Scholars Journal of Agriculture and Veterinary Sciences (SJAVS) e-ISSN 2348–1854 Abbreviated Key Title: Sch. J. Agric. Vet. Sci. p-ISSN 2348–8883

Abbreviated Key Title: Sch. J. Agric. Vet. Sci. ©Scholars Academic and Scientific Publishers (SAS Publishers) A Unit of Scholars Academic and Scientific Society, India www.saspublishers.com

Germplasm Diversity Of Banana (Musa Spp) in The City of Bandar Lampung, Indonesia by Type of Genome and Number of Chromosome

Eti Ernawiati^{*}, Rochmah Agustrina, Bambang Irawan, Eka Nurhasanah, Mohammad Kanedi

Department of Biology, Faculty of Mathematics and Sciences, University of Lampung, Bandar Lampung, Indonesia

Abstract: Bananas have a high diversity, so it can be a source of germplasm in plant breeding programs. This research aimed to investigate diversity of banana (Musa spp.) **Original Research Article** germplasm in the City of Bandar Lampung based on the number of chromosome and types of genome. The banana plant samples were taken randomly from 12 districts in *Corresponding author Bandar Lampung. Morphological characterization for genomic determination of the Eti Ernawiati plants was done using 15 characters from Simmonds and Shepherd (1955) and the expectation scores of genomes was adopted from Silayoi and Camchalow (1987). **Article History** Banana chromosomes number was determined using the squash method. The results Received: 07.04.2017 showed among 27 accessions of bananas collected from Bandar Lampung City, 26 Accepted: 17.04.2018 accessions belong to the genus Musa, namely Musa acuminata, Musa balbisiana, Musa Published: 30.04.2018 paradisiaca, and 1 accession belonging the genus Rhodhoclamys. The diversity of banana species in Bandar Lampung City is moderate. Variation of the genome obtained DOI: include AA (5 accessions), AAA (4 accessions), AAB (5 accessions), ABB (3 10.36347/sjavs.2018.v05i04.010 accessions), ABBB (1 accession), BBB (3 accessions), and 5 other accessions that cannot be determined. Chromosome number of the banana accessions found in Bandar Lampung is 22 (diploid), 33 (triploid) and 44 (tetraploid) Keywords: banana germplasm, Musa spp., genus Musa, chromosome number, genome type.

INTRODUCTION

Banana is a tropical crops abundant in Southeast Asia and has a high diversity. Bananas are much liked by many people around the world, because of their complete nutritional value. Every 100 g of bananas produces 90 calories. Bananas clearly contain no cholesterol, rich in vitamins A, B6, C, calcium, potassium, and phosphorus. One banana contains 380 mg of potassium, so it can meet the daily potassium requirement of an adult [1]. Thus, it is not surprising that bananas become one of the world's fourth most important foods after rice, wheat and corn [2].

In Indonesia bananas are included in superior fruit commodities, with plantation areas covering almost all regions in the country. According to the Indonesia Statistics, banana production in Indonesia increases every year. Lampung Province occupies the first position as the highest banana production center in Indonesia with production reaches 1,937,349 tons, followed by the Provinces of East Java, West Java, and Central Java., Considering that Bandar Lampung City is a home industry center of banana chips that has a high contribution in the community economy, the banana germplasm in this district is feasible to be preserved [3].

Available online: <u>www.saspublisher.com</u>

Bananas that are currently cultivated are bananas from natural crosses between bananas *Musa acuminata* and *Musa balbisiana* [4]. The genetic diversity of banana plants can be used as a source of germplasm that needs to be continuously evaluated, since it has not been fully revealed.

