

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

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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

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

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

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*]

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

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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,

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

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

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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