Oualitative Characteristics and Genetic Parameters of F3 Purple Corn Lines from Open Pollination Hybridization

10.18196/planta tropika.v11i1.12423

Rima Melati, Eries Dyah Mustikarini*, Gigih Ibnu Prayoga

Study Program Agricultural Science, Faculty of Agriculture, Universitas Bangka Belitung, Kampus Terpadu UBB, Gedung Semangat, Balunijuk, Merawang, Bangka, Province Kepulauan Bangka Belitung, 33172, Indonesia *Corresponding author, email: eriesdyah79@gmail.com

ABSTRACT

Purple corn is a cereal plant that has health benefits. Purple corn contains anthocyanins, which are useful as anticancer and antioxidants, and has high nutritional value. Improving the quality of corn production can be done through plant breeding. This study aimed to determine the value of variability, heritability and expected genetic advance of F₂ purple corn from open-pollinated hybridization. The research was conducted from December 2020 to April 2021 at Universitas Bangka Belitung. The study used a single plant design experimental method. The selection used was ear to row. The results showed that the color of the seeds of the F3 line was dominated by orange. The lines have different quantitative characters. The lines that have the most purple color were F3-PXU-11-25, F3-PXU-6-16, and F3-PXU-11-20. The value of phenotypic variability and genotypic variability of the broad criteria was found in the weight of corn cob with husk and without husk. The broad heritability was high on the weight corn cob with and without husk. The value of the expected genetic advance progress of the high criteria was on the characters of cob length, the weight of 100 seeds, the weight of corn cob with husk, and the weight of corn cob without husk.

Keywords: Expected genetic advance; Heritability; Purple corn; Qualitative; Variability

ABSTRAK

Jagung ungu merupakan tanaman yang memiliki manfaat bagi kesehatan. Jagung ungu memiliki kandungan antosianin yang bermanfaat sebagai anti kanker, antioksidan dan memiliki nilai gizi yang tinggi. Peningkatan kualitas produksi jagung dapat dilakukan melalui pemuliaan tanaman. Tujuan penelitian adalah mengetahui nilai variabilitas, heritabilitas dan keragaman genetik harapan jagung ungu pada generasi Fahasil persilangan bersari bebas. Penelitian dilaksanakan pada bulan Desember 2020 sampai dengan April 2021 di Universitas Bangka Belitung. Penelitian menggunakan metode eksperimen rancangan tanpa ulangan. Seleksi yang digunakan yaitu tongkol kebaris. Hasil penelitian didapatkan warna biji galur F3 didominasi warna orange. Galur memiliki perbedaan karakter kuantitatif. Galur yang memiliki warna ungu terbanyak yaitu F₃-PXU11-25, F₃-PXU6-16, F₃-PXU11-20. Nilai variabilitas fenotip dan variabilitas genotip kriteria luas terdapat pada karakter bobot tongkol dengan kelobot dan tanpa kelobot. Nilai heritabilitas arti luas kriteria tinggi pada karakter bobot tongkol dengan kelobotdan bobot tongkol tanpa kelobot. Nilai kemajuan genetik harapan kriteria tinggi pada karakter panjang tongkol, bobot 100 biji, bobot tongkol dengan kelobot, dan tanpa kelobot.

Kata Kunci: Kemajuan Genetik Harapan; Heritabilitas; Jagung Ungu; Kualitatif; Variabilitas

INTRODUCTION

Corn plants have a relatively high genetic diversity in seeds. The color of the corn kernels of plant genetic potential (Azrai, 2016). Plant breedis genetically controlled. The color of the corn ing can be done by crossbreeding plants to obtain kernels consists of purple, red, yellow, and white the desired characteristics (Sain, 2016). Previous (Pamandungan & Ogie, 2018). Purple corn con- researchers have carried out corn plant breeding. tains anthocyanins, which function as antioxidant Selected purple corn has no resistance to corn stem compounds beneficial to prevent cancer, diabetes, borer (Ostrianiafurnacalis G.) (Oktaviani, 2017). and coronary heart disease (Tumei et al., 2018). Improvement of resistance characters was carried Therefore, it is necessary to assemble purple corn, out by crossing 'Marassempulu' purple corn with which has a high anthocyanin content.

open

access

Plant breeding aims to increase the expression 'Magetan' white corn and 'Sungailiat' yellow corn



Article History Received: 28 August 2022 Accepted: 14 February 2023



Planta Tropika: Jurnal Agrosains (Journal of Agro Science) is licensed under a Creative Commons Attribution-NonCommercial 4.0 International License.

