

Population Distribution of Phosphate-solubilizing Microorganisms in Agricultural Soil

IRNANDA AIKO FIFI DJUUNA^{1*}, SARASWATI PRABAWARDANI², and MARIA MASSORA³

¹Department of Soil and Land Resources, Faculty of Agriculture The University of Papua, Gunung Salju St. Kampus UNIPA Amban Manokwari 98314 Papua Barat-Indonesia; ²Department of Agronomy Faculty of Agriculture The University of Papua, Gunung Salju St. Kampus UNIPA Amban Manokwari 98314 Papua Barat-Indonesia; and ³Department of Biology, Faculty of Mathematics and Sciences The University of Papua, Gunung Salju St. Kampus UNIPA Amban Manokwari 98314 Papua Barat-Indonesia

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Phosphorus (P) is an essential macronutrient for plant growth and is mainly present in agricultural soil in unavailable forms. Phosphate-solubilizing microorganisms (PSMs) increase soil P availability. The objective of the present study was to assess the population and type of PSMs and their relationships with soil characteristics in the agricultural soil of Manokwari. Twenty-one composite soil samples (0–20 cm) were collected at the rhizospheres of plants in the Prafi and Masni Districts. A dilution technique and plate count method on Pikovskayas agar medium were used to examine the PSM population, phosphate-solubilizing index (PSI), and various soil properties. The results obtained showed that the total population of phosphate-solubilizing bacteria ranged between 25×10³ and 550×10³ CFU g⁻¹ of soil, while that of phosphate-solubilizing fungi was between 2.0×10^3 and 5.0×10^3 CFU g⁻¹ of soil at all locations. The PSI of the isolates ranged between 1.1 to 3.6 mm, with the most efficient and highest PSI being obtained for Bacillus sp. (strain 8) and the lowest for Pseudomonas sp. (strain 15). Six isolates found at all locations were identified at the genus level: Chromobacterium sp., Pseudomonas sp., Bacillus sp., Micrococcus sp., Caulobacter sp., and Aspergillus sp. A correlation was observed between the number of PSMs and the level of soil P availability and moisture content, indicating an increase in soil P availability with a greater abundance of PSMs in soil.

Key words: biofertilizers, soil microorganisms, phosphorus, Manokwari-Papua Barat

Soil microorganisms play an important role in the decomposition of organic matter and the transformation of soil nutrients used for plant growth and development. They are also crucially involved in soil P dynamics and P availability for plants (Richardson, 2001) as well as in pollutant bioremediation and the maintenance of soil productivity (Dong et al., 2014; Luo et al., 2015). Many agricultural soils represent a phosphate sink in which this element is not readily available to plants, but may still be recovered. Groups of phosphate-solubilizing microorganisms (PSMs) are currently the most common among soil microorganisms because they are used as biofertilizers and as one of the alternatives to increase the efficiency of phosphate fertilizers in order to overcome low phosphate availability in soil (Alori et al., 2017; Kalayu, 2019; Nosheen et al., 2021; Tian et al., 2021). PSMs may dissolve unavailable phosphate, thereby increasing soil P availability. Since it is easily absorbed by plants, P may enhance crop yields if its previous level was a limiting factor. Furthermore, PSMs include different groups of microorganisms, which not only assimilate phosphorus from insoluble forms of phosphates, but also release a large portion of soluble phos-

