharide (EPS) derived from bacterial, fungal mycelium, and products synthesized by plants. However, the use of EPS-producing microorganisms as a biofertilizer has not been reported. This study was aimed to explore indigenous EPS-producing bacteria to solidify soil aggregation potential for biofertilizer. Bacterial strains were isolated from soils of three regions at Malang East-Java; two areas of green bean plantation in Kendal Payak and Jambe Gede, as well as forest land. Soil samples were derived from forest had has a total bacteria population of 9.3 x 10¹¹ CFU/mL. While soil samples from area Kendal Payak and Jambe Gede had total bacteria population of 1.5 x 10⁹ CFU/mL and 2.4 x 10⁹ CFU/mL, respectively. We selected three bacteria that could potentially produce abundant slime, namely as SPE-2, SPE-10 and SPE-20. The three selected bacteria are potential for biofertilizer because of their abundant slime, no antagonism and no symptoms as pathogen.

Keywords: bacteria, biofertilizer, exopolysaccharide, indigenous

Introduction

Soil fertility is the most important thing in agriculture because as a growing medium that is commonly used. Factors affecting soil fertility are texture, climate, organisms, altitude and water availability of the soil. Soil fertility is determined by physical, chemical and biological properties of the soil. The physical properties of the soil include effective depth, structure, soil texture, aggregation, field capacity of water, drainage, topography, humidity, climate and soil aeration.

Meanwhile, steady soil aggregates are very important for agriculture. The structure of soil with a steady aggregate will provide a positive influence on the growth of plant roots, water availability, and the movement of air and also able to enhance the resistance to disintegration when disruptive forces associated with tillage and water or wind erosion (Bronick and Lal, 2005). Soil aggregate is a group of sand, silt, clay and organic particles such as microbial cells that clot due to gum (adhesive), polysaccharides or other metabolites secreted by microbes (Hattori, 1998; Bertin et al., 2003). The formation of micro-aggregate becomes macro-aggregates is mediated by organic materials and various kinds of macro-organisms such as bacteria, fungi, worms, ants and insects. Organic agency that can increase soil aggregate stability is exopolysaccharide (EPS) derived from bacteria, fungi mycelium and plant products synthesis. The formation of soil aggregates is generally influenced by EPS which is the result of the activity of microorganisms (Lynch and Bragg, 1985; Goenadi, 1995).

Some exopolysaccharide-producing bacteria that have been reported include Pseudomonas aeruginosa, Erwinia, Ralstonia and Azotobacter vinelandii. Exopolysaccharides protect the bacteria from a variety of environmental stresses (Iqbal et al., 2002), protects cells from antimicrobial compounds, antibodies, and bacteriophages, or for sticking to other bacteria.
animal and plant tissues (Wingender et al., 1999). Indigenous bacteria are bacteria that naturally live freely in nature and known to have high capability to survive in their habitat. Some researches utilizing indigenous bacteria have been reported, for example as a bioremediation agent (Arfarita et al., 2016; Nuraini et al., 2015). However, the use of indigenous microorganisms producing EPS as a biofertilizer, especially in Indonesia has not been reported. The study was aimed to obtain indigenous bacteria in area of Malang-East Java, that are potential in producing exopolysaccharide (EPS).

**Materials and Methods**

The initial stage of exploration was to process a screening and isolation of the target bacteria around the rhizosphere of green bean plants in the three regions. Three bacteria had been selected and then observed its potential as a bacterial consortium that will be used as biofertilizer agent. Its potential was also supported by antagonism and pathogenicity test on germination of green beans.

**Soil samples**

Soil samples were taken in April 2016 from the rhizosphere of mung bean plants (*Vigna radiata* L.) in the area of Kendalpayak (SK) and Jambegede (SJ), as well as forest area in Junrejo (FS) Malang, East Java. Soil samples were taken by pulling out the plant along with its roots carefully. Plant shoot was cut and then the roots along the soil were added to plastic bags. They were kept in a cooling box and then taken to the laboratory or stored at 4°C to isolate the bacteria immediately.

**Total population of soil bacterial**

Ten grams of soil samples were suspended in physiological saline solution (0.85%) and then made serial dilution to 10⁻⁸. Calculation of the total population of soil bacteria was made by the Standard Plate Count (SPC) method. Certain dilution series of soil samples were made by taking 10 μL suspension and spreading to PCA 9Palte Count Agar) medium on a petridish. Petridishes were then incubated for 2 days at room temperature.

**Isolation and selection bacteria producing-EPS**

The initial selection of bacteria potential of producing exopolysaccharide was performed by a screening process. Soil samples at specific dilutions of suspensions in a solid ATCC no. 14 medium were processed using spread plate method. Media ATCC No.14 per liter medium was prepared as follows: 0.2 g KH₂PO₄; 0.8 g K₂HPO₄; 0.2 g MgSO₄·7H₂O; 0.1 g CaSO₄·2H₂O; 2.0 mg FeCl₃; Na₂MoO₄. 2H₂O (trace); 0.5 g Yeast Ektstrac; 20 g sucrose) and 15 g bakto agar with pH 7.2. Bacterial colonies appeared on selective medium were isolated based on differences of colony morphology for further study (Emtiazi et al., 2004). In the early stages of this exploration, three bacterial strains that produced abundant slime were selected based on morphological differences.