Banana plants have varying ploidy levels. It occurs naturally as a result of crossings between wild banana species, takes place continuously and is influenced by its environment [5]. The study result of chromosome number can be used for plant breeding because the number of chromosomes is constant, so it is important to maintain the character of a plant over time. Banana plants have the basic chromosome number x =11 with the size of the haploid genome ranging from 500-600 Mb. Size of the banana genome, one-quarter times larger than the rice genome [6]. It has suggested that the small size of the genome has the potential to provide a better picture of the function and structure of the genomics. This is useful for banana plant breeding purposes [7]. Determination of genome type of banana plant can be done using scoring system based on morphological characteristics. Information from the results of the study of the determination of the number of chromosomes and the genome type of banana plants will complement each other in the study of banana cultivation, and the results can be used as information in the plant breeding program. Therefore, the results of this study can reveal the biodiversity of banana germplasm based on the number of chromosomes and genomes in the city of Bandar Lampung.

MATERIALS AND METHODS

The study was conducted in three steps as follows: 1) collecting the banana plants, 2) characterization of the plant morphology to determine the genome of the banana, and 3) determination of chromosome number to determine the level of ploidy.

Colection of Banana germ plasm

Banana plant samples were collected randomly from 12 sub-districts in Bandar Lampung. The Diversity Index Value of the banana plants collection then determined using Shannon-Wiener Diversity Index, the Shannon-Wiener Diversity Index Value (H') is calculated by equation bellows [8].

$$H' = -\sum_{i=1}^{s} Pi * \ln Pi$$

$$Pi = \frac{number \ of \ individu \ for \ each \ species}{number \ of \ individu \ for \ all \ species}$$

With: H' = value of Shannon-Wiener diversity index Pi = proportion of total abundance represented by ith species s = number of species in community

If the index value (H') < 1, diversity is low; if the index value (H') is $1 < H' \leq 3$, the diversity is moderate; and if the index value (H') > 3, then the diversity is high.

Mhorphological characterization to determinate genome of banana plant

The type of genome was determined using table of morphological character based on 15 characters according to Simmonds and Shepherd which was then calculated using the expectation scores of Silayoi and Camchalow. The characters observed to differ the genome of *Musa acuminata* from *Musa balbisisana* are vegetative characters. The characters observed to distinguish the genome of *Musa acuminata* and *Musa balbisisana* are vegetative characters consisting of the color of the pseudo stems, the stem of the bunches, the petiole, the fruit stalk, and the generative characters consisting of number of seeds of per locus, bractea ratio, bractea tip, rate of bractea coloring (color fading), bractea marks, free tepal of male flowers, flower color of males, and stigma color [9].

Scores of each cultivar were then summed and matched with the expectation scores of the genome group from Silayoi and Camchalow [10]. The number of scores from the matching results of the expectation scores with the AA / AAA genome group was (15-25), with the AAB genome group was (26-46), with the AB / AABB genome group was (47-49), with the ABB genome group was (59-63), with the ABBB genome group was (67-69) and with the genomes group was BB / BBB (70-75).

Chromosome number determination

Determination of the chromosome number was performed using the squash method Darnaedi, 1991 with the following sequence of steps: cutting the meristem of the root tip of banana done at 08.30 - 10.00 am, the pre-treatment of banana root pieces was done by soaking it in 0.03% hydroxiquinoline for 3-5 hour at 18-20°C, root- pieces fixation with 45% acetic acid for 10 minutes, hydrolysis of the root pieces in a mixture of HCL 1 N and 45% acetic acid (3: 1) was carried out for 3-5 min, heating the root pieces on the surface of the hot plate at 60°C, staining the root pieces by immersing them in 2% aseto-orcein for 15 min, then squashing the root pieces using erased wood pencils, and observations of banana root chromosome preparations under a microscope with 1000x magnification.