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phates at quantities that are in excess of requirements. PSMs have been shown to increase the bioavailability of soil insoluble phosphorus for plant use (Zhu et al., 2011). This group of bacteria, fungi, and actinomycetes is normally found in agricultural soil, with larger populations of phosphate-solubilizing bacteria (PSB) being found in agricultural and rangeland soils (Yahya and Azawi, 1998). Among the phosphate-solubilizing fungi (PSF) commonly found in agricultural soil, such as Penicillium sp., Mucor sp. and Aspergillus sp. increased plant growth by 5–20% after inoculation (Gunes et al., 2009). PSB have been shown to enhance the solubilization of insoluble P compounds through the release of low-molecular-weight organic acids and phosphatases (Rossolini et al., 1998; Sahu and Jana, 2000). Zhang et al. (2020) reported that P availability increased with the amount of PSB in solubilizing organic P (or PSBop), while the number of PSF increased with the content of soil organic carbo (SOC) and produced organic acids by dissolving insoluble phosphate with a decrease in pH, the chelation of cations, and competition with phosphate on sorption sites in soil (Mardad et al., 2013; Anand et al., 2016). Many PSB are effective biofertilizers or biocontrol agents and are regarded as broad spectrum biofertilizers (Gupta, 2004). Due to the negative environmental impact of chemical fertilizers and the increasing cost of energy, the utilization of PSB is advantageous for sustainable agricultural practices (Gyaneshwar et al., 2002). The use of PSMs as biofertilizers may also overcome the excess of phosphate in acidic soil; therefore, they are expected to become an

^{*} Corresponding author. E-mail: i.djuuna@unipa.ac.id; Tel: +62-986-211899; Fax: +62-986-211455.

alternative that reduces dependence on the excessive use of phosphate fertilizers. The population of PSMs in soil ranges between 10⁴ and 10⁶ g⁻¹ of soil and they are mostly located in the rhizosphere. Among these microorganisms, bacteria are more effective at phosphorus solubilization than fungi (Alam et al., 2002). Previous studies reported that among the whole microbial population in soil, PSB were responsible for between 1 and 50% of the P solubilization potential, whereas PSF only accounted for between 0.1 and 0.5% (Chen et al., 2006). The population density of PSB ranged between 8×10^5 and 5.33×10^9 in the different rhizospheres of vegetable fields (Alia *et al.*, 2013), between 5.0×10^3 and 7.5×106 CFU g⁻¹ of soil in Wamena soil (Suliasih and Widawati, 2005), and between 0 and 107 cells g⁻¹ soil in the soil of North Iran, with PSB accounting for 3.98% of the total population of bacteria (Fallah, A. et al., 2006. Abundance and distribution of phosphate solubilizing bacteria and fungi in some soil samples from north of Iran. 18th World Congress of Soil Science. Philadelphia, Pennsylvania, USA July 9-15). However, their number is not sufficiently high to compete with other microbial species in the rhizosphere (Jain et al., 2012). The population or proportions of these microbes vary between ecosystems due to the influence of complex biological factors. Many soil factors, such as the soil nutrient status, soil pH, moisture content, organic matter, and soil enzyme activities, also contribute to these variations. Therefore, the objectives of the present study were to (1) isolate and assess the number and distribution of PSMs in agricultural soil samples, particularly the rhizosphere of agricultural plants; (2) characterize the isolated microorganisms; and (3) assess their relationships with soil characteristics.

Materials and Methods

Study area

The Prafi and Masni Districts are located in the Southern Manokwari Region, Papua Barat in the Province of Indonesia. These areas have been developed as a Central Agricultural Area based on the Transmigration Program since 1979 and are classified as plain areas called Arfak Flakte (the Arfak Alluvial Plain). These two areas are fertile, and the major crops are paddy rice, legumes, horticultures, corn, and oil palm. The land use history of this area is mostly food crops (paddy rice and tuber crops) and vegetables as well as some plantation crops. Soil developed from alluvial soil that was dominated by coarse fractions on the surface and fine fractions in the subsoil. There are four soil types in these areas: Entisols, Histosols, Inceptisols, and Ultisols (Darmawanto, 1994).

Collection of soil samples

Soil samples were collected at a depth of 0–20 cm in the rhizosphere at sample points in several locations in the Prafi and Masni Districts of the Manokwari Region in October 2018. A composite soil sample was obtained by mixing 10 sub-samples collected at the rhizosphere of each plant with soil auger (diameter of 2.5 inches). Samples were placed into sterile containers and transported to the laboratory, where they were air-dried and crushed. A portion of each sample was sieved with a 2-mm sieve mesh to remove pebbles and large organic debris, while the remainder was unsieved for a soil biological analysis. Samples were kept in sterile plastics bags and stored in a refrigerator at 4°C for further biological analyses. The coordinates of each sample collection point were obtained using a global positioning system (GPS). Twenty-one samples were collected from the rhizospheres of agricultural plants, mainly soybean (*Glycine max merril*), paddy rice (*Oryza sativa*), long bean (*Vigna sinensis*), corn (*Zea mays*), squash (*Luffa acutangula*), cacao (*Theobrema cacao*), chili (*Capsicum anuum*), kangkong (*Ipomoea aquatic*), eggplant (*Solanum tuberosum*), cassava (*Manihot utilisina*), and peanut (*Arachis hypogea*).

