

Review of: "First high-quality genome assembly data of sago palm (*Metroxylon sagu* Rottboll)"

zhang chuan-qing

Potential competing interests: The author(s) declared that no potential competing interests exist.

This ms. provided high-quality genome assembly data of sago palm by utilization of the Nanopore sequencing. This will be undoubtedly enhance functional genetic characterization. However, the genomic characteristics should be been further analyzed.