(Yahya, 2018). F1 selection results obtained 14 lines Universitas Bangka Belitung. (Safitri, 2019), and F2 selection obtained 9 corn lines (<u>Abadi et al., 2021</u>).

had high diversity. Genetic variance and diversity lines used, namely F2-PxU-11-14, F2-PxU-11-20, F2of F2 corn plant populations were found (Sudika PxU-11-21, F2-PxU-11-2, F2-PxU-6-15, F2-PxU-6-16, et al., 2022). The estimates of genetic effects traits F2-PxU-11-13, F2-PxU-11-11, and F2-PxU-11-25. were specific in eachgenotypes (Nabila et al., 2022). The control varieties were hybrid parents (purple According to Juita et al. (2022), the F2 and F3 lines corn Marassempulu from South Sulawesi and from free pollination crosses had better characters white Magetan corn from East Java). This research than their parents, in which 88.11% of the lines applied distance and time isolation. The study was had purple seed color.

row selection is used to select segregated plant pop- planting time was 2 weeks. Each plot was planted ulations. Segregated plants have more dominant with 30 plants. diversity between families (Hallauer et al., 1988). tion (Lasmono et al., 2018). High genetic progress x 20 cm; each planting hole was given 1 seed with a results from character improvement through selec- 2-3 cm planting depth. Fertilization was performed tion (Wulandari et al., 2016). The lines have differ- when the plants were 14 and 30 days after planting, genetic diversity.

It is expected that the results of this study will color of the corn husks was brown. produce F4 lines of purple corn that have wide variability, high hertability, and high genetic prog- and quantitative characters. Qualitative characters ress. The F4 lines with high anthocyanin content consist of leaf shape, leaf color, panicle shape, top will continue in the next generation selection. The seed surface shape, seed row shape and seed color aim of the study was to determine the diversity of distribution. Quantitative characters consist of qualitative characters, the variability value, the plant height, age of flowering plants, weight of heritability value and the genetic progress of the 100 seeds, stem diameter, weight of corncob with F3 purple corn lines as a result of open pollination. husks, harvest age, number of leaves per plant,

MATERIALS AND METHOD

The research was conducted from December

The research was arranged in an experimental method with a single plant design. The selection The lines produced from free-pollinated crosses method used was ear to row. There were 9 purple cord conducted in 4 locations with a distance of 20 m. Selection is made to get superior lines. Ear-to- A total of 11 maps were made. The difference in

The stages of the research included land prepa-The selection success is determined by genetic di- ration, making plots, selecting seeds, soaking the versity, heritability, and correlation between traits seeds, planting, watering, fertilizing, controlling (Privanto et al., 2017). High heritability values are pests, and harvesting. The research map was made important in increasing the effectiveness of selec- with a size of 3.4 x 1.5 m. Plant spacing was 75 cm ent quantitative characters indicate the presence of with the NPK fertilizer dose of 450 kg/ha. Harvesting was done when the corn cob looked dry, or the

> The observed characters included qualitative corncob length, weight of corncob without husk, and number of seed rows.

Qualitative data was written descriptively. 2020 to April 2021. The research location was in Quantitative data were used to calculate the value the Research and Experimental Garden (KP2) of of variability, heritability and expected genetic the Faculty of Agriculture, Fisheries and Biology, progress. The value of genetic variance, environmental variance and phenotype variance are three analysis of variance.

important components in finding variability values. According to <u>Suharsono et al. (2006)</u>, this value is obtained using the formula:

Genotype
variance
$$(\sigma_g^2) = \sigma_{f}^2 \sigma_e^2$$
 (1)

Phenotype variance $(\sigma_{f}^{2}) = \frac{\sum_{i=1}^{N} (xi - \mu)^{2}}{n}$ (2)

Environment variance $(\sigma_e^2) = \frac{\sigma_{2p1+\sigma_{2p2}+\dots+\sigma_{2pn}}}{n}$ (3)

Formula description:

 σ_{P1}^{2} = Variance of the female parent σ_{P2}^{2} = Variance of the male parent σ_{g}^{2} = Genotype Variance

Determining the phenotypic coefficient of diversity (PCD) and genetic coefficient of diversity (GCD):

%PCD =
$$\frac{\sqrt{\sigma_{\rm f}^2}}{\mu} x \, 100$$
 (4)

$$\% \text{GCD} = \frac{\sqrt{\sigma_{\text{g}}^2}}{\mu} x \ 100 \tag{5}$$

Formula description:

