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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

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

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

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climate change and the flowering time of annual - of the impacts of climate change on the flowering and cowpea (Vigna unguiculata)] and long In quantitative types, flowering is delayed but

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.

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

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qtl analysis of flowering time in arabidopsis - Quantitative trait loci (QTL) analyses based on restriction fragment length polymorphism maps have been used to resolve the genetic control of flowering time in a

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

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

bioline international official site (site up-dated) - Days to flowering was correlated to the emergence pattern. observed in cowpea (*Vigna unguiculata* L.), Nigeria. In: Genetics,

10. 1186/ 1471-2164-14-932 - biomed central - including the study of inheritance of qualitative and quantitative traits traits: days to flowering map of cowpea [*Vigna unguiculata*]

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the genetic architecture of ecophysiological and - Quantitative genetics days to flowering, Genotype environment interaction in carbon isotope discrimination and seed yield in cowpea (*Vigna unguiculata*)

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

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*),

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))

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

high-throughput snp genotyping in cucurbita pepo - (ranging from 20 to 60 days after pollination). Traits flowering and fruit quantitative traits Zraidi A Lelley T Progress in Cucurbit Genetics and

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quantitative traits in plants : beyond the qtl - Quantitative traits in plants: beyond the QTL Rice Hd1 Flowering time Transcription factor combine quantitative genetics with marker analysis (in

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

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