Isolation and identification of PSMs

PSMs were isolated from soil samples collected from each rhizosphere using serial dilutions and the agar pour plate method. Ten grams of soil samples from all rhizosphere sampling points was dispersed in 90 mL of sterile distilled water and thoroughly shaken. A 1-mL aliquot was transferred using a sterile pipette to 9 mL of sterile distilled water in a test tube and stirred for 10 s to form a 10^{-2} dilution. Serial dilutions to 10^{-7} were then prepared using the same method. A 0.1-mL of aliquot from each serial dilution was transferred to a sterile plate and covered with Pikovskayas agar medium (50°C) containing insoluble tricalcium phosphate, followed by an incubation at 27-30°C for 7 days. The composition of the medium was 5 g Ca₃(PO₄), 0.5 g (NH₄)₂SO₄, 0.2 g NaCl, 0.1 g MgSO₄.7H₂O, 0.2 g KCl, 10 g glucose, 0.5 g yeast extract, 20 g agar, small amounts of MnSO₄ and FeSO₄, and 1,000 mL distilled water (Subba Rao, 1982). Colonies with clear halos (a sign of solubilization) were considered to be phosphatesolubilizing colonies (Subba Rao, 1977; Vyas et al., 2007). The number of viable cells was calculated using the following formula: number of cells mL⁻¹ (CFU g⁻¹)=(number of colonies)×(dilution factor) (Cappuccino and Sherman, 1992). Colonies surrounded by a halo zone were then transferred to Pikovskayas agar medium to maintain the purity of the culture for morphological, physiological, and biochemical analyses as well as microbial identification. PSM isolates was identified based on colony and cell morphologies as well as microscopic observations using Bergey's Manual of Systematic Bacteriology (Krieg and Holt, 1984).

Phosphate-solubilizing index (PSI)

Halozone and colony diameters were successively measured during the incubation period to assess the PSI of PSMs. The PSI is the ratio of the total diameter, *i.e.* the clearance zone, including bacterial growth, and the diameter of the colony.

Soil analysis

Composite soil samples from all locations were air-dried, crushed, and passed through a 2-mm sieve and then analyzed for soil properties, such as pH related to soil-water 1:5 (w/v) (H₂O 1:5), available phosphorus (mg kg⁻¹) (Olsen, spectrophotometer), C-organic (g kg⁻¹) (Walkley and Black, 1934, spectrophotometer), and total nitrogen (g kg⁻¹) (Kjeldahl Method, spectrophotometer), while the soil moisture content (%) (gravimetric method) was measured using a fresh soil sample. Data on the population of PSMs and PSI were analyzed using the *t*-test. The relationships between PSMs and soil characteristics were examined using Pearson's correlation (SPSS).

Results

Population of PSMs

A significant difference was observed in the population of PSMs between plant rhizospheres in agricultural soil (Table 1). The total population of PSB in soil samples from the Prafi and Masni Districts ranged between 25×10^3 and 55×10^4 CFU g⁻¹ soil, while that of PSF ranged between 2.0×10^3 and 5.0×10^3 CFU g⁻¹ soil. The highest number of PSB was found in the rhizosphere of cacao (*Theobrema cacao*), followed by chili (*Capsicum anuum*) and corn (*Zea*

