ACCEPTED RESEARCH NOTE

Exploration And Characterization of Superior Locals Durian In Two

Subdistricts of Malang East Java Indonesia

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Durian production centers in East Java areas are quite numerous, two of which are Kasembon

and Ngantang Subdistricts, Malang District. In these areas, most durian trees are propagated

from seeds. This research aimed to determine the genetic diversity by exploration and

characterization of superior types, it was conducted by descriptive method. Determination of

samples in the study using a non probability sampling design with a purposive sampling

method. The research was conducted by characterizing the qualitative and quantitative

morphological characters from fruit, leaves, and trees. The characters identified are 81

characters. Characterization of the durian types was made based on the guidebook description

from Bioversity International. Quantitative data was used for descriptive statistical analysis.

Qualitative data analysis was carried out using the Unweighted Pair Group Method with

Arithmetic Averaging (UPGMA) method using NTSYS version 2.02 software. The results of

the research showed that local superior durians from Kasembon and Ngantang Subdistrict had

diverse characteristics. Diversity are in the character of fruits, leaves, and trees. The genetic

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ABSTRACT

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distance among superior durians is 0.109-0.343. Some of durian accessions observed such as 31 Vodca, and Kunir Amad have the potential to be registered as regional assets or as local 32

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ripe durian fruit owing to its unique taste and aroma is considered locally the king of fruits

superior cultivars.

INTRODUCTION

KEYWORDS: Characteristic, durian, local, morphology.

(Srianta et al., 2012), contain high in essential micro and macro nutrients, minerals and

Durian (Durio zibethinus) is a climacteric, seasonal tropical fruit of Southeast Asia. The

vitamins (Ho & Bhat, 2015), provide phytoestrogens, anti inflammatory, and antioxidants (Zannah et al., 2022). Durian is a type of annual plant, and evergreen leaves, but it has a period of flushing, which is a period of growing new leaves and generally occurs after the fruiting period is over (Ashari, 2017).

Durian production in Indonesia reaches 1.169 million tons annually, East Java being the largest durian producer with 289,000 tons (Agriculture Ministry, 2019), provides additional income for the community (Matius et al., 2018; Prasetyo et al., 2018). In Indonesia at least there are 84 varieties have been released by the Ministry of Agriculture, based on their morphological characteristics, and originate. The wide variety of cultivars is an important source of germplasm as selection material for breeding durian and becoming superior local fruit cultivars from each region (Yulita & Nurnianjari, 2010). Efforts to identify superior types of durian from this region to various areas carried out through inventory activities, includes exploration and identification, is an out of the field activity collect data on types of durian in the area, genotype morphology characteristic expected to reveal potential seed of plant to be developed in another region, also is used as a reference to introduce all types durian in this area to a wider scope (Yuniastuti, Anggita, et al., 2018; Yuniastuti, Nandariyah, et al., 2018).

Characterization and relationship among them are very important for plant breeding program (Hariyati et al., 2013). First select a good character edible fruit, flesh color, water content, attractive flesh aroma, soft aril texture, fair flesh creaminess, sweet and sweet with bitter after-taste flesh. These two subdistricts are known as centers of local durians in Malang Regency which have wide diversity of local cultivars with delicious taste and more expensive than other types but have not been reported yet. This research was conducted to determine the genetic diversity and characterization of local superior durians in Kasembon and Ngantang Subdistricts.

MATERIALS AND METHODS

The research was conducted in peak season from January to August 2018 in Subdistricts of Kasembon 7°47′02″, 112°18′32″ and altitude of 500-721 m asl and Ngantang -7°51′23″, 112°22′09″ and altitude of 199-539 m asl.



Figure 1. Research map location (Source: http://:www.malangkab.go.id).

Durian is known as an obligate cross pollinated plant (Santoso et al., 2014). The way to map the differences in durian plant types is by conducting exploration with farmers who own trees in their yards. Observed plant samples were determined based on the results of interviews through field surveys involving farmers, farmer groups, traders and local agricultural services as well as a durian fruit festival organized by the Government in East Java. Criteria for evaluating the superiority of each identified durian variety using the Durian Assessment Standard Guidelines (Direktorat Budidaya Tanaman Buah, 2010).

The characterization of the durian types was made based on the guidebook description from Bioversity International (Bioversity International, 2007). The characters identified are 81 characters, namely 48 fruit characters, 20 leaf characters, and 13 tree characters. The characteristics identified in the organoleptic test were aril texture, aril juiciness, presence of fiber, creaminess, flesh taste flesh aroma, and flesh stickiness.