 σ_{f}^{2} = Phenotype variance σ_{e}^{2} = Environment variance μ = mean

n = population

According to <u>Moedjiono and Mejaya (1994)</u>, the criteria for the PCD and GCD are divided into four criteria in percentage form, namely:

0-25% : Narrow 25-50%: Rather narrow 50-70%: Rather wide 75-100%: wide

according to <u>Syukur et al. (2012)</u>, estimated value of broad sense heritability h²(BS) is based on

h²(BS) =
$$\frac{\sigma^2 F3 - \sqrt{(\sigma^2 P1)(\sigma 2 P2)}}{\sigma^2 F3} X100\%$$
 (6)

Formula description:

h² (BS): Broad sense heritability σ_{F3}^{2} = Variance F₃ σ_{P1}^{2} = Variance of the female parent σ_{P2}^{2} = Variance of the male parent

The criteria for heritability values are as follows: Low = $h^2(BS) < 20\%$ Moderate = $20\% \le h^2(BS) \ge 50\%$ High = $h^2(BS) > 50\%$

Expected Genetic Advance value (EGA) can be calculated using the formula:

 $EGA = i.h^{2}(BS).\sigma p$ (7)

$$\% EGA = \frac{EGA}{\mu} \times 100\%$$
 (8)

Formula description:

EGA = Expected Genetic Advance

= Selection intensity (10% = 1,76)

h²(BS)= Heritability

 σp = Phenotypic standard deviation

μ = Mean

The expected genetic advance criteria are as follows:

 $0 < KGH \le 3.3\%$ = Low 3,3% < KGH $\le 6,6\%$ = Rather low 6,6% < KGH $\le 10\%$ = High enough KGH > 10% = High

RESULTS AND DISCUSSION

The F3 generation of the purple corn lines resulted from crossing purple and white corn, with 330 plants studied. Selection was made to get the selected F4 lines. The qualitative characters observed in the F3 line of corn plants consisted of

	Qualitative Characters							
Lines	Leaf Shape	Leaf Color	Panicle shape	Top Seed Surface Shape	Seed RowShape			
F3-PxU11-14	100% Spiky	100% Moderate Olive Green	5.6%Very Small	27.8%Wrinkled	22.3%Regular			
			27.8%Small	38.8%Serrated	33.3% Irregular			
			55.5%Medium	16.7%Flat	44.4%Straight			
			11.1%Large	16.7%Round				
F3-PxU11-20	100% Spiky	100% Moderate Olive Green	3.7%Very Small	3.7%Wrinkled	37.1%Regular			
			29.7%Small	51.8%Serrated	33.3% Irregular			
			33.3%Medium	37.1%Flat	25.9%Straight			
			33.3%Large	3.7%Round	3.7%Curved			
				3.8%Tapered				
F3-PxU11-21	100% Spiky	10% Moderate Olive Green	6.7%Very Small	40%Serrated	40%Regular			
		90% Greyish Olive Green	26.7% Small	40% Flat	30% Irregular			
			46.7% Medium	20% Round	30% Straight			
			16.7% Large					
			3.3% Very Large					
F3-PxU11-2	88.9% Spiky	94.4% Moderate Olive Green	5.5% Very Small	11.1% Wrinkled	27.8% Regular			
	11.1% Pointed Slightly rounded	5.6% Strong Yellow Green	61.1% Small	61.2% Serrated	50% Irregular			
			16.7% Medium	11.1% Flat	22.2% Straight			
	40001 0 1		16.7% Large	16.6% Round				
F3-PxU6-15	100% Spiky	89.5% Moderate Olive Green	5.3% Very Small	5.3% Wrinkled	15.8% Regular			
		10.5% Strong Yellow Green	10.5% Small	42.1% Serrated	57.9% Irregular			
			52.6% Medium	10.5% Flat	21.1% Straight			
			31.6% Large	36.8% Round 5.3% Tapered	5.2% Curved			
F3-PxU6-16	90% Spiky	70% Moderate Olive Green	45% Small	10% Wrinkled	50% Regular			
	10% Pointed and Slightly	30% Greyish Olive Green	30% Medium	25% Serrated	50% Irregular			
	Tounded		25% Large	30% Flat				
				30%Round				
				5%Tapered				
F3-PXU11-13	100% Spiky	96.4% Moderate Olive Green	3.6% Very Small	7.1% Wrinkled	35.7% Regular			
		3.6% Strong Yellow Green	28.7% Small	85.7% Serrated	10.7% Irregular			
			32.1% Medium	3.6% Flat	46.4% Straight			
			35.6% Large	3.6% Round	7.2% Curved			
F3-PxU11-11	100% Spiky	91.3% Moderate Olive Green	4.3% Very Small	30.4% Serrated	26.1% Regular			
		8.7% Strong Yellow Green	26.1% Small	39.2% Flat	43.5% Irregular			
			56.5% Medium	30.4% Round	30.4% Straight			
			13.1% Large					
F3-PxU11-25	69.2% Spiky	46.2% Moderate Olive Green	26.9% Small	61.6% Serrated	53.9% Regular			
	rounded	53.8% Greyish Olive Green	50% Medium	34.6% Flat	11.5% Irregular			
			23.1% Large	3.8% Round	34.6% Straight			
Parent	44.8% Spiky	82.8% Moderate Olive Green	3.4%Very Small	13.7%Serrated	37.8% Regular			
'Marassempulu'	55.2%Pointed Slightly rounded	17.2%Strong Yellow Green	31.1%Small	55.2%Flat	31.1% Irregular			
			37.9%Medium	31.1%Round	31.1% Straight			
			27.6%Large					
Parent	58.6% Spiky	89.7% Moderate Olive Green	6.9%Very Small	41.4%Serrated	34.5%Regular			
'Magetan'	41.4% Pointed and Slightly rounded	10.3% Greyish Olive Green	31.1%Small	48.3%Flat	34.5% Irregular			
			58.6%Medium 3.4% VeryLarge	10.3%Round	31% Straight			