RESULTS AND DISCUSSION

The results showed that from 12 sub-districts in Bandar Lampung city, 27 accessions of bananas were obtained. All the 27 of banana accessions fall into two groups of genera. First is the group of genus *Eumusa* consisting of *Musa acuminata* which has 9 accessions, *Musa paradisiaca* which has 9 accessions, *Musa balbisiana* which has 3 accessions, and second is genus of *Rhodhoclamys* which only has 1 accession. The remaining 5 accessions of bananas whose genome type cannot be determined yet because when taking the banana root preparation will be done, the generative period of the banana plants have passed (Table 1). The result of calculation of Diversity Index Value (H') is 2,8369. Thus the diversity of banana plants in Bandar Lampung City is moderate. The study results shows the genome types of banana plant obtained are AA, AAA, AAB, ABB, ABBB, and BBB with the number of basic chromosomes x = 11. The growth type of the upright to slightly erect leaves is characteristic of the banana cultivar from the descendants of *Musa acuminata* whereas in the descendants of *Musa balbisiana* has a drooped leaf growth type [11]. Another research results found 33 chromosome number from banana accession of morosebon (AAA), tanduk (AAB), raja sereh, kepok kuning, ambon kuning, and the number of chromosome 22 from banana accession of emas and lampung with AA genome type [12].

The results of characterisation above show that banana cultivar in Bandar Lampung City is dominated by the type of banana from the descendants of *Musa acuminata*. This shows that the type of banana *Musa acuminata* has a high crossing ability. The data obtained also indicate the presence of predominantly banana with triploid banana species. According to Simmonds (1962), the triploid banana cultivar has a sturdy and sturdy stout quasi-stature character, with leaves that are thicker and wider than those of diploid bananas, and in addition the triploid banana species are tolerant to minimal soil nutrients and more resistant to disease [13].

No.	Species	Accessions	Score	Types of Genome	Number of Chromosome
1	Musa acuminata	Lilin (Janten)	20	AA	22
2		Mas	18	AA	22
3		Mas Kuning	21	AA	22
4		Muli (Lampung)	20	AA	22
5		Rejang	21	AA	22
6		Ambon Kuning	17	AAA	33
7		Ambon Lumut	22	AAA	33
8		Cavendish	19	AAA	33
9		Papan	21	AAA	33
10	Musa paradisiaca	Kepok Abu	42	AAB	33
11		Morosebo	28	AAB	33
12		Raja Nangka	34	AAB	33
13		Seribu	45	AAB	33
14		Tanduk	47	AAB	33
15		Kepok Batu	58	ABB	33
16		Rabig	64	ABB	33
17		Raja Sajen	59	ABB	33
18		Batu	65	ABBB	44
19	Musa balbisiana	Kepok Kapas	71	BBB	33
20		Kepok Kuning	71	BBB	33
21		Kepok Manado	70	BBB	33
22	Hasn't been determined	Ambon Australi	-	-	44
23		Kepok Libanon	-	-	33
24		Kidang	-	-	33
25		Raja Bakar	-	-	33
26		Raja Sereh		-	33
27	Musa ornata	Musa ornata	*	*	22

Table-1. Types of genome and	chromosome number	of hanana in Ban	dar Lamnung	Indonesia
Table-1. Types of genome and	Chi omosome number	VI Danana in Dan	uai Lampung	, muunesia

*belong to genus Rhodhoclamys

In this study, the banana cultivars found to have 22 (diploid), 33 (triploid) and 44 (tetraploid) chromosomes. It has suggested that the type of diploid, triploid, and tetraploid bananas that exist today is the result of a naturally occurring plant crosses continuously and influenced by environmental factors [14]. Previously, it has reported that in Bandar Lampung there were 12 accessions of banana plants that have potential as table fruit, processed fruits, and ornamental plants [15]. Another study conducted in Lampung Province found 21 accessions of bananas in the district of South Lampung with variation of genomes are AA, AAA, AAB, ABB, BB [16]. From the district of Pesawaran, still in the province of Lampung, Puspitasari (2011) found 13 accessions of bananas with the variation genomes are AA, AAA, AAB, ABB, and BB [17].

