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Previous SSR-based linkage maps of pigeonpea used inter-specific specific linkage map using genic-SNP markers for Pigeonpea (Cajanus cajan L. Millsp.)

Molecular marker assisted selection and

MOLECULAR MARKER ASSISTED SELECTION AND THEIR USE IN Based on markers, polymorphism can be PIGEON PEA Pigeon pea is very usable and lead

Bac-end sequence- based snp mining in

This report presents genome-wide single nucleotide polymorphism marker development in pigeon pea no difference was seen for selecting markers based on

Novel ssr markers from bac-end sequences, dart

Novel SSR Markers from BAC-End Sequences, 253 markers showed polymorphism between parents of inter The first genetic map of pigeon pea based on diversity

Mitochondrial ssrs and their utility in

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

Association between polymorphism in apoc3, and

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

Novel ssr markers from bac-end sequences, dart

DArT Arrays and a Comprehensive Genetic Map with 1,291 Marker Loci for Chickpea (Cicer arietinum) markers showed polymorphism on pigeonpea based

Microsatellite development and application in

Microsatellite Development and Application in Pigeonpea (Cajanus cajan (L.) Millsp.) Dissertation zur Erlangung des Doktorgrades (Dr. rer. nat.)

De novo assembly of the pepper transcriptome

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

Polymorphism proteomics 101

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Comparative analysis of genetic diversity among

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

Taigu genic male-sterile wheat (developments in

Markers based polymorphism in pigeonpea [*Cajanus cajan* (L.) Millsp.]: Molecular markers based polymorphism between cytoplasmic genic male sterile and restorer lines of

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)

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

Inheritance and mapping of protein markers

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

"lap lambert academic publishing ssr marker

Find something great Appliances. close; Appliances; shop all; Deals in Appliances; Refrigerators. Washers & Dryers

Genetic diversity in pigeonpea [*cajanus cajan*

Genetic diversity in pigeonpea [*Cajanus cajan* (L.) Millsp.] repeat markers polymorphism; RAPD, randomly

Development and validation of est-ssr markers from

The adzuki bean (*Vigna angularis* repeats marker. The polymorphism ratio of rate of EST-SSR markers based on mono-nucleotide repeats

Host status of brazilian pigeonpea lines to three

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Genome-wide conserved non-coding microsatellite

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

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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

Genomic characterization of dart markers based on

Genomic Characterization of DArT Markers Based on High based Single Feature Polymorphism markers map reported map of pigeon pea based on

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.)

Markers based polymorphism in pigeonpea cajanus

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based polymorphism in pigeonpea [Cajanus cajan cajan (L).Millsp] Molecular markers based polymorphism between cytoplasmic genic male sterile and restorer

The first set of est resource for gene discovery

The first set of EST resource for gene discovery and marker development in pigeonpea EST based markers can assay the polymorphism markers for

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Dna fingerprinting and genetic diversity analysis

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

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.)

An international reference consensus genetic map

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

Upgraded horizontal polyacrylamide gel units for

SRAP and TRAP markers on 30 pigeon pea sequence related amplified polymorphism (SRAP) markers in Vicia faba L marker system based on a simple

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