Table 1. Qualitative characters of the F3 generation of corn lines and parents

.....

64

								·	· · ·		
	Lines Parents								ents		
Color Seed	F3-PxU11-14	F3-PxU11-20	F3-PxU11-21	F3-PxU11-2	F3-PxU6-15	F3-PxU11-16	F3-PxU11-13	F3-PxU11-11	F3-PxU11-25	'Marassempulu'	'Magetan'
Purple	5	8	0	4	4	11	7	2	14	29	0
Yellow	5	5	10	1	7	7	7	5	1	0	0
Orange	2	13	20	11	8	2	11	14	10	0	0
Red	6	1	0	2	0	0	3	2	1	0	0
White	0	0	0	0	0	0	0	0	0	0	29

Table 2. Qualitative characters of seed color distribution in the F3 generation of corn plant lines and parent

Table 3. Variance Value of Phenotypes, Genotypes and Environment

Characters	Variance Value of Phenotypic	Genotypes	Environment	
Plant Height	966.19	226.22	739.98	
Stem Diameter	9.63	1.95	7.68	
Number of Leaves per Plant	1.50	0.19	1.30	
Age of Male Flowering Plants	9.65	1.95	7.70	
Age of Female Flowering Plants	10.13	1.51	8.62	
Harvest Age	25.57	3.24	22.32	
Corncob Length	8.56	3.34	5.22	
Weight of Corncob with Husk	3060.61	2282.64	777.97	
Weight of Corncob without Husk	2370.67	1747.20	623.47	
Weight of 100 Seeds	82.28	27.44	54.83	
Number of Seed Rows	3.72	1.05	2.67	

6 characters: leaf shape, leaf color, panicle shape, wrinkled (13), serrated (104), flat (54), round (35), are identified based on CPVO (2020).

Leaf tip shape characters are grouped into five criteria. The shape of the leaves is divided into two seed surface, thought to be inherited from the criteria, namely sharp and slightly rounded, and white parents. The shape of the seed row shape is the ratio is 197:12. Leaf shape with pointed, slightly influenced by the various characteristics of the two rounded criteria has more dominant results. The parents. Differences in plant characteristics are due shape of the pointed leaves is based on the parent to the maternal effect. According to Pamandungan of white corn. Leaf color characters consist of three & Ogie (2018), the maternal effect occurs when the groups, namely moderate olive green (56), greyish female parent's nuclear genotype determines the olive green (47), and strong yellow-green (6). The offspring's phenotype. panicle shape characters consist of five criteria, namely very small (8), small (64), medium (87), various seed color distributions. The seed color large (49), and very large (1). The characters of the obtained was grouped into purple, orange, yellow, seed surface shape consist of five criteria, namely and red. The orange color showed the highest ratio

top seed surface shape, seed row shape and seed and tapered (3). The seed row shape characters concolor distribution (Table 2). Qualitative characters sist of four criteria, namely regular (74), irregular (70), straight (61), and curved (4) (Table 1).

