

CHARACTERIZATION AND CLUSTERING OF SOME GUAVA GERmplasm COLLECTIONS BASED ON LEAF AND FRUIT CHARACTERS

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ABSTRACT

Guava has several different accessions. Guava diversity needs to be studied and evaluated in order to determine the next steps in the guava breeding. The objective of this research was to characterize and cluster some guava germplasm collections. The study was conducted at Aripan and Subang experimental farm, Indonesian Tropical Fruit Research Institute from January 2012 to December 2012. Five fruits of each accession were randomly selected, sampled, and then characterized using UPOV guidelines. Obtained data were analyzed by NTSYS ver.2.1. The Similarity level of 19 accessions ranging from 70 – 90% or the genetic distance was between 0-20%. Dendrogram obtained could be clustered into two different groups, namely group I (ARP9406, ARP9407, ARP8653, ARP8742, ARP10.2, JBT001, JBT002, ARP8740, JBT003andJBT004) and group II (ARP10.7, ARP10.6, ARP10.1, ARP10.12, ARP10.9, ARP10.11, ARP8744, ARP8741 and ARP8743). The result of this research can be used for guava breeding. Species diversity and genetic resources are very important to produce new varieties. This is expected to be highly valuable in the future.

Keywords: clustering, characterization, guava germ- plasm

INTRODUCTION

Guava (*Psidium guajava* L.) is an important tropical fruit crops. Guava has long been known to be used by the community for healing. The plant has been extensively studied in terms of

pharmacological activity of its major components, and the result indicated potent anti-diarrheal, antihypertensive, hepatoprotective, antioxidant, antimicrobial, hypoglycemic and anti-mutagenic activities (Joseph, 2011). Owen *et al.* (2008) reported that the consumption of guava (*Psidium guajava* L.) and noni (*Morinda citrifolia* L.) may protect betel quid-chewing Papua New Guineans against diabetes. Guava leaves also have anti-oxidant (Daud *et al.*, 2011).

Guava has several different accessions so we need some activities to characterize them. Guava diversity needs to be studied and evaluated in order to determine the next steps in the guava breeding. Information about description and genetic distance are needed to get new hybrid. Species diversity and genetic resources are very important to get new varieties.

Morphological characterization is the easiest activity to be done because it is simple, inexpensive and useful to determine the relatedness between accessions. One of morphological characters easy to be observed is fruit. Fruit is noticeable without special tools. Fruit, an important component of production, is usually different in shape, weight and color.

The relatedness and genetic distance between accessions can be obtained through cluster analysis. Cluster analysis is grouping accessions which have the same characteristics in homogeneous categories of each stratum (Crossa *et al.*, 1995a, 1995b). The relatedness analysis based on morphological characters will be perfect when using the descriptions of the characters that have high heritability values and stability (Beer *et al.*, 1993, Lamadji, 1998). Besides the morphological characterization, to determine the related-

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ness between accessions can also be done with biotechnology. Some countries such as Cuba have used microsatellites to characterize the guava (Infante *et al.* 2007). In India, to see the genetic diversity of guava derived from somatic embryo-genesis, SSR and ISSR markers are used (Rai *et al.*, 2012). Liu and Yang (2012) used ISSR markers to assess clonal fidelity of micro-propagated guava (*Psidium guajava* L.) plants.

The aim of the research was to characterize and accession the relatedness or grouping of guava. This research is expected to be used for selection of parents to get new superior varieties. This study is expected to be highly valuable in the future.

Table 1. Accessions list of guava germplasm collection at Aripian and Subang experimental farm

| NO | Accession | Location |
|----|-----------|---------------------------|
| 1 | ARP9409 | Aripian experimental farm |
| 2 | ARP9407 | Aripian experimental farm |
| 3 | ARP8653 | Aripian experimental farm |
| 4 | ARP10.2 | Aripian experimental farm |
| 5 | ARP10.7 | Aripian experimental farm |
| 6 | ARP10.6 | Aripian experimental farm |
| 7 | ARP10.1 | Aripian experimental farm |
| 8 | ARP10.12 | Aripian experimental farm |
| 9 | ARP10.9 | Aripian experimental farm |
| 10 | ARP10.11 | Aripian experimental farm |
| 11 | ARP8740 | Aripian experimental farm |
| 12 | ARP8742 | Aripian experimental farm |
| 13 | ARP8744 | Aripian experimental farm |
| 14 | ARP8741 | Aripian experimental farm |
| 15 | ARP8743 | Aripian experimental farm |
| 16 | JBT001 | Subang experimental farm |
| 17 | JBT002 | Subang experimental farm |
| 18 | JBT003 | Subang experimental farm |
| 19 | JBT004 | Subang experimental farm |

