

## Genome constitution for *Musa beccarii* (Musaceae) varieties

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### 摘要

*Musa beccarii* N.W.Simmonds var. *beccarii* and *Musa beccarii* N.W.Simmonds var. *hottana* H? kkinen were described earlier in *Acta Phytotaxonomica et Geobotanica* based on their morphological characteristics. In order to distinguish the genomes between the two *M. beccarii* varieties and *M. coccinea* Andrews, we have now studied them with the Inter-Retrotransposon Amplified Polymorphism (IRAP) marker analyses. The high levels of IRAP polymorphism detected in this study showed that *M. beccarii* var. *beccarii* and *M. beccarii* var. *hottana* are two distinct varieties. Additional IRAP bands found in *M. beccarii* var. *beccarii* indicated that *M. beccarii* var. *hottana* might be evolutionarily more ancient than *M. beccarii* var. *beccarii* because LTR retrotransposons are not transpositionally removed.

关键词 [genome constitution](#) [inter-retrotransposon amplified polymorphism](#) [Musa beccarii var. beccarii](#) [Musa beccarii var. hottana](#) [Musa coccinea](#) [phylogenetic analysis.](#)

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