Serrated criteria dominate the shape of the top

The F3 generation of purple corn lines has

Karakter	Phenotypic Variability (%)	Criteria	Genotypic Variability (%)	Criteria	Heritability Broad Sense	Criteria	Expected Genetic Advance	Criteria
Plant Height	17.45	Rather narrow	8.44	Narrow	23.41	Medium	7.19	High enough
Stem Diameter	21.39	Rather narrow	9.61	Narrow	20.20	Medium	7.60	High enough
Number of Leaves Per Plant	10.47	Narrow	3.79	Narrow	13.15	Low	2.42	Slightly Low
Age of Male Flowering Plants	6.98	Narrow	3.14	Narrow	20.24	Medium	2.49	Slightly Low
Age of Female Flowering Plants	6.61	Narrow	2.55	Narrow	14.10	Low	1.76	Low
Harvest Age	6.16	Narrow	2.21	Narrow	12.69	Low	1.38	Low
Corncob Length	21.93	Rather narrow	13.70	Rather narrow	39.04	Medium	15.07	High
Corncob Weight with Husk	48.01	Broad	41.46	Wide	74.58	High	63.02	High
Corncob Weight without Husk	49.45	Broad	42.45	Wide	73.70	High	64.14	High
Weight of 100 Seeds	36.16	Broad Enough	20.88	Rather narrow	33.35	Medium	21.22	High
Number of Seed Rows	16.47	Rather narrow	8.76	Narrow	28.32	Medium	8.21	High enough

Table 4. Phenotypic variability values, genotypic variability values, heritability values, and expected genetic advance values

Remarks: 1. Phenotypic variability criteria: 0-12.4 (Narrow), 12.4-24.7 (Rather narrow), 24.7-37.1 (Broad Enough), 37.1-49.5 (Broad). Genotypic variability criteria: 0-10.6 (Narrow), 10.6-21.2 (Rather narrow), 21.2-31.8 (Broad Enough), 31.8-42.4(Broad).

2. Criteria for heritability value: $h^{2}(BS) < 20\%$ (Low), 20% $\leq h^{2}(BS) \geq 50\%$ (Medium), $h^{2}(BS) \geq 50\%$ (High).

3. Criteria for Expected Genetic Advance (EGA): 0 <EGA ≤ 3.3% (Low), 3.3% <EGA ≤ 6.6% (Slightly Low), 6.6% <EGA ≤ 10% (High enough) and EGA> 10% (high).

tions with ratios can be seen in Table 4. The lines flowering plants, and the age of harvest. The relawith the most purple color were F3-PXU11-25, tively narrow criteria were found in the character F3-PXU6-16, and F3-PXU11-20. The purple color of plant height, stem diameter, corncob length, came from one of the elders. The color that appears and the number of seed rows. The broad enough indicates the influence of the xenia effect. Accord- criteria were found in the character of the weight ing to Ishartati et al. (2020), the xenia effect shows of 100 seeds, while the broad criteria were found that the pollen of the male parent has a direct and in the character of the weight of corncob with and specific impact on the character of the seeds. The without husks (Table 4). emergence of colors such as yellow and orange in corn is thought to be due to open pollination in the variability value is divided into 4 criteria, namely previous generation. The pollination results often 0.25% (narrow), 25-50% (rather narrow), 50-70% get the same color in one corncob even though (broad enough), and 75-100% (broad). The phenothe cross parents differ. According to <u>Hariyanti et</u> typic and genotype variability value was adjusted by al. (2014), seed color is caused by dominance by setting 49.45% of the largest value as 100% diverdominant color control genes. The presence of one sity. Therefore, the criteria for phenotypic variabil-

from narrow to broad criteria. The narrow criteria Therefore, the criteria for genotypic variability are were found in the number of leaves per plant, the 0-10.6 (narrow), 10.6-21.2 (rather narrow), 21.2-31.8

compared to the other colors. Complete varia- age of male flowering plants, the age of female

According to Moedjiono & Mejaya (1994), the dominant gene can express color and dominance. ity are 0-12.4 (narrow), 12.4-24.7 (rather narrow), Phenotypic variability values were obtained 24.7-37.1 (broad enough), and 37.1-49.5 (broad).