Qualitative data analysis using the Unweighted Pair Group Method with Arithmetic Averaging (UPGMA) method using NTSYS (Numerical Taxonomy and Multivariate Analysis System) software. Qualitative data of characters are transformed into binary data and arranged in table form in Ntedit software version 1.07. After that, the data from the Ntedit result is processed with NTSYS version 2.02 software (Handayani & Rahayu, 2017). The results of processing data in this software produce a dendrogram and matrix.

RESULTS AND DISCUSSION

Genetic variation of local durians

In the exploration, there were 18 durian types (Table 1) were selected based on national durian idiotypes and consumer preference (Santoso et al., 2016). Most durian trees in these areas a were originated from seeds.

Table 1. Names and subdistricts of superior	aurian	touna.
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No.	Durian name	Initials	Origin subdistrict
1.	Kendi	KD	Ngantang
2.	Kunir Amad	KA	Ngantang
3.	Vodka	VK	Ngantang
4.	Manalagi Kuning	MK	Ngantang
5.	Duri Tajam	DT	Ngantang
6.	Getuk	GT	Ngantang
7.	Ranti	RT	Ngantang
8.	Selaput Udang	SU	Ngantang
9.	Toni	TN	Ngantang
10.	Aroma Kweni	AK	Ngantang
11.	Joko	JK	Ngantang
12.	Tarum	TR	Ngantang
13.	Ketan	KT	Ngantang
14.	Gipat	GP	Ngantang
15.	Klenting Kuning	KK	Kasembon
16.	Lima Ratus	LR	Kasembon
17.	Duri In	DI	Kasembon
18.	Brojo	BJ	Kasembon

Diversity exists in three levels, namely genetic, species, and ecosystem diversity. Morphological characteristics are based on the appearance of phenotypes, namely fruit shape, fruit spines, fruit flesh taste, flowers, leaf colour, and other characteristics (Rivero-Guerra, 2011; Sunaryo et al., 2015; Sundari et al., 2015; Daryono & Maryanto, 2017).

Durian fruit shape is globose (7 of 18 individuals) in general (Figure 2). Characters of shape of the fruit apex are: mammiform, convex, truncate, depressed, and pointed. Shape of the fruit base are: depressed, convex, truncate, and necked. Fruit spine shape are: convex, concave, pointed concave, and conical. Fruit spine densityare: intermediate and sparse. Fruit rind color, are: yellow, yellowish green, greenish yellow, and brown. Flesh color are: yellow, orange, and white (Figure 3). Local superior durians mostly have intermediate easiness of splitting, thin-rinded durian fruit has a longer shelf life, and has a more edible portion of fruit (Fitmawati et al. 2015).

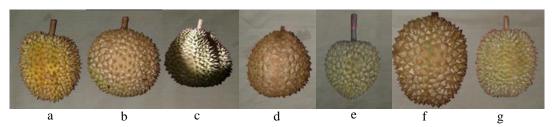


Figure 2. Fruit shape (a) globose; (b) oblate; (c) star; (d) obovoid; (e) oval; (f) oblong; (g) ovoid.

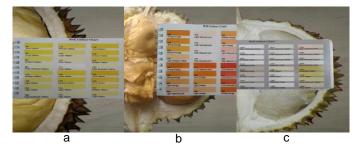


Figure 3. Flesh color, a. Yellow, b. Orange, and c. White.

Durian individuals with aborted and immature seeds have the potential to develop as superior durians without seeds and enlarge the edible portion. Durian had seed shape variations (Figure 4), with seed coat color intensity light and dark.



Figure 4. Seed Shape, a. Spheroid, b. Ellipsoid, c. Oblong, d. Ovoid, e. Obovoid, and f. Irregular.

The 18 durian types were varied in leaf form, generally leaf blade shape is elliptic (16 of 18 individuals). Most of the leaf attitude is drooping at 45° (17 of 18 individuals), the crown shape of the tree was 2, its growth habit variants of the durian tree were 3, most of the tree growth habit is intermediate (15 of 18 individuals). Durian tree age range from 11 until there are more than 100 years. Tree height range between 10.1-30.1 meters.

