

Chloroplast genome of acanthochlamys bracteata (China) and xerophyta (Africa) completed

August 31 2021, by Zhang Nannan



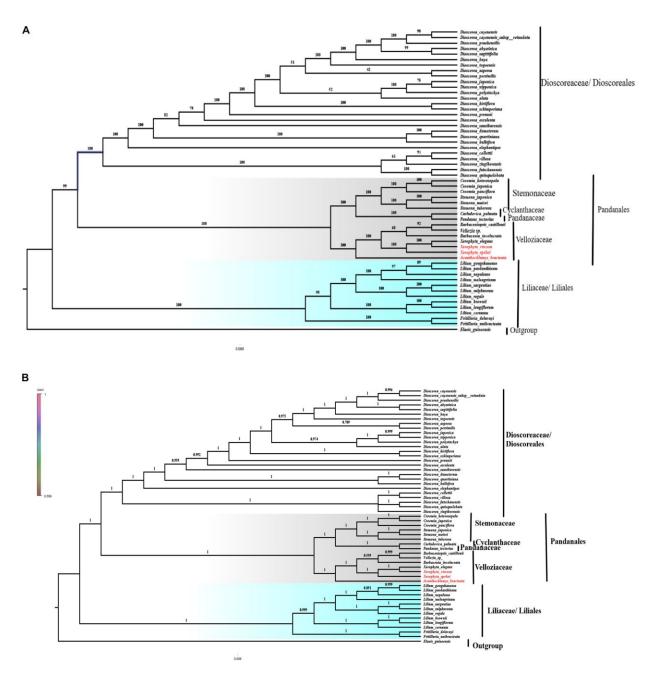


Fig. 1. Phylogenetic relationship of the three species of Velloziaceae based on 59 single-copy genes shared by all chloroplast genomes. A. The Maximum likelihood (ML) tree with bootstrap value on the branches; B. Bayesian inference (BI) tree with posterior probabilities values on the branches. Credit: WBG



Velloziaceae is a monocot family that is made up of five genera and c. 250 species with a disjunct distribution in Africa mainland, Madagascar, Arabian Peninsula, China and South America. They are recognized as the largest lineage of vascular plants that can tolerate desiccation, hence are resurrection plants.

The family is one of the classical examples of "taxonomic nightmares" among <u>plants</u>, due to its floral similarities and the huge variability in the morphological features in terms of leaf forms, sizes and life forms among others. Thus, relationship among the members of this family is still unclear.

Researchers at the Flora and Plant Taxonomy in Eastern Africa Research Group of the Wuhan Botanical Garden of the Chinese Academy of Sciences (CAS)analyzed the complete chloroplast genomes of Acanthochlamys bracteata collected from the Hengduan Mountains, China, and they did both molecular and morphological comparative analysis with two African Xerophytaspecies (Xerophyta spekei and Xerophyta viscosa).

The chloroplast genomes of Acanthochlamys bracteata, X.spekei, and X. viscosa are 153,843 bp, 155,235 bp, and 155,498 bp respectively. The chloroplast genomes consist of inverted repeats (IRa and IRb), each with a length ranging from 27,022 bp to 27,110 bp in the three species. The Large Single-Copy region in the three species shows length ranging from 81,919 bp to 83,813 bp and Small Single-Copy region (17,387 bp to 17,880 bp).





Fig. 2. Photo of Xerophyta spekei Baker in Chyulu Hills, Kenya. Credit: Prof. HU Guangwan

The <u>phylogenetic analysis</u> showed that Acanthochlamys bracteata P.C. Kao is the early diverged species in the family Velloziaceae because it is sister to the rest of the Velloziaceae <u>species</u>.

In terms of morphological and anatomical comparisons, Acanthochlamys is different from the rest of the family (Velloziaceae), with exception of sieve tube plastids. The <u>family</u> Velloziaceae currently is governed by four-character states: persistent leaves, presence of abscission zones, two phloem strands and violet tepals. Despite their geographical occurrences, they fit the classification into similar taxa.





Fig. 3. Photo of Acanthochlamys bracteata P.C. Kao from China. Credit: Prof. HU Guangwan

The research entitled "Complete Chloroplast Genomes of Acanthochlamys bracteata (China) and Xerophyta (Africa) (Velloziaceae): Comparative Genomics and Phylogenomic Placement"



was published in *Frontiers in Plant Science*, Plant Systematics and Evolution.

More information: Vincent Okelo Wanga et al, Complete Chloroplast Genomes of Acanthochlamys bracteata (China) and Xerophyta (Africa) (Velloziaceae): Comparative Genomics and Phylogenomic Placement, *Frontiers in Plant Science* (2021). DOI: 10.3389/fpls.2021.691833

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