MATERIALS AND METHODS

The study was conducted at Aripian and Subang experimental farm from January 2012 to December 2012. The plant materials were 19 germplasm accessions of guava (Table 1). Each accession was characterized in 5 pieces. The characterization was made based on qualitative and quantitative characters. Guidelines of UPOV (1987) were also applied. Qualitative characters

include leaves (fully developed leaf, leaf curvature in cross section, leaf twisting, leaf curvature of midrib, leaf shape of base, leaf shape of tip, color of upper leaf, color of under leaf), and fruit (fruit shape, fruit shape at stalk end, fruit width of neck in relation to that of fruit, color of fruit skin, color of flesh). Quantitative characters cover leaf length (cm), leaf width (cm), fruit weight (g), fruit width (cm), fruit length (cm), petiole length (cm), diameter core (cm), flesh thickness (cm), total soluble solid (TSS) (° brix).

To make dendrogram, results of quantitative data and qualitative characterization were grouped by category/ class and were converted into binary form. Binary data were analyzed with the program NTSYS version 2.1 with the final result of dendrogram.

RESULTS AND DISCUSSION

Morphological characterization results showed that 19 accessions of germplasm collections guava in experimental farm at Aripian and Subang have morphological diversity (Appendix 1).

Leaf shape of some plants are obtrullate, obovate, oblong, and ovate. There was no difference in color of upper and lower surface of the leaf. In general, leaf is green with 11.96 – 15.02 cm in length and 5.39 to 10.34 cm in width. According to Mani *et al.* (2011), leaf shape of some *Psidium* species in India was ovate, lanceolate, oblong, elliptical to oblong and oval. Leaf length was about 4.2 - 13.3 cm, and 1.5 - 7.4 cm in width.

Fruit shape from 19 accessions observed had difference (Figure 1). There are round oval, oval, rounded, symmetry, rounded inverted cone, and ovate.

What becomes the main concern among consumers is fruit weight. Fruit size is a character which serves as the first attribute considered by consumers. In papaya, Chan *et al.* (1992) reported that papaya of a medium size is able to boost the market. Now the market preference is for small to medium size papayas because one fruit is enough for one person. In this research, the average of fruit weight ranged from 106.95 - 300.91 grams. Accessions with small weight involved ARP9406,

ARP9407, ARP8653, ARP10.2, ARP10.11, ARP8742, JBT001, JBT002, and JBT003, while the accessions with heavy weight comprised ARP10.8, ARP8741, and ARP8743. In Taiwan, there are some cultivars of guava. Taiwanese guava cv. has weights about 400-700 grams in average. Century guava cv. weighs about 200-400 grams. Pearl guava cv. weighs about 200-400 grams (Anonymous, 2011).

The color of fruit flesh varied for some accessions. Some accessions had red color, yellow, red purple, white, orange red. Red guava fruit has high content of vitamin C and beta carotene efficacious as anti-oxidant (Astawan 2013). However white guava has one special quality. Adnyana *et al.* (2004) stated that the ethanol extracts of white guava leaves showed stronger antibacterial activity than that of ethanol extracts of red guava leaves against *Escherichia coli*, *Shigella dysenteria*, *Shigella flexneri* and *Salmonella typhi*.

Total soluble solid (TSS) as a measurement tool was used to measure the level of sweetness. In this study, the values of TSS ranged from 7.8 – 11.03° brix. In general, there is an assumption that red guava fruit is sweeter than others. But in this research, it is known that red and white guava fruit almost had the same TSS values. So this assumption cannot always be said true. Accessions having low TSS values were ARP8743, and JBT004, while the accessions having high TSS values were ARP9406, ARP9407, ARP10.2, ARP10.7, ARP10.6, ARP10.8, ARP10.1, ARP10.12, ARP10.11, ARP8740, ARP8744 and JBT001.

Cluster analysis of 19 accession numbers were based on morphological characters of leaf and fruit (Figure 2.) It shows that 19 accessions of guava had 70-90% of similarity level or genetic distance of 0-20%. According to Cahyarini *et al.* (2004), the plants had the value of similarity level less than 0.60 or 60%, meaning that they had far genetic distance. In this research, accessions

which had far genetic distance were ARP8744 with ARP8741 and JBT 003 with ARP 10.12. Both of these accessions had similarity distance on 0.59. Sharma *et al.* (2010) stated that the variation among genotypes for different morphological characters could be attributed to genetic differences of these genotypes.