(broad enough), and 31.8-42.4 (broad).

have very little or almost uniform diversity. The (Wulandari et al., 2016). According to Sari & Sugibroad phenotypic variability value was found in <u>harto (2018)</u>, a high heritability value will make the the characters of the weight of corncob with and without husks. According to Napitupulu & Damanhuri (2018), the low phenotypic variability value indicates that individuals in the population tested tend to be uniform. High phenotypic variability values indicate a high level of diversity. According values were found in the characters of the age of to Amoros et al. (2020), a high phenotypic variability value indicates a large environmental influence on this character.

environment influence the variety of phenotypes. the characters of plant height, stem diameter, and Genetic variety has a negative correlation with the number of seed rows. Meanwhile, the high environmental variation (Table 3). If plants have a was obtained in the characters of long length, the broad genetic variability value, the environmental weight of the corncob with and without husks, and variability value is narrow/low. Narrow genetic the weight of 100 seeds (Table 4). variability values were found in the characters of plant height, stem diameter, number of leaves tor of success in plant selection. High expected per plant, age of male flowering, age of harvest, and the number of seed rows. The value of broad genotypic variability was found in the characters According to Ibrahim et al. (2018), a good selection of the weight of corncob with and without husks (Table 4). Broad variability values can increase the ity values and genetic advance values. According chances of obtaining plants with good properties (Joshi et al., 2018). Variability with broad criteria can increase the selection response (Effendy et al., <u>2018</u>).

The low criteria of heritability values were obtained in the characters of the number of leaves per plant, the age of female flowering, and the advance values guide plant breeders in selecting. age of harvest. Medium criteria were obtained in Genetics influences inheritance and gene expresplant height, stem diameter, age of male flowering plants, corncob length, the weight of 100 seeds, and number of seed rows. The high criteria were useful for estimating how many stages of selection obtained in the characters of the weight of the corn must be carried out. According to Miftahorrachcob with and without husks (Table 4). Characters man (2010), characters that have low variability with high heritability values indicate that they are values indicate genetic homogeneity. Improving

more influenced by genetic factors. High heritabil-Narrow variability values indicate lines that ity values can be passed on to the next generation selection more effective. Selection is more effective because the influence of the environment is very small, so genetic factors are more dominant in the appearance of plant phenotypes.

The low criteria of the expected genetic advance female flowering plants and the age of harvest. The rather low criteria were obtained in the number of leaves per plant and the age of the male flowering The values of the variety of genotypes and the plants. The high enough criteria were found in

> Expected genetic advance value is an indicagenetic advance values are directly proportional to heritability values for several plant characters. of genotypes can use characters with high heritabilto Kristamtini et al. (2016), a high value of genetic advance in a character indicates that genetic factors support the character. The value of genetic advance can be a guideline for selection in the next generation.

> Variability, heritability, and expected genetic sion tends to be affected by the environment. The values of heritability and genetic advance are very

character through plant breeding tends to be difficult. According to <u>Hadiati et al. (2003)</u>, broad genetic variability will provide a higher chance of selection.

Characters with broad genetic variability and phenotypic variability will increase the value of genetic advance. Plant breeders have a high chance of obtaining the desired genotype (Sari & Susilo, 2013). According to Muin(2021), a high heritability value indicates that genetic factors have a large influence. Selection of corn plants can be stopped if they have high expected genetic advance values, broad heritability values, and narrow variability. The selected purple corn lines are expected to have characteristics that correlate well with production so that high-yielding purple corn can be obtained. Selection needs to be continued because the plants show broad criteria of variability values on characters that are correlated with production.

CONCLUSION

The results showed that the seeds of the F3 lines were dominated by orange color. The lines have different characters of leaf shape, leaf color, panicle shape, top seed surface shape, seed row shape and seed color distribution. The lines that had the most purple color were F3-PXU-11-25, F3-PXU-6-16, and F3-PXU-11-20. The weight of corncob with and without husk showed broad criteria of variability value and high criteria of broad sense heritability value. The high expected genetic advance value was found in the characters of corncob length, weight of corncob with and without husk, and weight of 100 seeds.

ACKNOWLEDGMENT

Acknowledgments are conveyed to the Universitas Bangka Belitung for the research facilities provided in experimental and research fields.