Sampling	Coord	linate	(Crops	Mean Popula (CFU g ⁻¹	tion of PSM soil)×10 ³
Sile	North (X)	East (Y)	Common Name	Botanical Name	Bacteria	Fungi
PRAFI						
Prafi 1	360157	9911265	Chili	Capsicum anuum	540	3.0
Prafi 1	359535	9911392	Kangkung	Kangkung Ipomoea aquatica		2.0
Prafi 1	358187	9911891	Paddy Rice	Oryza sativa	500	2.0
Prafi 1	363571	9906585	Squash	Luffa acutangula	25	5.0
Prafi 2	363700	9905520	Soybean	Glycine max	34	3.0
Prafi 2	364130	9904922	Eggplant	Solanum tuberosum	47	2.0
Prafi 2	364469	9904450	Cassava	Cassava Manihot utilisina		2.0
Prafi 3	368901	9902105	Corn	Zea mays	420	5.0
Prafi 3	368579	9901765	Cacao	Theobrema cacao	51	3.0
Prafi 3	368537	9901041	Cassava	Manihot utilisina	43	2.0
Prafi 4	370247	9901060	Peanut	Arachis hypogea	59	2.0
Prafi 4	370715	9901748	Long beans	Vigna sinensis	370	5.0
Prafi 4	371477	9901584	Cacao	Theobrema cacao	550	3.0
MASNI						
Masni 1	375397	9898650	Cacao	Theobrema cacao	16	3.0
Masni 1	376286	9899340	Corn	Zea mays	520	3.0
Masni 2	377516	9897641	Cacao Theobrema cacao		100	2.0
Masni 2	376932	9897807	Corn Zea mays		19	2.0
Masni 2	376832	9897370	Kangkong	Ipomoea aquatica	390	5.0
Masni 2	376902	9897444	Paddy rice	Oryza sativa	480	2.0
Masni 3	388221	9890357	Paddy rice	Oryza sativa	350	5.0

 Table 1.
 Population of phosphate-solubilizing microorganisms (bacteria and fungi) in rhizospheres of agricultural crops in Prafi and Masni Districts, Manokwari

t-test: PSB: t=5.523 df=19 Sign. (2 tailed)=0.001 PSF: t=11.168 df=19 Sign. (2 tailed)=0.001

mays) in Masni District, while the lowest was detected in the rhizosphere of squash (*Luffa acutangula*) in Prafi District. The total population of PSF was low in most rhizosphere samples. In comparisons between Prafi and Masni Districts, the total populations of PSB and PSF were low in the former. Prafi District is known as the center of agricultural production in Manokwari, most of the lands are cultivated with agricultural crops, and, thus, the use of chemical fertilizers and pesticides is more intense in this area.

Identification and morphological and physiological characteristics of isolates

Colony shapes were mostly circular/round (31 isolates) and whitish in color (32 isolates), while cell shapes were mainly rods (coccobacillus) (32 isolates) that were Gram negative (24 isolates) or Gram positive (12 isolates), and the majority of isolates were motile (Table 2). Six genera were identified among PSMs (5 PSBs and 1 PSF): *Chromobacterium* sp. (3 strains), *Pseudomonas* sp. (17 strains), *Bacillus* sp. (8 strains), *Micrococcus* sp. (5 strains), *Caulobacter* sp. (3 strains), and *Aspergillus* sp. (1 strain). The representative PSM types found in the agricultural soils of Prafi and Masni Districts are shown in Fig. 1.

PSI

Approximately 58 PSM isolates were found in all sampling locations; however, only 37 PSM isolates, which considered as best PSM isolates, were identified and tested for their PSI (Table 3). The PSI of the isolates ranged between 1.1 and 3.6 mm, with the most efficient and highest PSI being obtained for *Bacillus* sp. (strain 8) and the lowest for *Pseudomonas* sp. (strain 15).

However, among the isolates of *Pseudomonas* sp., six showed higher PSI than the others.

In the present study, an increase in the halo zone was not associated with a larger colony diameter.

