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(SSR) markers. Diversity analysis detected a total of 178 alleles ...

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the tropical and subtropical fruits of great nutritional value, basically due to their vitamin and mineral contribution

Genetic resources and breeding of guava (Psidium guajava L ... The study showed Page 31/47

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markers were
arranged into 11
linkage groups
possibly
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representing the 11 chromosomes (2n = 22) of the guava genome with 9 to 24 markers per linkage group and a total genome length of 1,349 cM. QTL analysis for vegetative characters identified a total of 15 loci.

Molecular Of characterization Tuban accessions of quava ... Guava (Psidium quajava L.), one of the most widely grown plants in the tropics, is very susceptible to disease which can decrease its marketability. Leaf and fruit spot Page 39/47

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Abstract. A (GA) n and (GT) n microsa tellite-enriched library was constructed and 23 nuclear simple sequence repeat (SSR) loci were characterized in the guava species (Psidium guajava L.). All SSR loci were found to be polymorphic after

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