REFERENCES

- Abadi, H. K., Mustikarini, E. D., & Prayoga, G. I. (2021). Parameter Genetik Hasil Persilangan Jagung Bersari Bebas untuk Mendapatkan Galur Berbiji Ungu. Jurnal Ilmu Pertanian Indonesia, 26(3), 450–458. <u>https://doi.org/10.18343/jipi.26.3.450</u>.
- Amoros, W., Salas, E., Hualla, V., Burgos, G., De Boeck, B., Eyzaguirre, R., zum Felde, T., & Bonierbale, M. (2020). Heritability and genetic gains for iron and zinc concentration in diploid potato. *Crop Science*, 60(4), 1884–1896. <u>https://doi.org/10.1002/ csc2.20170</u>
- Azrai, M. (2016). Pemanfaatan markah molekuler dalam proses seleksi pemuliaan tanaman. Jurnal AgroBiogen, 1(1), 26–37.
- CPVO[Community Plant Variety Office]. (2020). Plant Variety and Cultiver Identification: Advances and Prospects. <u>https://cpvo.</u> <u>europa.eu/</u>.
- Effendy, E., Respatijarti, R., & Waluyo, B. (2018). Keragaman genetik dan heritabilitas karakter komponen hasil dan hasil ciplukan (Physalis sp.). Jurnal Agro, 5(1), 30–38. <u>https://doi.org/10.15575/1864</u>.
- Hadiati, S., Mudaningsih, Baihaki, H.K.A., & Rostini, N. (2003). Parameter Genetik KarakterKomponenBuah padaBeberapa Aksesi Nanas. Zuriat, 14:47-52.
- Hallauer, A. R., Carena, M. J., & Filho, J. B. M. (1988). Quantitative Genetics in Maize Breeding. In *Lowa State University Press*, *Ames, Lowa*.
- Hariyanti, I. D., Soegianto, A., & Sugiharto, A. N. (2014). Hariyanti, I.D. (2014). Efek Xenia pada Beberapa Persilangan Jagung Manis (Zea mays L. Saccharata) Terhadap Karakter Biji. [Skripsi]. Jurusan Budidaya Pertanian. Universitas Brawijaya, Malang.
- Ibrahim, A. K., Dawadi, K. D., & Hassan, S. M. (2018). Genetic Variability, Heritability And Correlation Among Soybean [*Glycine* max. (L.) Merrill] Varieties. BAJOPAS: Bayero Journal of Pure and Applied Sciences, 11(2), 72–80.
- Ishartati, E., Sufianto, Fadjri, I. A., Jana, M., & Priadi, A. (2020). Efek Pollen Pada Persilangan Jagung Hitam, Pulut (*Zea mays* ceratina), dan Manis (*Zea mays saccharata*) Terhadap Karakter Biji. *Gontor AGROTECH Science Journal*, 6(3), 499–515.
- Joshi, D., Singh, K., & Adhikari, S. (2018). Study of Genetic Parameters in Soybean Germplasm Based on Yield and Yield Contributing Traits. Int. J. Curr. Microbiol. App. Sci, 7(01), 700–709. <u>https:// doi.org/10.20546/ijcmas.2018.701.085</u>
- Juita, R., Pamandungan, Y., & Lengkong, E. F. (2022). Karakterisasi Tanaman Jagung Ungu F2 Dan Biji F3 (Zea mays L.) Hasil Bersari Bebas Jagung Manado Kuning Dengan Jagung Ungu. Jurnal Agroekoteknologi Terapan, 3, 63–74.
- Kristamtini, Sutarno, Wiranti, E. W., & Widyayanti, S. (2016). Kemajuan Genetik dan Heritabilitas Karakter Agronomi Padi Beras Hitam pada Populasi F2. *Penelitian Pertanian Tanaman Pangan*, 35(2), 119–124.
- Lasmono, G, Sugiharto, A.N., & Respatijati. (2018). Pendugaan Nilai Haritabilitas, Keragaman Genetik dan Kemajuan Genetik Harapan pada Beberapa Genotipe F5 Cabai (*Capsicum annuum*L) *Jurnal Produksi Tanaman*, 4(6), 668-667.
- Miftahorrachman. (2010). Korelasi dan Analisis Sidik Lintas Karakter Tandan Bunga TerhadapBuah Jadi Kelapa Genjah Salak. *Buletin Palma*, 38, 60–66.

