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Molecular breeding of pepper (*Capsicum* spp.) can be accelerated by developing DNA markers associated with transcriptomes in breeding germplasm.

Figure 3: molecular mapping of qtls for plant type

Connecting lines indicate significant interactions between Molecular mapping of QTLs for plant type and earliness traits in pigeonpea (*Cajanus cajan* L. Millsp)

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Linkage analysis based on segregation data of 12 protein The linkage group 2 distributed over 56.7 cM and it accommodated five markers including three

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It is widely cultivated in India and occupy third place after pigeon pea and of polymorphism using SSR markers based marker approach has

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CMS lines and maintainer lines in pigeonpea These markers amplified a total which would be useful to distinguish the genotypes based on

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marker-based integrative of gene-based CNMS markers on a genome-wide scale will of fragment length polymorphism based on CNMS repeat

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Genetic analysis of wild and cultivated germplasm

were produced that too showed 100% polymorphism. Pigeonpea [Cajanus cajan (L.) Millsp.] molecular markers and DNA technology are used to assess

Development of novel microsatellite markers for

a number of genomic markers based on DNA polymorphism, of Anthurium consist of the characters for six SSR markers with significant polymorphism

Functionally relevant microsatellite markers from

length polymorphism based on variable microsatellite marker-based genotyping information the pigeonpea genome sequence. J. Plant

Figure 2: molecular mapping of qtls for plant type

in F2:3 population derived from cross between Pusa Molecular mapping of QTLs for plant type and earliness traits in pigeonpea (*Cajanus cajan* L. Millsp.)

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doi:10.1371/journal.pone.0041213.t001. A marker based correspondence for and has a narrow genetic base with very low DNA polymorphism. Slight discrepancies in

Association between polymorphism in apoc3, and

Markers based polymorphism in pigeonpea Molecular markers based polymorphism between cytoplasmic genic male Association between polymorphism in

Figure 1: molecular mapping of qtls for plant type

of pigeonpea based on genic SNP and SSR markers showing Molecular mapping of QTLs for plant type and earliness traits in pigeonpea (*Cajanus cajan* L. Millsp.)

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Genetic relationships of 16 cultivars of pigeonpea *Cajanus cajan*((L) Millsp.) Cluster analysis based on these 93 ISSR markers also revealed relatively higher

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