The accessions showing slight morphological similarity were good to be used as parent for crossing by considering of their characters. The clustering of this cultivar is useful in breeding activity to get new variety. The new variety will be better if the parent has far distance genetic relationship. Tatineni *et al.* (1996) said that parent plants having far genetic distance had opportunity to get higher heterosis hybrids than their parents. In this research, ARP8744 with ARP8741 and JBT003 with ARP10.12 were suitable as parent plant because these accessions had small similarity level.

Dendogram was obtained by 2 different groups at 70% of the similarity level, namely group I (ARP9406, ARP9407, ARP8653, ARP8742, ARP10.2, JBT001, JBT002, ARP8740, JBT003 and JBT004). Group I has the similarities in the character of the leaf shape of tip (acute), Color of under leaf (green), short petiole length (1.2-2.28 cm), and thin flesh thick(0.98-3.17 cm). At group I, the accessions which have the high similarity value (89%) are ARP9406 with ARP9407.

Group I involved ARP10.7, ARP10.6, ARP10.1, ARP10.12, ARP10.9, ARP10.11, ARP8744, ARP8741 and ARP8743. Group II had the same character in leaf twisting (absent), leaf curvature of midrib (present), and color of upper leaf surface (green). In group I, the accession which had the high similarity value (89%) was ARP10.1 with ARP10.2.

This dendogram was based on quantitative and qualitative characters. It described the cluster of every accession. It can be used to choose accession to be material selection in breeding.

