

Genetics Of Flowering And Quantitative Traits In Vigna Unguiculata: Days To Flowering In Cowpea By Williams Manggoel

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variation and heritability of sixteen characters - Plant Breeding and Seed Science Number of days to flowering, and Phenotypic variability for seed yield and other traits in cowpea (Vigna unguiculata (L)

quantitative genetic analysis of flowering time - 1. Genome. 2007 Mar;50(3):303-15. Quantitative genetic analysis of flowering time in tomato. Jim nez-G mez JM, Alonso-Blanco C, Borja A, Anastasio G, Angosto T

engineering plants for nematode resistance - - Serine Protease Inhibitors. The potential of PIs as antinematode effectors was first explored for the serine PI cowpea trypsin inhibitor (CpTI). CpTI has been shown

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resistance to pathogens and host developmental - This type of resistance may also be referred to as flowering Cowpea (*Vigna unguiculata* Cg1 mid-whorl leaves continue to display juvenile traits and

temperature responses of developmental processes - They also allow the study of quantitative genetics based on between days and plants to avoid any bicolor) and cowpea (*Vigna unguiculata*),

crop breeding and applied biotechnology - path - Cowpea (*Vigna unguiculata* and number of days to flowering may lead to decreases in for yield and some quantitative traits in Iran cowpea

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construction of a genetic linkage map and genetic - 34 were treated as quantitative traits and four shortening of days to flowering, *Vigna radiata* L. Wilczek) and cowpea (*V. unguiculata* L

genetics of flowering and quantitative traits in - The Nigerian cowpea accessions differ in phenology with respect to flowering and other quantitative traits.

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quantitative genetics - stanford university - There is a really cool paper by Buckler and colleagues in the current issue of Science. The basic gist of this paper is that flowering time in maize (*Zea mays*) is not

issuu - cowpeas by international institute of - This paper presents information on the major crops grown in the Savannah. These crops include cowpea, sorghum, millet, soybeans, maize, wheat and cotton.

genomics and plant breeding - sciencedirect - Much of our most basic understanding of genetics has its roots in plant genetics and crop breeding.

citeseerx l. - Mendelian factors underlying quantitative traits in tomato: A genetic linkage map of cowpea (*Vigna unguiculata*) Nature Genetics 14 - Frankel,

cowpea - springer - Cowpea (*Vigna unguiculata* (2013) Quantitative trait loci analysis of flowering time related traits identified in recombinant The genetics of cowpea:

genetic architecture of flowering time in maize - Abstract. Genetic architecture of flowering time in maize was addressed by synthesizing a total of 313 quantitative trait loci (QTL) available for this

genetic variability for yield and yield component - relatively less effort has been made to understand the genetics of important quantitative traits days to 50% flowering, cowpea [*Vigna unguiculata*]

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