Complete chloroplast genome sequences of *Hippophae neurocarpa*

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**ABSTRACT**

*Hippophae neurocarpa* are dioecious deciduous shrubs, diffused in the Qinghai–Tibetan Plateau at high altitude. Here, we report the complete chloroplast genome of *H. neurocarpa*. The chloroplast genome is found to be 156,316 bp in length with 36.65% GC contents. The chloroplast genome sequences contained 124 genes, including 78 protein-coding genes, 38 tRNA genes, and 8 rRNA genes. The complete chloroplast genome of *H. neurocarpa* will be beneficial for identifying molecular markers for further conservation and utilization of these multipurpose natural resources.
Disclosure statement
No potential conflict of interest was reported by the authors.

Funding
This work was supported by the project of Qinghai Science & Technology Department [2016-ZJ-Y01], the open project of State Key Laboratory of Plateau Ecology and Agriculture, Qinghai University [2018-ZZ-09], the Qinghai Provincial Science Foundation [2015-NK-509, 2019-ZJ-966Q, 2017-SF-A8].

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Figure 1. Maximum-likelihood phylogenetic tree inferred from 31 complete chloroplast genome sequences. The position of H. neurocarpa is marked in red and bootstrap values are listed for each branch.