Soil characteristics

An analysis of soil properties at each location examined (Table 4) revealed that soil pH varied between 5.20 and 6.40 (acidic to slightly acidic) in Prafi District and between 5.30 and 6.90 (acidic to neutral) in Masni District. The total nitrogen content ranged between 1.07 g kg⁻¹ (low) and 2.95 g kg⁻¹ (medium) in Prafi District, while that at all sampling locations in Masni District was low (0.14–0.16 g kg⁻¹). The phosphorus content in all areas was low to medium $(6.00-9.90 \text{ mg kg}^{-1})$, while the carbon organic content was very low to low (0.10 to 0.28 g kg⁻¹) in Prafi District and low (1.20 to 1.91 g kg⁻¹) in Masni District. Prafi and Masni Districts are the main centers of agricultural crop production for vegetables and food crops in Manokwari region and Papua Barat Province; therefore, most of the agricultural land is highly dependent on chemical fertilizers and pesticides, the high and continuous use of which without the application of organic matter has affected the physical, chemical, and biological properties of soil.

Relationship between PSMs and soil properties

Correlations were observed between the number of PSMs and soil P availability and moisture content, indicating an increase in soil P availability with a greater abundance of PSMs (Table 5). However, no correlations were noted with other soil characteristics, such as soil pH, N-total, and Corganic.

Isolate	Halozone Diameter	Colony Diameter	Phosphate-solubilizing Index	Species/Strain
	(mm)	(mm)	(mm)	
1	23	12	1.9	Chromobacterium sp. (1)
2	19	10	1.9	Pseudomonas sp. (1)
3	17	10	1.7	Pseudomonas sp. (2)
4	17	8	2.1	Pseudomonas sp. (3)
5	14	8	1.7	Bacillus sp. (1)
6	15	8	1.9	Pseudomonas sp. (4)
7	16	8	2.0	Bacillus sp 2
8	12	7	1.7	Pseudomonas sp. (5)
9	12	11	1.1	Pseudomonas sp. (6)
10	10	7	1.4	Bacillus sp. (3)
11	13	8	1.6	Pseudomonas sp. (7)
12	13	6	2.2	Pseudomonas sp. (8)
13	10	6	1.7	Bacillus sp. (4)
14	11	7	1.6	Bacillus sp. (5)
15	12	6	2.0	Bacillus sp. (6)
16	15	12	1.2	Chromobacterium sp. (2)
17	15	11	1.4	Pseudomonas sp. (9)
18	13	7	1.8	Pseudomonas sp. (10)
19	14	6	2.3	Micrococcus sp. (1)
20	11	8	1.4	Chromobacterium sp. (3)
21	21	9	2.3	Pseudomonas sp. (11)
22	22	9	2.4	Pseudomonas sp. (12)
23	3	2	1.5	Pseudomonas sp. (13)
24	8	5	1.6	Bacillus sp. (7)
25	11	7	1.6	Micrococcus sp. (2)
26	18	5	3.6	Bacillus sp. (8)
27	15	10	1.5	Pseudomonas sp. (14)
28	13	8	1.6	Caulobacter sp. (1)
29	11	6	1.8	Caulobacter sp. (2)
30	14	11	1.3	Micrococcus sp. (3)
31	9	7	1.3	Micrococcus sp. (4)
32	13	11	1.1	Pseudomonas sp. (15)
33	17	7	2.4	Pseudomonas sp. (16)
34	11	7	1.8	Micrococcus sp. (5)
35	14	11	1.3	Caulobacter sp. (3)
36	13	6	2.2	Pseudomonas sp. (17)
37	10	7	1.4	Aspergillus sp. (1)

 Table 2.
 Phosphate-solubilizing index of 37 PSM isolates from rhizosphere samples collected in Prafi and Masni Districts

t-test for PSI: t=22.654 df=36 Sign. (2 tailed)=0.001 Numbers in parentheses show the strains of PSB