CONCLUSION

• There are 27 accessions of banana in the city of Bandar Lampung. Based on the cytological analysis and the diversity of morphological characters it is found that the Banana Diversity Index is 2.83 and so it is classified as medium.

- Types of banana genome obtained are: *Musa acuminata* (AA, AAA), *Musa balbisiana* (BBB), *Musa paradisiaca* (AAB, ABB, ABBB).
- The chromosome number of the bananas are 22 (diploid), 33 (triploid), and 44 (tetraploid).

REFERENCES

- 1. Simmonds, NW. *Bananas*. Longmands. London. 1966.
- Frison EA, Escalant JV, Sharrock S, Jain SM, Swennen R. The global Musa genomic consortium: A boost for banana improvement. Banana improvement: cellular, molecular biology, and induced mutations. 2004:341-9.
- Statistik BP. Produksi Tanaman Pisang Seluruh Provinsi. Diakses dari www. bps. go. id pada tanggal. 2015;14.
- 4. Paull RE and Duarte O. *Tropical Fruits*. (2). (Crop production Science in horticulture series: No. 24). 2 nd Ed. The British Library London, U.K. 2011.
- Valmayor RV, Jamaluddin SH, Silayoi B, Kusumo S, Danh LD, Pascua OC, Espino RR. Banana cultivar names and synonyms in Southeast Asia. Advancing Banana and Plantain R & D in Asia and the Pacific. 1999.
- Escalant JV, Sharrock S, Frison E, Carreel F, Jenny C, Swennen R, Tomekpe K. The genetic improvement of Musa using conventional breeding, and modern tools of molecular and cellular biology. IPGRI, Rome, Italy. 2002 Dec:17.
- 7. Megia, R. 2005. *Musa* Sebagai Model Genom. *Hayati*. 12 (4): 167-170.
- Bengen DG. Pedoman teknis pengenalan & pengelolaan ekosistem mangrove. Pusat Kajian Sumberdaya Pesisir dan Lautan (PKSPL), Institut Pertanian Bogor (IPB); 2000.
- 9. Simmonds NW and K. Shepherd. *Bananas*. Longmans. London. 1959; 466 pp.
- Silayoi B, Chomchalow N. Cytotaxonomic and morphological studies of Thai banana cultivars. Proceedings of Banana and Plantain Breeding Strategies. 1987:157-60.
- Siddiqah, M. 2000. Biodiversitas dan Hubungan Kekerabatan berdasarkan Karakter Morfologi Berbagai Plasma Nutfah Pisang. *Skripsi*. Jurusan Biologi. Fakultas Matematika dan Ilmu Pengetahuan Alam. Institut Pertanian Bogor. Bogor. 21pp.
- 12. Damayanti F, Roostika I. Koleksi plasma nutfah pisang secara ex vitro dan in vitro serta kajian sitologi dan analisa keragaman antar karakter berdasarkan penanda fenotipe. Faktor Exacta. 2015 Jul 13;3(2):145-57.
- 13. Simmonds NW. *Evolution of the bananas*. Longmans, Green and Co. London. 1962.
- 14. Simmonds NW, Shepherd K. The taxonomy and origins of the cultivated bananas. Botanical Journal of the Linnean Society. 1955 Dec 1;55(359):302-12.

- Irawan, Bobby. Karakterisasi dan Potensi Pisang (Musa spp.) di Bandar Lampung. Skripsi. Jurusan Biologi. Universitas Lampung. 2011; 52 pp.
- 16. Rozyandra C, Darma K. Studi keragaman morfologi aksesi pisang Koleksi dari kabupaten lampung selatan.2006.
- Puspitasari R. Keanekaragaman dan karakterisasi tanaman pisang (*Musa* spp.) di Kabupaten Pesawaran Provinsi Lampung. *Skripsi*. Jurusan Biologi. Fakultas Matematika dan Ilmu Pengetahuan Alam. Universitas Lampung. Lampung. 2011.