The organoleptic test showed that most of the local durians had moderate flesh aroma, soft aril texture, non-juicy aril, medium presence of fibre, fair flesh creaminess, slightly sticky flesh, sweet and sweet with bitter after-taste flesh with total sugar range between 34.96 ± 0.33 – $48.38\pm0.46\%$. It caused by genetic differences, environmental factors and the possibility of fruit ages at different harvest times (Hadiati et al., 2016), the sweetness level is influenced by level of fruit maturity and water content (Belgis et al. 2016)

Propagation by seeds will produce offspring that vary with the properties of the parent plant, because durian is a cross pollinating plant (Bumrungsri et al., 2009). According to (Indriyani et al. 2012), maternal parent influence the character of fruit (length, skin thickness, number of seeds, seed weight per fruit) and percentage of aborted seeds. There are many hybrids which are suitable for selection because durian is an open pollination plant (Prihatini et al. 2016), this is consistent with the results of this study. Exploration results for local durians in the two sub-districts show a high level of diversity as most durians grow from

seeds that are the result of crossing different combinations of parent trees. This increases the genetic diversity of durian plants, giving us the opportunity to obtain superior accession with high heterosis to use as mother trees and develop through vegetative propagation. However, the consequences of seed propagation will make the durian population heterogeneous, resulting individual inferior fenotypes that are less profitable for farmers.

Clustering of local durians

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Cluster analysis was carried out based on qualitative data, showed that morphological similarity value of 18 durian species was 0.657-0.891 (Figure 5). Local superior durians in Kasembon and Ngantang Subdistricts based on the qualitative characters of fruit, leaves and trees, are divided into 2 Main Clusters, Ranti's accession was identified as the only individual with the lowest similarity coefficient (0.657) with the other 17 accessions, so it split from the two main cluster that formed, as it bears little morphological resemblance to other accessions. Grouping between accessions is not always related with origin, first group consists of various accessions originating from both subdistrict, Kendi and Duri In are the two accessions that show the highest morphological similarity (0.891). Grouping occurs because these accessions have similarities in morphological characters which can be caused by natural cross pollination of durian plants with the help of bats, birds and several species of beetles (Sritongchuay et al., 2016; Stewart & Dudash, 2017; Wayo et al., 2018). The highest similarity value was found between Kendi and Duri In durian (0.891), while the lowest similarity value was found between durian Ranti and 17 other durians (0.657) so it has not been identified. This is in accordance with the results of the study (Santoso et al., 2017) and (Sundari et al., 2021) that the fewer similarities, smaller similarity value or greater the genetic distance. Based on morphological observations (leaf and tree characters), genetic distance between durians is 0.109-0.343.

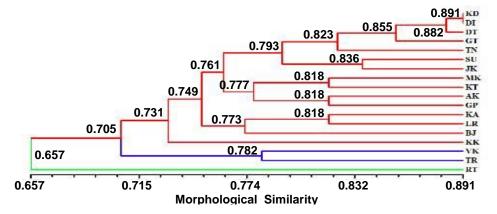


Figure 5. The dendrograms of Local superior durian Kasembon and Ngantang Subdistricts

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Figure 6. The resemblance matrix of local superior durian characters in Kasembon and Ngantang Subdistricts.

Differences and similarities in the emergence of outside morphology of a plant species can be used to determine the proximity of kinship. Environmental factors also influence the expression of these characteristics, even if they are only temporary (Hafizah et al., 2018).

CONCLUSIONS

The results of the research showed that local superior durians from Kasembon and Ngantang had diverse characters in the character of fruits, leaves and trees. The genetic distance between 18 durian accessions is 0.109–0.343 on the fruit, leaf and tree characters. Some of durian accessions observed such as Vodca, and Kunir Amad have the potential to be registered as regional assets or as local superior cultivars. This research was carried out observing agromorphology and organoleptics, so the results are still influenced by the environment.

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REFERENCES

- 194 Ashari, S. (2017). Durian: king of the fruit. UB Press.
- Belgis, M., Wijaya, C. H., Apriyantono, A., Kusbiantoro, B., & Yuliana, N. D. (2016).

 Physicochemical differences and sensory profilling of six lai (Durio kutejensis) and four
- durian (Durio zibethinus) cultivar indigenous indonesia. *International Food Research*
- 198 *Journal*, 23(4), 1466–1473.
- Bioversity International. (2007). *Descriptors for durian (Durio zibethinus Murr.)*. Bioversity International.