Discussion

PSM populations were abundant and varied in most of the agriculture soil samples collected from different plant and soil rhizospheres in Prafi and Masni Districts. This result is consistent with previous findings reported by Kucey (1983) and Baliah *et al.* (2016), showing that the population of PSMs varied within the soil rhizosphere and with soil characteristics. However, in the present study, PSM populations were smaller in Prafi and Masni agricultural soil samples than in other types of soil containing high organic matter and soil nutrients, in which the population ranged between 10^4 and 10^6 CFU g⁻¹. Although PSMs are found in all soils, their number depended on the soil climate as well as cropping history (Gupta *et al.*, 1986). The number of PSB was found to higher than that of PSF in Prafi and Masni Districts. Bacteria are more effective at phosphorus solubili-

zation than fungi (Alam et al., 2002). According to Chen et al. (2006), among the whole microbial population in soil, PSB are responsible for between 1 and 50% of the P solubilization potential and PSF for only 0.1 to 0.5%. Khan et al. (2009) reported that 1 g of fertile soil contained between 10¹ and 10¹⁰ bacteria and their live weight may exceed 2,000 kg ha⁻¹. Yahya and Azawi (1998) also showed that the abundance of PSB was generally large in agricultural and rangeland soils. PSB have been detected in the majority of soils (Chhonka and Taraedar, 1984), and their population is generally low in arid and semi-arid regions, which is attributed to the low level of organic matter and high temperature regime (Gupta et al., 1986). The PSB population was found to be higher in soils in mild and moist climates than those in dry climates (Subba Rao, 1982). Different plant species or genotypes are another factor that affected the number and activity of soil microorganisms in the present study. This is consistent with the findings reported by Katiyar and



b. Isolate 9 from the Paddy rice rhizosphere of Praf11: *Pseudomonas* sp. 6

Fig. 1. Isolates of phosphate-solubilizing microorganisms: (a) Isolate 10: *Bacillus* sp. 3 from Masni Districts in Manokwari. (b) Isolate 9: *Pseudomonas* sp. 6 from Prafi Districts in Manokwari.

Goel (2003) showing that the abundance of PSB in soil depended on the plant species, the microbial composition in soil, and soil conditions, in addition to the location of sampling (Kundu et al., 2009). Belimov et al. (2015) and Schreiter et al. (2014) demonstrated that the diversity of microorganisms in different plant species may be attributed to plant-microbe interactions being highly dependent on soil conditions and the plant genotype. Furthermore, Baliah et al. (2016) found that the population level of PSB varied in the rhizosphere soils of okra, chili, tomato, cotton, and eggplant. Ponmurugan and Gopi (2006) reported that the PSB population was the largest in the rhizosphere soil of groundnut and the lowest in the rhizosphere soils of ragi, sorghum, and maize, and suggested that this was due to high phosphatase activity in the rhizosphere. Another factor contributing to variations in the PSB population may be the development of microorganisms in soil that are strongly influenced by the metabolic activity of plant roots through root exudates. Kato et al. (1997) showed that metabolic activity and metabolite compounds released by plants through the roots had a marked impact on the soil microorganisms present in plant root areas; therefore, soil microorganism activity will increase in the rhizosphere. In addition, variations in the PSB population of different crops were attributed to soil factors, such as nutrients, pH, moisture content, organic matter, and some soil enzyme activities for each crop (Ponmurugan and Gopi, 2006).

However, another study reported no relationship between the number of PSB and the type of vegetation (Kucey, 1983) or sampled sites and soil management programs (Fernández *et al.*, 2015). Another factor affecting the population of PSMs in soil is soil properties, *i.e.* physical, chemical, and biological soil properties. In the present study, no correlations were observed between the population of PSMs and some of the soil characteristics analyzed (pH, total N, and C-organic). In contrast, a correlation was found between the number of PSB and total nitrogen and organic matter (Rfaki et al., 2017); differences in the organic carbon content in soil affected the distribution of PSB (Yadav and Singh, 1991); and a high content of soil organic matter increases productive soil microbes (John et al., 2001) including PSB. In addition, the intensive and excessive use of chemical fertilizers and pesticides in agricultural soils affects soil microorganisms and PSMs. Adul Rahman et al. (2021) reported that any external pressure, including chemicals, affects the chemistry and physics of soil and, thus, its living organisms. Among PSM types, the six PSM isolates identified in the present study have been widely reported for PSB in previous studies with isolation sources ranging from rhizosphere soils (Gupta et al., 2012; Singh and Prakash, 2012) to other common soils (Baliah et al., 2016). The genus Aspergillus was the most frequent PSF isolate reported (Gupta et al., 2007; Khan et al., 2010; Simfukwe and Tindwa, 2018).