Morphological characteristics were observed microscopically and macroscopically. At the macroscopic observation, the colony was observed the colours, shape of surface, edge, elevation, and the structure of colony. Microscopic was observed in gram staining properties and cell morphology.

**Antagonism test**

Antagonism test was performed by streaking selected bacteria on NA (Nutrient Agar) medium in the same petri dishes and incubated for 24 hours at room temperature. The test results of association among consortium isolates were to show negative results if it did not form an inhibition zone when cultured simultaneously. Thus isolates used in this consortium were not antagonistic that could be cultured and used in a formulation of biofertilizer.

**Phatogenicity test**

The pathogenicity test was performed using green bean sprouts as the test plants. Yosida medium was used for sprouts growth. Media stock was prepared by mixing 1000 mL solution consisting of 80 g NH₄NO₃, 40.3 g K₂SO₄, 71.4 g CaCl₂, 88.6 g MgSO₄, 32.4 g MnCl₂, 1.5 g (NH₄)₆Mo₇O₂₄, 0.074 g H₂BO₃, 0.93 g ZnSO₄, 0.035 g CuSO₄, 0.031 g FeCl₃, 7.7 g. The solution for planting was prepared by mixing 30 mL of stock solution with 4 liters of distilled water and heated the solution to a mixed material, divided into reaction tubes and sterilized.

Green bean sprouts (1-2 pieces) were inserted aseptically into tubes containing cotton and sterile Yosida medium and then inoculated 1 mL of pure cultures of bacteria into the tube (repeated 6-7 tubes). All of tube was incubated at the place with indirect sunlight. Observations were made every day for 7-10 days. Phatogenicity test was to observe the presence of necrosis, lesions or abnormalities of sprouts growth comparing with control. The data of sprouts growth were analyzed using analysis of variance.
(F test) with significant value (p<0.05) and if there were significant effects they were then followed by LSD test (p<0.05).

Results and Discussion

**Total population of soil bacteria**

Total population of bacteria on PCA medium (Table 1) shows the highest number of population is SF soil sample of $1.5 \times 10^{11}$ CFU/mL. This is due to the high content of organic matter in SF soil samples (Chemical soil analysis are presented in Table 2). These results are consistent with Suntoro (2003) that the organic matter content of soil affects biological properties that can increase the activity and microbiological populations in the soil, especially related to the activity of the decomposition of organic matter. As for SK and SJ soil samples have a total population of bacteria $2.4 \times 10^{9}$ CFU/mL and $9.3 \times 10^{9}$ CFU/mL lower than SF due to the organic matter content is lower than SF samples. SF soil samples are derived from forest area, where it has been known that forest soils are rich in organic material and have not been disturbed by chemical and human activities.

**Selected bacteria producing-EPS**

In this screening process had been obtained twenty-producing-exopolysaccharide isolates randomly. The presence of bacteria producing exopolysaccharide (EPS) on a selective medium were characterized by the ability of the growth and slimy colonies.

Exopolysaccharide-producing bacteria will grow well in a medium with a carbon source that is easily oxidized. ATCC medium which was used in screening process contained carbon sources used for optimization EPS bacterial growth medium consisting of sucrose, glucose, lactose, and 4-hydroxy-phenil acetic acid (4-HAA) and trace elements that were specific to the growth of EPS-producing bacteria. Name of exopolysaccharide (EPS) is the common name for all forms of bacterial polysaccharide that is found extracellular and one of the bioactive products produced by microorganisms (Flemming and Wingender, 2001). Exopolysaccharide is a high molecular weight polymer composed of long-chain sugar residues and secreted by the microorganisms into the surrounding environment. Bacterial exopolysaccharide is a complex mixture of electrolytes macro-polymer included polisakrida, proteins and nucleic acids, each of which is composed of variable molecular mass and structural properties (Kumon et al., 1994).

Of the 20 isolates of slimy bacteria were then selected three isolates based on the abundance of slime and morphological differences that significant for both macroscopic and microscopic morphology (Table 3).

| No | Soil Samples          | Dilution | The average of EPS bacteria population and other bacteria |
|----|-----------------------|----------|----------------------------------------------------------|
| 1  | FS (Jungejo Forest)   | $10^{-7}$| $1.5 \times 10^{11}$ CFU/mL                              |
| 2  | SK (Kendal Payak)     |          | $2.4 \times 10^{9}$ CFU/mL                               |
| 3  | SJ (Jambegede)        |          | $9.3 \times 10^{9}$ CFU/mL                               |

| Code | C- Organic (%) | N-Total (%) | C/N | Organic Material (%) | P. Bray I (mg/kg) |
|------|----------------|-------------|-----|----------------------|-------------------|
| Sample FS | 0.19          | 7           | 2.31| 4.67                 |
| Sample SJ  | 1.12          | 10          | 1.94| 10.56                |
| Sample SK  | 1.31          | 9           | 2.26| 27.44                |
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Table 3. Microscopic observation of 20 isolates of EPS bacteria present on the ATCC Media, isolated randomly based on differences of morphological colonies.

