

ABSTRAK

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KeragamanGenetikGalur-galurKacang Bambara
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Kacangbambaramerupakanantamankacang-kacangan ketigaterpenting di Indonesia setelah kacang tanah dan kacang tunggak, namun belum terlaludiperhatikan di Indonesia tetapi memilikiperan dalam program diversifikasi pangan. Salah satumasalah penting yang perludiketahuidalam peningkatan produksi kacang bambara di Indonesia adalah perlunya kajian tentang galur-galur kacang bambara sebagai bahan nutrimental dan tanaman.

Keragamangenetiksangatmempengaruhikeberhasilansuatu proses seleksidalam program pemuliaantanaman. Selainitu, perlujugadiketahuinilaiheritabilitasdankemajuanngeneticharapankarakter-karakter yang akandijadikan target seleksiguna mengetahuikemajuanseleksi. Penelitianinibertujuanmengetahuidanmenentukankeragamangeneticberdasarkansi fatpolongdanbijisebagaisumbergenetikuntukseleksilebihlanjutkacangbambara. PenelitiandilaksanakanselamaduabulanmulaidaribulanNopemberinggaDesember 2017.Penelitianinidilakukan di Laboratorium*Bambara Groundnut Research Centre* (BGRC) FakultasPertanianUniversitasMuhammadiyah Gresik.Rancanganpercobaan yang digunakanadalahRancanganAcakLengkap (RAL) dengansatufaktoryaitugenotipakacangbambarasebanyak 3 genotipadenganjumlahulangan 31. Variabelpenelitian yang diamatimeliputivariabelkuantitatifdankualitatif.Variabelkuantitatifterdiridarijumlah; bobotkereng; diameter polong, ketebalankulitpolong, diameter; bobotkereng; jumlahbiji, anova, ujilanjut BNT_{0.05}, ujiberabilitas, nilai koefisien keragamangenotip dan fenotip, keragamangenetikdankemajuanngeneticharapan; karakter kualitatifterdiridari bentuk; warna; tekstur polong, warna testa, bentuk biji, warna hilum, bentuk hilum, corak sekitar hilum, janggut pada biji. Hasil penelitian menunjukkan bahwa nilai heritabilitas menunjukkan sedang dan rendah. Rendahnya nilai heritabilitas pada variabel kuantitatif dipengaruhi oleh faktor lingkungan. Nilai keragamangenetik harapan tertinggi adalah 38% pada karakter bobot kereng biji. Dengan demikian karakter bobot kereng biji dapat dijadikan acuan untuk seleksi lebih lanjut pada tanaman kacang bambara.

Kata kunci: *heritabilitas; kacangbambara; kemajuangenetik; keragamangenetik; koefisienkeragaman.*

ABSTRACT: Bambara groundnut are the third most important legume crop in Indonesia after peanuts and cowpea, but have not been given much attention in Indonesia but have a role in food diversification programs. One important problem that needs to be known in increasing the production of bambaragroundnut in Indonesia is the need for a study of the strains of bambara beans as the main ingredient of plant breeding. Genetic diversity greatly influences the success of a selection process in a plant breeding program. In addition, it is also necessary to know the heritability value and genetic progress of the characters' expectations that will be targeted for selection in order to know the progress of the selection. This study aims to determine and determine genetic diversity based on the nature of the pods and seeds as a genetic source for further selection of bambaragroundnut. The study was conducted for two months from November to December 2017. The research was conducted at the Bambara Groundnut Research Center (BGRC) Laboratory of the Faculty of Agriculture Muhammadiyah University Gresik. The experimental design used was a Completely Randomized Design (CRD) with one factor, namely genotipabambaragroundnut totaling 3 genotipas with 31 replications. The observed research variables consisted of quantitative and qualitative variables. Quantitative variables consist of quantities; dry weight; pod diameter, pod thickness, diameter; dry weight; number of seeds, anova, 5% LSD test, heritability test, coefficient value of genotype and phenotype diversity, genetic diversity and genetic progress expectations; Qualitative character consists of forms; color; pod texture, testa color, seed shape, hilum color, hilum shape, style around hilum, beard on seeds. The results showed that the heritability was moderate and low. The low value of heritability on quantitative variables is influenced by environmental factors. The highest expected genetic advancement value is 38% in the dry weight character of the seeds. Thus the dry weight character of the seed can be used as a reference for further selection in the bambaragroundnut plant.

Keywords: *heritability; bambaragroundnut; genetic progress; genetic diversity; coefficient of diversity.*