Conclusions

The number and type of PSMs in the agricultural soils of Prafi and Masni Districts revealed that these areas are suitable habitats and these microorganisms may increase the P content in soil as well as its supply to crops. Correlations were observed between the number of PSMs and the level of soil P availability and moisture content, indicating an increase in soil P availability with a greater abundance of PSMs in soil.

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18 Wisish Gradie Red Neise -++++ - ++++ - -++++ -	8 Winks Creating Red Neglive + + + + + - + + + + - + + + + - + + + + + + + +	17	Whitish	Circular	Rod	Negative	 + + +	+ + +	+	+	+	I	I	+	I	+	I	I	+	+	+	+	I	+	I	Pseudomonas sp.
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		23	Whitish	Circular	Rod	Negative	 + + +	+++	+	+	+	1	I	+	I	+	ī	I	+	+	+	+	I	+	I	Pseudomonas sp. 13
		24	Whitish	Circular	Rod	Positive	+++++++	+++	+	+	+	+	I	+	I	+	I	+	+	+	I	I	I	+	+	Bacillus sp. 7
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21 Which Circler Rod Negative $+$ <th>$\begin{array}{c c c c c c c c c c c c c c c c c c c$</th> <th>26</th> <th>Whitish</th> <th>Circular</th> <th>Rod</th> <th>Positive</th> <td>+++++++++++++++++++++++++++++++++++++++</td> <td>+++</td> <td>+</td> <td>+</td> <td>+</td> <td>+</td> <td>I</td> <td>+</td> <td>I</td> <td>+</td> <td>ī</td> <td>+</td> <td>+</td> <td>+</td> <td>I</td> <td>I</td> <td>I</td> <td>+</td> <td>+</td> <td>Bacillus sp. 8</td>	$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	26	Whitish	Circular	Rod	Positive	+++++++++++++++++++++++++++++++++++++++	+++	+	+	+	+	I	+	I	+	ī	+	+	+	I	I	I	+	+	Bacillus sp. 8
38 Withish Circler Rod Negative $-+++$ $+++$ $++$	$ \begin{array}{c c c c c c c c c c c c c c c c c c c $	27	Whitish	Circular	Rod	Negative	 + + +	++++	+	+	+	I I	I	+	I	+	I	I	+	+	+	+	I	+	I	Pseudomonas sp. 14
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34 Which Irregular Rod Positive - + + + + + + + + Micrococcus sp.5 35 Transparent Circular Rod Negative - +	34 Whish Irregular Rod Positive - + <th>33</th> <th>Whitish</th> <th>Rhizoid</th> <th>Rod</th> <th>Negative</th> <td>+ + +</td> <td>+++</td> <td>+</td> <td>+</td> <td>+</td> <td>I I</td> <td>I</td> <td>+</td> <td>I</td> <td>+</td> <td>I</td> <td>I</td> <td>+</td> <td>+</td> <td>+</td> <td>+</td> <td>I</td> <td>+</td> <td>I</td> <td>Pseudomonas sp. 16</td>	33	Whitish	Rhizoid	Rod	Negative	+ + +	+++	+	+	+	I I	I	+	I	+	I	I	+	+	+	+	I	+	I	Pseudomonas sp. 16
35 Transparent Circlular Rod Negative - + <t< th=""><th>35 Transparent Circular Rod Negative - + <td< th=""><th>34</th><th>Whitish</th><th>Irregular</th><th>Rod</th><th>Positive</th><td> + </td><td>+++</td><td>I</td><td>+</td><td>+</td><td> +</td><td>I</td><td>I</td><td>I</td><td></td><td>I</td><td>+</td><td>+</td><td>I</td><td>+</td><td>I</td><td>I</td><td>I</td><td>+</td><td>Micrococcus sp. 5</td></td<></th></t<>	35 Transparent Circular Rod Negative - + <td< th=""><th>34</th><th>Whitish</th><th>Irregular</th><th>Rod</th><th>Positive</th><td> + </td><td>+++</td><td>I</td><td>+</td><td>+</td><td> +</td><td>I</td><td>I</td><td>I</td><td></td><td>I</td><td>+</td><td>+</td><td>I</td><td>+</td><td>I</td><td>I</td><td>I</td><td>+</td><td>Micrococcus sp. 5</td></td<>	34	Whitish	Irregular	Rod	Positive	 + 	+++	I	+	+	 +	I	I	I		I	+	+	I	+	I	I	I	+	Micrococcus sp. 5
36 Whitsh Circular Rod Negative + 5 Navorish Circular Circular Circular + 5 Navorish Circular	36 Whitsh Circular Rod Negative + <th>35</th> <th>Transparent</th> <th>Circular</th> <th>Rod</th> <th>Negative</th> <th> + </th> <th>+ + </th> <th>+</th> <th>+</th> <th>+</th> <th>+</th> <th>I</th> <th>I</th> <th>+</th> <th>+</th> <th>I</th> <th>I</th> <th>I</th> <th>+</th> <th>+</th> <th>+</th> <th>I</th> <th>+</th> <th>+</th> <th>Caulobacter sp. 3</th>	35	Transparent	Circular	Rod	Negative	 + 	+ +	+	+	+	+	I	I	+	+	I	I	I	+	+	+	I	+	+	Caulobacter sp. 3
37 Brownish Circular - + + + + + +	37 Bownisk Circular _ + + + + MR=Methyl Red; VP=Voges-Proskauer	36	Whitish	Circular	Rod	Negative	 + + +	+++	+	+	+	ı I	I	+	I	+	I	I	+	+	+	+	I	+	I	Pseudomonas sp. 17
	MR=Methyl Red; VP=Voges-Proskauer	37	Brownish	Circular			 + 	+ + 1																		Aspengillus sp. 1