| No | Isolate Code | Gram Staining | Morphology           | Slime Formation |
|----|--------------|---------------|----------------------|-----------------|
| 1  | SPE 1        | -             | Staphylococcus       | Slimy           |
| 2  | SPE 2        | -             | Monobasid            | Very Slimy      |
| 3  | SPE 3        | -             | Coccus-Basil         | Very Slimy      |
| 4  | SPE 4        | -             | Monobasid            | Very Slimy      |
| 5  | SPE 5        | -             | Staphylococcus       | Slimy           |
| 6  | SPE 6        | -             | Staphylococcus       | Slimy           |
| 7  | SPE 7        | -             | Palisade             | Rather Slimy    |
| 8  | SPE 8        | +             | Palisade             | Not Slimy       |
| 9  | SPE 9        | -             | Streptobasid         | Very Slimy      |
| 10 | SPE 10       | -             | Staphylococcus       | Very Slimy      |
| 11 | SPE 11       | -             | Palisade             | Rather Slimy    |
| 12 | SPE 12       | -             | Streptobasid         | Rather Slimy    |
| 13 | SPE 13       | -             | Coccus -Basil        | Rather Slimy    |
| 14 | SPE 14       | +             | Coccus -Basil        | Very Slimy      |
| 15 | SPE 15       | -             | Coccus -Basil        | Very Slimy      |
| 16 | SPE 16       | -             | Staphylococcus       | Very Slimy      |
| 17 | SPE 17       | -             | Coccus               | Rather Slimy    |
| 18 | SPE 18       | -             | Streptokokus         | Not Slimy       |
| 19 | SPE 19       | -             | Palisade             | Not Slimy       |
| 20 | SPE 20       | -             | Coccus               | Very Slimy      |

Antagonism activities

Figure 1(a) shows the result of antagonism test of three isolates (SPE2, SPE10, and SPE20), which were then cultured simultaneously on NA media. It shows negative antagonism because it did not form a zone of inhibition. Test antagonism in all selected three isolates were not antagonistic so that it can be cultured in fertilizer formulations as a potential bacterial consortium.

Phatogenicity activities

Based on the results of pathogenicity tests showed that three isolates of bacteria producing EPS that had been selected, namely SPE2, SPE10, SPE20 were not pathogenic. As shown in Figure 2(b), there is no symptoms of necrosis, lesions or abnormalities in growth compared to controls.

Observations on the growth of green bean sprouts were also performed by analysis of variance with a randomized block design (CRD) with 7 replicates. The results indicated that the length of the plant, total root length, and fresh weight showed a significant effect on the growth of green bean sprouts (Table 4). Average in the growth of green bean sprouts show that there is no inhibition on growth. With a growth rate greater than control, the possibility of selected
three bacteria were to produce growth hormone. In this study, exploration of bacterial EPS production potential as a biological fertilizer limited only to characterization of the antagonism test and pathogenicity test.

Biochemical analysis and identification of bacteria to the molecular level will be observed in the next study. These observations are to determine the bacterial species are really not a pathogen and to determine other potential of bacteria that had been selected in this study. Analysis of growth hormone is also necessary, because the growth in the average observation of green bean sprouts showed a higher growth than the control. As reported by Mu’minah et al. (2015), of 34 EPS bacterial isolates, 4 isolates produced high value of IAA.

Table 4. Average of the growth of green bean sprouts

| Treatments | Long Sprouts (cm) | Total length Roots (cm) | Fresh weights (g) |
|------------|------------------|-------------------------|------------------|
| Control    | 14.33 ab         | 37.42 c                 | 0.96 c           |
| SPE2       | 12.92 a          | 31.67 b                 | 0.84 a           |
| SPE10      | 17.5 b           | 23.29 b                 | 0.85 b           |
| SPE20      | 19.00 a          | 35.00 a                 | 0.84 a           |
| F. Table 5%| Significant      | Significant             | Significant      |
| LSD        | 4.06             | 9.65                    | 0.08             |

Conclusion

The total population of bacteria producing exopoly-saccharide of each soil sample was different. Soil sample from Forest had has a total bacteria population of 9.3 x 10\(^{11}\) CFU/mL, while soil samples from area Kendal Payak and Jambe Gede were much lower than that of forest land, with total population of 1.5 x 10\(^{9}\) CFU/mL and 2.4 x 10\(^{9}\) CFU/mL, respectively. Three potential bacteria that produced EPS had been selected, namely SPE2, SPE10, and SPE20. Antagonism tests on three selected isolates showed no inhibitory zone occurs simultaneously when cultured on NA. Pathogenicity test did not indicate the nature of the pathogen against green bean sprouts. All three isolates are suspected growth hormone-producing, as germination was observed in green bean sprouts. Biochemical analysis and identification of bacteria to the molecular level need to be performed to determine the bacterial species are really not a pathogen and to determine the other potential of bacteria that had been selected in this study. Test aggregation on the ground also needs to be done in the laboratory and field conditions.

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