- Muin, A. (2021). Evaluasi Pertumbuhan Uji Keturunan Meranti Tembaga (*Shorea leprosula*) Umur12 Tahun diAreal IUPHHK-HA PT. Erna Djuliawati Kalimantan Tengah. *Jurnal Tengkawang*, 11(2), 87–97.
- Moedjiono, & Mejaya M.1994. Variabilitas Genetik beberapa Karakter Plasma Nutfah Jagung Koleksi Balittan Malang. *Zuriat*, 5(2): 27-32.
- Nabila, N., Wahyu, Y., & Widodo, W. D. (2022). Determination of Genetic Parameters and Inbreeding Depression of Half-Sib and Selfing Families to Developed as Baby Corn Variety. *Current Topics in Agricultural Sciences Vol. 6, March*, 58–65. <u>https:// doi.org/10.9734/bpi/ctas/v6/3525e</u>
- Napitupulu, M., & Damanhuri. (2018). Keragaman Genetik, Fenotipe Dan Heritabilitas Pada Generasi F2 Hasil Persilangan Tanaman Padi (Oryza sativa L.). Jurnal Produksi Tanaman, 6(8), 1844–1850.
- Oktaviani. (2017). Serangan Hama Penggerek Batang Jagung (Ostrinia furnacalis Guenee) (Lepidoptera : Crambidae). Program Studi Agroteknologi. [Skripsi]. Universitas Bangka Belitung.
- Pamandungan, Y., &Ogie, T.M. (2018). Pewarisan Sifat Warna dan Tipe Biji Jagung ManadoKuning. *Eugenia*, 24(1), 1-8.
- Priyanto, S. B., Azrai, M., & Syakir, M. (2017). Analisis Ragam Genetik, Heritabilitas, dan Sidik Lintas Krakter Agronomik Jagung Hibrida Silang Tunggal.. *Informatika Pertanian*, 27(1), 1–8.
- Safitri, F. (2019). Analisis Heritabilitas dan Variabilitas Generasi F₁ Hasil Persilangan Jagung Bersari Bebas. [Skripsi]. Fakultas Pertanian Perikanan dan Biologi.Universitas Bangka Belitung.
- Sain, A. (2016). Keragaman Genetik Empat Varietas Jagung (Zea Mays. L) Bersari Bebas Menggunakan Marka Ssrs (Simple Squence Repeats). [Skripsi]. Universitas Islam NegeriAlauddin Makasar 1–91.
- Sari, E. N., & Sugiharto, A. N. (2018). Keragaman Beberapa Galur Jagung Pakan (*Zea mays* L.) Generasi S₇. Jurnal Produksi Tanaman, 6(1), 56–65.
- Sari, I. A., & Susilo, A. W. (2013). Pengembangan Kriteria Seleksi Karakter Berat Biji pada Tanaman Kakao (Theobroma cacao L.) melalui Pendekatan Analisis Sidik Lintas Development of Selection Criteria on Bean Weight Character of Cocoa (Theobroma cacao L.) through Path Analysis Approach. *Pelita Perkebunan*, 29(3), 174–181.
- Sudika, I. W., Sutresna, I. W., & Anugrahwati, D. R. (2022). Estimation of Genetic Variance and Populations of Corn Plants in Dry Land Heritability of. 8, 117–123. <u>https://doi.org/10.29303/jppipa.</u> v8iSpeciallssue.2481
- Suharsono, Jusuf, M., &Paserang, A.P. (2006). Analisis Ragam, Heritabilitas, danPendugaanKemajuanSeleksi Populasi F₂ dari Persilangan KedelaiKultivar Slamet danNokonsawon. Jurnal Tanaman Tropika 11(2): 86-93.
- Syukur, M., Sujiprihati, S., &Yuniati, R. (2012). *Teknik Pemuliaan Tanaman*. Jakarta: Penebar Swadaya.
- Tumei, O. D., Marjam Toding, & Pamandungan., Y. (2017). Karakterisasi Tanaman Jagung Ungu F1 Hasil Bersari Bebas Jagung Manado Kuning dengan Jagung Ungu. *Pendidikan Kimia PPs* UNM, 1(1), 91–99.
- Wulandari, J. E., Yulianah, I., & Saptadi, D. (2016). Heritabilitas dan Kemajuan Genetik Harapan Empat Populasi F2 Tomat (Lycopersicum esculentum Mill.) PADA BUDIDAYA ORGANIK. Jurnal

Produksi, 4(5), 361-369.

Yahya, R. (2018). Karakterisasi Tetua dan Hibridisasi Jagung Tahan Penggerek Batang Melalui Penyerbukan Bersari Bebas. [Skripsi]. Jurusan Agroteknologi, Universitas Bangka Belitung. Balunijuk.