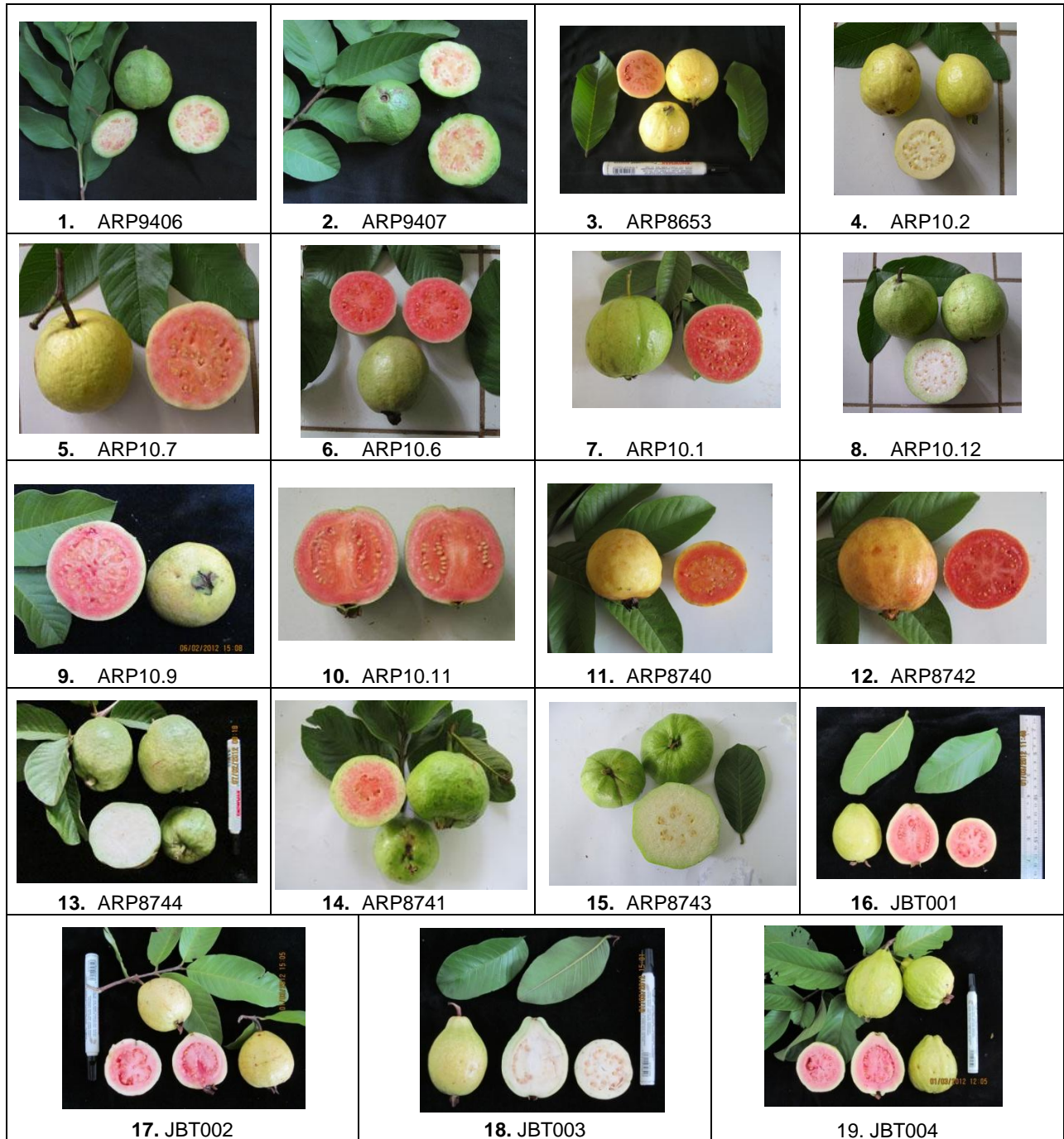


Figure 1. Fruit shape of nineteen accessions of guava

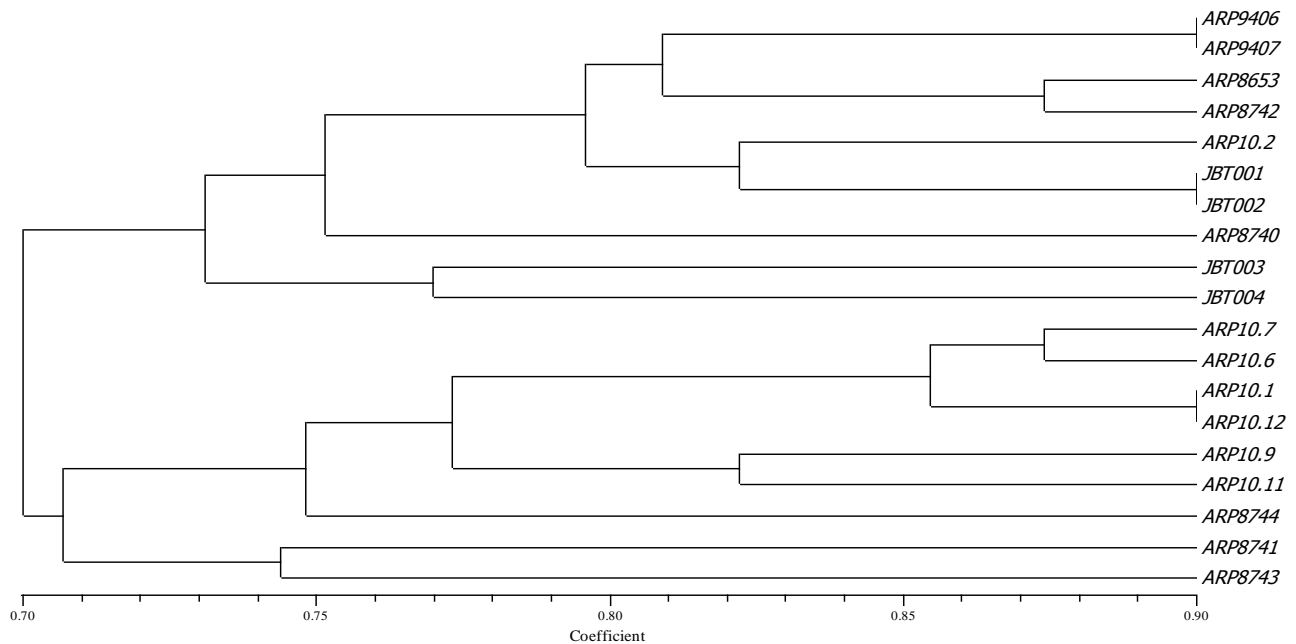


Figure 2. The dendrogram of clustering some accessions of guava based on morphological leaf and fruit characters

CONCLUSIONS

Similarity level from 19 of guava accessions was 70 – 90% or genetic distance was 0 – 20%. All of accessions can be clustered into 2 clusters at 70% genetic similarity level. Accession which had the heaviest fruit weight was ARP8743 (300.91 grams) and accession which had the highest value of TSS was ARP10.1 (11.03°brix)

