

CORRECTION

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Correction: Complete chloroplast genomes provide insights into evolution and phylogeny of *Zingiber* (Zingiberaceae)

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Correction: BMC Genomics 24, 30 (2023)
<https://doi.org/10.1186/s12864-023-09115-9>

Following the publication of the original article [1], the authors reported that the 2nd author was assigned affiliation 1 erroneously. Xiaodong Cai's affiliation is presented correctly in this correction article.

Moreover, the authors identified an error in Fig. 3. The correct figure is given below.

The original article [1] has been updated.

Reference

1. Jiang D, Cai X, Gong M, et al. Complete chloroplast genomes provide insights into evolution and phylogeny of *Zingiber* (Zingiberaceae). *BMC Genomics*. 2023;24:30. <https://doi.org/10.1186/s12864-023-09115-9>.

Published online: 14 July 2023

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The original article can be found online at <https://doi.org/10.1186/s12864-023-09115-9>.

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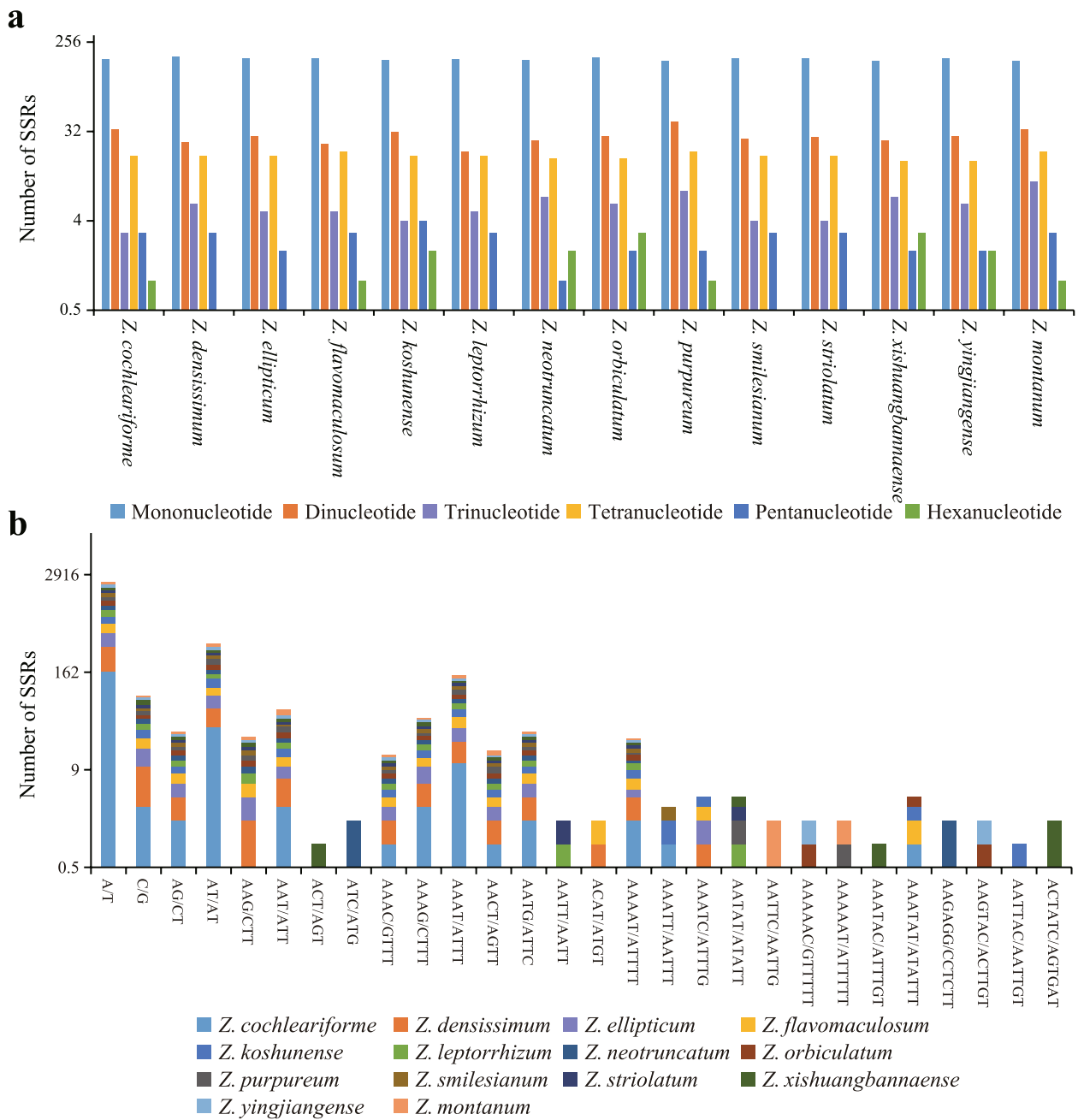


Fig. 3 Comparison of the simple sequence repeats (SSRs) among fourteen *Zingiber* species. **a** The number of different SSR types. **b** The frequency of the identified SSRs in different repeat class types