		S	oil Characteri	stics	
Sampling site	pH H ₂ O (1:5)	Moisture Content (%)	N-Total (g kg ⁻¹)	Available P (mg kg ⁻¹)	C-Org (g kg ⁻¹)
PRAFI					
Prafi 1 (1)	6.3 SA	31.0	12 L	7.6 M	1.4 VL
Prafi 1 (2)	6.4 SA	24.0	12 L	6.6 L	1.3 VL
Prafi 2 (1)	6.3 SA	26.9	21.5 L	7.5 M	2.0 VL
Prafi 3 (1)	5.2 A	20.8	29.5 M	9.9 M	2.8 VL
Prafi 3 (2)	6.4 SA	28.3	10.7 L	6.3 L	1.0 VL
Prafi 4 (1)	5.8 SA	15.9	13.5 L	7.1 M	1.4 VL
Prafi 4 (2)	6.1 SA	20.7	24.7 M	8.4 M	2.5 VL
MASNI					
Masni 5 (1)	5.3 A	27.5	15 L	6.0 L	17.5 L
Masni 5 (2)	5.9 SA	17.8	16 L	7.1 M	19.1 L
Masni 6 (1)	6.1 SA	18.4	16 L	7.9 M	16.0 L
Masni 6 (2)	6.9 N	35.3	15 L	7.6 M	13.5 L
Masni 7 (1)	6.0 SA	32.6	14 L	6.8 L	12.0 L

Table 4. Soil characteristics at each sampling site of agricultural soil in Prafi and Masni Districts, Manokwari

VL=Very Low; L=Low; M=Medium; A=Acid; SA=Slightly Acidic; N=Neutral

Table 5. Correlation matrices between populations of PSMs and soil characteristics

Variables	Water Content	рН	C organic	N total	Р	PSB	PSF
Water Content	1						
pН	0.546**	1					
С	-0.371	0491	1				
N	-0.301	-0.523*	0.971**	1			
Р	-0.067	0.291	0.134	0.145	1		
PSB	0.694*	0.220	0.139	0.184	0.641**	1	
PSF	0.084	-0.100	0.676**	0.190	0.070	0.891**	1

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