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Appendix 1. Morphological characterization of some guava germplasm collections at Aripan and Subang Experimental Farm

| Accessions | ARP9406 | ARP9407 | ARP8653 | ARP10.2 | ARP10.7 |
|--|----------------|----------------|----------------|----------------|----------------|
| Shape fully developed leaf | Obtrullate | Obtrullate | Obtrullate | Obovate | Oblong |
| Leaf curvature in cross section | Medium | Medium | Medium | Weak | Medium |
| Leaf twisting | Absent | Present | Absent | Absent | Absent |
| Leaf curvature of midrib | Absent | Present | Present | Absent | Absent |
| Leaf shape of base | Rounded | Rounded | Rounded | Cordate | Rounded |
| Leaf shape of tip | Obtuse | Obtuse | Obtuse | Obtuse | Apiculate |
| Leaf length (cm) | 11.96 | 12.80 | 15.02 | 12.00 | 13.37 |
| Leaf width (cm) | 5.68 | 5.76 | 6.52 | 6.03 | 8.10 |
| Color of upper leaf | Green | Green | Green | Green | Green |
| Color of under leaf | Green | Yellow green | Green | Green | Green |
| Fruit shape | Round oval | Round oval | Oval | Rounded | Rounded |
| Fruit shape at stalk end | Pointed | Pointed | Pointed | Rounded | Rounded |
| Fruit width of neck in relation to that of fruit | Very broad | Very broad | Very broad | Very broad | Very broad |
| Color of fruit skin | Green | Green | Yellow | Yellow | Yellow green |
| Color of flesh | Red | Red | Red | Yellow | Red purple |
| Fruit weight (gram) | 115,82 | 148.43 | 162.04 | 118.31 | 191.09 |
| Fruit width (cm) | 6.32 | 6.29 | 6.50 | 5.92 | 6.86 |
| Fruit length (cm) | 6.45 | 6.23 | 7.07 | 6.40 | 7.11 |
| Petiole length (cm) | 1.57 | 1.67 | 1.84 | 1.55 | 2.21 |
| Diameter core (cm) | 4.26 | 4.28 | 4.30 | 3.74 | 4.43 |
| Flesh thick (cm) | 1.21 | 1.13 | 1.05 | 1.01 | 1.30 |
| TSS (° brix) | 11.00 | 10.60 | 9.20 | 10.20 | 9.86 |

| Accessions | ARP10.6 | ARP10.1 | ARP10.12 | ARP10.9 | ARP10.11 |
|--|----------------|----------------|-----------------|-----------------|-----------------|
| Shape fully developed leaf | Obovate | Obovate | Obovate | Obovate | Obovate |
| Leaf curvature in cross section | Medium | Medium | Medium | Medium | Medium |
| Leaf twisting | Absent | Absent | Absent | Absent | Absent |
| Leaf curvature of midrib | Absent | Absent | Absent | Absent | Absent |
| Leaf shape of base | Rounded | Rounded | Rounded | Rounded | Rounded |
| Leaf shape of tip | Obtuse | Obtuse | Obtuse | Apiculate | Obtuse |
| Leaf length (cm) | 14.82 | 13.02 | 14.39 | 11.97 | 12.03 |
| Leaf width (cm) | 8.08 | 6.30 | 6.05 | 5.39 | 7.94 |
| Color of upper leaf | Green | Green | Green | Green | Green |
| Color of under leaf | Green | Green | Green | Green | Green |
| Fruit shape | Round oval | Round | Round | Round | Round |
| Fruit shape at stalk end | Rounded | Rounded | Rounded | Broadly rounded | Rounded |
| Fruit width of neck in relation to that of fruit | Very broad | Very broad | Very broad | Very broad | Very broad |
| Color of fruit skin | Yellow green | Yellow green | Yellow green | Yellow green | Yellow green |
| Color of flesh | Red | White | White | Orange red | Red |
| Fruit weight (gram) | 185.59 | 177.70 | 223.51 | 185.44 | 128.98 |
| Fruit width (cm) | 6.92 | 6.90 | 7.58 | 6.25 | 5.09 |
| Fruit length (cm) | 7.34 | 6.93 | 7.71 | 5.40 | 5.25 |
| Petiole length (cm) | 2.36 | 4.43 | 2.49 | 3.46 | 2.31 |
| Diameter core (cm) | 4.44 | 4.32 | 4.59 | 4.16 | 4.30 |
| Flesh thick (cm) | 1.20 | 1.21 | 1.04 | 1.37 | 0.98 |
| TSS (° brix) | 10.15 | 11.03 | 10.83 | 9.38 | 10.27 |

