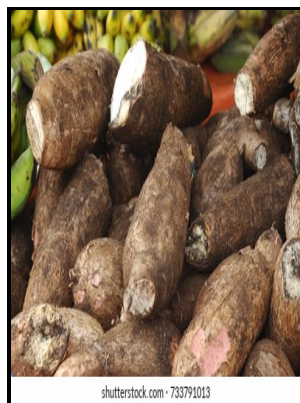


2.000 abstracts on cassava (manihot esculenta Crantz).

Cassava Information Center, Centro Internacional de Agricultura Tropical - 2,000 abstracts on cassava (manihot esculenta Crantz)



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Cassava genome from a wild ancestor to cultivated varieties

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Histological analysis of embryo

The sequence integrated 26- and 0.

Histological analysis of embryo

Keating BA, Evenson JP, Fukai S 1982. Proceedings of the Second International Scientific Meeting, CBN II, held at Bogor, Indonesia, 22—26 August 1994. Total yield losses occur in heavily infested fields.

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Methods in Molecular Biology, vol 1224. Gene family analysis in Euphorbiaceae BlastP was used on all the protein sequences against a database containing a protein data set of M. Eds SS Narwal, L Szajdak, DA Sampietro pp.

Characterization of hemicellulose in Cassava (Manihot esculenta Crantz) stem during xylogenesis

The genotypes showed similar leaf water potential Ψ_w in both the predawn period and the noon period Table 3.

Characterization of hemicellulose in Cassava (Manihot esculenta Crantz) stem during xylogenesis

These results show that the W14 draft genome sequence assembly is of high quality in spite of its high heterozygosity ; and complexity.

Cassava genome from a wild ancestor to cultivated varieties

From the total of 31,396 genes expressed in W14 or cultivated varieties, 749 show significantly differential expression in leaves and 2,732 in storage roots , ,. Taken together, these results suggest that transposon activity may have played a role in the reduction of cyanogenic glucoside content in the domesticated cassava. .

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