- 201 Bumrungsri, S., Sripaoraya, E., Chongsiri, T., Sridith, K., & Racey, P. A. (2009). The
- pollination ecology of durian (Durio zibethinus, Bombacaceae) in southern Thailand.
- 203 *Journal of Tropical Ecology*, 25(1), 85–92. https://doi.org/10.1017/S0266467408005531
- 204 Daryono, B. S., & Maryanto, S. D. (2017). Keanekaragaman dan Potensi Sumber Daya
- 205 Genetik Melon. Gadjah Mada University Press.
- 206 Direktorat Budidaya Tanaman Buah. (2010). Pedoman Standar Penilaian Durian. Direktorat
- 207 Jendral Hortikultura Kementerian Pertanian.
- 208 Fitmawati, Aisyah, L., & Iriani, D. (2015). Comparative study of anatomical structure of
- durian perikarp (Durio zibethinus Murr.) shelf life and not shelf life from Bengkalis
- Island, Riau Province. In A. Soemargono, Mulyati, S. Hadiati, Martias, A. Sutanto, N. L.
- P. Indriyani, & Jumjunidang (Eds.), Nusantara II. Bukittinggi 23–25 September 2015
- 212 (pp. 631–640).
- 213 Hadiati, S., Nasution, F., & Kuswandi. (2016). Karakterisasi dan Evaluasi Koleksi Sumber
- Daya Genetik Durian Berdasarkan Karakter Morfologi Buah. Bul. Plasma Nutfah, 22(1),
- 215 1–10.
- Hafizah, R. A., Adawiyah, R., Harahap, R. M., Hannum, S., & Santoso, P. J. (2018). Aplikasi
- Marka SSR Pada Keanekaragaman Genetik Durian (Durio zibethinus Murr.) Di
- Kabupaten Deli Serdang, Sumatra Utara. Al-Kauniyah: Jurnal Biologi, 11(1), 49–56.
- 219 https://doi.org/10.15408/kauniyah.v11i1.5668
- 220 Handayani, F., & Rahayu, S. P. (2017). Short Communication: Assessment of genetic
- diversity in Lai (Durio kutejensis) local cultivars of Batuah (Indonesia) using ISSR
- marker. Biodiversitas Journal of Biological Diversity, 18(2), 525–529.
- 223 https://doi.org/10.13057/biodiv/d180212
- Hariyati, T., Kusnadi, J., & Arumingtyas, E. L. (2013). Genetic diversity of hybrid durian
- resulted from cross breeding between Durio kutejensis and Durio zibethinus based on
- 226 random amplified polymorphic DNAs (RAPDs). American Journal of Molecular
- 227 Biology, 03(03), 153–157. https://doi.org/10.4236/ajmb.2013.33020
- 228 Ho, L.-H., & Bhat, R. (2015). Exploring the potential nutraceutical values of durian (Durio
- zibethinus L.) An exotic tropical fruit. Food Chemistry, 168, 80–89.
- 230 https://doi.org/10.1016/j.foodchem.2014.07.020
- Indriyani, N. L. P., Hadiati, S., Nasution, F., Sudjijo, E., & Irawati, Y. (2012). Maternal and
- Paternal Effect on the Characters of Durian (Durio Zibethinus Murr.) Fruit from Cross-
- Pollination. Journal of Fruit and Ornamental Plant Research, 20(2), 23–33.
- 234 https://doi.org/10.2478/v10290-012-0012-x