| Accessions | ARP8740 | ARP8742 | ARP8744 | ARP8741 | ARP8743 |
|--|----------------|----------------|----------------|-----------------|-----------------|
| Shape fully developed leaf | Obovate | Obovate | Ovate | Obovate | Obtrullate |
| Leaf curvature in cross section | Strong | Medium | Weak | Strong | Medium |
| Leaf twisting | Absent | Absent | Absent | Absent | Absent |
| Leaf curvature of midrib | Present | Absent | Absent | Absent | Absent |
| Leaf shape of base | Rounded | Rounded | Rounded | Rounded | Obtuse |
| Leaf shape of tip | Obtuse | Obtuse | Obtuse | Rounded | Obtuse |
| Leaf length (cm) | 12.96 | 14.18 | 12.36 | 14.60 | 12.86 |
| Leaf width (cm) | 7.20 | 6.94 | 7.46 | 10.34 | 6.34 |
| Color of upper leaf | Green | Green | Green | Green | Green |
| Color of under leaf | Yellow green | Green | Green | Green | Green |
| Fruit shape | Oval | Oval | Symmetry | Round | Round |
| Fruit shape at stalk end | Rounded | Rounded | Rounded | Broadly rounded | Broadly rounded |
| Fruit width of neck in relation to that of fruit | Very broad | Very broad | Very broad | Very broad | Very broad |
| Color of fruit skin | Yellow | Yellow | Yellow green | Yellow green | Yellow green |
| Color of flesh | Red | Red | White | Red | White |
| Fruit weight (gram) | 227.08 | 158.39 | 175.82 | 287.54 | 300.91 |
| Fruit width (cm) | 7.33 | 6.06 | 7.56 | 8.40 | 8.93 |
| Fruit length (cm) | 7.31 | 5.58 | 8.29 | 7.27 | 6.65 |
| Petiole length (cm) | 1.25 | 1.97 | 2.26 | 4.31 | 1.50 |
| Diameter core (cm) | 4.03 | 4.30 | 7.56 | 2.67 | 3.89 |
| Flesh thick (cm) | 1.60 | 1.125 | 7.56 | 2.78 | 2.34 |
| TSS (° brix) | 10.10 | 9.00 | 10.02 | 8.80 | 7.10 |

| Accessions | JBT001 | JBT002 | JBT003 | JBT004 |
|--|---------------|-----------------------|---------------|---------------|
| Shape fully developed leaf | Oblong | Oblong | Oblong | Oblong |
| Leaf curvature in cross section | Weak | Weak | Medium | Medium |
| Leaf twisting | Absent | Absent | Present | Absent |
| Leaf curvature of midrib | Absent | Absent | Absent | Absent |
| Leaf shape of base | Rounded | Rounded | Rounded | Rounded |
| Leaf shape of tip | Obtuse | Obtuse | Obtuse | Obtuse |
| Leaf length (cm) | 12.45 | 12.96 | 13.15 | 15.04 |
| Leaf width (cm) | 5.82 | 5.51 | 7.07 | 7.42 |
| Color of upper leaf | Green | Green | Green | Green |
| Color of under leaf | Green | Green | Green | Green |
| Fruit shape | Round oval | Rounded inverted cone | Ovate | Oval |
| Fruit shape at stalk end | Necked | Pointed | Necked | Necked |
| Fruit width of neck in relation to that of fruit | Broad | Broad | Medium | Medium |
| Color of fruit skin | Yellow | Yellow | Yellow | Yellow |
| Color of flesh | Red | Red | Yellow | Red |
| Fruit weight (gram) | 122.64 | 106.95 | 139.82 | 215.008 |
| Fruit width (cm) | 5.93 | 5.74 | 5.96 | 7.14 |
| Fruit length (cm) | 6.69 | 6.32 | 7.54 | 8.60 |
| Petiole length (cm) | 1.48 | 1.67 | 1.29 | 1.20 |
| Diameter core (cm) | 3.20 | 3.69 | 3.99 | 4.41 |
| Flesh thick (cm) | 1.23 | 1.13 | 1.21 | 1.33 |
| TSS (° brix) | 10.75 | 9.71 | 9.13 | 7.80 |