

High-quality reference plastomes in *Tillandsia* species living at the dry limits

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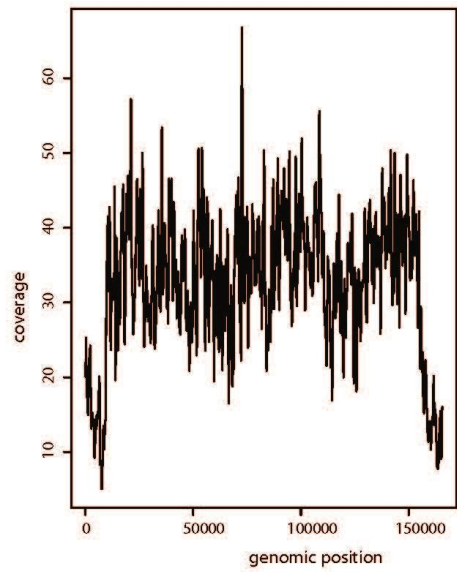
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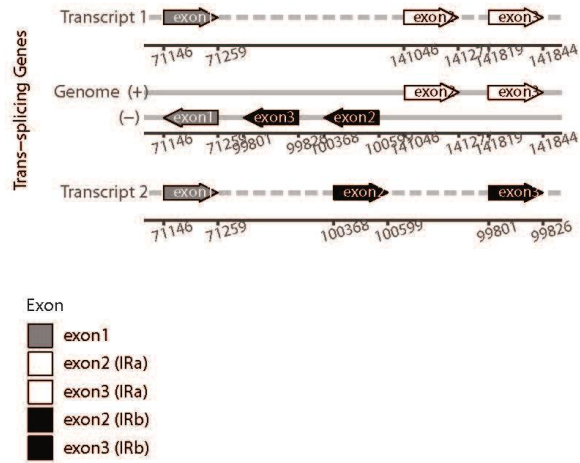
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Online Resource 6. (a) Sequencing coverage for sample TL10611 obtained by back-mapping the sequencing reads to the completed plastid genome. A sliding window approach was used (window size 500 bp, step size 100 bp), and the resulting mean values over the 500 bp windows were plotted in R. **(b)** Trans-splicing genes as annotated by the CPGView package (Liu et al. 2023). **(c)** Cis-splicing genes were plotted as annotated by the CPGView package (Liu et al. 2023).

A



B



C