- 235 Kementerrian Pertanian. (2019). Produksi Durian Menurut Provinsi , Tahun 2015-2019.
- Www. Pertanian.Go.Id.
- Matius, P., Tjwa, S. J. M., Raharja, M., Sapruddin, S., Noor, S., & Ruslim, Y. (2018). Plant
- diversity in traditional fruit gardens (munaans) of Benuaq and Tunjung Dayaks tribes of
- West Kutai, East Kalimantan, Indonesia. Biodiversitas Journal of Biological Diversity,
- 240 19(4), 1280–1288. https://doi.org/10.13057/biodiv/d190414
- 241 Prasetyo, B., Chikmawati, T., Walujo, eko baroto, & Amzu, E. (2018). Ethnoecology: The
- traditional landscape of Osing Tribe in Banyuwangi, Indonesia. *Biodiversitas Journal of*
- 243 Biological Diversity, 19(6), 2003–2009. https://doi.org/10.13057/biodiv/d190604
- 244 Prihatini, R., Ihsan, F., & Indriyani, N. L. P. (2016). Genomic Profiling of F 1 Hybrids of
- Durian (Durio zibethinus) Revealed by RAPD-PCR. Journal of Horticultural Research,
- 24(2), 69–76. https://doi.org/10.1515/johr-2016-0022
- 247 Rivero-Guerra, A. O. (2011). Morphological Variation within and Between taxa of the
- Santolina rosmarinifolia L. (Asteraceae: Anthemideae) Aggregate. Systematic Botany,
- 249 *36*(1), 171–190. https://doi.org/10.1600/036364411X553261
- 250 Santoso, P. J., Granitia, A., Indriyani, N. L. P., & Pancoro, A. (2016). Analisis Lokus dan
- 251 Keragaman Sumber Daya Genetik Durian (Durio sp.) Berdasarkan Marka Mikrosatelit.
- 252 *Jurnal Hortikultura*, 26(1), 9. https://doi.org/10.21082/jhort.v26n1.2016.p9-20
- Santoso, P. J., Pancoro, A., Suhandono, S., & Aryantha, I. N. P. (2017). Development of
- Simple-Sequence Repeats Markers from Durian (Durio zibethinus Murr. cultv. Matahari)
- Genomic Library. AGRIVITA Journal of Agricultural Science, 39(3), 257-265.
- 256 https://doi.org/10.17503/agrivita.v39i3.1171
- 257 Srianta, I., Hendrawan, B., Kusumawati, N., & Blanc, P. J. (2012). Study on durian seed as a
- new substrate for Angkak production. International Food Research Journal, 19(3), 941-
- 259 945.
- 260 Sritongchuay, T., Kremen, C., & Bumrungsri, S. (2016). Effects of forest and cave proximity
- on fruit set of tree crops in tropical orchards in Southern Thailand. *Journal of Tropical*
- 262 *Ecology*, 32(4), 269–279. https://doi.org/10.1017/S0266467416000353
- Stewart, A. B., & Dudash, M. R. (2017). Flower- visiting bat species contribute unequally
- 264 toward agricultural pollination ecosystem services in southern Thailand. *Biotropica*,
- 265 49(2), 239–248. https://doi.org/10.1111/btp.12401
- Sunaryo, W., Hendra, M., Rudarmono, Suprapto, H., Pratama, A. N., & Rahman. (2015).
- Exploration and identification of Lai Durian, new highly economic potential cultivars
- derived from natural crossing between Durio zibethinus and Durio kutejensis in East

269	Kalimantan. Asian Journal of Microbiology, Biotechnology and Environmental Sciences
270	<i>17</i> (2), 365–371.
271	Sundari, Arumingtyas, E. L., Hakim, L., & Azrianingsih, R. (2015). Exploration and
272	Morphological Character Identification of Local Durian (Durio zibethinus Murr.) from
273	Tidore Island, North Maluku. Proceeding of 6th ICGRC, 1-4.
274	Sundari, Mas'Ud, A., Wahyudi, D., Arumingtyas, E. L., Hakim, L., & Azrianingsih, R
275	(2021). Genetic diversity of local durian from Tidore Island based on morphological and
276	moleculer data for tropical fruit conservation in North Maluku. IOP Conference Series.
277	Earth and Environmental Science, 739(1). https://doi.org/10.1088/1755-
278	1315/739/1/012073
279	Wayo, K., Phankaew, C., Stewart, A. B., & Bumrungsri, S. (2018). Bees are supplementary
280	pollinators of self-compatible chiropterophilous durian. Journal of Tropical Ecology
281	34(1), 41–52. https://doi.org/10.1017/S0266467418000019
282	Yulita, K. S., & Nurnianjari, M. (2010). Keragaman Genetik Beberapa Klon Durian (Duric
283	zibethinus Murray) asal Jawa Barat berdasarkan Sidik Random Amplified Polimorphic
284	DNA. Jurnal Berita Biologi, 10(3), 269–275.
285	Yuniastuti, E., Anggita, A., Nandariyah, & Sukaya. (2018). Local durian (Durio zibethinus
286	murr.) exploration for potentially superior tree as parents in Ngrambe District, Ngawi
287	IOP Conference Series: Earth and Environmental Science, 142(1)
288	https://doi.org/10.1088/1755-1315/142/1/012029
289	Yuniastuti, E., Nandariyah, N., & Bukka, S. R. (2018). Karakterisasi Durian (Durio
290	zibenthinus) Ngrambe di Jawa Timur, Indonesia. Caraka Tani: Journal of Sustainable
291	Agriculture, 33(2), 136. https://doi.org/10.20961/carakatani.v33i2.19610
292	Zannah, F., Kamaliah, Pramudiyanti, Ayatusaadah, & Hidayati, N. (2022). Exploration of the
293	Potential of Local Plants of Melastoma malabatchricum Fruit for Food Fortification
294	Journal of Tropical Life Science, 12(3), 333–338. https://doi.org/10.11594/jtls.12.03.06